

# ***MICROACTIVE GEMINI<sup>®</sup> VII***

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***SURFACE AREA ANALYZER***



## ***OPERATOR MANUAL***

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239-42828-01  
July 2018  
(Rev F)

## ***TRADEMARKS***

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## ***CONTACT US***

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## ABOUT THIS MANUAL

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Log in to your [customer portal](#) to access the following:

- Calculations
- Error Messages
- MicroActive Gemini VII Operator Manual in PDF format
- MicroActive Report Tutorials
- Smart VacPrep Operator Manual in PDF format

Parts and accessories can be found online at [www.Micromeritics.com](http://www.Micromeritics.com).



All references to Gemini VII in this document encompass the Gemini VII models 2390a, 2390p, and 2390t unless otherwise noted. It also encompasses all references to the MicroActive Gemini VII and MicroActive Gemini VII Confirm unless otherwise noted.



This manual contains instructions for both standard installations and installations in 21CFR11 environments.

The following icons may be found in this manual:



**NOTE** — Notes contain important information applicable to the topic.



**CAUTION** — Cautions contain information to help prevent actions that may damage the analyzer or components.



**WARNING** — Warnings contain information to help prevent actions that may cause personal injury.



**NOTE** — Notes that apply to 21CFR11 environments only (Confirm applications).

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## 1 ABOUT THE ANALYZER

The Gemini VII series analyzers provide single and multipoint surface area and pore size measurements. Three models are available — the Gemini 2390a, 2390p, and 2390t.

The Gemini 2390a and 2390p are enclosed in the same size cabinet. The only physical difference is the  $P_0$  (saturation pressure) tube which is installed on the Gemini 2390p, allowing continuous measurement of the saturation pressure.

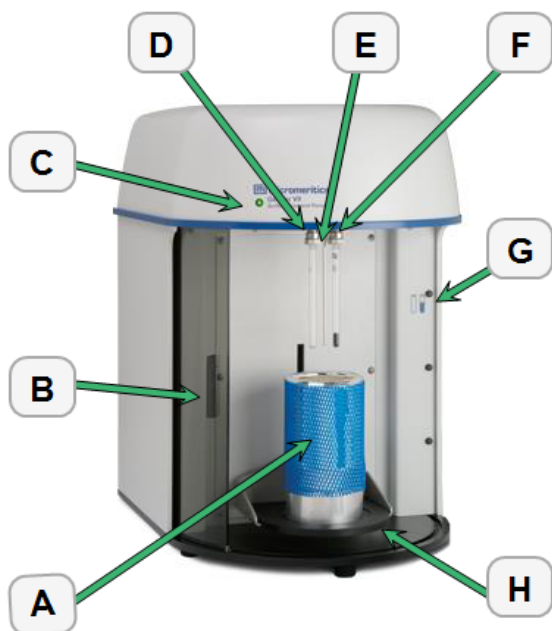
The Gemini 2390t is in a slightly larger cabinet allowing the use of a larger dewar and longer sample tubes for extended analyses. This model also is equipped with a  $P_0$  tube.

Most application features are available on all three models, with the exception of a continuous  $P_0$  measurement and a few reports. Refer to the following table for a brief summary of some of the model reports and features.

### Gemini Model Features

Feature	2390a	2390p	2390t
Single and multipoint BET surface area	✓	✓	✓
Langmuir surface area	✓	✓	✓
Automatically generated pressure tables with user-selected points	✓	✓	✓
Total pore volume	✓	✓	✓
t-Plot method micropore volume and area	✓	✓	✓
Analysis log	✓	✓	✓
Statistical Process Control (SPC) reports	✓	✓	✓
Horvath-Kawazoe data reduction	✓	✓	✓
BJH pore size distribution using adsorption isotherm		✓	✓
BJH pore size distribution using desorption isotherm			✓
Continuous measurement of saturation pressure ( $P_0$ )		✓	✓
Dubinin reports	✓	✓	✓
MP-Method reports	✓	✓	✓
DFT reports	✓	✓	✓

## FRONT COMPONENTS



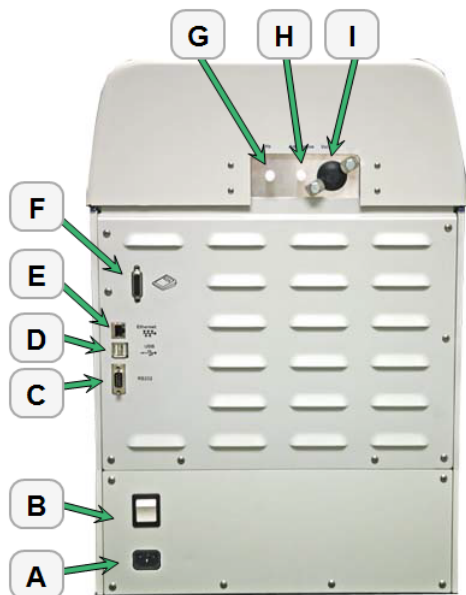
### Front Components

Component		Description
A	Dewar	For housing cryogenics used for analysis. The dewar rests on an elevator.
B	Sample compartment door	Closes to cover the dewar.
C	Power indicator light	Blinks when power is applied to the analyzer; illuminates when the analysis program is initiated and ready for operation.
D	Balance port and balance tube	The balance tube is identical to the sample tube. This design negates free-space errors introduced by thermal gradient variations or by initial mis-measurement of free space.
E	P <sub>0</sub> port and tube	Models 2390p and 2390t only. For measuring the saturation pressure.

## Front Components (continued)

Component		Description
<b>F</b>	Sample port and sample tube	For installing one sample tube and one balance tube.
<b>G</b>	Tube position diagram	Diagram of port numbers and locations as well as the positions of the balance and sample tubes. The empty tube represents the balance tube installed on the left port. The filled tube represents the sample tube installed on the right port.
<b>H</b>	Elevator	Allows placement of the dewar around the sample and $P_0$ tubes. The elevator is raised automatically when the analysis is started and lowers automatically upon completion. During analysis, the elevator optionally lowers after the free space measurement to allow evacuation, then is raised and continues the analysis.

## BACK PANEL COMPONENTS



### Front Components

Component	Description
<b>A</b>	Power connector.
<b>B</b>	Power switch.
<b>C</b>	RS232 connector - for service personnel use.
<b>D</b>	USB ports. Not used.
<b>E</b>	Ethernet port for Ethernet cable for communication between the analyzer and the computer.
<b>F</b>	Keypad connection. Not used.
<b>G</b>	Helium inlet port - for helium gas supply.
<b>H</b>	Adsorptive inlet port - for analysis gas supply.
<b>I</b>	Vacuum pump connector - for vacuum pump hose.



## DEGASSER OPTIONS

Parts and accessories can be found online at [www.Micromeritics.com](http://www.Micromeritics.com).

Degasser Option	Description
<b>FlowPrep</b>	The FlowPrep applies both heat and a stream of inert gas to the sample to remove adsorbed contaminants from the surface and pores in preparation for analysis for up to six samples. Choose the temperature, gas, and flow rate best suited for your sample material. The FlowPrep is an independent unit and not controlled by the analyzer.
<b>Smart VacPrep</b>	<p>The Smart VacPrep prepares samples by heating and evacuation. It contains six sample ports in which up to five temperatures, ramp rates, and soak times per sample are individually controlled by the analyzer program so that all degas information is integrated into the sample data file for future reference. Samples can also be prepared, started, and completed independently. There is no need to wait for samples on other ports to finish. Front panel buttons allow a QuickStart operation with preprogrammed conditions.</p> <p>Up to three additional Smart VacPrep degassers can be connected to one computer permitting 24 preparation ports to be used. The Smart VacPrep is the recommended degassing unit.</p>
<b>SmartPrep</b>	The SmartPrep passes flowing-gas over the sample at elevated temperatures. It contains six sample ports in which temperature, ramp rates, and soak times are individually controlled by the analyzer program so that all degas information is integrated into the sample data file for future reference. It contains two serial ports — one for connecting to the computer and the other for connection of up to three additional SmartPrep degassers permitting 24 preparation ports to be used.
<b>VacPrep</b>	The VacPrep offers two methods for removing contaminants. In addition to flowing gas, it provides vacuum to prepare samples by heating and evacuation of up to six samples. This combination provides preparation method options best suited to your material or application. Needle valves are also provided for introducing the vacuum slowly to prevent fluidization of samples. The VacPrep is an independent unit and not controlled by the analyzer.

## EQUIPMENT OPTIONS AND UPGRADES

Parts and accessories can be found online at [www.Micromeritics.com](http://www.Micromeritics.com).

Equipment Option	Description
<b>Chiller Dewar</b>	<p>For Gemini model 2390t.</p> <p>A closed loop recirculating system that utilizes a high surface area copper coil to provide excellent heat transfer between the dewar and the recirculating liquids. Log in to your <a href="#">customer portal</a> to access the Chiller Dewar Quick Start Guide [part number 025-42801-00].</p>
<b>Vacuum Pump</b>	<p>The analyzer requires a vacuum pump for sample analysis. Vacuum pumps used must meet the following criteria:</p> <ul style="list-style-type: none"> <li>• achieve vacuum levels of 20 <math>\mu</math>mHg at the analyzer inlet</li> <li>• contain an anti-suckback valve to prevent vacuum pump oil from back streaming into the analyzer in the event of a vacuum pump failure</li> <li>• contain an NW16 inlet port for connection to the analyzer</li> </ul> <hr/> <div>  <p>A device to reduce oil vapor backstreaming is recommended.</p> </div> <hr/> <div>  <p>The vacuum pump must have an anti-suckback valve to prevent oil from being admitted into the analyzer should the power fail while the system is under vacuum. Pumps available from Micromeritics are equipped with an anti-suckback valve.</p> </div> <hr/> <p>An oil-based or oil-free vacuum pump can be used with the analyzer. Appropriate vacuum pumps are available from Micromeritics.</p>

## ***GAS REQUIREMENTS***

Compressed gases are required for analyses. Gas cylinders or an outlet from a central source should be located near the analyzer.

Appropriate two-stage regulators which have been leak-checked and specially cleaned are required. Pressure relief valves should be set to no more than 30 psig (200 kPag). All gases should be of a purity of 99.999% or better. Gas regulators can be ordered from Micromeritics. Parts and accessories can be found online at [www.Micromeritics.com](http://www.Micromeritics.com).

When helium is used for differential free-space measurement and nitrogen is used as the adsorbate gas, they should be of the following purity or better:

- Helium — purity of 99.9%. For analysis of materials with very low surface areas, Micromeritics recommends use of helium with purity of 99.995%.
- Nitrogen — purity of 99.9%. For analysis of materials with very low surface areas, Micromeritics recommends use of nitrogen with purity of 99.995%.

(CGA 580) N <sub>2</sub>	99.999%
(CGA 580) He	99.999%

## ***CRYOGEN REQUIREMENTS***

Liquid nitrogen is used as the cryogen to cool the sample during analysis. A liquid nitrogen transfer system eliminates the need to pressurize storage dewars. The Model 021 liquid nitrogen transfer system is available from Micromeritics. Log in to your [customer portal](#) to access parts and accessories.

## SPECIFICATIONS FOR THE GEMINI VII

Specification	Description
<b>Analysis</b>	
<b>Surface Area</b>	From 0.1 m <sup>2</sup> /g, total From 0.01 m <sup>2</sup> /g, specific
<b>Pore Volume</b>	From 4 × 10 <sup>-6</sup> cm <sup>3</sup> /g
<b>Electrical</b>	
<b>Frequency</b>	50 to 60 Hz
<b>Power</b>	150 VA, maximum
<b>Voltage</b>	85 to 265 VAC
<b>Environment</b>	
<b>Humidity</b>	20 to 80% relative, noncondensing
<b>Temperature</b>	10 °C and 35 °C (50 °F to 95 °F), operating 0 °C to 50 °C (32 °F to 122 °F), nonoperating
<b>Gases</b>	
<b>Adsorbate</b>	Optimized for nitrogen in a liquid nitrogen sample bath. Gemini may be used with non-corrosive adsorbate gases having vapor pressures at both room and bath temperatures that are acceptably high relative to the resolution of the 1,000 mmHg pressure transducer. Typically, oxygen, argon, carbon dioxide, butane, methane, and other light hydrocarbons will produce useful data above absolute pressures of a few mmHg.
<b>Physical</b>	
<b>Height</b>	23 in. (59 cm) - Gemini VII 2390a and 2390p 29 in. (74 cm) - Gemini VII 2390t
<b>Width</b>	16 in. (40 cm)
<b>Depth</b>	20 in. (51 cm)
<b>Weight</b>	75 lbs. (32 kg) - Gemini VII 2390a and 2390p 78 lbs. (35 kg) - Gemini 2390t
<b>Pressure Measurement</b>	
<b>Range</b>	0 to 950 mmHg
<b>P/P<sub>0</sub> Resolution</b>	<10 <sup>-4</sup>
<b>Relative Pressure Range</b>	0 to 1.0 P/P <sub>0</sub> (adsorption only)



Specification	Description
Pressure Resolution	<0.1 mmHg
Accuracy and Linearity (transducer manufacturer specification)	Better than $\pm 0.5\%$ full scale
<b>Sample Tube and Dewar</b>	
Standard Tube	<ul style="list-style-type: none"> <li>• <b>Gemini VII 2390a and 2390p.</b> 0.95 cm (3/8 in.) OD, 15.5 cm (6.1 in.) long with 6.5 cm<sup>3</sup> of volume. Sample capacity is approximately 2.0 cm<sup>3</sup>.</li> <li>• <b>Gemini VII 2390t.</b> 0.95 cm (3/8 in.) OD <math>\times</math> 20.5 cm (8.1 in.) long with 8.9 cm<sup>3</sup> of volume. Sample capacity is approximately 2.0 cm<sup>3</sup>.</li> </ul>
Dewar	<ul style="list-style-type: none"> <li>• <b>Gemini VII 2390a and 2390p.</b> 8 hours</li> <li>• <b>Gemini VII 2390t.</b> &gt;24 hours</li> </ul>
<b>Vacuum System</b>	
<p>An external vacuum source achieving <math>20 \times 10^{-3}</math> mmHg (or better) at the instrument inlet for oil-based or oil-free pumps.</p> <p>For oil-based pumps, an anti-suckback valve is required to prevent oil from being admitted into the analyzer should there be a power failure. A device to reduce oil vapor backstreaming is also recommended.</p>	
<b>Computer Requirements</b>	
<ul style="list-style-type: none"> <li>• <b>Operating System.</b> Windows 7 Professional or higher operating system is recommended for the best user experience. For 21CFR11 environments, Windows 10 Professional or Windows 10 Enterprise or higher is required.</li> <li>• <b>Desktop Installation Required.</b> The application should not be installed on a network drive with shared access. Multiple users cannot operate the application at the same time.</li> <li>• <b>10 Base T or 100 Base T Ethernet Port.</b> If the computer is to be connected to a network, two Ethernet ports are required. If more than one Ethernet based unit is connected to the same computer, an Ethernet switch will also be required.</li> <li>• <b>Read / Write Permissions.</b> All users of the application will need Read/Write permission to all directories and subdirectories where the application is installed.</li> <li>• <b>Drives.</b> CD-ROM drive and USB port.</li> </ul>	
<p><i>In keeping with a policy of ongoing product improvement, specifications are subject to change without notice.</i></p>	

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## 2 ABOUT THE SOFTWARE

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**See also:**

[About the Software in 21CFR11 Environments on page 2 - 24](#) for CFR11 environments.

The analyzer allows other computer programs to run while an automatic operation is in progress. The *Help* menu provides access to the online operator manual.

Report options can be specified when creating the sample file. When running an analysis, data gathered during the analysis process are compiled into the predefined reports. Reports can also be defined and generated after an analysis has been run. Each selected report is displayed on its own tab and reflects data collected during the analysis.

The MicroActive feature offers a Windows interface with an easy way to collect, organize, archive, reduce raw data, and store sample files for later use. Scalable and editable graphs, and copy and paste graphics, are easily generated. Customized reports can be generated to computer monitor, paper, or exported for use in other programs.

Data can be manipulated and displayed interactively using MicroActive reports.

### SOFTWARE SETUP



If the computer is to be connected to a network, a second Ethernet port on the computer must be used for that purpose.

The *Setup* program is located on the installation media and is used to reinstall the software and make analyzer changes — such as adding or removing a unit, etc.



If the IP address needs to be changed on the computer connected to the analyzer, refer to the computer's operating system manual or the internet for instructions. The IP address for the computer and the IP address specified in the setup program must match. The IP address must be 192.168.77.100.

## MENU STRUCTURE

All program functions use standard Windows menu functionality. The title bar contains a *Unit [N]*. If multiple analyzers are installed, ensure the appropriate unit is selected before continuing.

### Main Menu Bar Options

Option	Description
<b>File</b>	Use to manage files used by the application — such as sample files, analysis conditions files, report options files, etc.
<b>Unit [n]</b>	Use to perform analyses, calibrations, and other analyzer operations. <i>Unit [n]</i> displays on the menu bar for each analyzer attached to the computer.
<b>Smart VacPrep</b>	Use to access the menu for each installed Smart VacPrep. This menu does not display if a Smart VacPrep is not installed.
<b>SmartPrep</b>	Use to access the menu for each installed SmartPrep. This menu does not display if a SmartPrep is not installed.
<b>Reports</b>	Use to run reports and view the results.
<b>Options</b>	Use to edit the default method, specify system configuration, specify units, and change presentation options.
<b>Window</b>	Use to manage open windows and display a list of open windows. A checkmark appears to the left of the active window.
<b>Help</b>	Provides access to the online operator manual, the Micromeritics web page, the analyzer web page, and information about the application.

## COMMON FIELDS AND BUTTONS

The fields and buttons in the following table are located in multiple windows throughout the analyzer application and have the same description or function. Fields and button descriptions not listed in this table are found in tables in their respective sections.

Common Fields and Buttons Table

Field or Button	Description
<b>Add</b>	Add an item to the list.
<b>Add Log Entry</b>	Use to enter information to appear in the sample log report that cannot be recorded automatically through the application. Click the button again to enter multiple log entries.
<b>Append</b>	Use to insert one row at the end of a table.
<b>Autoscale</b>	When enabled on report parameters windows, allows the x- and y-axes to be scaled automatically. <i>Autoscale</i> means that the x- and y-ranges will be set so that all the data is shown. If <i>Autoscale</i> is not selected, the entered range is used.
<b>Axis Range</b>	On report parameters windows, the <i>From / To</i> fields are enabled when <i>Autoscale</i> options are not selected. Enter the starting and ending values for the x- and/or y-axes.
<b>Bar Code</b> (default field label name)	Go to <i>Options &gt; Default Method</i> to rename this field to something suitable for your environment. A bar code reader may be used to enter text into many of the fields on the <i>Sample Description</i> window. The bar code reader must be connected to the computer's USB port. To use a bar code reader, use a mouse to click in the field first where information is to be entered then scan the bar code with the bar code reader.  If a bar code reader is not used, this alphanumeric field can be used to enter additional information about the sample, such as a sample lot number, sample ID, etc.
<b>Browse</b>	Searches for a file.
<b>Cancel</b>	Discards any changes or cancels the current process.
<b>Clear</b>	Use to clear the table entries and display only one default value.
<b>Close</b>	Closes the active window.
<b>Close All</b>	Closes all active windows. If changes were made and not yet saved, a prompt displays for each changed file providing the option to save the file.

## Common Fields and Buttons Table (continued)

Field or Button	Description
<b>Comments</b>	Enter comments about the sample or analysis. Comments display in the report header.
<b>Copies</b>	Select the number of copies to print. This field is only enabled when <i>Print</i> is selected.
<b>Delete</b>	When working with pressure tables and pressures, <b>Delete</b> removes the selected information.
<b>Destination</b>	Select the report destination.
<b>Edit</b>	When working with report parameters, highlight the item in the <i>Selected Reports</i> list box and click <b>Edit</b> to modify the report details.
<b>Exit</b>	Exits the application. If a file is open with unsaved changes, a prompt displays providing the option to save the changes and exit or to exit the application without saving the changes.
<b>Export</b>	Exports data in a sample file as a .REP, .TXT or .XLS file. When saved to a file, the data can be imported into other applications.
<b>File</b>	Select the destination directory. Enter a new file name in the <i>File name</i> field, or accept the default. Select to save the file as a report system (.REP), a spreadsheet (.XLS), a portable document format (.PDF), or an ASCII text (.TXT) file format.
<b>File name</b>	Select a file name from the list shown or enter a file name. If the required file type is not shown, select the type of file from the list.
<b>From / To</b>	When working with report parameters windows, enter the <i>From</i> and <i>To</i> range for x- and/or y-axes.
<b>Insert</b>	Use to insert one row above the selected row in the table.
<b>List</b>	Creates a list of sample or other type files. The list will contain file name, date / time the file was created or last edited, file identification, and file status.
<b>Name</b>	The <i>Name</i> column is a list of files in the selected directory or library.
<b>Next</b>	Click to move to the next window or next step.
<b>OK</b>	Saves and closes the active window.
<b>Open</b>	Opens the selected file. Alternatively, double click the file name in the <i>Name</i> column to open the file.
<b>Prev</b>	Click to move to the previous window.
<b>Preview</b>	Previews predefined reports. Click the tabs at the top of the window to preview each selected report. When an analysis has not been run on a sample, this button is disabled.

Common Fields and Buttons Table (continued)

Field or Button	Description
<b>Print</b>	Sends the report to the selected destination (screen, printer, or file).
<b>Remove</b>	Remove the selected file or files from the list.
<b>Replace</b>	Click to select another file where the values will replace the current file's values.
<b>Replace All</b>	Click to select another .SMP file where the values will replace all values for the active sample file. The original file will remain unchanged.
<b>Report</b>	Click to display a window to specify report output options.
<b>Save</b>	Saves changes to the active window.
<b>Save As</b>	Saves a file in the active window under a different file name.
<b>Start</b>	Starts the report, test, analysis, or operation.
<b>Start Date</b>	Displays a calendar to select the start date for the report.
<b>View</b>	<p><b>Operation.</b> Use to display the data from the current analysis.</p> <p><b>Instrument Log.</b> Use to display recent analyses, calibrations, errors, or messages.</p> <p><b>Instrument Schematic.</b> Use to display a schematic of the analyzer system.</p>

## FILE STATUS, DESCRIPTION, EXTENSION, AND LOCATION

In the *File Selector* window, the *Mic Description* column and the *Mic Status* column display file description and file status. The *File Selector* incorporates standard Windows features for resizing windows, reordering and repositioning columns, and right clicking an entry to display a menu of standard Windows functions.

### File Status and Description Table

File Status	Description
Analyzing	Sample files that are currently being used for analysis.
Complete	Sample files used in an analysis that has been completed.
Entered	Sample files containing manually entered data.
No Analysis	Sample files which have not been used to perform an analysis.
Prepared	Sample files that have been used in an automatic degas operation but have not been analyzed. This status is applicable only if using the Smart VacPrep or SmartPrep degasser.
Preparing	Sample files that are currently being used in an automatic degas operation. This status is applicable only if using the Smart VacPrep or SmartPrep degasser.

### File Type, Extension, and Location Table

File Type	File Name Extension	Default Location
Alpha-s Curve *	.ALS	Param directory
Adsorptive Properties	.ADP	Param directory
Analysis Conditions	.ANC	Param directory
Degas Conditions	.DEG	Param directory
Heat of Adsorption Report	.HOA	Param directory
Methods	.MTH	Param directory
Report Options	.RPO	Param directory
Sample Information	.SMP	Data directory
Thickness Curve **	.THK	Param directory

\* **Alpha-S Curve file.** Saves the relative pressures and resulting quantities adsorbed as an ASCII text file. These data are derived by dividing the isotherm by the quantity adsorbed at 0.4 relative pressure.



**\*\* Thickness Curve file.** Saves the relative pressures and corresponding thicknesses as an ASCII text file. These data are derived by dividing the condensed volume of adsorptive by the selected surface area. The density conversion factor in the adsorptive properties file is used to convert quantity adsorbed to volume of condensed adsorptive.

File types available when printing or exporting reports:	
Portable document format	.PDF
Report	.REP
Spreadsheet	.XLS
Unicode	.TXT
Extensible markup language	.XML

## KEYBOARD SHORTCUTS

Shortcut keys can be used to activate some menu commands. Shortcut keys or key combinations (when applicable) are listed to the right of the menu item.

Certain menus or functions can also be accessed using the **Alt** key plus the underlined letter in the menu command. For example:

- to access the *File* menu, press **Alt + F**, then press the underlined letter on the submenu
- **Alt + F** opens the *File* menu, then press **O** to access the *File Selector*



If the underscore does not display beneath the letter on the menu or window, press the **Alt** key on the keyboard.

### Keyboard Shortcut Table

Field or Button	Description
<b>Alt + [Unit n]</b>	Opens the <i>Unit [n]</i> menu.
<b>Alt + F</b>	Opens the <i>File</i> menu.
<b>Alt + F4</b>	Exits the program. If files are open with unsaved changes, a prompt to save changes displays.
<b>Alt + H</b>	Opens the <i>Help</i> menu.
<b>Alt + I</b>	Opens the <i>Options</i> menu.
<b>Alt + R</b>	Opens the <i>Reports</i> menu.
<b>Alt + V</b>	Opens the <i>Smart VacPrep</i> menu.
<b>Alt + W</b>	Opens the <i>Window</i> menu.
<b>Shift + F9</b>	Opens the shortcut menu of (1) selected component on analyzer schematic when manual control is enabled or (2) onscreen reports.
<b>Ctrl + N</b>	Opens a new sample file.
<b>Ctrl + O</b>	Opens the <i>File Selector</i> window.
<b>Ctrl + P</b>	Opens the <i>File Selector</i> to start a report from a selected .SMP file.
<b>Ctrl + S</b>	Saves the open file.
<b>F1</b>	Opens the online help operator manual.
<b>F2</b>	Opens the <i>File Selector</i> window.
<b>F3</b>	When in the <i>File Selector</i> window, opens the file search box.
<b>F4</b>	When in the <i>File Selector</i> window, opens the address bar.

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**Keyboard Shortcut Table (continued)**

Field or Button	Description
<b>F6</b>	Cascades open windows.
<b>F7</b>	Tiles all open application windows.
<b>F8</b>	Opens the <i>File Selector</i> to start a report from a selected .SMP file.
<b>F9</b>	Closes all open reports.
<b>F10</b>	Opens the <i>Heat of Adsorption</i> window.

## OPTION PRESENTATION

### Options > Option Presentation

#### CFR Note

For 21CFR11 environments, see [Option Presentation for 21CFR11 Environments on page 2 - 12](#).

Use to change the way sample files and parameter files display: *Advanced*, *Basic*, or *Restricted*. Each display option shows sample information and options differently.

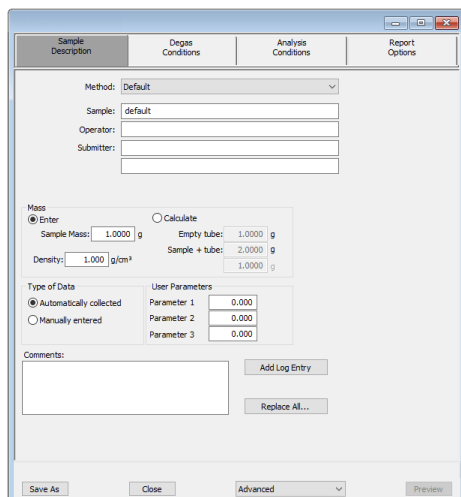
#### Option Presentation Display Table

Presentation Display	Description
<b>Advanced</b>	Displays all parts of sample and parameter files. Navigate to parameter windows by selecting the tabs across the top of the window.
<b>Basic</b>	Displays sample information in a single window. This display option is used after the parameter files have been created. The previously entered or default parameter files are then accessible using drop-down lists.
<b>Restricted</b>	Displays the sample file in a single window similar to the <i>Basic</i> display option with certain functions disabled. A password is set when the <i>Restricted</i> option is selected. That same password must be entered to change to the <i>Basic</i> or <i>Advanced</i> display option. This display type is typically used in laboratories where analysis conditions must remain constant — such as the pharmaceutical industry. The <i>Advanced</i> option is not available at the bottom of the window when using the <i>Restricted</i> display option.
<b>Always Open Edit View</b>	Opens files with a <i>Complete</i> status in tabbed file editor rather than in the Peak Editor view.
<b>Show Splash Screen</b>	Enables (or disables) the splash screen upon application startup.
<b>Show Degas Conditions</b>	When enabled, displays the <i>Degas Conditions</i> tab when using <i>Advanced</i> option presentation and the Degas Conditions drop-down list when using <i>Basic</i> or <i>Restricted</i> option presentation. This option may be deselected to hide the <i>Degas Conditions</i> tab if not using a SmartPrep or Smart VacPrep.

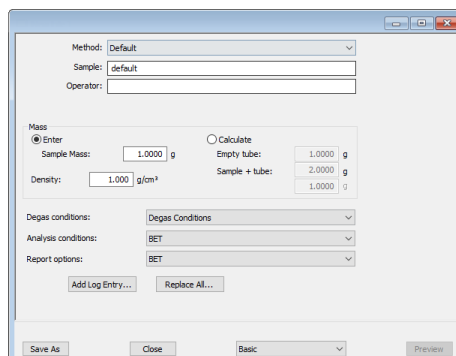


Specify or change the default option presentation by selecting **Options > Option Presentation**, or select *Basic* or *Advanced* from the drop-down list at the bottom of the window.

The following examples show the same sample file in *Advanced* and *Basic* display. *Basic* and *Restricted* displays will look the same.



**Advanced  
option presentation**



**Basic / Restricted  
option presentation**



A sample file must be created for each analysis. The file can be created prior to or at the time of analysis. The sample file identifies the sample, guides the analysis, and specifies report options.

## OPTION PRESENTATION FOR 21CFR11 ENVIRONMENTS

### Options > Option Presentation

#### Option Presentation Display Table

Presentation Display	Description
<b>Show Degas Conditions</b>	When enabled, displays the <i>Degas Conditions</i> tab when using <i>Advanced</i> option presentation and the Degas Conditions drop-down list when using <i>Basic</i> option presentation. This option may be deselected to hide the <i>Degas Conditions</i> tab if not using a SmartPrep or Smart VacPrep.
<b>Show Splash Screen</b>	Enables (or disables) the splash screen upon application startup.

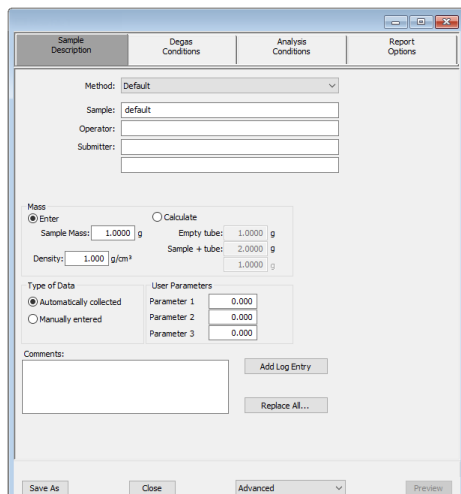
#### CFR Note

**For members of the Developer group only.** To change the view from *Advanced* (for Developers) to *Basic* (for Analysts), click the drop-down arrow at the bottom of the window. Select either *Advanced* (when in *Basic* view) or *Basic* (when in *Advanced* view).

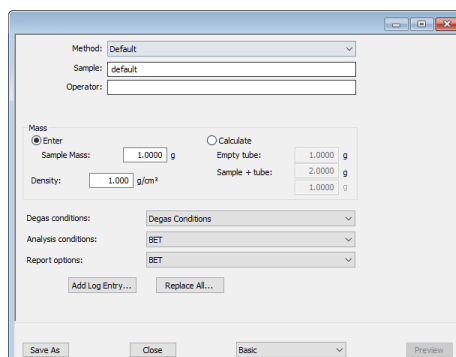
#### Option Presentation Display Table

Presentation Display	Description
<b>Advanced</b>	Displays all parts of sample and parameter files. Navigate to parameter windows by selecting the tabs across the top of the window.
<b>Basic</b>	This display option is used after the parameter files have been created by Developer group members. The previously entered or default parameter files are then accessible using drop-down lists.

The following examples show the same sample file in *Advanced* and *Basic* display.



**Advanced view  
for the Developer group**



**Basic view  
for the Analyst group**



A sample file must be created for each analysis. The file can be created prior to or at the time of analysis. The sample file identifies the sample, guides the analysis, and specifies report options.

The **Save** button is disabled on sample files with a *Complete* status. When the **Preview** button is used to view reports for sample files with an unsaved status, the report will have a *Preview* watermark. The **Save As** and **Print** buttons on the report window are also disabled.

## ***LIBRARIES***

### ***Options > Manage Libraries***



This feature is available only to Windows 7 and higher operating systems.

The library provides an easy way to locate and open specific analyzer files. Libraries are located within the *File Selector* window and can be viewed only within the application.

The library gathers sample and parameter files that are stored in multiple locations — such as folders on a C: drive, a network location, a connected external hard drive, or a connected USB flash drive — providing access to all files. Even though libraries do not store actual sample and parameter files, folders can be added or removed within each library.

One library can include up to 50 folders. Other items — such as saved searches and search connectors — cannot be included.

When removing a folder from a library, the folder and its contents are not deleted from the original file storage location. However, when deleting files or folders from within a library, they are deleted from their original file storage location.



## METHODS

**File > New Method**

**Options > Default Method**

**File > Open > [.MTH File]**



**See also:**

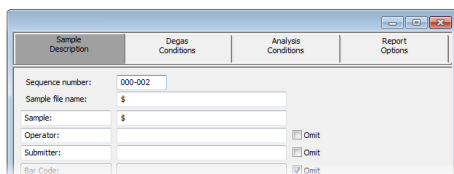
- [About Sample Files on page 3 - 1](#)
- [About Parameter Files on page 4 - 1](#)

### CFR Note

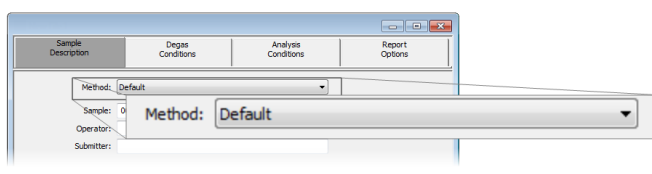
For 21CFR11 environments, this section is applicable only to members of the Developer group, however, members of the Analyst group may find information in this section helpful. Sample file information that is available to Analysts is created by a member in the Developer group using information in this section.

A *Method* determines the default sample identification format and sequence number. A *Method* is a template of specifications that go into a newly created sample file. It allows for the definition of complete sets of parameters for each type of sample commonly analyzed, so that only a single selection is required for each new sample file created.

The *Method* drop-down list displays only those methods applicable to the open sample file type.

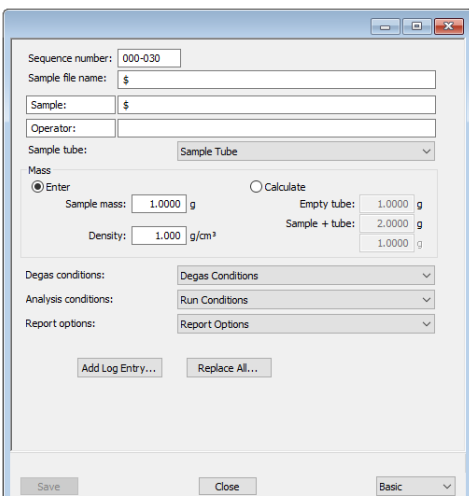


**Default Method**



**Sample Information File**

**Developer group view in a 21CFR11 environment**



**Analyst group view in a 21CFR11 environment**

### Default Method Fields and Buttons Table

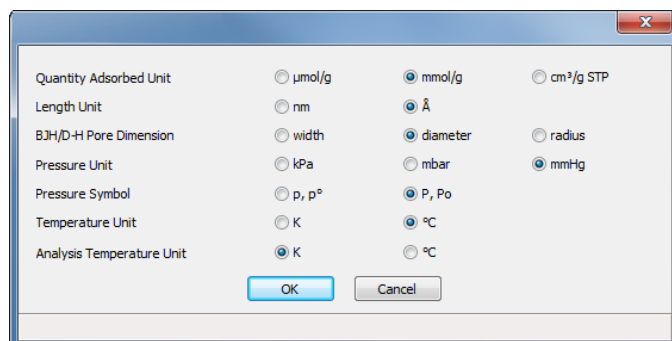
Field or Button	Description
<b>Sequence Number</b> [text box]	Specify a default numeric string to be used as a prefix in the <i>Sample</i> field when a new sample file is created. This number increments with each sample file created.
<b>Sample file name</b> [text box]	Enter a format for the sample identification. The entry in this field becomes a part of the saved sample file name. Include the \$ symbol to have the sample file number included as part of the identification.
<b>Sample Operator Submitter Bar Code</b> [text box]	These field labels may be renamed and the new label becomes a part of all new sample files.

## CONFIGURE THE ANALYZER

### SPECIFY UNIT SELECTIONS

#### Options > Units

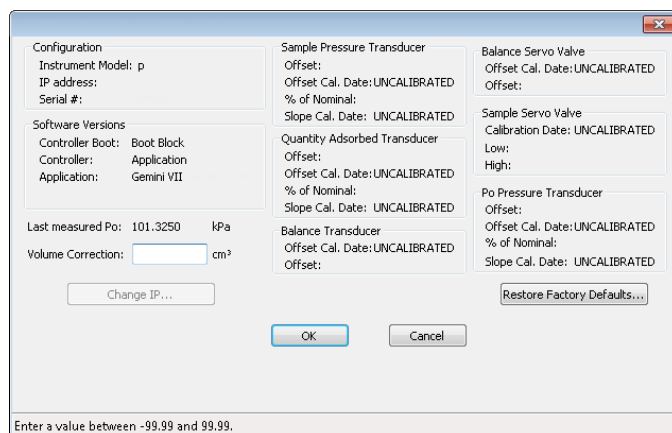
Use to specify how data should appear on the application windows and reports. This menu option is not available if using *Restricted* option presentation in a standard installation environment.



### UNIT CONFIGURATION

#### Unit [n] > Unit Configuration

Use to display and confirm hardware / software configurations and calibrations of the analyzer.



- **Volume Correction.** Displays the volume correction of the analyzer. If a different value is entered in this field, it is stored with sample data at the beginning of the analysis. The value in this field is used in calculating free space.
- **Change IP.** Click to display the *Unit IP Setup* window. Use to change the IP address and subnet mask assigned during installation. Do not edit these fields unless instructed by a Micromeritics Service Representative.

## INSTRUMENT STATUS

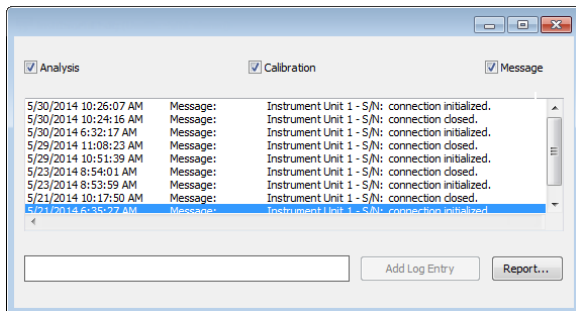
### SHOW INSTRUMENT LOG

#### Unit [n] > Show Instrument Log

#### CFR Note

In 21CFR11 environments, see [System Audit Trail on page 2 - 25](#).

Use to display a log of recent analyses, calibrations, errors, or messages.



#### Instrument Log Fields and Buttons Table

Field	Description
<b>Add Log Entry</b> [button]	Use to enter information to appear in the sample log report that cannot be recorded automatically through the application. Click the button again to enter multiple log entries.
<b>Analysis/ Calibration/ Message</b> [check box]	Select the logs to display.

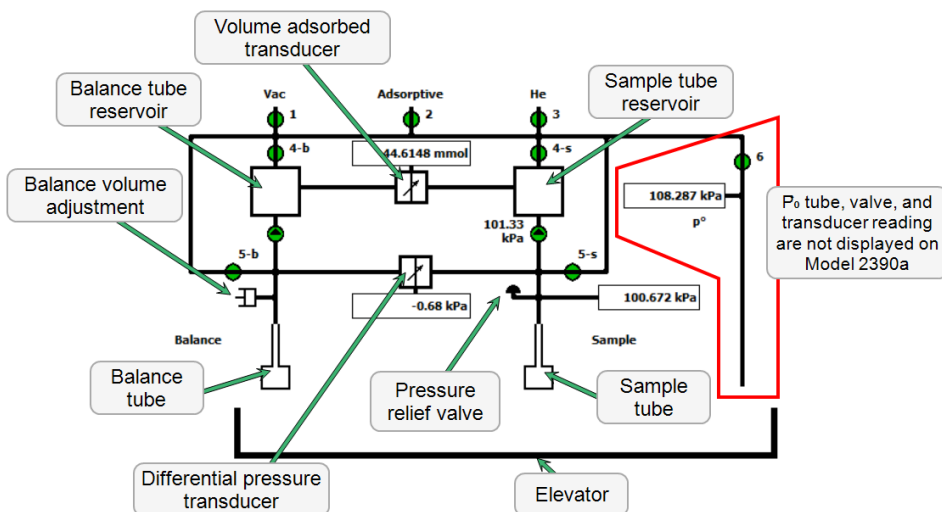


For fields and buttons not listed in this table, see [Common Fields and Buttons on page 2 - 3](#).

## SHOW INSTRUMENT SCHEMATIC

### Unit [n] > Show Instrument Schematic


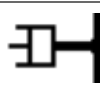


Use to enable the manual control of certain system valves and elevator components on the analyzer schematic. When this option is enabled, a checkmark appears to the left of **Unit [n] > Enable Manual Control**. If the analyzer schematic is not immediately visible, go to **Unit [n] > Show Instrument Schematic**.



### Analyzer Schematic Icon Table

Icon or Symbol	Description
	<b>Open Valve.</b> Green indicates an open valve.
	<b>Closed Valve.</b> Yellow indicates a closed valve. When manual control is disabled, closed valves appear white.
	<b>Servo Valve.</b> Closed.
	<b>Servo Valve.</b> Open.
	<b>Elevator.</b>
	<b>Reservoir.</b> For the balance tube and sample tube.

### Analyzer Schematic Icon Table

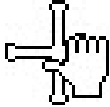

Icon or Symbol	Description
	<p><b>Volume adsorbed transducer</b> (upper).</p> <p><b>Differential pressure transducer</b> (lower).</p> <p>The reading for the volume adsorbed transducer is displayed just above the symbol. The reading for the sample pressure transducer is displayed just below the analysis servo valve and valve 5-s.</p>
	<b>Balance volume adjustment.</b> Can be adjusted by a Micromeritics Service Representative.
	<b>Pressure relief valve.</b> Prevents excessive pressure build-up in the event of abnormal operation.
	<b>Balance tube and sample tube.</b>

### Analyzer Schematic Components Table

Schematic Components	Description
<b>1</b>	Vacuum valve
<b>2</b>	Adsorptive valve
<b>3</b>	Helium valve
<b>4-b</b>	Balance reservoir valve
<b>4-s</b>	Analysis reservoir valve
<b>5-b</b>	Balance port valve
<b>5-s</b>	Analysis port valve
<b>6</b>	P <sub>0</sub> port valve (models 2390 and 2390t only)

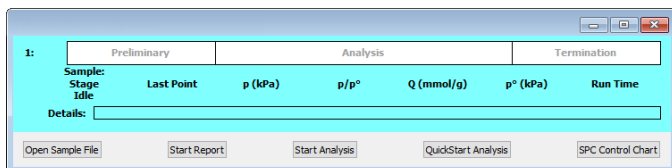
## Instrument Schematic Shortcut Menus

Schematic Shortcuts Table

Schematic Shortcut Icon	Description
<b>Valve options</b> 	<p>For valves:</p> <ul style="list-style-type: none"> <li>• <b>Close.</b> Closes the selected valve.</li> <li>• <b>Close both.</b> Closes valves 4b and 4s or valves 5b and 5s.</li> <li>• <b>Open.</b> Opens the selected valve.</li> <li>• <b>Open both.</b> Opens valves 4b and 4s or valves 5b and 5s.</li> <li>• <b>Pulse.</b> Use to quickly turn the valve on and off allowing the operation to proceed in small increments.</li> </ul> <p>For servo valve:</p> <ul style="list-style-type: none"> <li>• <b>On/Off.</b> Turns the servo valve on or off.</li> <li>• <b>Increase / Decrease.</b> Increases pressure by dosing the sample. Decreases pressure by evacuating the sample.</li> <li>• <b>Enter target.</b> Displays the servo valve settings window. The entered target pressure display to the left of the sample servo valve icon.</li> </ul> <p>For balance servo valve:</p> <ul style="list-style-type: none"> <li>• <b>Disable.</b> Select to disable the balance servo valve. Operations for the sample servo will also be performed by the balance servo.</li> <li>• <b>Enable.</b> Select to enable the balance servo valve. Operations for the sample servo will be ignored by the balance servo.</li> </ul>
<b>Elevator options</b> 	<p><b>Raise.</b> Select <i>Raise</i> to raise the elevator. When it is moving, press the keyboard space bar to stop the movement (or right click and select <i>Stop</i> from the menu).</p> <p><b>Lower.</b> Select <i>Lower</i> and press the keyboard space bar to lower the elevator.</p> <p><b>Stop.</b> Stops the elevator from moving.</p>

## SHOW STATUS

### *Unit [n] > Show Status*



Sample Stage	Last Point	p (kPa)	p/p°	Q (mmol/g)	p° (kPa)	Run Time
Idle						

If multiple units are attached to the computer, select *Show Status* on each *Unit [n]* menu. The status for all units display.



## EXPORT FILES

### File > Export



See also:

[Exported Data Example on page C - 1](#)

Provides the option to print the contents of one or more sample or parameter files to either the screen, a printer, or to a file. Data can be exported as a .PDF, .REP, .TXT, or .XLS file format. The type of data to include or exclude can be selected during the export process. When exported to a file, the data can be imported into other applications that read these file formats.

## LIST FILES

### File > List

Provides the option to create a list of sample file information —such as file name, date, time the file was created or last edited, file identification, and file status.

Select one or more files from the file selector, then click [List](#).

File Listing					
No.	File Name	Date	Time	Description	Status
1	co_pulse.smp	9/26/2017	5:12:26 PM	CO pulse chemisorption on 0.5 Pt Alumina	Complete
2	h2_o2_h2.smp	9/26/2017	5:12:26 PM	H2 O2 H2 titration on Pt Alumina	Complete
3	mfi-fit2.smp	9/26/2017	5:12:27 PM	Heat of desorption - fit peaks - strong chemi	Complete
4	mfi-fit1.smp	9/26/2017	5:12:26 PM	Heat of desorption - fit peaks - weak sites	Complete
5	mfi-hod1.smp	9/26/2017	5:12:27 PM	NH3 - heat of desorption - weak sites	Complete
6	mfi-hod2.smp	9/26/2017	5:12:27 PM	NH3 - Heat of desorption - strong chemisorption	Complete
7	mfi-01.smp	9/26/2017	5:12:26 PM	NH3 TPD	Complete
8	mfi-fit.smp	9/26/2017	5:12:26 PM	NH3 TPD - with peak deconvolution	Complete
9	bet.smp	9/26/2017	5:12:26 PM	Silica Alumina BET 2X	Complete
10	000-193.SMP	9/26/2017	11:28:12 AM	Silver Oxide TPR	Complete
11	ago_tpr.smp	9/26/2017	5:12:26 PM	Silver oxide TPR	Complete
12	n2o_puls.smp	9/26/2017	5:12:27 PM	Silver surface area using N2O	Complete
13	v2o5.smp	9/26/2017	5:12:27 PM	TPR of Vanadia	Complete

## CONVERT FILES

### File > Convert > [*\*.MGD file*]

Converts StarDriver files (.MGD extension) to a sample file with a .SMP file extension. Only those files with a .MGD file extension display in the *Name* column.

## ABOUT THE SOFTWARE IN 21CFR11 ENVIRONMENTS



### See also:

[Option Presentation for 21CFR11 Environments on page 2 - 12](#)



The Micromeritics Confirm applications for 21CFR11 environments require an operating system of Windows 10 Professional or Windows 10 Enterprise or higher. Management of users and groups is performed in Windows Users and Groups.

The Micromeritics Confirm application enables laboratory managers to develop analysis methods, enforce industry standards, and produce audit trails. It also enables laboratory analysts to perform analyses and produce reports.

## USER PERMISSIONS

### Confirm Groups

Group Name	Description
<b>Developer Group</b>	<p>The default Developer group name is <i>mic_[analyzer model number]_developer</i>. Members of the Developer group:</p> <ul style="list-style-type: none"><li>• have rights to all functions of the Micromeritics application - including Advanced option presentation which allows the creation and modification of methods, sample files, and parameter files.</li><li>• can run an analysis.</li><li>• can also be assigned Administrator rights which control the user profiles.</li></ul>
<b>Analyst Group</b>	<p>The default Analyst group name is <i>mic_[analyzer model number]_analyst</i>. Members of the Analyst group:</p> <ul style="list-style-type: none"><li>• have access to the Basic presentation option only.</li><li>• may create sample files from pre-defined methods and can modify only a limited number of input fields.</li></ul>

## CREATE NEW FOLDER

### File > Create New Folder

**CFR  
Note**

For 21CFR11 environments only.

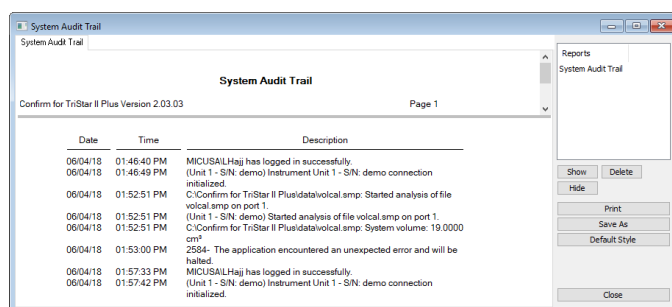
Provides the option to create and name a new folder in the Confirm application folder. This option may not be available depending on how the IT administrator configures Windows permissions.

## SYSTEM AUDIT TRAIL

**CFR  
Note**

For 21CFR11 environments only.

### File > System Audit Trail



Lists the current user, successful and failed application user login attempts, and contains a description of all the changes made to sample files.

Contains an audit trail of all system initializations, user login attempts, and sample analyses.

## SOFTWARE UNINSTALL

When the software is uninstalled using *uninstall.exe*, only the files required to run the application are removed. Parameter files, sample files, reports, calibration files, and data files are not removed.

To uninstall the software, double click the *uninstall.exe* file located in the software installation directory, then follow the prompts.

### CFR Note

In a 21CFR11 environment, upon uninstalling the Confirm application, the system administrator should go into Windows Users and Groups to remove the Confirm users and groups. See the Confirm Administrator Guide (part number 004-42821-00).

## SOFTWARE UPDATES

Log in to your [customer portal](#) to access and download the latest analyzer software version.

When performing a software update, existing data files are not overwritten. There are three types of subsequent installation:

- a later version than the current installation
- the same version as the current installation
- an earlier version than the current installation

Insert the setup media into the media drive. The setup program starts automatically. If the program does not start automatically, navigate to the installation media drive, locate and double click the *setup.exe* file.

### CFR Note

Existing Confirm application users and groups are not affected by software updates. Any changes to Confirm users and Confirm groups must be made using Windows Users and Groups.

## 3 ABOUT SAMPLE FILES

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Sample files include the information required by the analyzer to perform analyses and collect data. A sample file identifies the sample, guides the analysis, and specifies report options and may be displayed in either *Advanced*, *Basic*, or *Restricted* presentation display mode. See [Option Presentation on page 2 - 10](#) or [Option Presentation for 21CFR11 Environments on page 2 - 12](#).

A sample file consists of parameter sets; however, parameter sets can also stand alone. A sample file may be created either prior to or at the time of analysis.

Parameter files allow for repeated use of parameter sets. For example, if the same analysis conditions exist for multiple analyses, an *Analysis Conditions* file containing the recurring conditions can be created. When the sample file is created, the *Analysis Conditions* file can be selected for the analysis conditions. Once it becomes part of the new sample file, the new file can be edited as needed without affecting the original *Analysis Conditions* file.

The analysis application contains a default method. A method is a template for sample files that contains the parameters to be used for an analysis. When a new sample file is created, all the parameters are filled with the values in the default method.



Specify or change the default option presentation by selecting **Options > Option Presentation**, or select *Basic* or *Advanced* from the drop-down list at the bottom of the window.

## CREATE SAMPLE FILES

**Options > Option Presentation > Show Degas Conditions**

**File > New Sample > [.SMP File]**

**File > Open > [.SMP File]**

### CFR Note

For 21CFR11 environments, this section is applicable only to members of the Developer group, however, members of the Analyst group may find information in this section helpful. Sample file information that is available to Analysts is created by a member in the Developer group using information in this section.

Each analysis must be linked with a sample file before the analysis can proceed. A sample file can consist of parameter files; however, parameter files can also stand alone.

Specify or change the default display option by selecting **Options > Option Presentation** or select *Basic* or *Advanced* from the drop-down list at the bottom of the window.

Sample files must be initially created in *Advanced* option presentation. After the sample files are saved, the files are accessible in *Basic* and *Restricted* option presentation.

The values specified in the parameter portions of the default method are the defaults for new sample files. To navigate from one set of parameters to another, select the parameter tab across the top of the window.



The *Degas Conditions* tab displays only if enabled in **Options > Option Presentation > Show Degas Conditions**.

**Advanced view  
for the Developer group**

**Basic view  
for the Analyst group**




A bar code reader may be used to enter text into many of the fields on the *Sample Description* window. Use a mouse to click in the field first where information is to be entered then scan the bar code with the bar code reader.

## Sample File Fields and Buttons Table

Field or Button	Description
<b>Add Log Entry</b> [button]	Use to enter information to appear in the sample log report that cannot be recorded automatically through the application. Click the button again to enter multiple log entries.
<b>Comments</b> [text box]	Enter comments about the sample or analysis. Comments display in the report header.
<b>Mass</b> [group box]	<p>Enter a value for sample mass. Mass can be changed any time before, during, or after analysis.</p> <p><b>Enter.</b> Enables the <i>Sample mass</i> field. Enter a value for the sample mass.</p> <p><b>Calculate.</b> Enables the <i>Empty tube</i> and <i>Sample + tube</i> fields. Enter the values necessary to calculate the sample mass. Equation used to calculate sample mass:</p> $Mass_{sample} = Mass_{sample+tube} - Mass_{tube}$
<b>Method</b> [drop-down box]	Select a method from the drop-down list.

Sample File Fields and Buttons Table (continued)

Field or Button	Description
<b>Operator</b> [text box]	Enter operator identification information. This field label may have been renamed or may not display if modified in <b>Options &gt; Default Methods</b> .
<b>Sample</b> [text box]	Enter a sample description.
<b>Submitter</b> [text box]	Enter submitter identification information. This text box may have been renamed or may not display if modified in <b>Options &gt; Default Methods</b> .
<b>Type of Data</b> [group box]	<p><b>Automatically collected.</b> Select if the type of data will be automatically collected by the system while an analysis is running.</p> <p><b>Manually entered.</b> Use to enter data manually that was collected from another source. If <i>Manually entered</i> is selected, the Isotherm Report becomes available in the <i>Basic/Advanced</i> drop-down list for pasting or importing data into the file.</p> <p>See <a href="#">Manually Enter Data on page 3 - 7</a>.</p>
<b>User Parameters</b> [group box]	These fields are primarily used for the SPC (Statistical Process Control) reporting to specify sample characteristics or its manufacturing process but may be used for other data by entering specific analysis conditions or sample criteria. The entered parameters display on the <i>Summary Report</i> . Some fields may not display (or may have a different field label) if modified in the method from which the sample file was created, either through <b>Options &gt; Default Method</b> or <b>File &gt; Open &gt; Method</b> . See <a href="#">SPC Report on page 7 - 4</a> .
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	



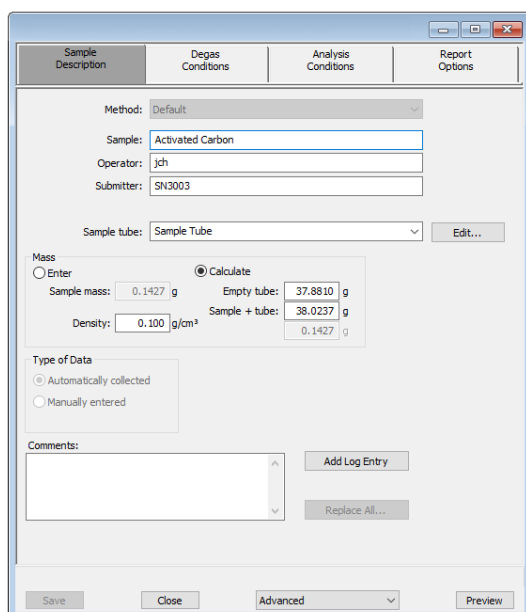
## OPEN A SAMPLE FILE

**File > Open > [.SMP File]**

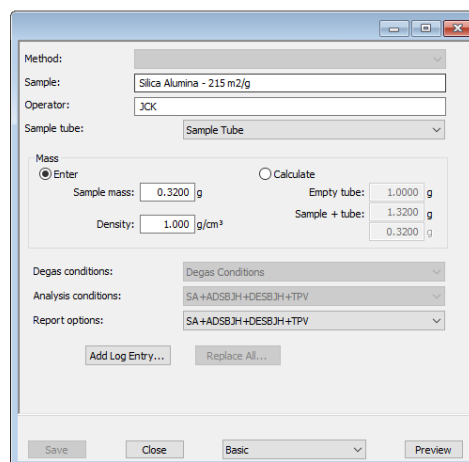


When working with an existing sample file, consider making a copy of the sample file to maintain the original configuration options.

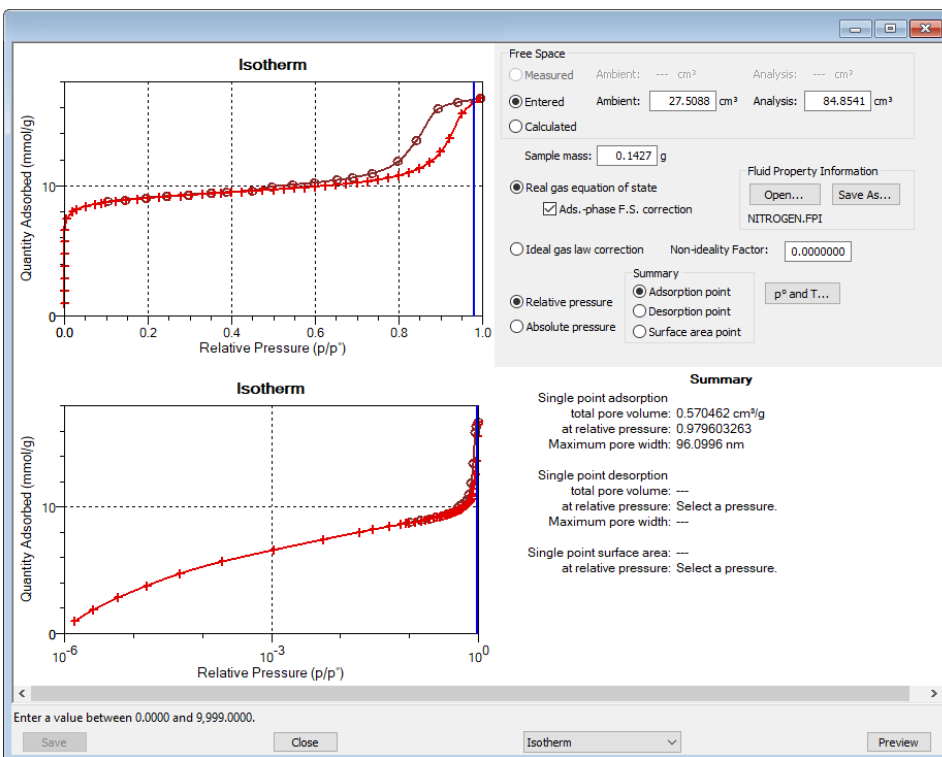
File Status	Displays
Preparing Prepared No Analysis	Tabbed file editor
Complete Analyzing Entered	MicroActive report window



Example of tabbed file editor for  
Advanced or Developer (Cfr) view



Example of file editor for  
Basic or Analyst (Cfr) view



**Example of MicroActive Report**

If a sample file with a *Complete* status is opened, to return to the tabbed file editor, select *Advanced* or *Basic* from the drop-down at the bottom of the window. **NOTE:** In a 21CFR11 environment, these selections are available only to members of the Developer group.

## MANUALLY ENTER DATA

### CFR Note

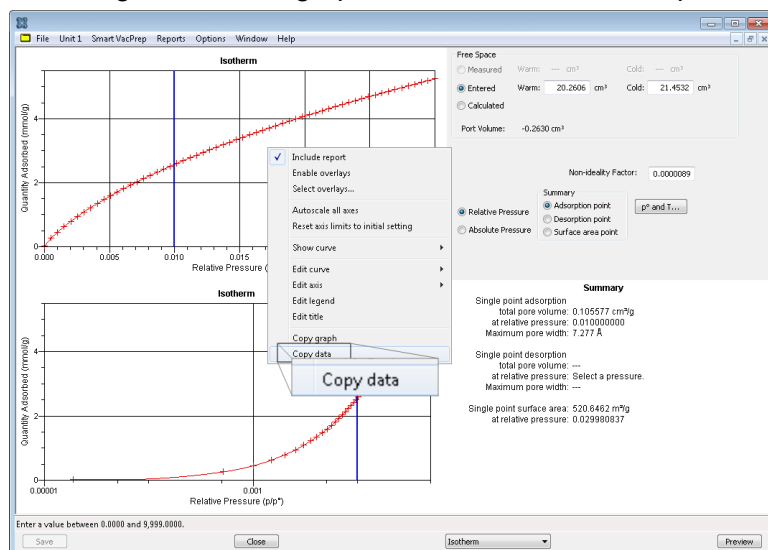
In 21CFR11 environments, this feature is applicable to members of the Developer group only.

This process allows the manual entry of pressure data from a sample file with a *Complete* status. There are two methods for manually entering data into a sample file:

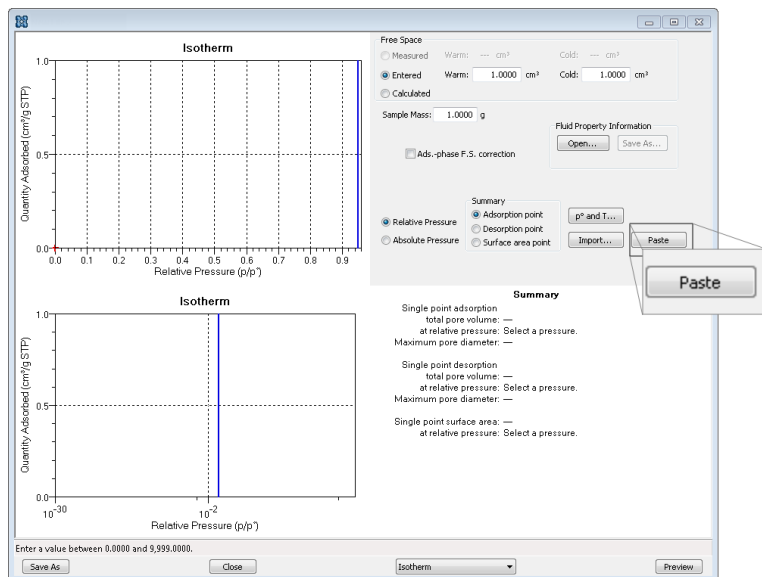
- Copy and paste onto the graph area of the interactive window
- Import data into the interactive window

## COPY AND PASTE MANUALLY ENTERED DATA

1. Open a sample file with a *Complete* status. The file will open to the interactive reports window.
2. Right click in the graph area of the interactive reports window, then select *Copy data*.



3. Open another sample file using *Advanced* option presentation.
4. On the *Sample Description* tab, select *Manually entered* in the *Type of Data* group box.
5. Click the *Advanced* down arrow at the bottom of the window, then select *Isotherm*.



6. Ensure that all parameter fields are set appropriately, then click the **Paste** button.

## IMPORT MANUALLY ENTERED DATA

When importing isotherm data from an external ASCII text file using the **Import** button on the interactive window, the ASCII text file must use the following rules:

### ASCII text file format rules

Data must be in two columns and separated by a comma or white-space. Acceptable column headings are:

- Relative Pressure
- Absolute Pressure (mmHg)
- Absolute Pressure (kPa)
- Absolute Pressure (mBar)
- Quantity Adsorbed (mmol/g)
- Quantity Adsorbed (cm<sup>3</sup>/g STP)
- Quantity Adsorbed (cm<sup>3</sup>/g STP)

### Sample Physical Adsorption ASCII Text File

Silica Alumina : Adsorption

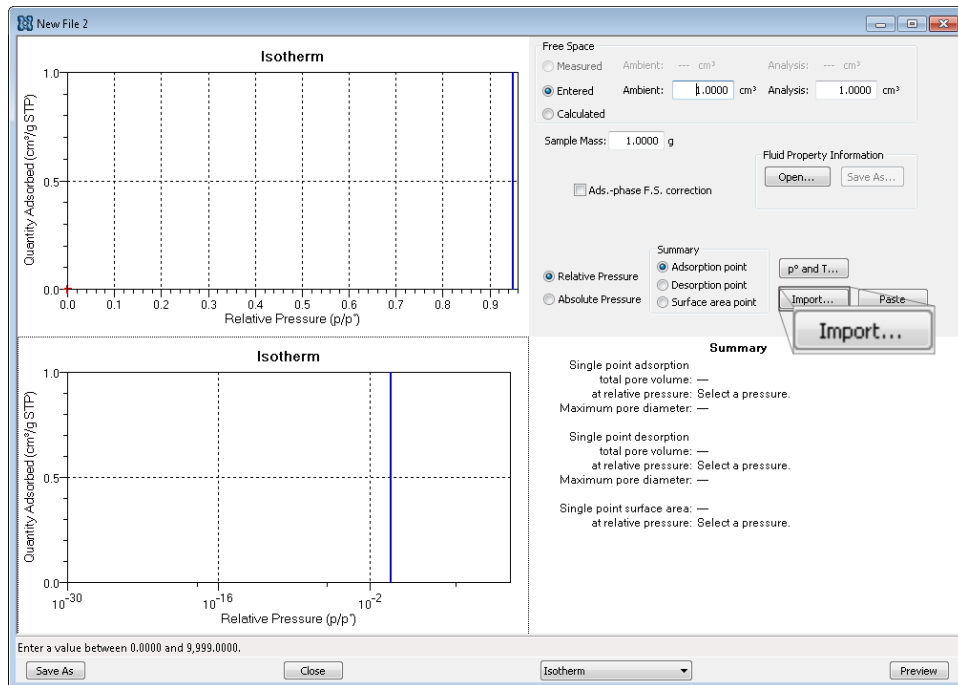
Relative Pressure	Quantity Adsorbed (cm <sup>3</sup> /g STP)
0.108629	50.6657
0.22288	60.7813
0.339909	71.3095
0.459512	84.4172
0.577447	102.672
0.654583	121.707
0.760074	179.096
0.855713	334.565
0.958511	394.675
0.996251	403.793

Silica Alumina : Desorption

Relative Pressure	Quantity Adsorbed (cm <sup>3</sup> /g STP)
0.996251	403.793
0.86016	389.626
0.753567	256.264
0.664418	133.099
0.542416	96.7366
0.422295	79.7351
0.346371	71.5994
0.2519	62.8256
0.152718	54.2336
0.103389	49.5803

### To import the ASCII text file

1. Open a new sample file in *Advanced* option presentation.
2. On the *Sample Description* tab, select *Manually entered*.
3. Click the *Advanced* down arrow at the bottom of the window, then select *Isotherm*.



4. Ensure that all parameter fields are set appropriately, then click **Import**.
5. Open the .TXT file. The data from the original sample file is imported and displayed. If an error message appears instead, verify that the .TXT file format is correct.

## 4 ABOUT PARAMETER FILES

### CFR Note

In 21CFR11 environments, this section is applicable only to members of the Developer group, however, members of the Analyst group may find information in this section helpful. Parameter file information that is available to Analysts is created by a member in the Developer group using information in this section.

Parameter files allow for repeated use of parameter sets. For example, if the same analysis conditions exist for multiple analyses, an *Analysis Conditions* file containing the recurring conditions can be created. When the sample file is created, the *Analysis Conditions* file can be selected for the analysis conditions. Once it becomes part of the new sample file, the new file can be edited as needed without affecting the original *Analysis Conditions* file.

Methods include both analysis conditions and report options, offering the most convenient way to repeat most analyses. See [Methods on page 2 - 15](#).

Predefined parameter files are included with the program and can be edited as needed or new parameter files can be created.

The following file types can exist as part of the sample file as well as individual parameter files:

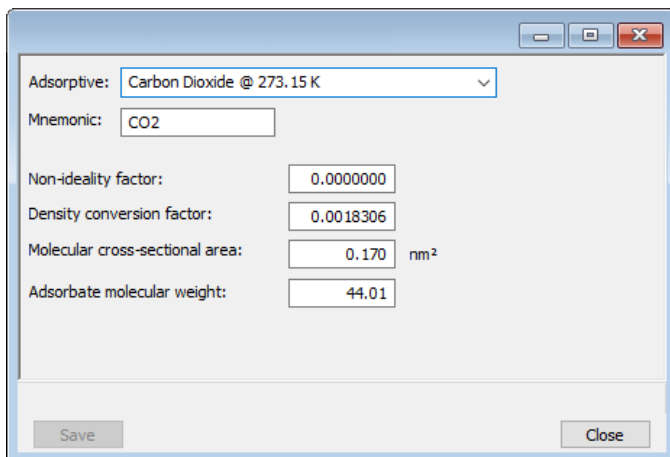
File Type	File Extension
Adsorptive Properties	.ADP
Analysis Conditions	.ANC
Degas Conditions	.DEG
Method	.MTH
Report Options	.RPO

## ADSORPTIVE PROPERTIES


### File > Open > [.ADP File]

(or click **Edit** next to the *Adsorptive* selection on the *Analysis Conditions* tab when in *Advanced* option presentation)

Adsorptive properties provide the adsorptive (analysis gas) characteristics for the analysis.



### Adsorptive Properties Fields and Buttons Table

Field or Button	Description
<b>Adsorbate molecular weight</b> [text box]	The molecular mass is used for the weight % column of the isotherm tabular report and for the pressure composition isotherm plot.
<b>Adsorptive</b> [text box]	Name of the adsorptive gas whose properties are being defined.
<b>Density Conversion Factor</b> [text box]	Factor determined by obtaining the ratio of the gas density (STP) to the liquid volume.
<b>Mnemonic</b> [text box]	Enter the mnemonic name for the adsorptive..
<b>Molecular cross-sectional area</b> [text box]	The area that a single adsorbed molecule occupies on the surface of the sample. It is used in surface area calculations.
<b>Non-ideality factor</b> [text box]	Compensates for the forces of attraction between molecules in a real gas. This value is used for a calculated free space.
 <b>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</b>	

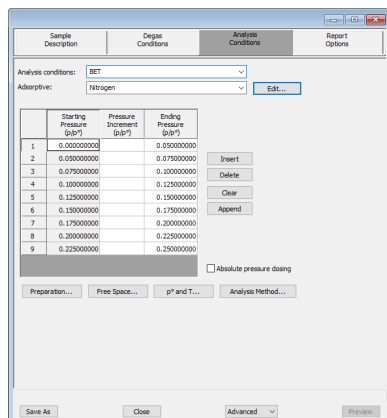


## ANALYSIS CONDITIONS

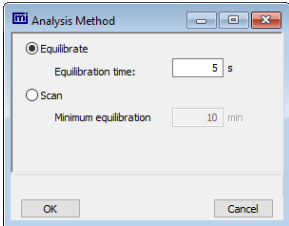
### File > Open > [.ANC File]

(or click the *Analysis Conditions* tab when in *Advanced* option presentation)

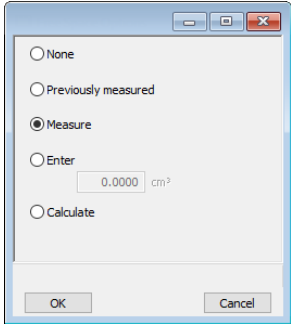
Analysis conditions specify the parameters used to guide an analysis.



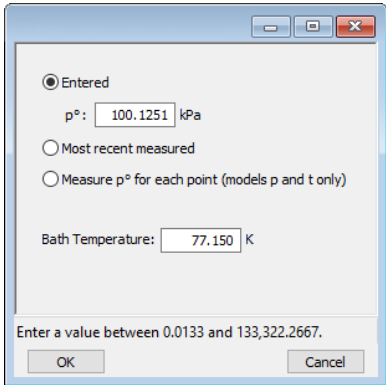
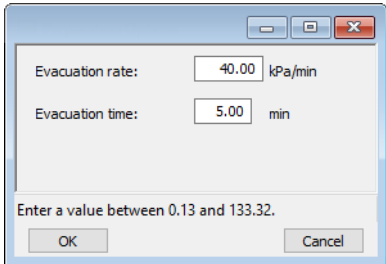
### Analysis Conditions Buttons Table

Button	Use to Specify...
<b>Analysis Method</b> [button]	Use to enter equilibration time in seconds or enter the minimum equilibration time for scanning in minutes  
<b>Free Space</b> [button]	How the free space is to be measured.
<b>P<sub>0</sub> and T</b> [button]	How the saturation pressure (P <sub>0</sub> ) is to be measured or calculated and the analysis bath temperature.
<b>Preparation</b> [button]	Evacuation rate / time / level, leak test and time values, elevator prompts, and in situ degassing or activation.


## Analysis Conditions Fields and Buttons Table

Field or Button	Description
<b>Absolute pressure dosing</b> [check box]	Specifies pressure targets in mmHg, mbar, or kPa instead of relative pressure. If this option is selected, the <i>Relative Pressure</i> labels and entries change to <i>Absolute Pressure</i> in the selected pressure units. This option is typically selected when using adsorptives at analysis conditions above the critical point of the gas; for example, H <sub>2</sub> adsorption on carbon at liquid nitrogen temperature.
<b>Adsorptive</b> [text box]	Select an <i>Adsorptive Properties</i> file from the list of defined gases to be used for analysis. After selection, click <b>Edit</b> to modify adsorptive properties.
<b>Analysis Conditions</b> [drop-down box]	Use to browse for an <i>Analysis Conditions</i> file that contains analysis condition parameters to be used in the analysis.
<b>Free Space</b> [button]	<p>Use to enter the free space measurement type.</p>  <p><b>None.</b> Select if free space will not be used.</p> <p><b>Previously measured.</b> Select if free space has been previously measured. Recommended if measuring a series of samples with similar volumes.</p> <p><b>Measure.</b> Select if the free space is to be measured before the analysis begins.</p> <p><b>Enter.</b> Allows manual entry of free space.</p> <p><b>Calculate.</b> Select to have the free-space calculated using the value entered for <i>Volume Correction</i> (entered on the <b>Unit [n] &gt; Unit Configuration</b> window) and the non-ideality factor (entered on the <i>Adsorptive Properties</i> window).</p>

## Analysis Conditions Fields and Buttons Table (continued)

Field or Button	Description
<b>P<sub>0</sub> and T</b> [button]	 <p>Use to select options for obtaining the saturation pressure (P<sub>0</sub>) and analysis bath temperature.</p> <p><b>Entered.</b> Select to manually enter the P<sub>0</sub>.</p> <p><b>Most recent measured.</b> Select to use the most recently measured saturation pressure.</p> <p><b>Measure P<sub>0</sub> for each point (models p and t only).</b> Select if P<sub>0</sub> is to be measured for each point.</p> <p><b>Bath temperature.</b> Calculates P<sub>0</sub> using the entered bath temperature.</p>
<b>Preparation</b> [button]	<p>Use to enter analysis preparation details.</p>  <p><b>Evacuation rate.</b> Rate at which unrestricted sample evacuation should begin.</p> <p><b>Evacuation time.</b> The length of time for preliminary evacuation which takes place prior to the free space measurement or sample analysis if free space is to be entered or calculated.</p>

## Analysis Conditions Fields and Buttons Table (continued)

Field or Button	Description
	For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .

## DEGAS CONDITIONS

**File > Open > [.DEG File]**

(or click the *Degas Conditions* tab when using *Advanced* option presentation)



**See also:**

[Degassing on page 5 - 1](#)

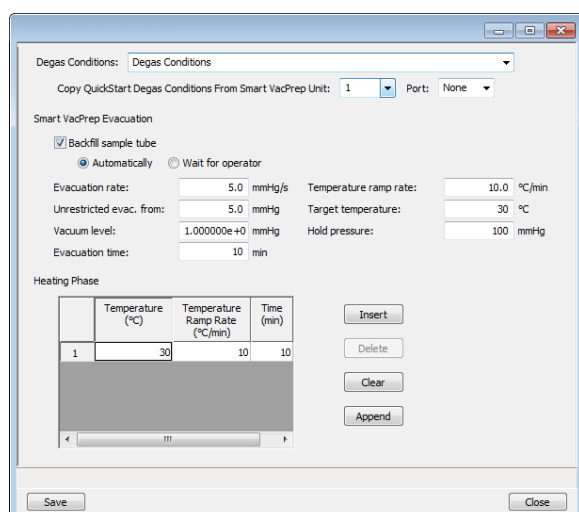


Use this option only when the SmartPrep or Smart VacPrep is installed. Degassing is a required step in preparation for an analysis. The *Degas Conditions* tab provides settings that will be automatically applied during the degassing procedure when using the Smart VacPrep.

If using a FlowPrep or a VacPrep use the heating phase section for reference only.

The *Degas Conditions* tab displays only if enabled in **Options > Option Presentation > Show Degas Conditions**.

Log in to your [customer portal](#) to access the Smart VacPrep Operator Manual.



Degas Conditions: Degas Conditions

Copy QuickStart Degas Conditions From Smart VacPrep Unit: 1 Port: None

Smart VacPrep Evacuation

☒ Backfill sample tube

☒ Automatically ☐ Wait for operator

Evacuation rate: 5.0 mmHg/s Temperature ramp rate: 10.0 °C/min

Unrestricted evac. from: 5.0 mmHg Target temperature: 30 °C

Vacuum level: 1.000000e+0 mmHg Hold pressure: 100 mmHg

Evacuation time: 10 min

Heating Phase

	Temperature (°C)	Temperature Ramp Rate (°C/min)	Time (min)
1	30	10	10


Buttons: Insert, Delete, Clear, Append

Buttons: Save, Close

### Degas Conditions Fields and Buttons Table

Field or Button	Description
<b>Copy QuickStart Degas Conditions from Smart VacPrep unit</b> <i>[drop-down box]</i>	Use to copy the degas conditions settings from the selected Smart VacPrep unit and port.
<b>Degas Conditions</b> <i>[drop-down box]</i>	Use to browse for a .DEG file that contains degas condition parameters to be used in the analysis.
<b>Heating Phase</b> <i>[table]</i>	<p>This option is applicable when degassing with either a Smart VacPrep or a SmartPrep.</p> <p>Enter up to five stages of degas conditions.</p> <p><b>Temperature.</b> Temperature at which the sample is to be held while degassing.</p> <p><b>Time.</b> How long the sample is to be held at the specified temperature before beginning to cool down.</p> <p><b>Temperature Ramp Rate.</b> The rate at which the temperature will change while advancing to the hold temperature.</p>
<b>Smart VacPrep Evacuation</b> <i>[group box]</i>	<p><b>Backfill sample tube.</b> Indicate if the sample tube should be backfilled automatically or wait for operator response.</p> <p><b>Evacuation rate.</b> Rate used for evacuation.</p> <p><b>Evacuation time.</b> Length of time for preliminary evacuation before proceeding with the <i>Heating Phase</i> temperature schedule. The timer starts when the vacuum level is reached.</p> <p><b>Hold pressure.</b> Pressure at which heating will stop and hold the sample temperature approximately constant until the pressure falls below the <i>Hold</i> pressure. This prevents damage to the sample structure due to 'steaming' and /or elutriation due to excessive escaping gas velocity.</p> <p><b>Target temperature.</b> Targeted temperature for evacuation.</p> <p><b>Temperature ramp rate.</b> Rate at which the temperature is to change when advancing to the target pressure.</p> <p><b>Unrestricted evac. from.</b> Pressure at which the unrestricted evacuation is to begin.</p>

Degas Conditions Fields and Buttons Table (continued)

Field or Button	Description
	<b>Vacuum level.</b> Evacuation time starts when the vacuum level is reached.
 For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .	

## REPORT OPTIONS

### **File > Open > [.RPO File]**

(or click the *Report Options* tab when in *Advanced* option presentation)



#### See also:

- [About Reports on page 7 - 1](#)
- [Selected Reports on page 8 - 1](#)

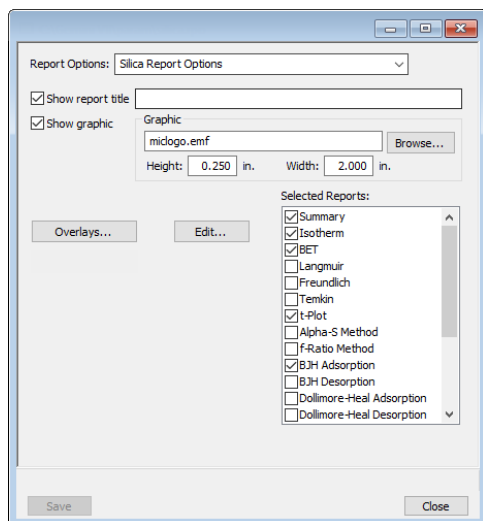
Log in to your [customer portal](#) to access MicroActive Report Tutorials and the Calculations document.

Additional reports are available using the *Reports* menu item.

Use to specify report options for data collected from an analysis or manually entered data. *Report Options* files also help in customizing report details such as axis scale, axis range, column headings, and components of thickness curve equations. These files may contain tabular reports, plots, or both, as well as advanced report tables.


Customized report options files can be created then loaded into a sample file, allowing quick generation of reports.

*Report Options* files may be defined to include overlay options. This system allows the overlay of up to 25 plots of different samples onto a plot of the same type or overlay one plot type onto a different plot type from the same analysis.





## Report Options Fields and Buttons Table

Field or Button	Description
<b>Overlays</b> [button]	See <a href="#">Graph and Sample Overlays on page 7 - 26</a> .
<b>Report Options</b> [drop-down box]	Browse for a .RPO file that contains report options parameters to be used in the report.
<b>Selected Reports</b> [group box]	Select the report names to include in the report.
<b>Show graphic</b> [text box]	Use to show a graphic on the report header.  <b>Height / Width.</b> Enter the height and width of the selected graphic. These values determine the graphic appearance on the generated report.
<b>Show report title</b> [check box]	Select and enter a report title to appear on the report header.
 For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .	

**Blank Page**

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## 5 DEGASSING

Most solid materials absorb moisture and other contaminants when exposed to the atmosphere. The sample must be clean when an analysis is performed. The degas process heats the sample to remove the moisture and contaminants.

Log in to your [customer portal](#) to access the Smart VacPrep Operator Manual or the SmartPrep Operator Manual.

### DEGAS ON THE SMARTPREP



**See:**

[Degasser Options on page 1 - 5](#)



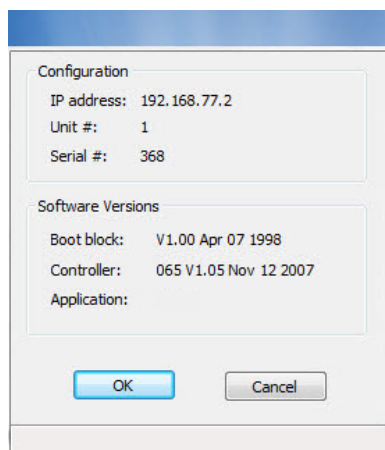
If a SmartPrep is not connected to the analyzer, the SmartPrep menu items are not available.

Log in to your [customer portal](#) to access the SmartPrep Operator Manual.

### SMARTPREP CONFIGURATION

#### Unit [n] > Degas > SmartPrep Configuration

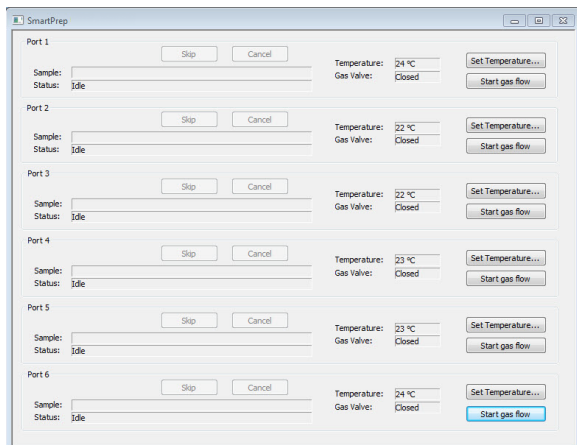
Displays the SmartPrep configuration and software versions.



## SHOW SMARTPREP STATUS

### Unit [n] > Degas > Show SmartPrep Status

The SmartPrep Status window allows the monitoring of degas operations and to stop the gas flow after samples are degassed.



The screenshot shows the 'SmartPrep' window with six ports (Port 1 to Port 6). Each port has a 'Sample' field, a 'Status' field (all set to 'Idle'), and three buttons: 'Skip', 'Cancel', and 'Set Temperature...'. To the right of each port, there are two readouts: 'Temperature' and 'Gas Valve'. For Port 1, Temperature is 24 °C and Gas Valve is Closed. For Port 2, Temperature is 22 °C and Gas Valve is Closed. For Port 3, Temperature is 22 °C and Gas Valve is Closed. For Port 4, Temperature is 23 °C and Gas Valve is Closed. For Port 5, Temperature is 23 °C and Gas Valve is Closed. For Port 6, Temperature is 24 °C and Gas Valve is Closed. Below each 'Gas Valve' readout is a 'Start gas flow' button. The 'Start gas flow' button for Port 6 is highlighted in blue.

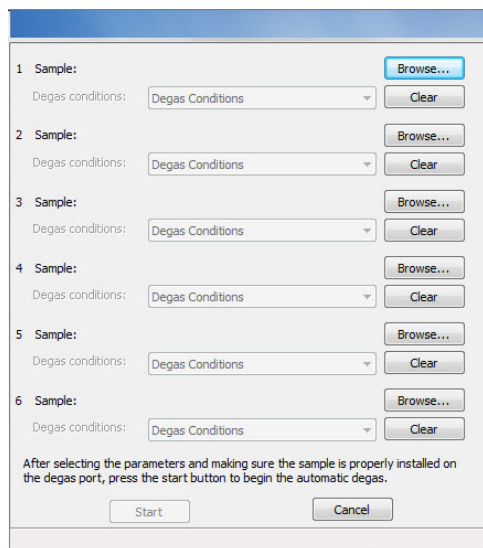
### Show SmartPrep Status Fields and Buttons Table

Button	Description
<b>Cancel</b> [button]	Discards any changes or cancels the current process.
<b>Set Temperature</b> [button]	Use to set the temperature of the selected port.
<b>Skip</b> [button]	Use to bypass the current stage.
<b>Stop Gas flow</b> [button]	Stops the gas flow to the selected port.

## START SMARTPREP DEGAS

### Unit [n] > Degas > Start SmartPrep Degas

The six SmartPrep heating stations are represented by row numbers on the *Automatic Degas* window.



The screenshot shows a software window titled 'Automatic Degas'. It contains six rows, each for a sample. Each row has a 'Sample:' label, a 'Degas conditions:' dropdown menu (all set to 'Degas Conditions'), a 'Browse...' button, and a 'Clear' button. At the bottom of the window, there is a 'Start' button and a 'Cancel' button. A note at the bottom states: 'After selecting the parameters and making sure the sample is properly installed on the degas port, press the start button to begin the automatic degas.'

### Start SmartPrep Status Fields and Buttons Table

Button	Description
<b>Browse</b> [button]	Searches for a file.
<b>Cancel</b> [button]	Discards any changes or cancels the current process.
<b>Clear</b> [button]	Clears the sample file selection for a port.
<b>Start</b> [button]	Starts degassing.

## DEGAS ON THE SMART VACPREP



See:

[Degasser Options on page 1 - 5](#)

Log in to your [customer portal](#) to access the Smart VacPrep operator manual.

## TRANSFER A DEGASSED SAMPLE TO AN ANALYSIS PORT



**See also:**

[Worksheets on page E - 1](#)

When degassing on a separate degasser such as a Smart VacPrep or SmartPrep, the sample tube must be removed from the degas port, weighed, and then installed onto the analysis port for analysis.



If the sample tube is not mounted on the analysis port immediately, leave it on the degas port. If it is necessary to remove the sample tube, insert a rubber stopper into the sample tube.

1. Allow the sample tube to cool to room temperature (approximately 15 minutes).



Do not touch the sample tube or the heating mantle until they have reached room temperature. Touching the sample tube, heating mantle, or heating mantle clip before they have cooled could result in burns.

2. Carefully remove the heating mantle clip and the heating mantle from the sample tube.
3. While holding the sample tube, loosen the port connector nut and remove the sample tube from the degas port. Immediately insert a rubber stopper into the sample tube.
4. Complete the sample data worksheet.
5. If a rubber stopper was used, remove it from the sample tube.
6. Place the connector nut, ferrule, and O-ring onto the sample tube stem.
7. On the analyzer, loosen the connector nut on the Psat tube and rotate it out of the way.
8. Position the dewar cover so that the slot for the Psat tube is on the left.
9. Insert the sample tube through one of the holes in the dewar cover.
10. Insert the sample tube into the analysis port and ensure it is completely in the port. Securely hand tighten the sample port nut onto the analysis port.
11. Position the dewar cover approximately 3/4 in. (19 mm) below the sample port nut.
12. Slide the Psat tube into the Psat slot in the dewar cover and retighten the Psat tube connector nut.
13. Attach the sample tube to the analysis port, pushing it fully up. Turn the connector nut clockwise and hand tighten.
14. Install the dewar onto the elevator and close the sample compartment door.
15. Begin the analysis.

## 6 PERFORM AN ANALYSIS

### CFR Note

In 21CFR11 environments, users are required to login to the Confirm application to start an analysis. Once the analysis window is opened, manual control is disabled until the analysis has completed. During analysis, pausing and resuming is allowed, however, steps cannot be skipped.

### DEWAR PRECAUTIONS



Always handle glass dewars with care. Any product incorporating a vacuum is a potential safety hazard and should be treated with caution. If in doubt, contact your safety officer.

When handling dewars containing liquefied gases or cryogenic liquids:

- Wear protective equipment:
  - goggles or face shield
  - an insulated or rubber apron
  - insulated gloves
- When pouring liquefied gases from one container to another:
  - cool the receiving container gradually to minimize thermal shock
  - pour the liquified gas slowly to prevent splashing
  - vent the receiving container to the atmosphere

### FOR GLASS DEWARS

- Use a plastic stirring rod when stirring substances in a dewar containing liquefied gases (or other materials of extremely low temperature). Do not use a glass or metal stirring rod unless it has a protective coating.
- Do not handle heavy objects above the dewar. If unavoidable, place a protective cover over the dewar opening. If an object of sufficient weight is accidentally dropped into the dewar, shattering may occur.
- If the dewar has a protective mesh covering, do not remove it. This cover minimizes the risk of flying particles should the dewar be knocked over, dropped, or broken.

## PREPARE FOR ANALYSIS

The steps in this topic properly prepare the equipment for an analysis. It is recommended to perform the tasks in the provided order.



### See also:

- [Dewar Precautions on the previous page](#)
- Step 1 - [Clean and Label Sample Tubes below](#)
- Step 2 - [Create the Sample File on page 6 - 4](#)
- Step 3 - [Determine the Sample Mass for Physical Adsorption on page 6 - 4](#)
- Step 4 - [Degas the Sample on page 6 - 6](#)
- Step 5 - [Sample Tube Installation on page 6 - 7](#)
- Step 6 - [Fill and Install the Dewar on page 6 - 9](#)

## CLEAN AND LABEL SAMPLE TUBES



The equipment images in this topic may differ slightly from your equipment, however the instructions are the same unless otherwise noted.

Sample tubes and filler rods must be clean and dry before samples are added and weighed. The following table indicates which materials are supplied by Micromeritics and which are supplied by the user. The procedures following the materials table are recommended.

Supplied by Micromeritics	Supplied by User
<ul style="list-style-type: none"> <li>• Filler rod</li> <li>• Funnel</li> <li>• Sample data worksheet</li> <li>• Sample tube</li> <li>• Sample tube brush</li> <li>• Sample tube rack</li> <li>• Sample weighing support</li> <li>• Stopper for sample tube</li> </ul>	<ul style="list-style-type: none"> <li>• Acetone or isopropyl alcohol</li> <li>• Analytical balance</li> <li>• Detergent (such as Alconox)</li> <li>• Drying oven</li> <li>• Forceps</li> <li>• Insulated gloves</li> <li>• Pipe cleaners</li> <li>• Rubber gloves or lint-free cloth</li> <li>• Safety glasses</li> <li>• Ultrasonic cleaning unit</li> <li>• Waste container</li> </ul>

1. Preheat drying oven to 110 °C.
2. Verify that the ultrasonic cleaning unit is clean.



3. Use 5 grams of Alconox (or other suitable detergent) per 500 mL of warm water and fill the ultrasonic unit with enough water to cover the sample tubes and filler rods (if used). If too much detergent is used, it may be difficult to rinse from the sample tubes. Ensure the detergent is dissolved before placing the sample tubes and filler rods into the water.
4. Fill the sample tubes with warm water and place them in the ultrasonic cleaning unit, then place the filler rods in the unit. Turn on the ultrasonic cleaning unit for approximately 15 minutes.



5. Use rubber gloves to ensure no oils or residue are transferred to the clean tubes and filler rods, then remove the sample tubes and filler rods from the unit.
6. Clean the interior of the sample tubes with the brush supplied with the analyzer.
7. Rinse the sample tubes and filler rods thoroughly with hot water. Rinse again with isopropyl alcohol or acetone. If isopropyl alcohol or acetone is not available, deionized water may be used.



8. Stand the sample tubes on the sample tube rack and place the filler rods in a basket or in the rack. Bake in a vacuum oven for two hours at 110 °C.



Samples tubes can also be cleaned with high purity acetone or isopropyl alcohol and dried for about 10 minutes under heat. If using this method, continue with step 10.

9. Remove the sample tubes and filler rods from the oven and allow to cool.



Do not insert the filler rods at this time. Filler rods are inserted before the sample tube is installed on the analysis port.

10. Blow out the sample tubes with oil-free compressed air.
11. Rinse the sample tube closure with isopropyl alcohol, then wipe the sample tube closure dry with a clean, lint-free cloth.
12. Label the sample tube and stopper for identification.
13. Replace the rubber stopper.

### **CREATE THE SAMPLE FILE**



**See:**

[Create Sample Files on page 3 - 2](#)

### **DETERMINE THE SAMPLE MASS FOR PHYSICAL ADSORPTION**



**See also:**

[Sample Data Worksheet for Gas Adsorption on page E - 2](#)

Clean, dry sample tubes are essential for accurate results. How much sample to use can be determined best by experiment. In general, a sample providing 40 to 120 square meters of total surface area is recommended for nitrogen analysis. Less than 40 square meters may cause unreliable results. More than 120 square meters will extend analysis time.

Smaller quantities are required for samples having high surface areas. These samples require careful weighing after degassing because a small error may represent a considerable percent of total weight. Proper weighing techniques are most important in this case. Use no less than 100 mg to reduce the effect of weighing errors.

Care should be taken when loading powders — the accessory funnel is useful for this purpose. Large granules or chunks may be loaded with forceps.

Analysis results are expressed in units of surface area per gram of sample; therefore, it is important the true sample mass be known.

Follow the instructions on the *Sample Data Worksheet* and complete all fields to find the true sample mass.

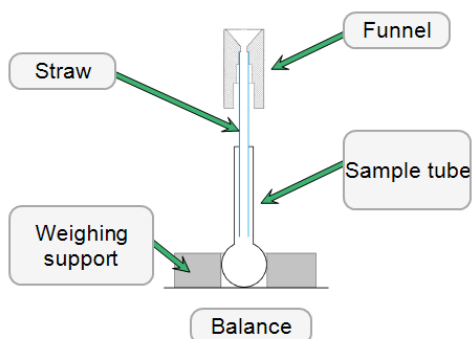
1. Record the sample tube identification on the *Sample Data Worksheet*.

2. Place the sample weighing support on the balance. Tare the balance and allow it to stabilize at zero (0).
3. Place the empty sample tube set (empty sample tube and stopper) on the sample weighing support and place it on the balance.
4. Record the stabilized mass on the *Sample Data Worksheet*. Remove the sample tube set from the balance.



Do not touch the sample with bare hands while performing the following steps. Doing so could affect the accuracy of results.

5. Place a sample container on the balance. Tare the balance and allow it to stabilize to zero (0).
6. Slowly pour the specified amount of sample into the sample container.
7. Remove either the rubber stopper from the sample tube.
8. Use the sample tube funnel (provided in the accessories kit) and pour the sample from the weighing container into the sample tube.



**TIP:**  
Use a drinking straw tube (approx 6mm [1/4 in.] diameter) that fits inside the sample tube to keep the inside wall of the sample tube clean.

9. Replace the rubber stopper.
10. On the Sample Data worksheet, record the following:
  - a. Mass of the sample tube set with the sample.
  - b. Subtract the *Mass of empty sample tube set* from the *Mass of sample tube set plus sample*.

## DEGAS THE SAMPLE



### See also:

- [Degas Conditions on page 4 - 7](#)
- [Degassing on page 5 - 1](#)
- [Sample Data Worksheet for Gas Adsorption on page E - 2](#)



If using the Smart VacPrep degasser, go to **Smart VacPrep > Unit [n] > Start Degas**, then degas the sample using menu commands and information entered on the *Degas Conditions* tab. Log in to your [customer portal](#) to access the Smart VacPrep Operator Manual.

For instructions on degassing on the SmartPrep, see [Degas on the SmartPrep on page 5 - 1](#).

After the sample has been weighed, use a degassing unit to remove any contaminants which may have adsorbed to the surface or pores. Appropriate degassing units are available from Micromeritics.

After degassing is complete, perform the following steps:

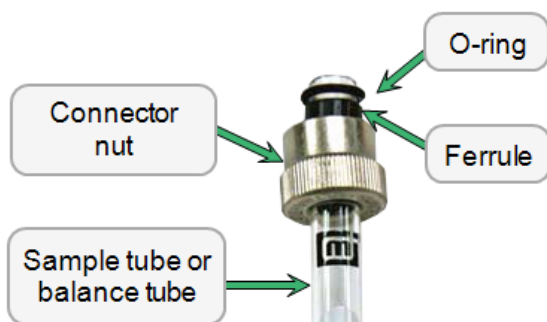
1. Weigh the sample tube set containing the sample. Record the mass on the [Sample Data Worksheet for Gas Adsorption on page E - 2](#) as *Mass of Sample tube set plus sample (After Degas)*.
2. Subtract the *Mass of empty sample tube set (Before Degas)* from the *Mass of Sample tube set plus sample (After Degas)* to obtain the sample's mass. Record this value as *Mass of sample (After Degas)*.

## SAMPLE TUBE INSTALLATION



The equipment images in this topic may differ slightly from your equipment, however the instructions are the same unless otherwise noted.

1. Remove the sample tube stopper, if used.
2. Place the connector nut, ferrule, and O-ring onto the sample tube stem. The sample tube ferrule is tapered slightly on one end, however it may be installed with the tapered end in the up or down position.



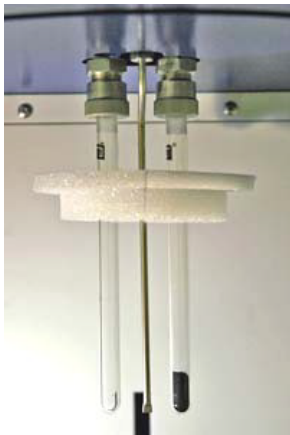
If analyzing samples with a surface area of  $1.0 \text{ m}^2$  or less, use filler rods in the sample and balance tubes. See [Analyze Samples with a Total Surface Area of  \$1.0 \text{ m}^2\$  or Less on page A - 1](#).

3. Attach the sample tube to the analysis port. Ensure it is fully in the port. Secure it in place by screwing the connector nut into the analysis port. Hand-tighten the connector nut.
4. If the balance tube is not installed, attach it to the balance port following the above procedures. If a filler rod was used in the sample tube, one must be used in the balance tube.



It is not necessary to remove and replace the balance tube between analyses unless it has been contaminated or if using a different size sample tube.

5. Place the dewar cover under the sample and balance tubes and slide upward until it is three inches from connector nuts.



## FILL AND INSTALL THE DEWAR



**See also:**

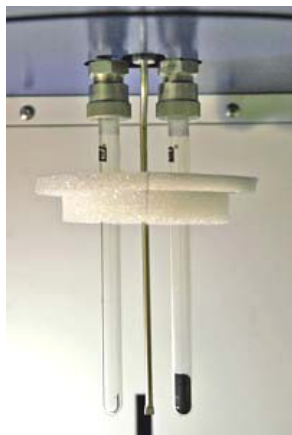
[Dewar Precautions on page 6 - 1](#)

1. Fill the dewar with the analysis bath liquid (such as liquid nitrogen) to no higher than 1 in. (2.5 cm) from the top for a large dewar or no higher than 3/4 in. (2 cm) from the top for a smaller dewar. Filling the dewar higher than this will cause an error in the free space measurement.



Incorrect fluid levels can lead to measurement errors. Check the level of the bath liquid before each analysis.

2. Check the level of the analysis bath liquid. For best results, if the dewar has not been used for a while, allow approximately 30 minutes for the temperature of the dewar to stabilize with the bath liquid, then recheck the level of the bath liquid. Add additional liquid if necessary.
3. Slide the dewar cover to approximately 3 in. from the sample port nuts to ensure a proper seal on the top of the dewar.



4. Place the dewar on the elevator. A dewar support is not necessary for the larger dewar used with the Gemini 2390t. Place the dewar directly on the elevator.
5. Close the safety shield on the front of the analyzer.

## PERFORM A SAMPLE ANALYSIS

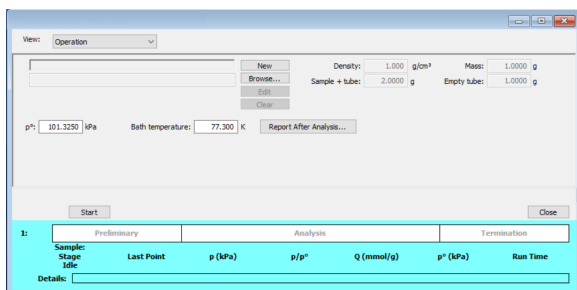
### Unit [n] > Start Analysis

#### CFR Note

In 21CFR11 environments, users are required to login to the Confirm application to start an analysis. Once the analysis window is opened, manual control is disabled until the analysis has completed. During analysis, pausing and resuming is allowed, however, steps cannot be skipped.



Ensure the analyzer safety compartment door is closed before beginning an analysis. If the analyzer is operated at an excessive pressure, the sample or balance tube could become dislodged from its port, possibly causing personal injury or damage to the equipment.



The screenshot shows a software window titled 'Operation'. It contains input fields for 'Density' (1.000 g/cm³), 'Mass' (1.0000 g), 'Sample + tube' (2.0000 g), and 'Empty tube' (1.0000 g). There are buttons for 'New', 'Browse...', 'Exit', and 'Clear'. Below these are fields for 'pH' (101.3250 kPa) and 'Bath temperature' (77.300 K), with a 'Report After Analysis...' button. At the bottom, there is a 'Start' button and a 'Close' button. Below the main window, there is a table with columns: 'Sample Stage', 'Last Point', 'p (kPa)', 'p/p\*', 'Q (mmol/g)', 'p\* (kPa)', and 'Run Time'. The table has one row labeled '1:' and a 'Details' link.

### Analysis Fields and Buttons Table

Field or Button	Description
<b>Density / Mass / Sample + Tube / Empty Tube</b> [text box]	Enter values for the sample's mass and density. These values may be edited after analysis.
<b>New</b> [button]	Creates a new sample file.
<b>Sample tube</b> [drop-down box]	Select the sample tube to be used for this analysis.
<b>Start</b> [button]	Click to start the analysis.



For fields and buttons not listed in this table, see [Common Fields and Buttons on page 2 - 3](#).





Sample Analysis Graph Fields and Buttons Table

Field or Button	Description
<b>Live Graph Settings</b> [button]	Select Thermal transpiration, x-axis Quantity (relative or absolute pressure) and the x-Axis Scale (linear or logarithmic).
<b>Report after analysis</b> [button]	Generates reports to the selected destination when the analysis is complete.
<b>Display Port</b> [button]	Generates a report on data being collected. The reports are displayed on the computer monitor only.
<b>Resume</b> [button]	Restarts the suspended analysis.
<b>Skip</b> [button]	Moves to the next step. Select the ports to skip. In 21CFR11 environments, steps cannot be skipped.
<b>Status window</b>	Displays the last point pressure and relative pressure for each port.
<b>Suspend</b> [button]	Suspends an analysis in progress.
For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .	

## PERFORM AN EMPTY TUBE ANALYSIS

### Unit [n] > Empty Tube Analysis

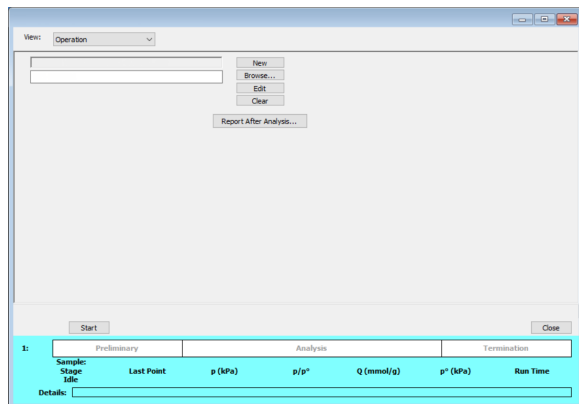
#### CFR Note

In 21CFR11 environments, users are required to login to the Confirm application to start an analysis. Once the analysis window is opened, manual control is disabled until the analysis has completed. During analysis, pausing and resuming is allowed, however, steps cannot be skipped.

Use to obtain sample tube parameters when using a calculated free space. An empty tube analysis can also be used for diagnostic purposes. The empty tube analysis should result in zero adsorptive. An amount other than zero could indicate a possible leak.



Ensure the analyzer safety compartment door is closed before beginning an analysis. If the analyzer is operated at an excessive pressure, the sample or balance tube could become dislodged from its port, possibly causing personal injury or damage to the equipment.



1. Click **Browse** to select a sample file.
2. Click **Report after analysis** to automatically generate reports when the analysis is complete. On the *Report Settings* window, select the report destination. Click **OK**.
3. Click **Start** to start the analysis. A window displays data as they are collected. A short delay is encountered before the port status changes from the Idle state at the bottom of the window.

When analysis is complete, remove the sample tube and store (or dispose) of the sample material as applicable.

## PERFORM A REFERENCE MATERIAL ANALYSIS

### CFR Note

In 21CFR11 environments, users are required to login to the Confirm application to start an analysis. Once the analysis window is opened, manual control is disabled until the analysis has completed. During analysis, pausing and resuming is allowed, however, steps cannot be skipped.

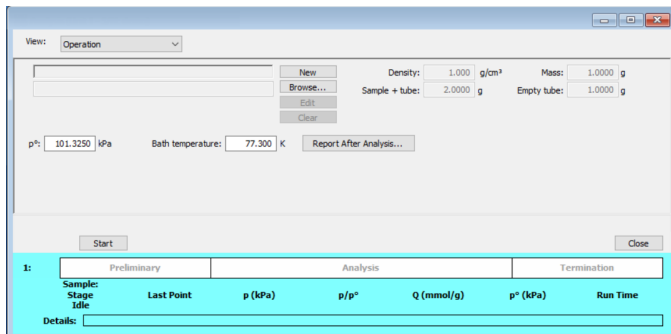
A reference material analysis is used to verify the analyzer is operating properly and producing optimum results. These methods provide specifications for critical report quantities and reporting of whether quantities are in or out of specification.

When running a reference material analysis, use the appropriate reference material provided in the accessories kit to perform this analysis. The results should match those shown on the label of the reference material bottle, within the tolerance level.



Ensure the analyzer safety compartment door is closed before beginning an analysis. If the analyzer is operated at an excessive pressure, the sample or balance tube could become dislodged from its port, possibly causing personal injury or damage to the equipment.

1. Go to **File > New Sample > [.SMP]** to create a new sample file.
2. Click **Replace All**. Locate and select the file *Carbon.smp*. Click **Load**. This file contains the appropriate conditions for analyzing the carbon reference material. All values will be copied into the new sample file. Edit fields as needed.
3. Click **Save As**, enter a new file name if necessary, click **Save**, then click **Close**.
4. Clean and label a sample tube. If installing a new balance tube, clean and label it also.
5. Determine the sample mass.
6. Degas the sample using a degassing unit.
7. Install the sample tube with degassed sample on the sample port.
8. Install the balance tube (if not installed) on the balance port.
9. Fill the dewar with liquid nitrogen to about 5 cm (2 in.) from the top. Allow the dewar to equilibrate to ambient conditions (approximately 30 minutes).
10. Close the sample compartment door.
11. Go to **Unit [n] > Start Analysis**.



12. Click **Browse**. Locate and select the sample file created previously for this analysis. Click **Open**.
  - a. Click **Report After Analysis**.
  - b. Select *Report after analysis* and *Preview* as the destination. Click **OK**.
13. Click **Start**.
14. Compare the BET Surface Area in the report with the BET Surface Area on the reference material bottle.
  - If results are within tolerance, the analyzer is operating properly.
  - If results are not within tolerance, refer to the following table for possible causes and actions. After performing the action, perform the reference material analysis again.

Cause	Action
The sample was not degassed properly.	Degas the sample again.
The gas lines are not clean.	Clean and verify the gas lines. See <a href="#">Clean and Verify the Gas Line on page 11 - 5</a> .
The measured free space is too high. This indicates the helium may not be pure enough.	Helium must be 99.9999% pure.

## PERFORM A QUICKSTART ANALYSIS

### Unit [n] > QuickStart Analysis

#### CFR Note


In 21CFR11 environments, users are required to login to the Confirm application to start an analysis. Once the analysis window is opened, manual control is disabled until the analysis has completed. During analysis, pausing and resuming is allowed, however, steps cannot be skipped.

The *QuickStart Analysis* option automatically creates a sample file and assigns the default parameters. Click **Edit** to modify the parameters or **Browse** to select another file.



Ensure the analyzer safety compartment door is closed before beginning an analysis. If the analyzer is operated at an excessive pressure, the sample or balance tube could become dislodged from its port, possibly causing personal injury or damage to the equipment.

### QuickStart Analysis Fields and Buttons

Field or Button	Description
<b>Sample tube</b> [drop-down box]	Click the drop-down arrow to select a sample tube file. This option is applicable only if the calculated method for measuring free space is selected for analysis conditions.
 For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .	

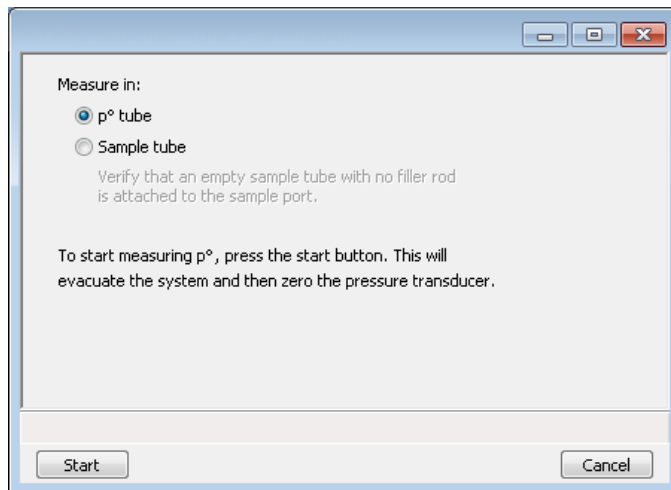
## MEASURE SATURATION PRESSURE

### Unit [n] > Start $P_0$ Measurement

#### CFR Note

In 21CFR11 environments, users are required to login to the Confirm application to start an analysis. Once the analysis window is opened, manual control is disabled until the analysis has completed. During analysis, pausing and resuming is allowed, however, steps cannot be skipped.

Evacuates the system and zeroes the pressure transducer.



Gemini Model	Description
2390a	$P_0$ can only be measured in the sample tube, therefore selections are not provided. Verify that a sample tube is attached to the sample port and that it does not contain a filler rod. Click <b>Start</b> .
2390p and 2390t	A $P_0$ tube is installed on these models. $P_0$ can be measured in the $P_0$ tube or in the sample tube. If using a sample tube, ensure that a sample tube is attached to the sample port and that it does not contain a filler rod. Click <b>Start</b> .

After analysis, go to **Unit [n] > Unit Configuration**. The recorded value now displays in the *Last Measured  $P_0$*  field.

## 7 ABOUT REPORTS

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### **Reports > Open Report > [.REP file]**

Opens a saved report.

### **Reports > Close Reports**

Closes all open reports. This option is unavailable if reports are being generated.

### **Reports > Start Report**

Generates a report on a sample analysis.

Reports can be generated for data collected on a sample that has completed analysis, collected on a sample currently being analyzed, or manually entered.

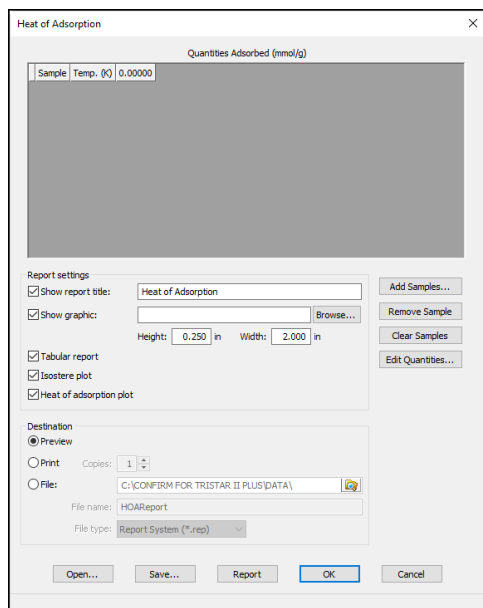
## **START REPORTS**

### **Reports > Start Report**

Starts the selected report. Select a file from the *Files* list. Ensure the selected file has a status of either *Complete* or *Analyzing*.

## HEAT OF ADSORPTION REPORT

### Reports > Heat of Adsorption



The dialog box titled "Heat of Adsorption" contains the following elements:

- Quantities Adsorbed (mmol/g):** A table with columns "Sample" and "Temp. (K)". The first row shows "0" and "0.00000".
- Report settings:**
  - ☒ Show report title: Heat of Adsorption
  - ☒ Show graphic: (with a "Browse..." button)
  - Height: 0.250 in Width: 2.000 in
  - ☒ Tabular report
  - ☒ Isotherm plot
  - ☒ Heat of adsorption plot
- Buttons:** Add Samples..., Remove Sample, Clear Samples, Edit Quantities...
- Destination:**
  - ☒ Preview
  - ☐ Print: Copies: 1
  - ☐ File: C:\CONFIRM FOR TRISTAR II PLUS\DATA\ (with a file name field containing "HQAReport" and a file type dropdown set to "Report System (\*.rep)")
- Bottom Buttons:** Open..., Save..., Report, OK, Cancel

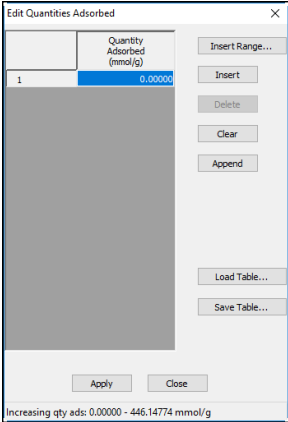

Use to select sample files, define quantities, and generate a *Heat of Adsorption* report. The isosteric heat of adsorption is an important parameter for characterizing the surface heterogeneity and for providing information about the adsorbent and the adsorption capacity. Multiple adsorption isotherms are obtained on the same sample using the same adsorptive but at different temperatures to obtain the heat of adsorption.

### Heat of Adsorption Fields and Buttons Table

Field or Button	Description
<b>Add Samples</b> [button]	Adds a sample file to the table.
<b>Clear Samples</b> [button]	Removes all entries from the table.
<b>Edit Quantities</b> [button]	Use to specify the range of surface coverage to include in the report.



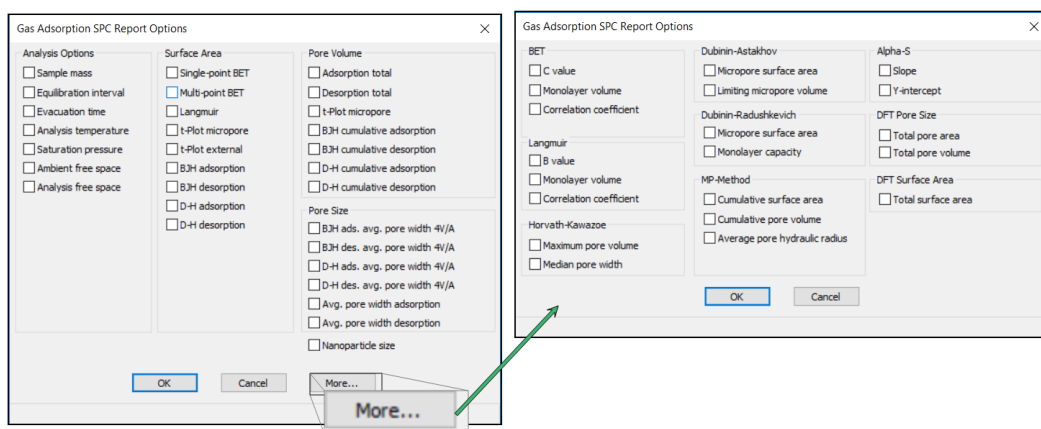
## Heat of Adsorption Fields and Buttons Table (continued)

Field or Button	Description
	 <p><b>Insert Range.</b> Click to specify the starting and ending quantities adsorbed and number of points to insert.</p> <p><b>Load Table.</b> Imports values from another file.</p> <p><b>Save Table.</b> Saves the current table as a .QNT file.</p> <p><b>Apply.</b> Applies all table changes.</p>
<b>Heat of adsorption plot</b> [selection]	Generates the <i>Heat of Adsorption</i> data in a graphical format.
<b>Isostere plot</b> [selection]	Generates a graph showing quantities of gas adsorbed versus the temperature.
<b>Remove Sample</b> [button]	Removes the selected sample from the list.
<b>Show graphic</b> [check box]	Use to show a graphic on the report header.  <b>Height / Width.</b> Enter the height and width of the selected graphic. These values determine the graphic appearance on the generated report.
<b>Show report title</b> [check box]	Select and enter a report title to appear on the report header.
<b>Tabular report</b> [check box]	Generates a tabular report of the included samples. A tabular report contains the numeric values contributed by each sample.
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	

## SPC REPORT

### Reports > SPC Report Options

Use to generate reports with various *SPC* (Statistical Process Control) options. All selected variables must be computed for each sample file used in an SPC report; therefore, it is more efficient to select only the necessary variables.

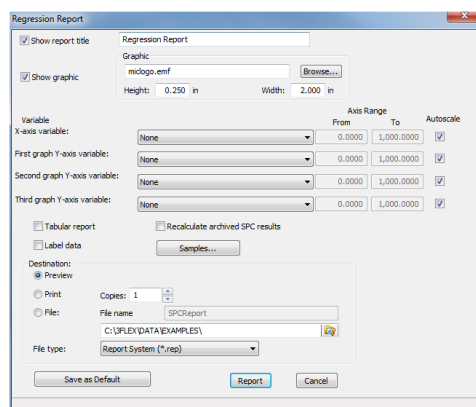


The selected items display as graph variable selections in **Reports > Regression Report** and graph selections in **Reports > Control Chart**. If additional report options are required, click [More](#).


## REGRESSION REPORT

### Reports > Regression Report



Use to generate a Statistical Process Control (SPC) Regression report to determine the interdependency between two variables. Up to three dependent variables (y-axis) may be plotted against a single independent variable (x-axis). The degree of correlation between the variables is also reported.



### Regression Report Fields and Buttons Table

Field or Button	Description
<b>Autoscale</b> [check box]	When enabled, allows the x- and y-axes to be scaled automatically.
<b>Axis Range</b> [text box]	Enter the beginning and ending values for the x- and y-axis ranges. These fields are disabled if <i>Autoscale</i> is selected.
<b>Label data</b> [check box]	Use to label the points on the plot to correspond with the values in the sample files.
<b>Recalculate archived SPC results</b> [check box]	<p>Use to have archived SPC values recalculated ensuring any changes made to the SPC Report Options are included in the new report. This option lengthens the time required to generate the report.</p> <div>  <p>If this recalculation option is enabled and sample files from an earlier application version are selected, it is recommended that copies of the archived sample files be used rather than the original. Selecting this option will make some archived sample files unreadable by the original application. If an earlier application version will not be reused, this warning message can be safely disregarded.</p> </div> <p>When this option is selected, this message will display:</p>

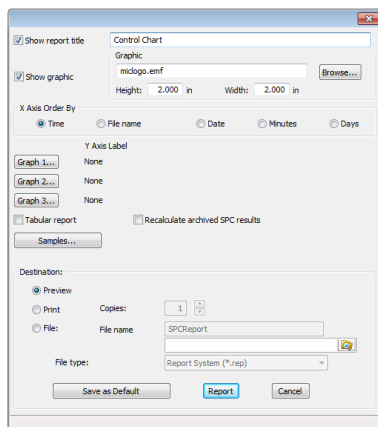
Regression Report Fields and Buttons Table (continued)

Field or Button	Description
	<p><b>Saving the recalculated SPC data may render some files unreadable by the original application. Saving the SPC data speeds up future SPC reports.</b></p> <hr/> <p> If <i>Do not show me this message again</i> is selected, the message cannot be redisplayed without Micromeritics assistance.</p> <hr/> <p>The first time this option is used, the time it takes to generate the report is lengthened. The second time the report is generated, if using the same sample files used in the initial calculation, it is recommended that this option not be selected since the data was recalculated previously. If a sample file is added or removed from the report after the initial recalculation, this option should be selected again to ensure the data from the newly added or removed sample file is recalculated.</p>
<b>Samples</b> [button]	Select additional sample files to add to the report.
<b>Save as Default</b> [button]	Click to save selected report options as default report settings.
<b>Show graphic</b> [check box]	Use to show a graphic on the report header.  <b>Height / Width.</b> Enter the height and width of the selected graphic. These values determine the graphic appearance on the generated report.
<b>Show report title</b> [check box]	Select and enter a report title to appear on the report header.
<b>Tabular report</b> [check box]	Generates a tabular report of the included samples. A tabular report contains the numeric values contributed by each sample.
<b>X- and Y-Axis Variable</b> [drop-down box]	Use to designate the x- and y-axes variables. The variables in the drop-down lists are those selected in the <b>Reports &gt; SPC Report Options</b> window. Use these options to plot the regression of up to three y-axis variables against the x-axis variable.
	For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .

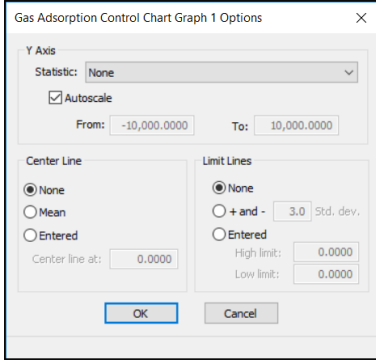
## CONTROL CHART REPORT

### Reports > Control Chart



Generates a Statistical Process Control (SPC) control chart report which plots the changes in a statistic.




### Control Chart Fields and Buttons Table

Field or Button	Description
<b>Graph [n] [button]</b>	<p>Defines the y-axis of each graph.</p>  <p><b>Statistic.</b> [drop-down box] Displays the SPC variables selected on the <b>Reports &gt; SPC Report Options</b> window. The selected variable will be plotted for each selected sample. This selection also becomes the y-axis label.</p> <p><b>Autoscale.</b> Allows the y-axis to be scaled automatically. To specify a range, deselect this option and enter a range in the <i>From</i> and <i>To</i> fields.</p>

Control Chart Fields and Buttons Table (continued)

Field or Button	Description
	<p><b>Center Line.</b> Displays placement options for the center line in the graph. Select <i>Entered</i> to specify placement of the line or <i>Mean</i> to place the center line at the calculated mean value for the selected samples.</p> <p><b>Limit Lines.</b> Displays limiting lines options. Lines can be placed at some multiple of the standard deviation or at specified positions (<i>Entered</i>). When <i>Entered</i> is selected, enter the <i>High limit</i> and <i>Low limit</i> fields with appropriate values.</p>
<b>Recalculate archived SPC results</b> [check box]	<p>Use to have archived SPC values recalculated ensuring any changes made to the SPC Report Options are included in the new report. This option lengthens the time required to generate the report.</p> <hr/> <div>  <p>If this recalculation option is enabled and sample files from an earlier application version are selected, it is recommended that copies of the archived sample files be used rather than the original. Selecting this option will make some archived sample files unreadable by the original application. If an earlier application version will not be reused, this warning message can be safely disregarded.</p> </div> <hr/> <p>When this option is selected, this message will display:</p> <div> <p><b>Saving the recalculated SPC data may render some files unreadable by the original application. Saving the SPC data speeds up future SPC reports.</b></p> </div> <hr/> <div>  <p>If <i>Do not show me this message again</i> is selected, the message cannot be redisplayed without Micromeritics assistance.</p> </div> <hr/> <p>The first time this option is used, the time it takes to generate the report is lengthened. The second time the report is generated, if using the same sample files used in the initial calculation, it is recommended that this option not be selected since the data was recalculated previously. If a sample file is added or removed from the report after the initial recalculation, this option should be selected again to ensure the data from the newly added or removed sample file is recalculated.</p>
<b>Report</b> [button]	Generates the report.
<b>Samples</b> [button]	Select additional sample files to add to the report.

## Control Chart Fields and Buttons Table (continued)

Field or Button	Description
<b>Show graphic</b> <i>[check box]</i>	Use to show a graphic on the report header.  <b>Height / Width.</b> Enter the height and width of the selected graphic. These values determine the graphic appearance on the generated report.
<b>Show report title</b> <i>[check box]</i>	Select and enter a report title to appear on the report header.
<b>Tabular report</b> <i>[check box]</i>	Generates a tabular report of the included samples. A tabular report contains the numeric values contributed by each sample.
<b>X Axis Order by</b> <i>[group box]</i>	Select the order in which x-axis statistics are placed. Sort by:  <b>Time.</b> Time the files were analyzed.  <b>File name.</b> Alphanumeric order.  <b>Date.</b> Date the files were analyzed.  <b>Minutes.</b> Minutes elapsed from the first file placed on the list, which is the earliest-analyzed file.  <b>Days.</b> Number of days elapsed from the first file placed on the list, which is the earliest-analyzed file.
 For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .	

## ***MICROACTIVE REPORTS***

MicroActive reports are generated automatically after an analysis is performed. This feature provides a quick and easy way to investigate and manipulate analysis data using a variety of reporting methods.

When a sample file with a status of *Complete*, *Analyzing* or *Entered* is opened, a linear plot and log plot of the data collected during analysis are displayed as well as a summary of the analysis giving the total pore volume. Numerous reports are accessible from a drop-down menu.

When a report is opened, plots and summary data are displayed, and in some reports certain parameters (for example, thickness curve type, pore geometry, and interaction parameters) are also displayed. Plots may be edited by selecting the data points or data point range to be included in the plots and modifying the parameters. When a report is edited, the results are immediately reflected in the plots and summary data.

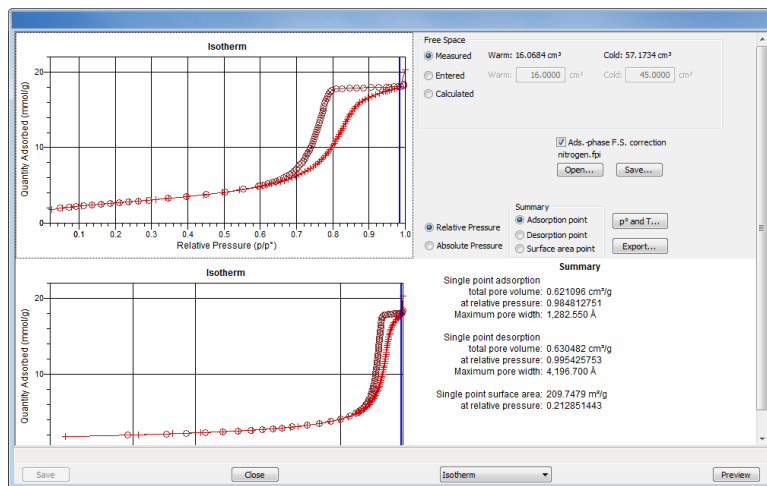
Log in to your [customer portal](#) to access MicroActive Report Tutorials.



## INTERACTIVE REPORTS

When opening a sample file that contains data from a complete or in-progress analysis, the interactive reporting feature is enabled.

1. When opening a sample file that contains analysis data, a window with the following information will display:
  - a linear plot and log plot of the data collected during analysis
  - a summary of the analysis giving a single total pore volume and surface area



2. To view the plots in either relative or absolute pressure, select either the *Relative Pressure* or *Absolute Pressure* option.
3. To view the reports selected for generation during the analysis, click **Preview**.
4. From the drop-down list at the bottom of the window:
  - change the option presentation of the sample information window to either *Basic* or *Advanced* to modify certain file parameters, or
  - select another plot from the list and edit the data contained in the plot.
5. When ranges are edited, the changes are reflected immediately in the plots and the summary data displayed in the window. Some editing options are:
  - Drag the blue bars to increase or decrease the range of data included in the plot.
  - Edit the Isotherm Linear Plot to include or omit the data point from the BET plot.
  - Right click to display a popup menu to include reports; enable or select overlays; edit curves, axes, legends, titles; and copy and paste the data in a graph or in tabular format.
6. Click **Save**.

## ***EVALUATE REPORT RESULTS***

Analysis reports provide a record of test conditions, experimental data, and information extracted from the experimental data by application of various reduction methods. This topic discusses the elements of various reports presented by Micromeritics' analyzers and suggests ways by which the merit of the reported information may be evaluated.

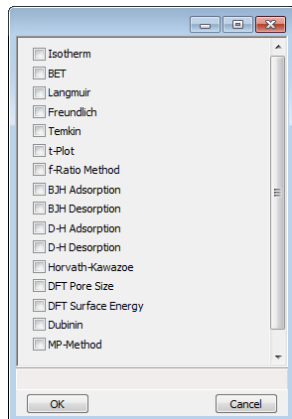
Regardless of the precautions exercised before the analysis, problems still may occur during the analysis, or as a result of using inappropriate parameters or even inappropriate methods. The analysis data should be inspected for evidence of experimental error. The traditional method of confirming the quality of the experiment is to repeat the analysis. Toward that end, Micromeritics' analyzers log and report the exact conditions of each analysis.

Analysis data can be evaluated by:

- Viewing the Validation Report
- Inspecting the Isotherm Plot
- Evaluating the Isotherm Tabular Data Set
- Reviewing Reduced Data

## VIEW THE VALIDATION REPORT

The *Validation* report shows whether the data collected during an analysis are within typical ranges. Select the types of reports to include by selecting the report in the *Validation Report Options* window.



When a selected report is generated, if errors occur, a message is displayed across the top portion of the report and a unique symbol displays on the graph.

## INSPECT THE ISOTHERM PLOT

Evaluation of data should begin with a visual inspection of the isotherm plot. The plot should be composed of data which have not been subjected to mathematical smoothing as far as possible. If the data describe a Type I isotherm, then the plot is best shown on a logarithmic pressure axis so that details of the low pressure region are revealed. Data in this region are important particularly for micropore studies. Examine the plot to determine if any points are outliers or if a region of the isotherm exhibits characteristics (spikes, steps, etc.) which are inconsistent with the physical process being monitored. The philosophical question of whether or not these suspected extraneous data points should be removed from the raw data is not considered here, but it may be appropriate to exclude an outlier from reduced data. Too many outliers can cause the integrity of the total data set to come under suspicion.

Examine specific reported values to confirm that the isotherm data were collected under reasonable conditions and using reasonable parameters. For example, confirm that the free space values reported are typical for the sample holder and bath in use. A problem with either ambient or analysis free space values may indicate a free space measurement error and affect all calculations of quantity adsorbed.

The raw data should be carefully examined before it is reduced. Errors that occur in raw data will only be exacerbated in reduced data.<sup>1)</sup>

## ***EVALUATE THE ISOTHERM TABULAR DATA SET***

Another place to look for reasonableness of the data is the adsorptive uptake by the sample in the BET range ( $P/P_0 = 0.05$  to  $0.30$ ). Total uptake is the specific quantity adsorbed ( $\text{cm}^3/\text{g STP}$ ) times the sample mass (g). As an example, the level of uncertainty in this range typically is less than  $0.1 \text{ cm}^3 \text{ STP}$  for a high performance system. Total uptake quantities should be some multiple of this level of uncertainty. Otherwise, an unfavorable signal-to-noise ratio and unreliable data result. The solution is to use a greater quantity of sample to increase adsorptive uptake.

Another valuable bit of information resides in the tabulated saturation pressure. This pressure is expected to change somewhat over the duration of an analysis, but it is not expected to do so with large or abrupt transitions. Unreasonable saturation pressures or unusual changes may indicate that a gas different from the adsorptive was used in determining  $P_0$ , that the level of the cryogen fell too far, or that the cryogen is impure or inappropriate.

With experience, obvious signs of problems can be detected by a quick inspection of the tabular and graphical data. If the data appear satisfactory, the next step is to evaluate the reduced data.<sup>2)</sup>

## ***REVIEW REDUCED DATA***

Isotherm data may be analyzed by any one of several reduction methods depending on the analyzer model and pressure range employed. The quality of the results depends on the quality of the isotherm, the congruity of the data reduction parameters with experimental conditions, the agreement of the theoretical model with the physical gas-solid system, and compliance to the pressure range over which the method is valid. Typically, results can be appraised by examining a few salient areas of the report as described in the following topics.<sup>3)</sup>

## ***PHYSICAL PARAMETERS***

The value of physical parameters which are used only in data reduction routines should be reviewed to assure that they agree with experimental conditions. These parameters can be changed and the experimental data recalculated if an error is discovered or if exploring an alternate value is desired. Analysis condition values used in the calculation of quantity adsorbed can be changed also. These are typically the manually entered free space(s), nonideality correction factor, and bath temperature.

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<sup>1)</sup> The information in this article is extracted from Analytical Methods in Fine Particle Technology, Webb, P. and Orr, C., (1997).

<sup>2)</sup> Most of the information in this article is extracted from Analytical Methods in Fine Particle Technology, Webb, P. and Orr, C., (1997).

<sup>3)</sup> Most of the information in this article is extracted from Analytical Methods in Fine Particle Technology, Webb, P. and Orr, C., (1997).

The area occupied by a single adsorbed molecule is a required parameter in the calculation of surface area by the BET and Langmuir methods. The software provides a default value, but other values are found in the literature. McClellan and Harnsberger<sup>1)</sup> provide a comprehensive review of such values.

The volume of pores of a specific size range is calculated from the gas quantity adsorbed in them by converting the quantity to its liquid equivalent volume. This is achieved through use of a density conversion factor calculated from the ratio of molar densities of the condensed adsorbate at bath temperature to the gaseous phase at STP. The necessary information is found in handbooks. The software contains default values for common adsorptives; values for other adsorptives must be calculated.

The terms for liquid surface tension  $\gamma$ , contact angle between solid and liquid phase  $\theta$ , molar volume of the adsorbate  $n$ , gas constant  $R$ , and sample temperature  $T$  are treated as one constant, the adsorbate property factor  $A$  expressed by:

$$A = \frac{\gamma n \cos \theta}{RT}$$

using which, the Kelvin equation<sup>2)</sup> reduces to

$$\ln \frac{p}{p_0} = \frac{A}{r_h}$$

Either surface tension, contact angle, or molar volume can be revised individually to give a new value for the factor  $A$ , or  $\theta$  can simply be altered arbitrarily for exploratory purposes.

The thermal transpiration correction requires two parameters which may be adjusted from those of the default values. The first is the inside stem (neck) diameter of the sample holder, and the second is the hard-sphere diameter of the adsorptive molecule. The sample holder inside diameter is available from the documentation provided with it or is measurable. Information on hard-sphere diameters of molecules may be obtained from handbooks.

For terms such as the interaction parameter found in the Horvath-Kawazoe calculation<sup>3)</sup>, the Dubinin affinity coefficient or Astakhov exponent<sup>4)</sup>, the default values as provided by the software generally are adequate. A search of the technical literature is required if the analysis involves a gas-solid system other than that covered by the default values.

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1) McClellan, A.L., and Harnsberger, H.F., Journal of Colloid and Interface Science, 23, 577 (1967).

2) Thomson, W., Phil. Mag. S., 42, 448 (1871).

3) Everett, D.H. and Powl, J.C., J. Chem Soc., Faraday Trans. 1, 72, 619 (1976).

4) Dubinin, M. and Radushkevich, L.V., Proc. Acad. Sci. USSR, 55, 331 (1947).

The t-Plot method plots quantity adsorbed ( $V_a$ ) against thickness ( $t$ ) derived from a thickness equation, and the Dubinin transform plots quantity adsorbed against  $\log(P/P_0)^n$ . All of these data reduction methods were first proposed for specific applications. The user must make a judgment as to the applicability of the method to a gas-solid system.

If applied appropriately, all transform plots will exhibit a linear range and the regression analysis must be applied only over the linear range and within the range of application. Fitting a regression line to surface area transformation plots should yield a correlation coefficient of 0.9999 or better and for t-plots and Dubinin plots the correlation coefficient should be 0.99 or better.

If the data reduction model does not apply to the gas-solid system under examination, then it may be that either no linear range exists within the pressure range of validity, or that solutions derived from the regression line of the linear range are intuitively incorrect, that is, they have no relevance to the physical situation, such as a negative C-value from a BET transform.

## ***BET C-VALUE***

BET theory assumes uniform surface coverage with no favored adsorption sites and it also assumes that the gas is more strongly attracted to the surface than to other gas molecules. The typical range of BET C-values is from about 5 to well over 100. Values much less than 5 imply that the gas-to-gas affinity is competing with the gas-to-solid affinity which conflicts with the basic assumptions of BET theory. C-values much greater than 100 indicate very strong attraction for the surface or preferential adsorption.

Provided the isotherm was determined with negligible error and the regression line to the BET transformation data was fit properly, then an out-of-range C-value probably indicates that the gas-solid interaction for the particular sample material does not conform to the BET model. An inappropriate adsorption model may be indicated also by the coefficient of correlation of the regression line, 0.999 being about the minimum value expected with five more or less equally spaced points. In the case of indications of poor conformance to the BET model, the Langmuir data reduction method should be examined.

## ***DATA ANALYSES BY THE BJH METHOD***

In general, this method visualizes the incremental decomposition of an experimental isotherm, starting at the highest relative pressure or pore size. At each step the quantity of adsorptive involved is divided between pore-emptying and film-thinning processes and is accounted for totally. This computational algorithm frequently leads to inconsistencies when carried to small mesopore sizes. If the thickness curve used is too steep, ultimately it will predict a larger increment of adsorptive for a given pressure increment than is actually observed. The algorithm must stop since a negative pore volume is nonphysical. Accumulated error results in the calculation of a too large volume of (possibly nonexistent) small pores if the thickness curve used underestimates film thinning.

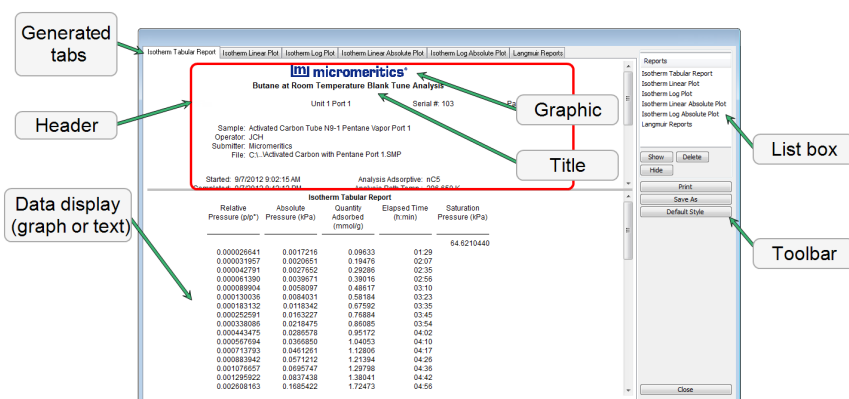
## REPORT FEATURES AND SHORTCUTS

### CFR Note

In 21CFR11 environments, members of the Analyst group must click [Preview](#) on the sample file window to access this screen.

Reports can be customized and manipulated using the toolbar, shortcut menus, the zoom feature, or axis cross-hairs.

- After analysis, reports can be viewed, printed, and/or copied and pasted into other documents.
- The report zoom feature provides the viewing of fine graph details and the ability to shift the axes.
- All reports contain a header displaying file statistics.

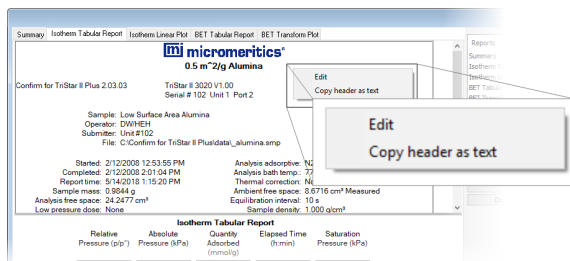


If configured, the report header can also contain a graphic and/or a title.

- Tabular and graphical reports contain sample and analyzer statistics such as analysis date/time, analysis conditions, etc.
- The headers contain notes of sample file changes occurring after analysis.
- Summary report headers contain the same information as tabular and graphical reports with the exception of notes.

## REPORT HEADER SHORTCUTS

Display header shortcuts by right clicking in the report header.



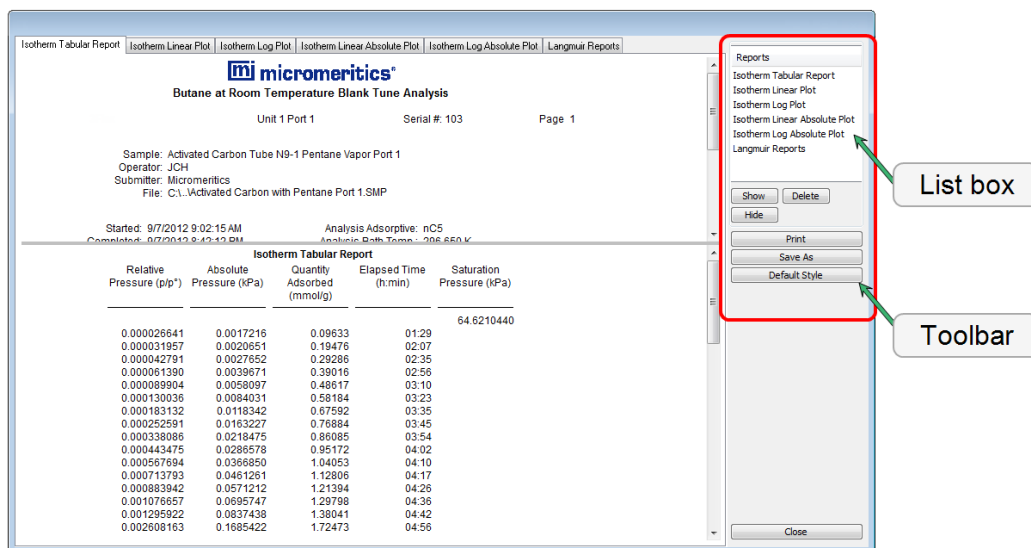
### Report header Shortcut Field and Button Table

Field or Button	Description
Copy header as text	Use to copy the report header as text. Text is copied to the clipboard and then can be pasted into other documents.
Edit	Use to edit the report title and/or graphic in the report header.



## REPORT TOOLBAR

The *Report* window has a toolbar on the right portion of the window and selectable tabs at the top of the report header. To view a specific report, either select the tab or the report in the *Reports* list box, then click **Show**.

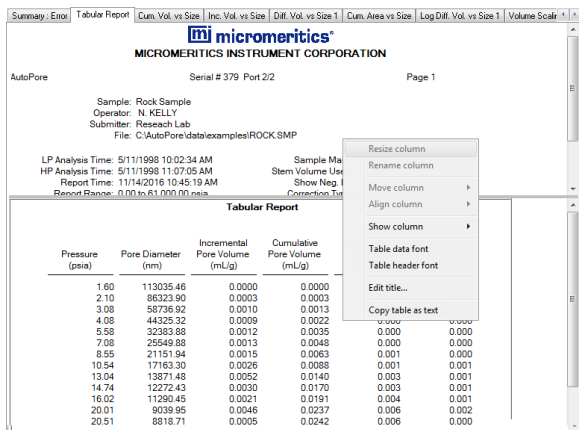


## Report Toolbar Fields and Buttons Table


Field or Button	Description
<b>Default Style</b> [button]	Specify default report parameters for fonts and curve properties.
<b>Delete</b> [button]	Deletes the selected report in the <i>Reports</i> list box. Deleted reports will have to be regenerated if deleted in error.
<b>Hide</b> [button]	Hides (or temporarily removes) the selected report from the tabbed view. The report name remains in the <i>Reports</i> list box.
<b>Print</b> [button]	Displays the <i>Print</i> window for report output.
<b>Reports</b> [group box]	Contains a list of all generated reports. The same reports display as tabs at the top of the report header unless the report has been hidden using the <b>Hide</b> button.
<b>Show</b> [button]	Displays the selected or hidden report in the <i>Reports</i> list box.
For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .	

## TABULAR REPORT FEATURES AND SHORTCUTS

Display tabular report shortcuts by right clicking in the body of the tabular report. Column shortcuts require right clicking on the column to be modified.

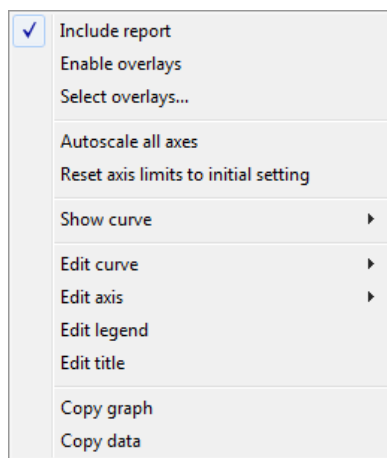


Tabular Reports Shortcut Options and Descriptions Table

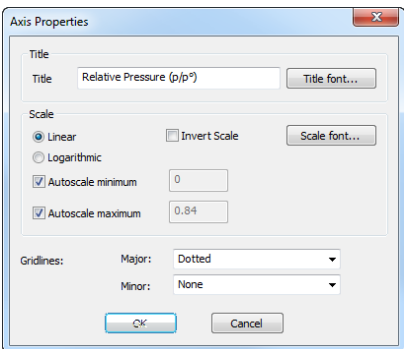
Field or Button	Description
<b>Align column</b>	Select to change the column alignment to either left, right, or centered.
<b>Copy table as text</b>	Use to copy the report contents to the clipboard as tab-delimited text. It can then be pasted into another document.
<b>Edit title</b>	Use to edit the report title and/or title font attributes. Click <a href="#">Font</a> to modify font attributes.
<b>Move column</b>	Right click the column to be moved. Select <i>Move column</i> on the shortcut menu and select <i>Left</i> or <i>Right</i> for the move.
<b>Rename column</b>	Right click the column to be renamed. Select <i>Rename column</i> on the shortcut menu and enter the new column name.
<b>Resize column</b>	Right click the column to be resized. Select <i>Resize column</i> on the shortcut menu and enter the new column width in inches.
<b>Show column</b>	Displays a list of all columns. Click a column to add a checkmark to show the column or remove the checkmark to hide the column.
<b>Table data font</b>	Right click in the report data. Select <i>Table data font</i> on the shortcut menu.
<b>Table header font</b>	Right click in the report data. Select <i>Table header font</i> on the shortcut menu.
 <b>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</b>	

## GRAPH FEATURES AND SHORTCUTS

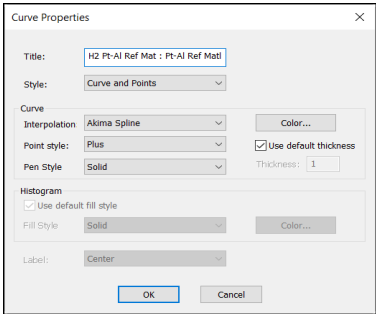
Right click in the graph area to display graph report shortcuts.



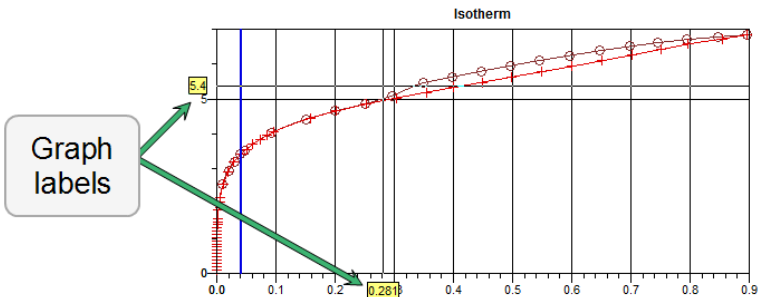
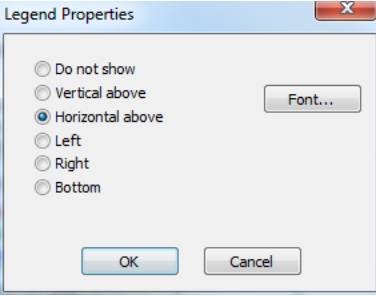

### Graph Shortcut Options and DescriptionsTable

Field or Button	Description
<b>Autoscale all axes</b>	Returns the report to full view after using the zoom feature.
<b>Copy Data</b>	Copies the report data to the clipboard. It can then be pasted into other software programs as tab-delimited columns or copied as an overlay onto another graph.
<b>Copy Graph</b>	Copies the graph to the clipboard. It can then be pasted into other software programs.
<b>Edit axis</b>	<p>Use to edit the selected axis properties.</p>  <p><b>Grid lines.</b> Use to change how to display major / minor grid lines.</p> <p><b>Scale.</b> [group box]</p>

## Graph Shortcut Options and DescriptionsTable (continued)

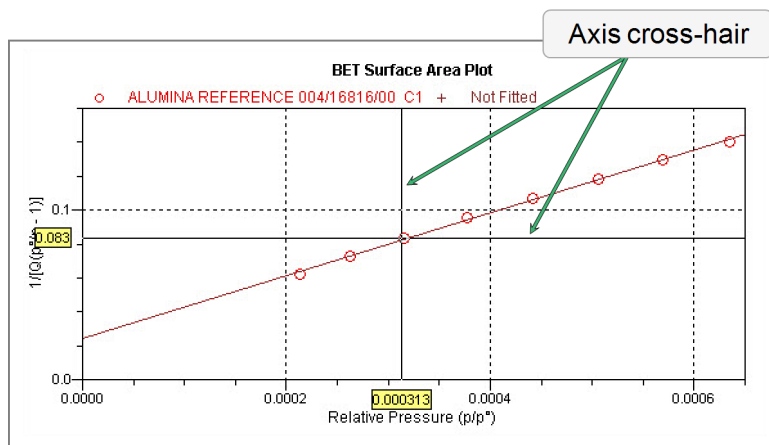
Field or Button	Description
	<ul style="list-style-type: none"> <li>• <b>Autoscale minimum / maximum.</b> To manually specify minimum / maximum autoscale, deselect the option and enter the new amount in the text box.</li> <li>• <b>Invert scale.</b> Use to invert the scale.</li> <li>• <b>Linear / Logarithmic.</b> Select the option to scale the graph as linear or logarithmic.</li> <li>• <b>Scale font.</b> Use to modify the font for the scale label. Deselect <i>Use default font</i> to enable font options.</li> </ul> <p><b>Title.</b> Use to edit the selected axis label.</p> <p><b>Title font.</b> Use to modify the font for the selected axis label. Deselect <i>Use default font</i>. Select new font attributes for report data. Enable <i>Use default font</i> to reset default fonts.</p>
<b>Edit curve</b>	<p>Use to edit selected curve properties.</p>  <p><b>Color.</b> Click to change the curve color.</p> <p><b>Curve.</b> <i>[group box]</i> Use to change the interpolation, point style, and pen style for the selected curve. These options are disabled if <i>Use default fill style</i> is selected in the <i>Histogram</i> group box.</p> <p><b>Histogram.</b> <i>[group box]</i> Enabled only if <i>Histogram</i> is selected in the <i>Style</i> drop-down list. Use to specify the type of fill, fill color, and label position for the selected curve.</p> <p><b>Label.</b> Select where the graph point labels will display (left, right, center, etc.) on the SPC report.</p>

Graph Shortcut Options and DescriptionsTable (continued)

Field or Button	Description
	 <p><b>Style.</b> Use to select another style for the collected data curve.</p> <p><b>Title.</b> Use to change the title of the selected curve.</p> <p><b>Use default thickness.</b> Uses the default curve thickness. Deselect to enter a new thickness number in the <i>Thickness</i> text box.</p>
<b>Edit legend</b>	<p>Use to change the legend location and font.</p> 
<b>Edit title</b>	Select to change the report title.
<b>Enable Overlays</b>	If overlays have been selected, this option displays (or hides) the overlays.
<b>Include report</b>	When selected, places a checkmark to the left of the report in the <i>Select Reports</i> list box on the <i>Report Options</i> tab.
<b>Reset axis limits to initial setting</b>	Removes the cross-hair and returns the graph back to the initial setting.
<b>Select overlays...</b>	Displays the option to select files to overlay onto the active graph. To view the overlays, click <i>Enable Overlays</i> on the shortcut menu.
<b>Show curve</b>	Displays a list of all curves. Select the curve(s) to display.
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	

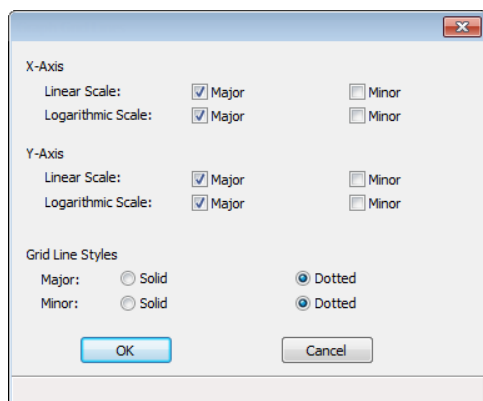
### Axis Cross-Hair

Left click on the graph to view the cross-hair coordinates.




## Graph Grid Lines

### Options > Graph Grid Lines



Use to select how grid lines appear on reports. This menu option is not available if using *Restricted* option presentation.

### Graph Grid Lines Fields and Buttons Table

Field or Button	Description
<b>Grid Line Styles</b> [selection]	Select if the major and/or minor grid lines should appear as solid or dotted lines.
<b>X-Axis / Y-Axis</b> [selection]	Select major and/or minor lines to display in reports for the logarithmic and linear scales. Deselect this option to remove the grid lines.
 <b>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</b>	

## Zoom Feature

Use the zoom feature to examine graph details. Click, hold, and drag the left mouse button on the graphical area to be enlarged. A box will display in the area to be enlarged. To return to normal view, right click in the graph and select *Autoscale all axes*.

## GRAPH AND SAMPLE OVERLAYS

### CFR Note

In 21CFR11 environments, this feature is applicable to members of the Developer group only.

Use the graph overlay functions to compare multiple graph options. Graphical lines are differentiated by the use of varying colored symbols outlined on a legend. Overlays may be generated in two ways:

- **Multiple Graph Overlays.** Overlay two different types of graphs from one sample.
- **Multiple Sample Overlays.** Overlay graphs of the same type with that of the current plot. This type of overlay is available only for:
  - BJH Adsorption / Desorption
  - DFT Pore Size / Surface Energy
  - Dollimore-Heal Adsorption / Desorption
  - Horvath-Kawazoe
  - M-P Method



This feature is available only when using *Advanced* option presentation. Go to **Options > Option Presentation > Advanced**.



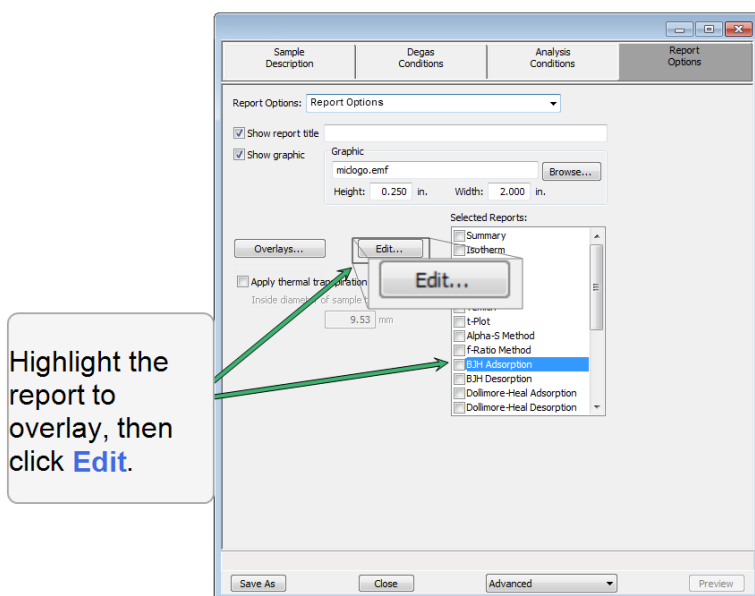
## OVERLAY MULTIPLE SAMPLE FILES

### CFR Note

In 21CFR11 environments, this feature is applicable to members of the Developer group only.

To overlay the same type of graph on multiple samples:

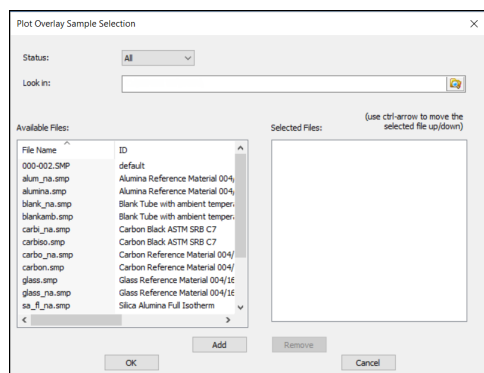
1. Go to **File > Open**.
2. Select a .SMP file, then click **Open**. If the Isotherm plot displays, select *Advanced* from the drop-down list at the bottom of the window to display the tabbed window view.
3. Click the *Report Options* tab.
4. In the *Selected Reports* list box, highlight a report then click **Edit**. Use the following table to complete the process for the selected report.



If overlaying this type of report...	Then...
<ul style="list-style-type: none"> <li>• Isotherm</li> </ul>	<ol style="list-style-type: none"> <li>On the <i>Isotherm Report Options</i> window, select one or more plots in the <i>Selected Reports</i> group box, then click <b>Options</b> to the right of the selected plot.</li> <li>On the <i>Plot Options</i> window, select <i>Plot curve</i> and/or <i>Plot points</i> if they are to be included in the overlay. If the x- and/or y-axes are to be autoscaled, enable <i>Autoscale</i>; otherwise, enter the <i>From</i> and <i>To</i> points for the axes. Click <b>OK</b>.</li> <li>On the <i>Isotherm Report Options</i> window, in the <i>Plot Options</i> group box, select <i>Plot overlays</i>. Click <b>OK</b>.</li> <li>Continue to Step 5.</li> </ol>
<ul style="list-style-type: none"> <li>• Alpha-S Method</li> <li>• BET Surface Area</li> <li>• <i>f</i>-Ratio Method</li> <li>• Freundlich</li> <li>• Langmuir Surface Area</li> <li>• <i>t</i>-plot</li> <li>• Temkin</li> </ul>	<ol style="list-style-type: none"> <li>On the pop-up window, select <i>Overlay samples</i>. Verify other fields. Click <b>OK</b>.</li> <li>Continue to Step 5.</li> </ol>
<ul style="list-style-type: none"> <li>• BJH</li> <li>• Dollimore-Heal</li> <li>• MP-Method</li> </ul>	<ol style="list-style-type: none"> <li>Select the report variable from the <i>Selected Reports</i> group box, then click <b>Edit</b>.</li> <li>Click the down arrow on the <i>Overlay</i> field, then select the <i>Samples</i> option. Verify other fields. Click <b>OK</b>.</li> <li>Click <b>OK</b> again.</li> </ol>

5. On the *Report Options* tab, click **Overlays**.


6. On the *Plot Overlay Sample Selection* window, move up to 25 files from the *Available Files* box to the *Selected Files* box:



7. Click **OK**.

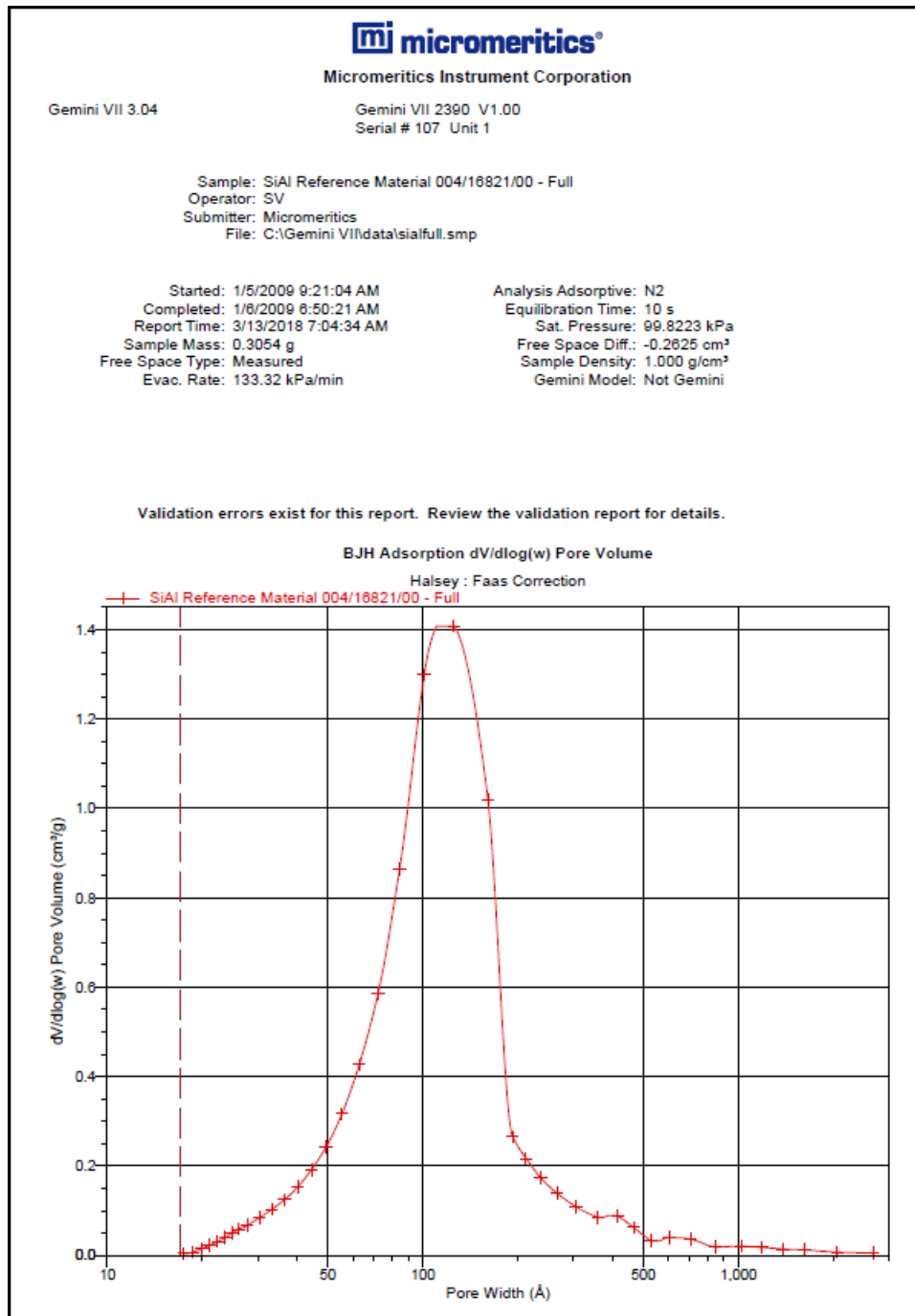
8. To view the report, click **Preview**.

Plot Overlay Sample Selection Fields and Buttons Table

Field	Description
<b>Status</b> <i>[drop-down box]</i>	Select the status of files to be combined.
<b>Look in</b> <i>[button]</i>	Click the <b>Browse</b> icon to change the file folder location.
<b>Available Files</b> <i>[selection]</i>	Lists files that meet the selected criteria. Select the files to be combined, then click <b>Add</b> . The selected files are moved to the <i>Selected Files</i> list box.
<b>Selected Files</b> <i>[selection]</i>	Lists the files selected to be combined. Click <b>OK</b> to combine the files.
 For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .	

## REPORT EXAMPLES

### BJH ADSORPTION REPORT



## HORVATH-KAWAZOE REPORT



Micromeritics Instrument Corporation

Gemini VII 3.04

Gemini VII 2390 V1.00

Serial # 107 Unit 1

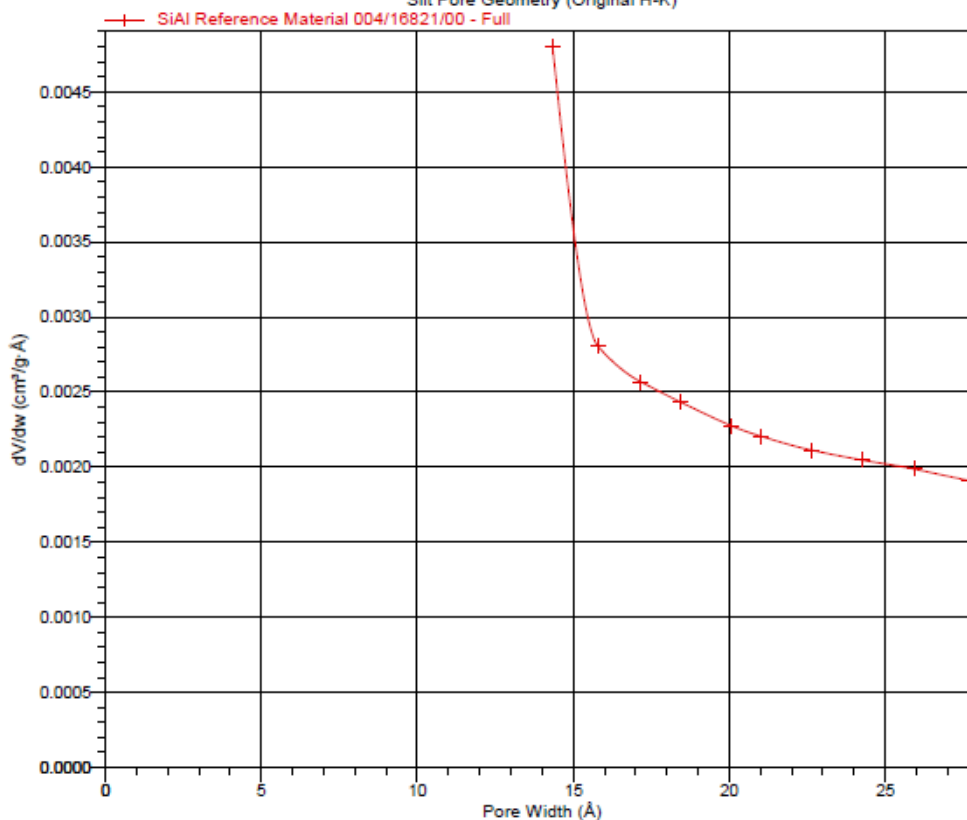
Sample: SiAl Reference Material 004/16821/00 - Full  
Operator: SV  
Submitter: Micromeritics  
File: C:\Gemini VII\data\sialfull.smp

Started: 1/5/2009 9:21:04 AM  
Completed: 1/6/2009 6:50:21 AM  
Report Time: 3/13/2018 7:04:34 AM  
Sample Mass: 0.3054 g  
Free Space Type: Measured  
Evac. Rate: 133.32 kPa/min

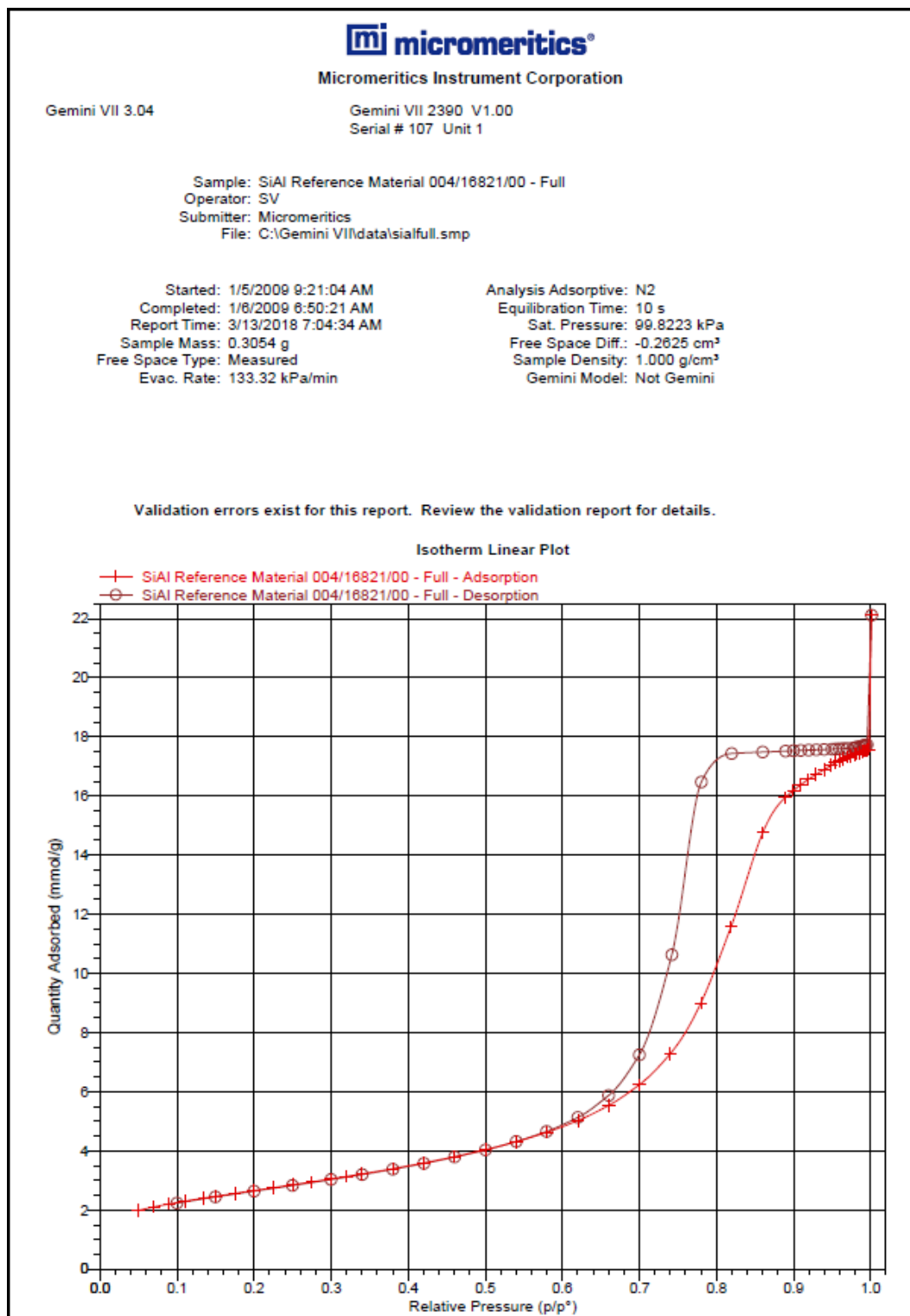
Analysis Adsorptive: N2  
Equilibration Time: 10 s  
Sat. Pressure: 99.8223 kPa  
Free Space Diff.: -0.2625 cm<sup>3</sup>  
Sample Density: 1.000 g/cm<sup>3</sup>  
Gemini Model: Not Gemini

### Horvath-Kawazoe Differential Pore Volume Plot


Slit Pore Geometry (Original H-K)



## ISOTHERM LINEAR PLOT REPORT



**SAMPLE LOG REPORT**

			
Micromeritics Instrument Corporation			
Gemini VII 3.04		Gemini VII 2390 V1.00 Serial # 107 Unit 1	
Sample: SiAl Reference Material 004/16821/00 - Full			
Operator: SV			
Submitter: Micromeritics			
File: C:\Gemini VII\data\sialfull.smp			
Started: 1/5/2009 9:21:04 AM		Analysis Adsorptive: N2	
Completed: 1/6/2009 6:50:21 AM		Equilibration Time: 10 s	
Report Time: 3/13/2018 7:04:34 AM		Sat. Pressure: 99.8223 kPa	
Sample Mass: 0.3054 g		Free Space Diff.: -0.2625 cm <sup>3</sup>	
Free Space Type: Measured		Sample Density: 1.000 g/cm <sup>3</sup>	
Evac. Rate: 133.32 kPa/min		Gemini Model: Not Gemini	
<b>Sample log</b>			
Date	Time	Log Message	
1/5/2009	9:21:04 AM	Analysis Started - Serial #: 107.	
1/6/2009	6:50:21 AM	Analysis Done - Serial #: 107.	

**T-PLOT REPORT**

Micromeritics Instrument Corporation

Gemini VII 3.04

Gemini VII 2390 V1.00

Serial # 107 Unit 1

Sample: SiAl Reference Material 004/16821/00 - Full  
 Operator: SV  
 Submitter: Micromeritics  
 File: C:\Gemini VII\data\sialfull.smp

Started: 1/5/2009 9:21:04 AM      Analysis Adsorptive: N2  
 Completed: 1/6/2009 8:50:21 AM      Equilibration Time: 10 s  
 Report Time: 3/13/2018 7:04:34 AM      Sat. Pressure: 99.8223 kPa  
 Sample Mass: 0.3054 g      Free Space Diff.: -0.2625 cm³  
 Free Space Type: Measured      Sample Density: 1.000 g/cm³  
 Evac. Rate: 133.32 kPa/min      Gemini Model: Not Gemini

**t-Plot Report**

Micropore Volume: 0.002373 cm³/g  
 Micropore Area: 9.0368 m²/g  
 External Surface Area: 205.5550 m²/g  
 Slope: 0.592888 ± 0.001717 mmol/g·Å  
 Y-Intercept: 0.068458 ± 0.007276 mmol/g  
 Correlation Coefficient: 0.999971  
 Surface Area Correction Factor: 1.000  
 Density Conversion Factor: 0.0015468  
 Total Surface Area (BET): 214.5918 m²/g  
 Thickness Range: 3.5000 Å to 5.0000 Å  
 Thickness Equation: Harkins and Jura

**Thickness Curve**


$$t = [ 13.99 / ( 0.034 - \log(p/p^*) ) ]^{0.5}$$

**t-Plot Report - Data**

Relative Pressure (p/p*)	Statistical Thickness (Å)	Quantity Adsorbed (mmol/g)	Fitted
0.050127557	3.2385	1.98507	
0.069963864	3.4300	2.10309	
0.090012579	3.5998	2.20441	*
0.109862177	3.7532	2.29511	*
0.134871644	3.9337	2.40085	*
0.149810730	4.0369	2.46094	*
0.174903377	4.2050	2.55921	*
0.199872020	4.3680	2.65619	*
0.224829880	4.5287	2.75168	*
0.249831907	4.6888	2.84903	*
0.275004731	4.8504	2.94747	*
0.300001339	5.0122	3.04804	
0.320088875	5.1439	3.13155	
0.339990182	5.2763	3.21636	
0.380129171	5.5507	3.39563	
0.420146589	5.8371	3.58895	
0.460280414	6.1408	3.80047	
0.500427370	6.4656	4.03753	
0.540155815	6.8121	4.30652	
0.579603932	7.1867	4.62301	
0.620178490	7.6114	5.02313	
0.660380872	8.0815	5.53708	
0.699950841	8.6051	6.23398	
0.740071237	9.2157	7.28941	



**VALIDATION REPORT**

  
**Micromeritics Instrument Corporation**

Gemini VII 3.04

Gemini VII 2390 V1.00  
Serial # 107 Unit 1

Sample: SiAl Reference Material 004/16821/00 - Full  
Operator: SV  
Submitter: Micromeritics  
File: C:\Gemini VII\data\sialfull.smp

Started: 1/5/2009 9:21:04 AM  
Completed: 1/6/2009 6:50:21 AM  
Report Time: 3/13/2018 7:04:34 AM  
Sample Mass: 0.3054 g  
Free Space Type: Measured  
Evac. Rate: 133.32 kPa/min

Analysis Adsorptive: N2  
Equilibration Time: 10 s  
Sat. Pressure: 99.8223 kPa  
Free Space Diff.: -0.2625 cm<sup>3</sup>  
Sample Density: 1.000 g/cm<sup>3</sup>  
Gemini Model: Not Gemini

**Validation Report**  
**Isotherm Reports**  
Free Space: Low free space values may be  
observed when using liquid argon or ice baths.  
  
p<sup>0</sup> Passed  
  
Pressure/Quantity adsorbed: Passed  
  
Desorption Not within limits. Increase the sample mass, increase  
the equilibration interval, or check the free space.  
  
**BJH Adsorption Reports**  
Pressure range: Passed  
  
Desorption pressures .45 to .40: Phantom 40-angstrom pores may be observed due to the  
properties of nitrogen at relative pressures 0.4 to 0.45.  
  
Minimum Pore Width: Passed  
  
Thickness curve: Passed

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## 8 *SELECTED REPORTS*

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### CFR Note

In 21CFR11 environments, this feature is applicable to members of the Developer group only.

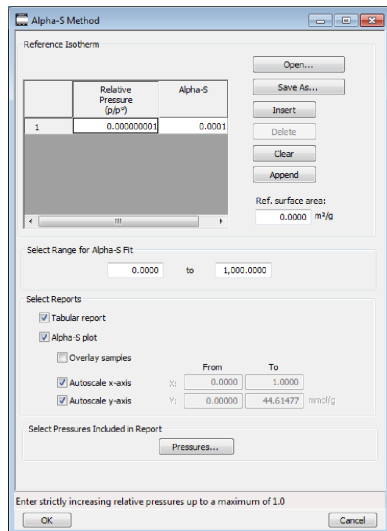


To edit reports, open the *Sample* file then select the *Report Options* tab. Highlight the report name in the *Selected Reports* list box and click **Edit**.

Log in to your [customer portal](#) to access MicroActive Report Tutorials.

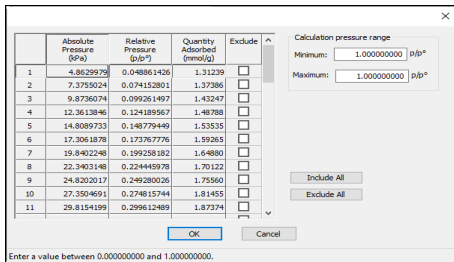
## ALPHA-S METHOD REPORT OPTIONS

The *Alpha-S* plot converts the standard adsorption isotherm into a dimensionless isotherm using the quantity adsorbed at a relative pressure of 0.4.




One predefined curve is located in the *Reference* file directory. Use the table buttons to enter relative pressure and the alpha-s values.

### Alpha-S Method Report Options Fields and Buttons Table

Field or Button	Description
<b>Open</b> [button]	Use to import values from an existing thickness curve (.ALS). The table to be imported must be saved as ASCII text with a .ALS file extension. It must have a two-column format with the relative pressures in the first column and the alpha-s values in the second column. Columns must be separated by a space or a tab.
<b>Pressures</b> [button]	Use to select a pressure range for report calculations and points for exclusion from calculations. <div data-bbox="555 1413 1006 1673" data-label="Image">  </div> <p><b>Calculation pressure range.</b> Enter the minimum and maximum pressures to be used in the pressure table. To exclude a point from the</p>

Alpha-S Method Report Options Fields and Buttons Table (continued)

Field or Button	Description
	<p>calculations used to generate the report, select <i>Exclude</i>.</p> <p><b>Exclude All.</b> Select to exclude all pressure points in the table.</p> <p><b>Include All.</b> Select to include all pressure points in the table.</p>
<b>Ref. surface area</b> [text box]	Enter the surface area from the reference curve. This value is used to calculate the sample surface area.
<b>Select Range for Alpha-S Fit</b> [group box]	Enter minimum and maximum relative pressures to determine the fit.
<b>Selected Reports</b> [group box]	<p><b>Alpha-S Plot.</b> Use to plot data in graph format.</p> <ul style="list-style-type: none"> <li>• <b>Autoscale x-axis.</b> The x-axis field shows the relative pressure.</li> <li>• <b>Autoscale y-axis.</b> The y-axis field shows the quantity of gas adsorbed.</li> <li>• <b>Overlay samples.</b> Use to overlay sample files on the plot.</li> </ul> <p><b>Tabular Report.</b> Use to have a tabular report of data generated.</p>
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	


## BET SURFACE AREA REPORT OPTIONS

The BET calculation obtains the sample surface area value by determining the monolayer volume of adsorbed gas from the isotherm data. BET uses a multilayer model.

### BET Report Options Fields and Buttons Table

Field or Button	Description
<b>Pressures</b> [button]	<p>This option is available when the sample file has a status of <i>Analyzing</i> or <i>Complete</i>. Use to enter a range of pressure points to be included in the report or to modify table values for pressure points.</p> <p><b>Calculation pressure range.</b> Enter the minimum and maximum pressures to be used in the pressure table if not using the <i>Use Interpolation</i> option.</p> <p><b>Exclude All.</b> Select to exclude all pressure points in the table.</p>

BET Report Options Fields and Buttons Table (continued)

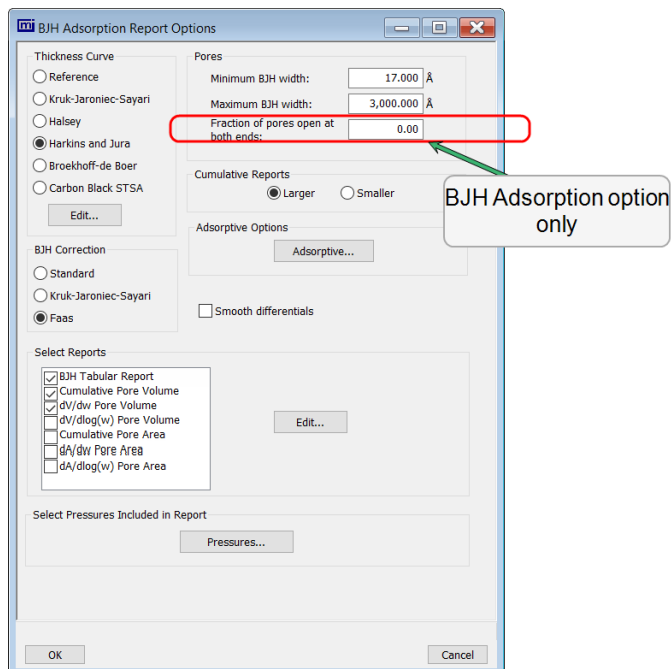
Field or Button	Description
	<p><b>Include All.</b> Select to include all pressure points in the table.</p> <p><b>Insert Predefined.</b> Click to insert a predefined (default) set of points into the report. <i>Use Interpolation</i> must be selected to enable this button. This button displays for BET reports only.</p> <p><b>Use Interpolation.</b> Use to indicate if the system should use the table or interpolated data. This option is available for BET and Langmuir reports only.</p>
<b>Select Pressure Range for BET fit</b> [text box]	Enter values to indicate the fitted pressure range.
<b>Selected Reports</b> [group box]	<p><b>BET Isotherm plot.</b> Uses BET monolayer volume and constant to produce an isotherm.</p> <ul style="list-style-type: none"> <li>• <b>Autoscale x-axis.</b> Linear x-axes begin at zero. The x-axis field shows the relative pressure for BET.</li> <li>• <b>Autoscale y-axis.</b> The y-axis field shows the quantity of gas adsorbed.</li> <li>• <b>Overlay samples.</b> Use to overlay sample files on the BET isotherm plot.</li> </ul> <p><b>BET Transform plot.</b> Use to generate a traditional BET surface area plot used to determine monolayer volume and BET C constant.</p> <ul style="list-style-type: none"> <li>• <b>Autoscale x-axis.</b> Linear x-axes begin at zero. The x-axis field shows the relative pressure for BET.</li> <li>• <b>Autoscale y-axis.</b> The y-axis field shows BET transformation.</li> <li>• <b>Overlay samples.</b> Use to overlay sample files on the BET transform plot.</li> </ul> <p><b>Tabular report.</b> Use to have a table of measured and calculated values generated.</p>
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	

## BJH ADSORPTION / DESORPTION REPORT OPTIONS

The BJH calculation determines the mesopore volume / area distribution, which accounts for both the change in adsorbate layer thickness and the liquid condensed in pore cores. Reports can be generated from both adsorption and desorption data. The fields for both *BJH Adsorption Report Options* and *BJH Desorption Report Options* are identical unless otherwise specified.



An incomplete pore distribution may be generated if a thickness curve selection is not a good match for the sample being analyzed.

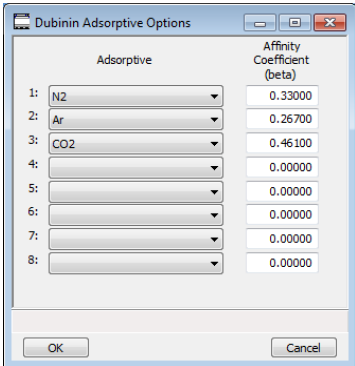


### BJH Adsorption / Desorption Report Options Fields and Buttons Table

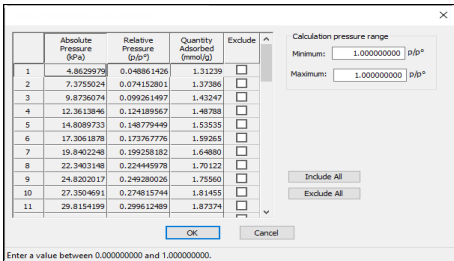
Field or Button	Description
<b>Adsorptive</b> [button]	Displays the <i>Adsorptive Options</i> window. The recommended adsorptives and their values are shown. Up to eight adsorptive and adsorbate property factor combinations may be specified.



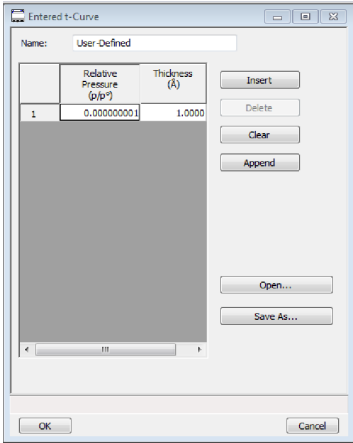

BJH Adsorption / Desorption Report Options Fields and Buttons Table (continued)

Field or Button	Description
	
<b>BJH Correction</b> <i>[group box]</i>	<p>Select the type of correction to apply to calculations. The selected type will display in the report header.</p> <p><b>Faas.</b> Good for statistical thickness curves.</p> <p><b>Kruk-Jaroniec-Sayari.</b> Good for reference thickness curves.</p> <p><b>Standard.</b> Uses original BJH models.</p>
<b>Cumulative Reports</b> <i>[group box]</i>	<p><b>Larger.</b> Use to report the total volume found in pores larger than the current pore size.</p> <p><b>Smaller.</b> Use to report the total volume found in pores smaller than the current pore size.</p>
<b>Pores</b> <i>[group box]</i>	<p>Enter the minimum and maximum diameter (radius or width) of pores to include in the BJH reports.</p> <p><b>Fraction of pores open at both ends.</b> This field is not available for the <i>BJH Desorption Report Options</i> window.</p> <p>During adsorption calculations, the analysis program assumes that all pores are closed at one end. Occasionally, a percentage of pores may be open at both ends causing disagreement in the adsorption and desorption data or in the values for total volume and total BJH pore volume. Enter the fraction of pores open at both ends to compensate for this error.</p>

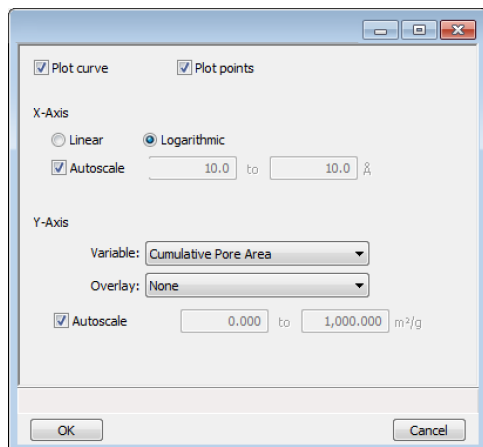
## BJH Adsorption / Desorption Report Options Fields and Buttons Table (continued)

Field or Button	Description
<b>Pressures</b> [button]	<p>Use to select a pressure range for report calculations and points for exclusion from calculations.</p>  <p><b>Calculation pressure range.</b> Enter the minimum and maximum pressures to be used in the pressure table. To exclude a point from the calculations used to generate the report, select <i>Exclude</i>.</p> <p><b>Exclude All.</b> Select to exclude all pressure points in the table.</p> <p><b>Include All.</b> Select to include all pressure points in the table.</p>
<b>Select Reports</b> [group box]	Select the report names to include in the report. Highlight the report name, then click <b>Edit</b> to modify report parameters.
<b>Smooth differentials</b> [check box]	Use to smooth all differential calculations, thus eliminating variations in the differential computation caused by noise in the input data.
<b>Thickness Curve</b> [group box]	<p>Select the thickness curve, then click <b>Edit</b> to modify the values in the equation for the selected curve. The Frenkel-Halsey-Hill thickness curve can be applied using the Halsey option.</p> <p><b>Kruk-Jaroniec-Sayari / Halsey / Harkins and Jura / Broekhoff-de Boer / Carbon Black STSA.</b> Select the thickness curve option, then click <b>Edit</b>. Modify the equation for the selected curve as needed.</p> <p><b>Reference.</b> Select <i>Reference</i>, then click <b>Edit</b> to define a t-curve by entering both the relative pressure and thickness values. One predefined curve is shipped with the analysis program and is found in the <i>Reference</i> directory.</p>

## BJH Adsorption / Desorption Report Options Fields and Buttons Table (continued)


Field or Button	Description
	 <p>To import values from an existing thickness curve (.THK file), click <a href="#">Open</a>, then select the file containing the values. The table to be imported must have a .TXT or .THK file extension and have a two-column format with the relative pressures in the first column and the thickness values in the second column. Columns must be separated by a space or a tab.</p>
	For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .

## BJH PLOT OPTIONS

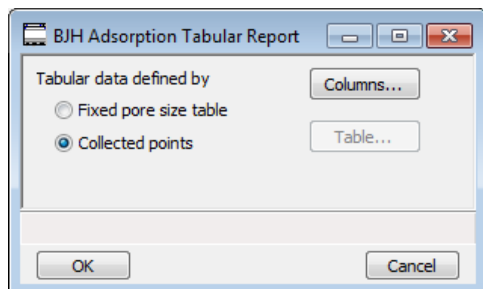


The fields for all plot options are identical for specifying plotting methods and customizing plots. Highlight any plot option in the *Selected Reports* list box in the *BJH Report Options* window, then click [Edit](#).

### BJH Plot Options Fields and Buttons Table

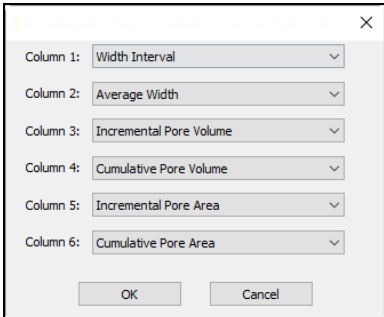

Field or Button	Description
<b>Autoscale</b> [check box]	When enabled on the report parameters windows, allows the x- and y- axes to be scaled automatically. <i>Autoscale</i> means that the x- and y- ranges will be set so that all the data is shown. If <i>Autoscale</i> is not selected, the entered range is used.
<b>Plot curve / Plot points</b> [check box]	Select to plot points on the graph.
<b>X-Axis</b> [group box]	Use to have the x-axis on a logarithmic or linear scale.
<b>Y-Axis</b> [group box]	<b>Overlay.</b> Select an option to overlay onto the current report. <b>Variable.</b> Select a variable.
 For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .	

## BJH TABULAR REPORT OPTIONS



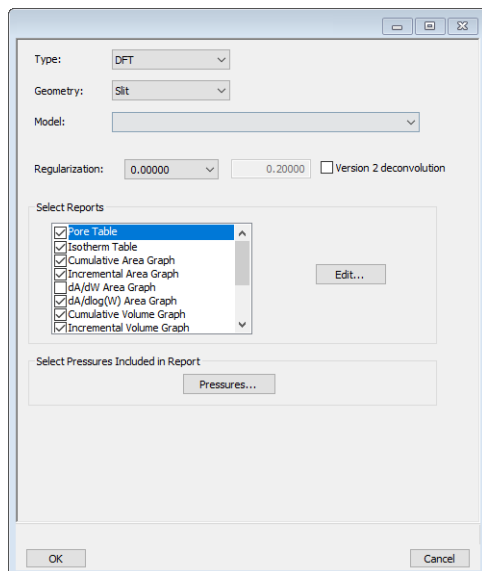
Highlight *BJH Tabular Report* in the *Selected Reports* list box on the *BJH Adsorption Report Options* window, then click **Edit** to specify the method of data reduction.

### BJH Tabular Report Options Fields and Buttons Table

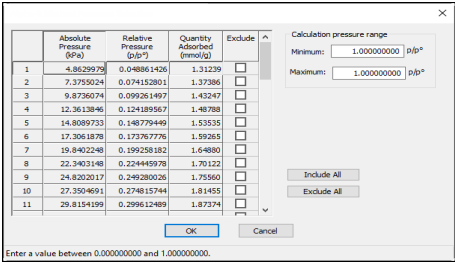
Field or Button	Description
<b>Collected points</b> [selection]	Use to include all relative pressure points collected. Refer to the <b>Columns</b> button below.
<b>Columns</b> [button]	Select the data types to include in the report. <i>Column [n]</i> indicates the column order and data contents for the report. 
<b>Fixed pore size table</b> [selection]	Use to specify exact pore sizes for volume or area data. Click <b>Table</b> to modify the fixed pore size table. Refer to <b>Table</b> and <b>Columns</b> buttons elsewhere in this table.
<b>Table</b> [button]	The fixed pore size table must contain a minimum of two points. The points must be strictly decreasing. Enabled only when <i>Fixed pore size table</i> is selected.
 <b>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</b>	

## DFT PORE SIZE REPORT OPTIONS

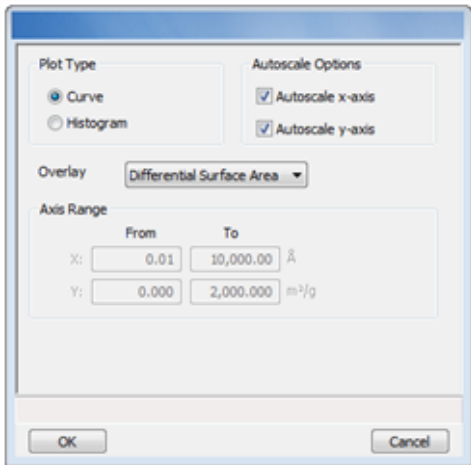
The *DFT Pore Size* report contains the results of pore size distribution analyses using a non-local DFT range of micro and mesopore ranges.




DFT Pore Size Report Options Fields and Buttons Table

Field or Button	Description
<b>Geometry</b> [drop-down box]	Select the pore shape.
<b>Model</b> [drop-down box]	Lists the models that meet the specified criteria and match the adsorbate and temperature of the sample data. If no models appear, no models meet the selected criteria. One model must be selected.
<b>Pressures</b> [button]	Use to select a pressure range for report calculations and points for exclusion from calculations.
	 <p><b>Calculation pressure range.</b> Enter the minimum and maximum pressures to be used in the pressure table. To exclude a point from the calculations used to generate the report, select <i>Exclude</i>.</p>

DFT Pore Size Report Options Fields and Buttons Table (continued)

Field or Button	Description
	<p><b>Exclude All.</b> Select to exclude all pressure points in the table.</p> <p><b>Include All.</b> Select to include all pressure points in the table.</p>
<b>Regularization</b> <i>[drop-down box]</i>	<p>Select the extent of smoothing to apply to the data. If <i>0.20000 (user)</i> is selected, enter a number in the text box giving a relative weight for the smoothing during deconvolution. Larger values produce more smoothing.</p>
<b>Select Reports</b> <i>[group box]</i>	<p>Select the reports to generate. To edit graph details, highlight the graph option and click <b>Edit</b>. The <i>Log Goodness of Fit</i> and <i>Goodness of Fit</i> graphs cannot be edited.</p>  <p><b>Autoscale Options.</b> Use to autoscale the x-axis and/or y-axes.</p> <p><b>Axis Range.</b> <i>From / To</i> fields are enabled when <i>Autoscale</i> options are not selected. Enter the starting and ending values for the x- and/or y-axes.</p> <ul style="list-style-type: none"> <li>• <b>x-axis.</b> Shows the pore size.</li> <li>• <b>y-axis.</b> Shows the area.</li> </ul> <p><b>Overlay.</b> Select an overlay for the report.</p> <p><b>Plot Type.</b> Select the method for data display.</p>

DFT Pore Size Report Options Fields and Buttons Table (continued)

Field or Button	Description
<b>Type</b> [drop-down box]	<b>Classical.</b> Model based on the Kelvin equation and thickness for determining the pore size distribution. See <a href="#">DFT Models on page B - 1</a> .  <b>DFT.</b> Model based on the density functional theory.
<b>Version 2 deconvolution</b> [check box]	If enabled, adds improved performance for nonzero values of the DFT regularization. The DFT results will be slightly better than when the checkbox is not selected and when using a regularization parameter. When the checkbox is not selected, the legacy behavior of the DFT calculation used by earlier versions of the application is produced
 For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .	

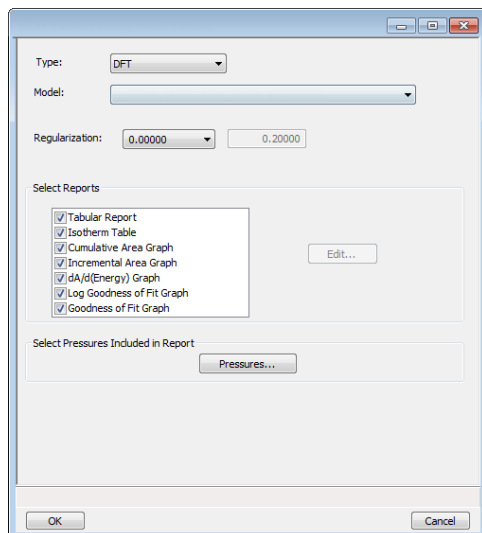
## DFT SURFACE ENERGY REPORT OPTIONS



See also:

[DFT Pore Size Report Options on page 8 - 12](#)

The *DFT Surface Energy* report contains the results of surface energy distribution analyses.



*DFT Surface Energy Report Options* fields and buttons are identical to the *DFT Pore Size Report Options*.



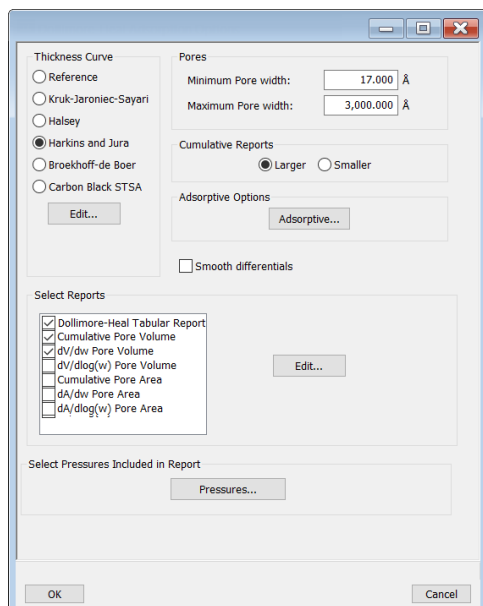
## DOLLIMORE-HEAL ADSORPTION / DESORPTION REPORT OPTIONS



**See also:**

[BJH Adsorption / Desorption Report Options on page 8 - 6](#) for additional information on field and buttons for this report.

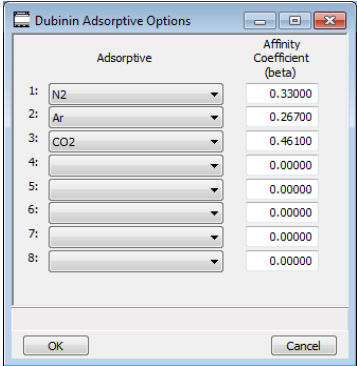
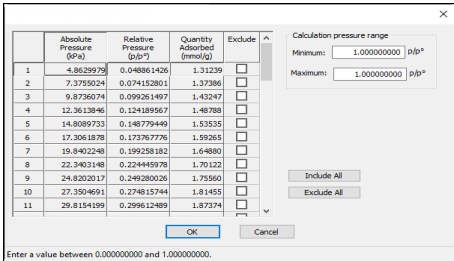
The *Dollimore-Heal Adsorption Report Option* and the *Dollimore-Heal Desorption Report Option* generate reports from both adsorption and desorption data.



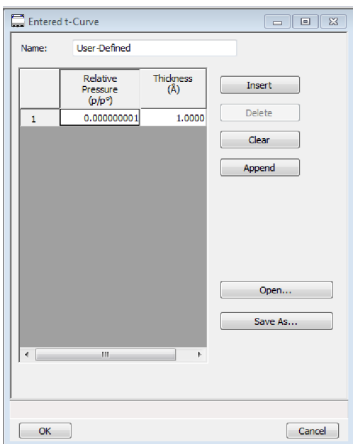

### BJH Adsorption / Desorption Report Options Fields and Buttons Table

Field or Button	Description
<b>Adsorptive</b> [button]	Displays the <i>Adsorptive Options</i> window. The recommended adsorptives and their values are shown. Up to eight adsorptive and adsorbate property factor combinations may be specified.

## BJH Adsorption / Desorption Report Options Fields and Buttons Table (continued)

Field or Button	Description
	
<b>Cumulative Reports</b> <i>[group box]</i>	<p><b>Larger.</b> Use to report the total volume found in pores larger than the current pore size.</p> <p><b>Smaller.</b> Use to report the total volume found in pores smaller than the current pore size.</p>
<b>Pores</b> <i>[group box]</i>	Enter the minimum and maximum diameter (radius or width) of pores to include in the BJH reports.
<b>Pressures</b> <i>[button]</i>	<p>Use to select a pressure range for report calculations and points for exclusion from calculations.</p>  <p><b>Calculation pressure range.</b> Enter the minimum and maximum pressures to be used in the pressure table. To exclude a point from the calculations used to generate the report, select <i>Exclude</i>.</p> <p><b>Exclude All.</b> Select to exclude all pressure points in the table.</p> <p><b>Include All.</b> Select to include all pressure points in the table.</p>
<b>Select Reports</b> <i>[group box]</i>	Select the report names to include in the report. Highlight the report name, then click <b>Edit</b> to modify report parameters.
<b>Smooth differentials</b> <i>[check box]</i>	Use to smooth all differential calculations, thus eliminating variations in the differential computation caused by noise in the input data.
<b>Thickness Curve</b> <i>[group box]</i>	Select the thickness curve, then click <b>Edit</b> to modify the values in the equation for the selected curve. The Frenkel-Halsey-Hill thickness

## BJH Adsorption / Desorption Report Options Fields and Buttons Table (continued)

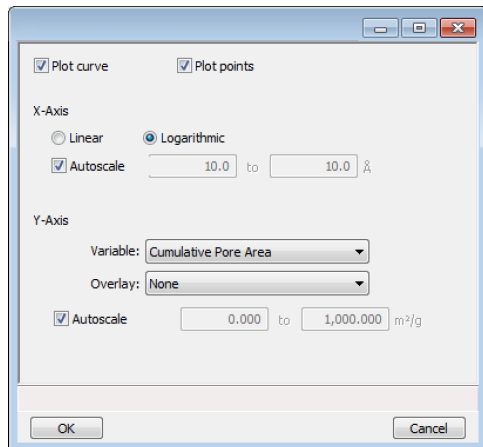
Field or Button	Description
	<p>curve can be applied using the Halsey option.</p> <p><b>Kruk-Jaroniec-Sayari / Halsey / Harkins and Jura / Broekhoff-de Boer / Carbon Black STSA.</b> Select the thickness curve option, then click <b>Edit</b>. Modify the equation for the selected curve as needed.</p> <p><b>Reference.</b> Select <i>Reference</i>, then click <b>Edit</b> to define a t-curve by entering both the relative pressure and thickness values. One predefined curve is shipped with the analysis program and is found in the <i>Reference</i> directory.</p>  <p>To import values from an existing thickness curve (.THK file), click <b>Open</b>, then select the file containing the values. The table to be imported must have a .TXT or .THK file extension and have a two-column format with the relative pressures in the first column and the thickness values in the second column. Columns must be separated by a space or a tab.</p>
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	

## DOLLIMORE-HEAL PLOT OPTIONS



See also:

[BJH Plot Options on page 8 - 10](#)



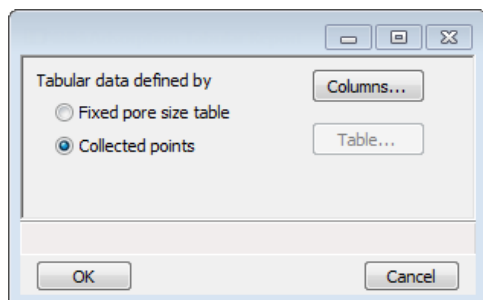
The fields for all plot options are identical for specifying plotting methods and customizing plots. Highlight any plot option in the *Selected Reports* list box in the *Dollimore-Heal Report Options* window, then click **Edit**. The fields and buttons for these reports are identical to the *BJH Plot Report Options*.

## ***DOLLIMORE-HEAL TABULAR REPORT OPTIONS***



**See also:**

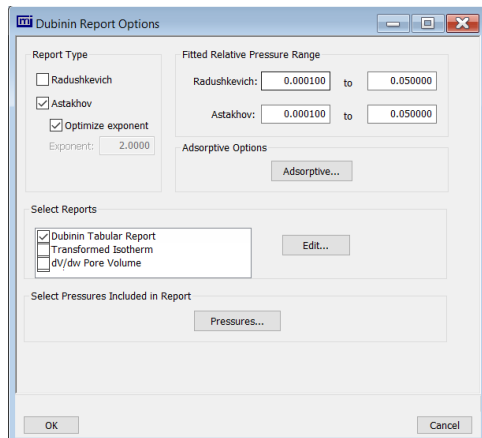
[\*BJH Tabular Report Options on page 8 - 11\*](#)



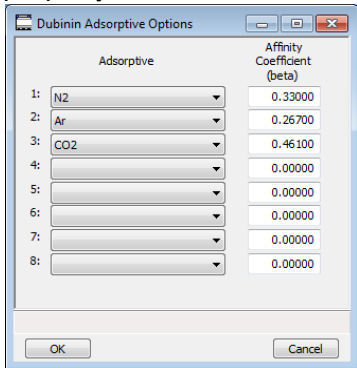
*Dollimore-Heal Tabular Report Options are identical to the BJH Tabular Report Options.*

## DUBININ REPORT OPTIONS

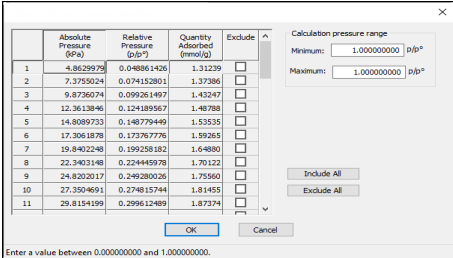

The *Dubinin* method provides pore volume distributions for microporous materials by making use of an expression for the adsorption potential.



Dubinin Report Options Fields and Buttons Table

Field or Button	Description																		
<b>Adsorptive</b> [button]	Displays the <i>Adsorptive Options</i> window. The recommended adsorptives and their values are shown. Up to eight adsorptive and adsorbate property factor combinations may be specified. <div data-bbox="555 1073 909 1438" data-label="Form">  <table border="1"> <thead> <tr> <th>Adsorptive</th><th>Affinity Coefficient (beta)</th></tr> </thead> <tbody> <tr><td>1: N2</td><td>0.33000</td></tr> <tr><td>2: Ar</td><td>0.26700</td></tr> <tr><td>3: CO2</td><td>0.46100</td></tr> <tr><td>4:</td><td>0.00000</td></tr> <tr><td>5:</td><td>0.00000</td></tr> <tr><td>6:</td><td>0.00000</td></tr> <tr><td>7:</td><td>0.00000</td></tr> <tr><td>8:</td><td>0.00000</td></tr> </tbody> </table> </div>	Adsorptive	Affinity Coefficient (beta)	1: N2	0.33000	2: Ar	0.26700	3: CO2	0.46100	4:	0.00000	5:	0.00000	6:	0.00000	7:	0.00000	8:	0.00000
Adsorptive	Affinity Coefficient (beta)																		
1: N2	0.33000																		
2: Ar	0.26700																		
3: CO2	0.46100																		
4:	0.00000																		
5:	0.00000																		
6:	0.00000																		
7:	0.00000																		
8:	0.00000																		
<b>Fitted Relative Pressure Range</b> [group box]	Enter the minimum and maximum limits for Radushkevich or Astakhov relative pressures included in the line fit.																		

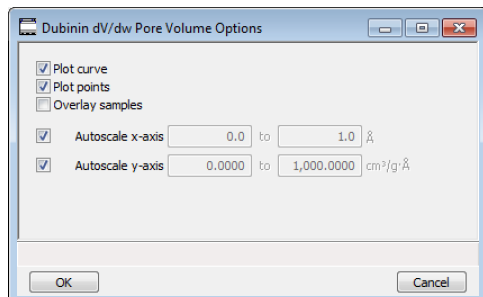
## Dubinin Report Options Fields and Buttons Table (continued)

Field or Button	Description
<b>Pressures</b> [button]	<p>Use to select a pressure range for report calculations and points for exclusion from calculations.</p>  <p><b>Calculation pressure range.</b> Enter the minimum and maximum pressures to be used in the pressure table. To exclude a point from the calculations used to generate the report, select <i>Exclude</i>.</p> <p><b>Exclude All.</b> Select to exclude all pressure points in the table.</p> <p><b>Include All.</b> Select to include all pressure points in the table.</p>
<b>Report Type</b> [group box]	Select report types. If <i>Astakhov</i> is selected, either select <i>Optimize exponent</i> or enter an appropriate exponent value in the text box.
<b>Select Reports</b> [group box]	Select the reports to generate. Highlight the report, then click <b>Edit</b> to modify report options.
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	


## DUBININ PORE VOLUME REPORT OPTIONS

In the *Dubin Report Options* window, highlight *dV/dw Pore Volume* in the *Selected Reports* list box, then click [Edit](#).

This option plots differential pore volume as a function of pore width.



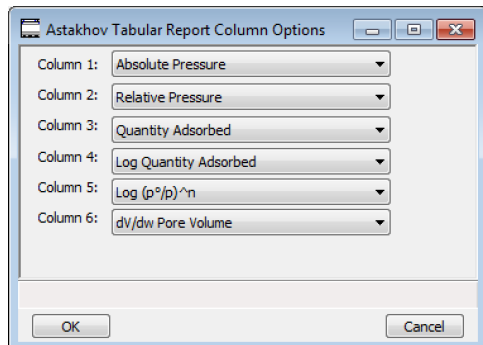
### Dubin Pore Volume Report Fields and Buttons Table

Field or Button	Description
<b>Autoscale x-axis / Autoscale y-axis</b> [check box]	Select an option to have the x- and/or y-axes scaled automatically. Both axes begin at 0; the system uses the highest values collected during analysis as the ending points for axis ranges.  Enable to enter beginning and ending values manually.
<b>Overlay samples</b> [selection]	Use to overlay sample files on the plot.
<b>Plot curve / Plot points</b> [selection]	Select to plot points on the graph.
 For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .	



## DUBININ TABULAR REPORT OPTIONS

In the *Dubinin Report Options* window, highlight *Dubinin Tabular Report* in the *Selected Reports* list box, then click **Edit**. *Column [n]* indicates the column order and data contents for the report.

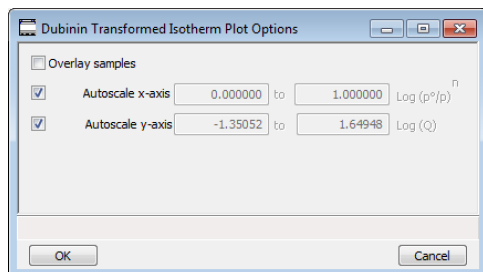


**Log (p°/p )^n.** The value for *[n]* is the optimized exponent if *Optimize exponent* is selected on the *Dubinin Report Options* window. If not, then the value for *[n]* is the entered exponent value.


## DUBININ TRANSFORMED ISOTHERM PLOT OPTIONS

Highlight *Transformed Isotherm* in the *Selected Reports* list box in the *Dubinin Report Options* window, then click **Edit**.

The transformed Dubinin isotherm is the logarithm of quantity adsorbed as a function of the log of relative pressure raised to a power. Isotherms for which the Dubinin method is applicable produce straight lines when transformed in this way.

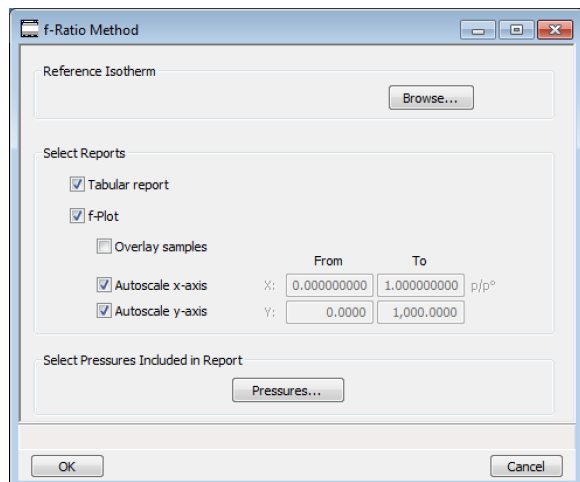


Dubinin Transformed Isotherm Plot Options Fields and Buttons Table

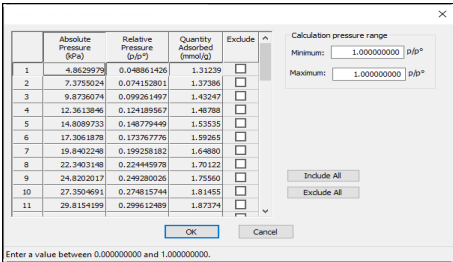
Field or Button	Description
<b>Autoscale x-axis / Autoscale y-axis</b> [check box]	<p>Select an option to have the x- and/or y-axes scaled automatically. Both axes begin at 0; the system uses the highest values collected during analysis as the ending points for axis ranges.</p> <p>Deselect to enter beginning and ending values manually.</p> <p><b>Autoscale x-axis.</b> Shows the quantity of gas adsorbed at standard temperature and pressure.</p> <p><b>Autoscale y-axis.</b> Shows the log of relative pressure.</p>
<b>Overlay Samples</b> [check box]	Use to overlay sample files on the plot.
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	

## F-RATIO METHOD REPORT OPTIONS


The *f*-Ratio report uses the measured isotherm and normalizes it using a reference isotherm.



### f-Ratio Fields and Buttons Table

Field or Button	Description
<b>Pressures</b> [button]	<p>Use to select a pressure range for report calculations and points for exclusion from calculations.</p>  <p><b>Calculation pressure range.</b> Enter the minimum and maximum pressures to be used in the pressure table. To exclude a point from the calculations used to generate the report, select <i>Exclude</i>.</p> <p><b>Exclude All.</b> Select to exclude all pressure points in the table.</p> <p><b>Include All.</b> Select to include all pressure points in the table.</p>

## f-Ratio Fields and Buttons Table (continued)

Field or Button	Description
<b>Reference isotherm</b> <i>[group box]</i>	Browse to select a sample file to use as a reference for the isotherm. Select a file containing an isotherm measured from a non-porous sample of the same material as the current sample.
<b>Selected Reports</b> <i>[group box]</i>	<p><b>Tabular Report.</b> Use to have a tabular report of data generated.</p> <p><b>f-Plot.</b> Use to generate a normalized isotherm.</p> <ul style="list-style-type: none"> <li>• <b>Autoscale x-axis.</b> The x-axis field is dimensionless in units of f-ratio.</li> <li>• <b>Autoscale y-axis.</b> The y-axis field shows the quantity of gas adsorbed.</li> <li>• <b>Overlay samples.</b> Use to overlay sample files on the f-plot.</li> </ul>
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	


## FREUNDLICH REPORT OPTIONS

The *Freundlich Isotherm* is an empirical isotherm used to model low-pressure adsorption data. It can also be applied to model some micropore isotherms. In the *Selected Reports* list box, highlight *Freundlich*, then click **Edit**.

### Freundlich Report Options Fields and Buttons Table

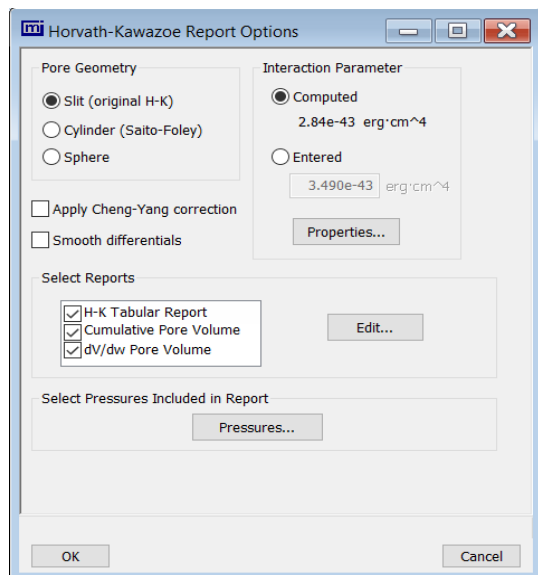
Field or Button	Description
<b>Pressures</b> [button]	<p>Use to select a pressure range for report calculations and points for exclusion from calculations.</p> <p><b>Calculation pressure range.</b> Enter the minimum and maximum pressures to be used in the pressure table. To exclude a point from the calculations used to generate the report, select <i>Exclude</i>.</p> <p><b>Exclude All.</b> Select to exclude all pressure points in the table.</p> <p><b>Include All.</b> Select to include all pressure points in the table.</p>

Freundlich Report Options Fields and Buttons Table (continued)

Field or Button	Description
<b>Select Reports</b> <i>[group box]</i>	<p><b>Freundlich Isotherm plot.</b> Plots the absolute pressure vs quantity adsorbed. Shows best fit line.</p> <ul style="list-style-type: none"> <li>• <b>Autoscale x-axis.</b> Linear x-axes begin at zero. The x-axis field shows the absolute pressure.</li> <li>• <b>Autoscale y-axis.</b> Y-axes begin at zero. The y-axis field shows the quantity of gas adsorbed.</li> <li>• <b>Overlay samples.</b> Use to overlay sample files on the Freundlich isotherm plot.</li> </ul> <p><b>Freundlich Transform plot.</b> Plots the <math>\log(P)</math> vs <math>\log(Q)</math> and the best fit.</p> <ul style="list-style-type: none"> <li>• <b>Autoscale x-axis.</b> The x-axis field shows the absolute pressure.</li> <li>• <b>Autoscale y-axis.</b> The y-axis field shows the quantity of gas adsorbed.</li> <li>• <b>Overlay samples.</b> Use to overlay sample files on the Freundlich transform plot.</li> </ul> <p><b>Tabular report.</b> Select to include pressure points included in the report.</p>
<b>Specify monolayer capacity</b> <i>[selection]</i>	Select and enter the monolayer capacity of the sample.
<b>Tabular report</b> <i>[selection]</i>	Use to have a report of the pressure points generated.
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	

## HORVATH-KAWAZOE REPORT OPTIONS

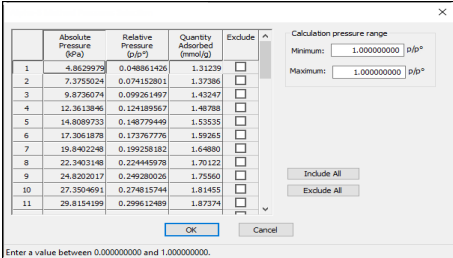
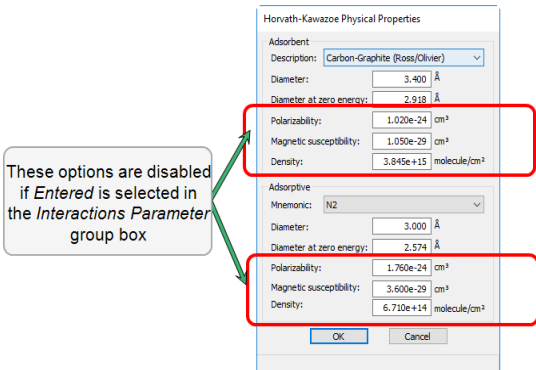
The *Horvath-Kawazoe* method plots individual peaks for different pore sizes even if the difference between one pore size and the next is only one angstrom (0.10 nm) or less.



### Horvath-Kawazoe Report Fields and Buttons Table


Field or Button	Description
<b>Apply Cheng-Yang correction</b> [selection]	Use to apply the Cheng-Yang correction to the pore size analysis. This correction substitutes the Langmuir equation of state for Henry's Law in the Horvath-Kawazoe derivation.
<b>Interaction Parameter</b> [group box]	<p>Use to determine which interaction parameter will be used in the report. These options are disabled if <i>Sphere</i> is selected in the <i>Pore Geometry</i> group box.</p> <p><b>Computed.</b> Use to calculate using the parameters on the <i>Horvath-Kawazoe Physical Properties</i> window (click <b>Properties</b> to display the <i>Physical Properties</i> window). The interaction parameter is recalculated each time a parameter in the <i>Physical Properties</i> window is edited.</p> <p><b>Entered.</b> Calculates using the value entered in the text box.</p>
<b>Pore Geometry</b> [group box]	Select the option that best represents the physical geometry of the micropores in the sample material. When <i>Sphere</i> is selected, options in the <i>Interaction Parameter</i> group box are disabled.

## Horvath-Kawazoe Report Fields and Buttons Table (continued)

Field or Button	Description
<b>Pressures</b> [button]	<p>Use to select a pressure range for report calculations and points for exclusion from calculations.</p>  <p><b>Calculation pressure range.</b> Enter the minimum and maximum pressures to be used in the pressure table. To exclude a point from the calculations used to generate the report, select <i>Exclude</i>.</p> <p><b>Exclude All.</b> Select to exclude all pressure points in the table.</p> <p><b>Include All.</b> Select to include all pressure points in the table.</p>
<b>Properties</b> [button]	<p>Click to view or edit the constants describing the physical properties of the adsorbent and adsorptive.</p>  <p><b>Adsorbent.</b> Contains the parameters for the sample. If using <i>Computed</i> for the interaction parameter, all fields are enabled. If using <i>Entered</i>, only the values in the <i>Diameter</i> and <i>Diameter at zero energy</i> text fields may be edited.</p> <ul style="list-style-type: none"> <li>• <b>Density.</b> Enter the density per unit area of the sample.</li> <li>• <b>Description.</b> Select the name of the sample used in the analysis.</li> <li>• <b>Diameter.</b> Enter the diameter of the sample atom.</li> <li>• <b>Diameter at zero energy.</b> Enter the diameter of an atom at zero interaction energy: <math>(2/5)^{1/6} \times \text{diameter}</math>.</li> </ul>



Horvath-Kawazoe Report Fields and Buttons Table (continued)

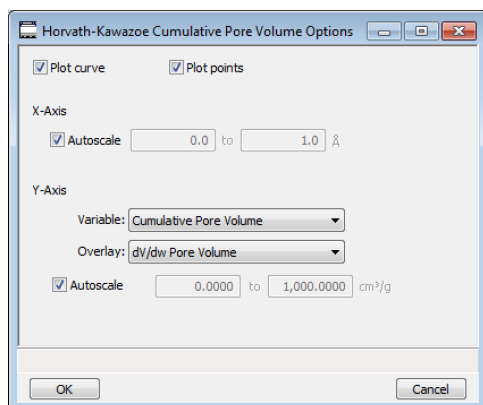
Field or Button	Description
	<ul style="list-style-type: none"> <li>• <b>Magnetic susceptibility.</b> Enter the magnetic susceptibility of the sample.</li> <li>• <b>Polarizability.</b> Enter the polarizability of the sample.</li> </ul> <p><b>Adsorptive.</b> Contains the parameters for the adsorptives. If using <i>Computed</i> for the interaction parameter, all fields are enabled. If using <i>Entered</i>, only the values in the <i>Diameter</i> and <i>Diameter at zero energy</i> text fields may be edited.</p> <ul style="list-style-type: none"> <li>• <b>Density.</b> Enter the density per unit area of the adsorptive.</li> <li>• <b>Diameter.</b> Enter the diameter of the gas phase atom.</li> <li>• <b>Diameter at zero energy.</b> Enter the diameter of an atom at zero interaction energy: <math>(2/5)^{1/6} \times \text{diameter}</math>.</li> <li>• <b>Magnetic susceptibility.</b> Enter the magnetic susceptibility of the adsorptive.</li> <li>• <b>Mnemonic.</b> Select the mnemonic of the adsorptive gas in use.</li> <li>• <b>Polarizability.</b> Enter the polarizability of the adsorptive.</li> </ul>
<b>Select Reports</b> [group box]	Select the types of reports to generate. Highlight the report, then click <b>Edit</b> to modify report parameters.
<b>Smooth Differentials</b> [selection]	Use to smooth all differential calculations, thus eliminating variations in the differential computation caused by noise in the input data.
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	

## HORVATH-KAWAZOE PLOT OPTIONS



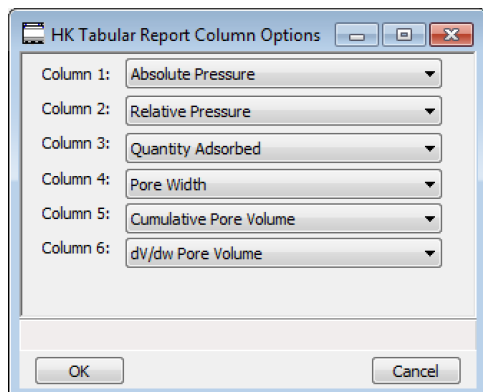
**See also:**

[BJH Plot Options on page 8 - 10](#) for additional information on fields and buttons for this report.



Highlight a plot option in the *Selected Reports* list box in the *Horvath-Kawazoe Report Options* window, then click **Edit** to customize the plotting method.

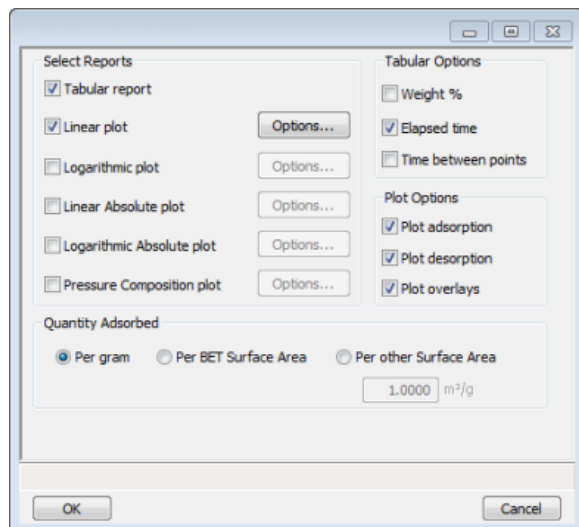
## HORVATH-KAWAZOE TABULAR REPORT OPTIONS



Highlight *H-K Tabular Report* in the *Selected Reports* list box in the *Horvath-Kawazoe Report Options* window, then click **Edit**. Select the data types to include in the report. *Column [n]* indicates the column order and data contents for the report.

## ISOTHERM REPORT OPTIONS


The *Isotherm* report indicates adsorption (up to saturation pressure) and desorption (down from saturation pressure) of a gas by a solid held at constant temperature.



### Isotherm Report Options Fields and Buttons Table

Field or Button	Description
<b>Options</b> [button]	Click to display related linear plot options. All plot windows contain identical fields.  <b>Autoscale x-axis.</b> Linear x-axes begin at zero. Logarithmic x-axes begin at an appropriate value. The x-axis field shows the relative or absolute pressure.  <b>Autoscale y-axis.</b> The y-axis field shows the quantity of gas adsorbed.  <b>Plot curve / Plot points.</b> Select to plot points on the graph.
<b>Plot Options</b> [group box]	Select the types of isotherm to plot.
<b>Quantity Adsorbed</b> [group box]	Select how to report the quantity adsorbed.  <ul style="list-style-type: none"> <li>per gram (cm<sup>3</sup>/g) STP</li> <li>per BET Surface Area (cm<sup>3</sup>/m<sup>2</sup>) STP or mmol/g</li> <li>per other Surface Area (cm<sup>3</sup>/m<sup>2</sup>) STP or mmol/m<sup>2</sup></li> </ul>
<b>Selected Reports</b>	Select each option to include on the final report. Click the Options button of a selected item to include plot curve, plot points, and to

Isotherm Report Options Fields and Buttons Table (continued)

Field or Button	Description
[group box]	autoscale x- and y-axes.
<b>Tabular Options</b> [group box]	Select the options to include on the report.  <b>Elapsed time.</b> Time elapsed during the analysis.  <b>Time between points.</b> Time elapsed between points during the analysis.  <b>Weight %.</b> Enter the mass percentage when plotting pressure composition.
	For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .


## LANGMUIR REPORT OPTIONS

The Langmuir calculation determines the surface area of a sample by relating the surface area to the volume of gas adsorbed as a monolayer. Langmuir uses a single layer model.

### Langmuir Report Options Fields and Buttons Table

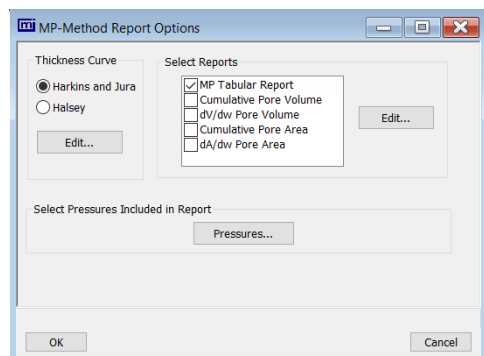
Field or Button	Description
<b>Pressures</b> [button]	<p>This option is available when the sample file has a status of <i>Analyzing</i> or <i>Complete</i>. Use to enter a range of pressure points to be included in the report or to modify table values for pressure points.</p> <p><b>Calculation pressure range.</b> Enter the minimum and</p>

Langmuir Report Options Fields and Buttons Table (continued)

Field or Button	Description
	<p>maximum pressures to be used in the pressure table. To exclude a point from the calculations used to generate the report, select <i>Exclude</i>.</p> <p><b>Exclude All.</b> Select to exclude all pressure points in the table.</p> <p><b>Include All.</b> Select to include all pressure points in the table.</p> <p><b>Use Interpolation.</b> Use to indicate if the system should use the table or interpolated data. This option is available for BET and Langmuir reports only.</p>
<b>Select Pressure Range for Langmuir fit</b> <i>[group box]</i>	Enter values to indicate the fitted pressure range.
<b>Selected Reports</b> <i>[group box]</i>	<p><b>Langmuir Isotherm Plot.</b> Uses the Langmuir monolayer volume and constant to produce an isotherm.</p> <ul style="list-style-type: none"> <li>• <b>Autoscale x-axis.</b> Linear x-axes begin at zero. The x-axis field shows the absolute pressure for Langmuir.</li> <li>• <b>Autoscale y-axis.</b> The y-axis field shows the quantity of gas adsorbed.</li> <li>• <b>Overlay samples.</b> Use to overlay sample files on the Langmuir isotherm plot.</li> </ul> <p><b>Langmuir Transform Plot.</b> Use to generate a traditional Langmuir surface area plot used to determine monolayer volume constant.</p> <ul style="list-style-type: none"> <li>• <b>Autoscale x-axis.</b> Linear x-axes begin at zero. The x-axis field shows the absolute pressure for Langmuir.</li> <li>• <b>Autoscale y-axis.</b> The y-axis field shows Langmuir transformation.</li> <li>• <b>Overlay samples.</b> Use to overlay sample files on the Langmuir transform plot.</li> </ul>
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	

## MP-METHOD REPORT OPTIONS


The *MP-Method Report Options* provides pore volume distributions for microporous materials by correlating quantity adsorbed with the thickness of the adsorbed layer as determined from a user-selected thickness curve. Pore size can be expressed in angstroms or nanometers. Go to **Options > Units** to specify the unit.



### MP-Method Report Options Fields and Buttons Table

Field or Button	Description
<b>Pressures</b> [button]	<p>Use to select a pressure range for report calculations and points for exclusion from calculations.</p> <p><b>Calculation pressure range.</b> Enter the minimum and maximum pressures to be used in the pressure table. To exclude a point from the calculations used to generate the report, select <i>Exclude</i>.</p> <p><b>Exclude All.</b> Select to exclude all pressure points in the table.</p> <p><b>Include All.</b> Select to include all pressure points in the table.</p>

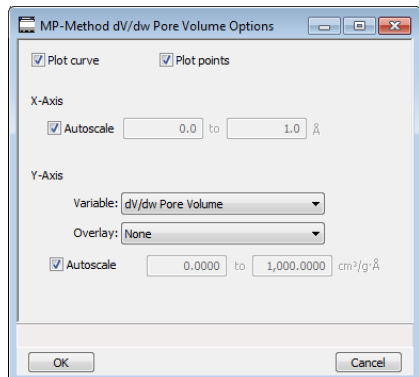
MP-Method Report Options Fields and Buttons Table (continued)

Field or Button	Description
<b>Select Reports</b> <i>[group box]</i>	Select the reports to generate. Highlight the report, then click <b>Edit</b> to modify report options.
<b>Thickness Curve</b> <i>[group box]</i>	Select the thickness curve, then click <b>Edit</b> to modify the values in the equation for the selected curve.
	For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .




## MP-METHOD PLOT REPORT OPTIONS

In the *MP-Method Report Options* window, highlight a plot option in the *Selected Reports* list box, then click **Edit** to customize the plotting method.

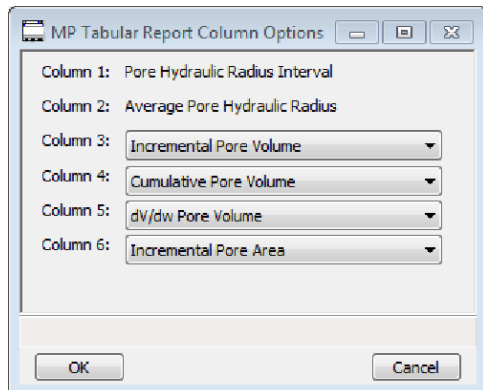


### MP Method Plot Options Fields and Buttons Table

Field or Button	Description
<b>Overlay</b> [drop-down box]	Select an option to overlay on the current report.
<b>Plot curve / Plot points</b> [selection]	Select to plot points on the graph.
<b>Thickness Curve</b> [group box]	Select the thickness curve, then click <b>Edit</b> to modify the values in the equation for the selected curve.
<b>X-Axis</b> [check box]	Use to have the x-axis autoscaled or enter beginning and ending values.
<b>Y-Axis</b> [group box]	<p><b>Autoscale.</b> Use to have the y-axis autoscaled or enter beginning and ending values.</p> <p><b>Overlay.</b> Select an option to overlay on the current report.</p> <p><b>Variable.</b> Select a variable.</p>
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	

## MP-METHOD TABULAR REPORT OPTIONS

In the *MP-Method Report Options* window, highlight *MP Tabular Report* in the *Selected Reports* list box, then click **Edit**. *Column [n]* indicates the column order and data contents for the report.



The MP Method reports hydraulic radius only. If Pore size in diameter is selected on the Unit Selection window, pore size in radius will be reported.

## OPTIONS REPORT

Lists the conditions used to perform the analysis— such as:

- Adsorptive properties
- Analysis conditions
- Analysis method
- Degas conditions
- Free space
- Saturation pressure ( $P_0$ ) and temperature



Options reports cannot be edited.

## ***SAMPLE AUDIT TRAIL REPORT***

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**CFR  
Note**

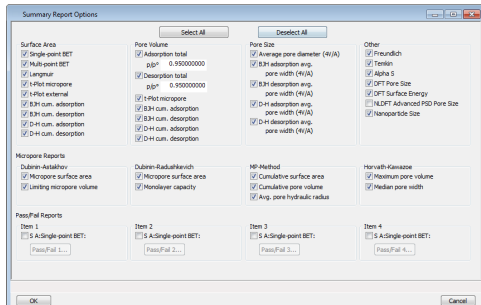
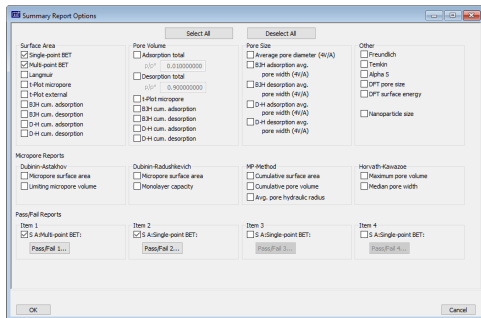
For 21CFR11 environments only.

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This report lists all changes and comments that been applied to sample files with a *Completed* status.

## SUMMARY REPORT OPTIONS

The *Summary Report* for physical adsorption analyses provides a condensed summary of selected data results.

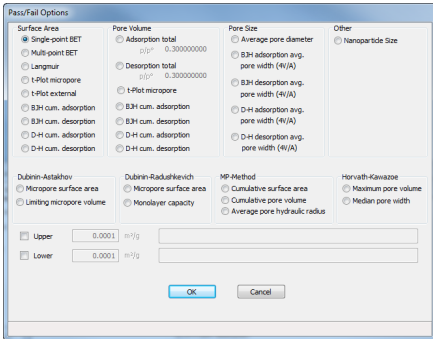




In the *Pore Volume* group box, if *Adsorption total* or *Desorption total* is selected, the  $p/p^0$  field is enabled. Enter the relative pressure used to calculate the total pore volume.

### Summary Report Fields and Buttons Table

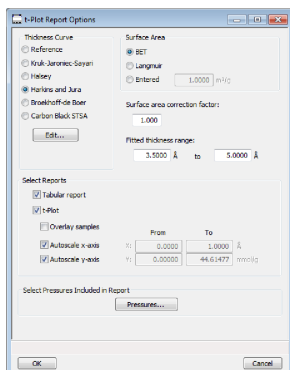
Field or Button	Description
<b>Item [n] [selection]</b>	Use to enable the first <i>Pass/Fail</i> item. Until the <i>Summary Report</i> is selected, <i>S A Single-point BET</i> will be displayed by default. When selected, click <b>Pass/Fail</b> , then select pass/fail criteria options.
<b>Pass/Fail [n].</b>	Click to display the <i>Pass/Fail Options</i> window for selection of pass/fail criteria.

## Summary Report Fields and Buttons Table (continued)

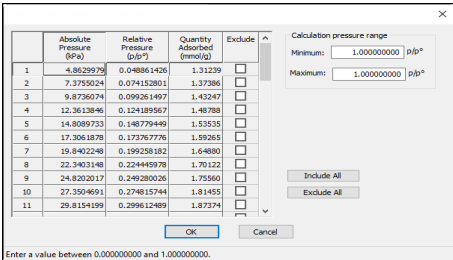
Field or Button	Description
	 <p><b>S A: Single-point BET.</b> Use to enable <b>Pass/Fail [n]</b> in the <i>Item [n]</i> group box.</p> <p><b>Upper / Lower.</b> Specify upper and lower limits for the selected parameter. A range can be left open by not selecting the limit. In the text box to the right of <i>Upper / Lower</i>, enter operator instructions to be displayed if a failure is encountered.</p>
<b>Select All / Deselect All [button]</b>	Selects (or deselects) all options.
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	

## T-PLOT REPORT OPTIONS

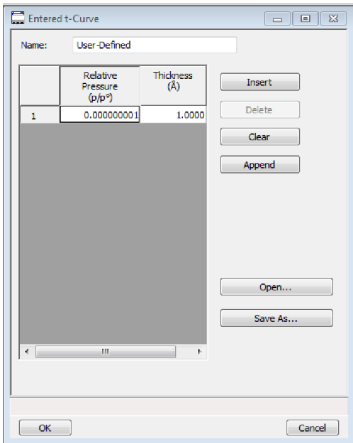
The *t*-Plot calculation allows quantitative analysis of the area and total volume ascribed to micropores. Matrix area (the area external to micropores) is directly determined and often proves to be a valuable way of characterizing complex mixed materials.




### t-Plot Report Options Fields and Buttons Table

Field or Button	Description
<b>Fitted thickness range</b> [text box]	Enter the minimum and maximum thicknesses (in angstroms or nanometers) to include in the thickness curve. Go to <b>Options &gt; Units</b> to specify default units.
<b>Pressures</b> [button]	<p>Use to select a pressure range for report calculations and points for exclusion from calculations.</p>  <p><b>Calculation pressure range.</b> Enter the minimum and maximum pressures to be used in the pressure table. To exclude a point from the calculations used to generate the report, select <i>Exclude</i>.</p> <p><b>Exclude All.</b> Select to exclude all pressure points in the table.</p> <p><b>Include All.</b> Select to include all pressure points in the table.</p>

t-Plot Report Options Fields and Buttons Table (continued)

Field or Button	Description
<b>Selected Reports</b> <i>[group box]</i>	<p><b>Tabular Report.</b> Use to have a tabular report of data generated.</p> <p><b>t-Plot.</b> Use to have a graphical representation of data generated.</p> <ul style="list-style-type: none"> <li>• <b>Autoscale x-axis.</b> The x-axis field shows the statistical thickness of the adsorbed film.</li> <li>• <b>Autoscale y-axis.</b> The y-axis field shows the quantity of gas adsorbed.</li> <li>• <b>Overlay samples.</b> Use to overlay sample files on the t-plot.</li> </ul>
<b>Surface area correction factor</b> <i>[text box]</i>	Enter the value to correct for surface areas that are not smooth. This brings the values for BET surface area and micropore surface area into accordance. For most samples, the default value of 1.000 is adequate.
<b>Surface Area</b> <i>[group box]</i>	Select the surface area value used for thickness calculations. BET is the most commonly used option.
<b>Thickness Curve</b> <i>[group box]</i>	<p>Select the thickness curve, then click <b>Edit</b> to modify the values in the equation for the selected curve. The Frenkel-Halsey-Hill thickness curve can be applied using the Halsey option.</p> <p><b>Kruk-Jaroniec-Sayari / Halsey / Harkins and Jura / Broekhoff-de Boer / Carbon Black STSA.</b> Select the thickness curve option, then click <b>Edit</b>. Modify the equation for the selected curve as needed.</p> <p><b>Reference.</b> Select <i>Reference</i>, then click <b>Edit</b> to define a t-curve by entering both the relative pressure and thickness values. One predefined curve is shipped with the analysis program and is found in the <i>Reference</i> directory.</p> 

t-Plot Report Options Fields and Buttons Table (continued)

Field or Button	Description
	To import values from an existing thickness curve (.THK file), click <b>Open</b> , then select the file containing the values. The table to be imported must have a .TXT or .THK file extension and have a two-column format with the relative pressures in the first column and the thickness values in the second column. Columns must be separated by a space or a tab.
<b>t-Plot</b> <i>[check box]</i>	<p>Use to have a graphical representation of data generated.</p> <p><b>Autoscale x-axis.</b> The x-axis field shows the statistical thickness of the adsorbed film.</p> <p><b>Autoscale y-axis.</b> The y-axis field shows the quantity of gas adsorbed.</p> <p><b>Overlay samples.</b> Use to overlay sample files on the <i>t</i>-plot.</p>
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	




## TEMKIN REPORT OPTIONS

The *Temkin* isotherm is used to model adsorption data where the heat of adsorption drops linearly with increasing coverage.

Temkin Report Options Fields and Buttons Table

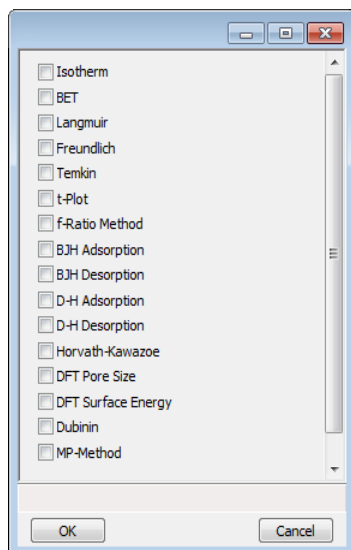
Field or Button	Description
<b>Pressures [button]</b>	<p>Use to select a pressure range for report calculations and points for exclusion from calculations.</p> <p><b>Calculation pressure range.</b> Enter the minimum and maximum pressures to be used in the pressure table. To exclude a point from the calculations used to generate the report, select <i>Exclude</i>.</p> <p><b>Exclude All.</b> Select to exclude all pressure points in the table.</p> <p><b>Include All.</b> Select to include all pressure points in the table.</p>

Temkin Report Options Fields and Buttons Table (continued)

Field or Button	Description
<b>Selected Reports</b> <i>[group box]</i>	<p><b>Tabular Report.</b> Generates a tabular report of the included samples. A tabular report contains the numeric values contributed by each sample.</p> <p><b>Temkin Isotherm plot.</b> Overlays the Temkin isotherm with the analysis data.</p> <ul style="list-style-type: none"> <li>• <b>Autoscale x-axis.</b> Linear x-axes begin at zero. The x-axis field shows the absolute pressure.</li> <li>• <b>Autoscale y-axis.</b> y-axes begin at zero. The y-axis field shows the quantity of gas adsorbed.</li> <li>• <b>Overlay samples.</b> Use to overlay sample files on the isotherm plot.</li> </ul> <p><b>Temkin Transform plot.</b> Plots a linear form of the Temkin transform plot.</p> <ul style="list-style-type: none"> <li>• <b>Autoscale x-axis.</b> The x-axis field shows the logarithm of pressure (ln).</li> <li>• <b>Autoscale y-axis.</b> The y-axis field shows the quantity of gas adsorbed.</li> <li>• <b>Overlay samples.</b> Use to overlay sample files on the transform plot.</li> </ul>
<b>Specify differential heat of adsorption</b> <i>[check box]</i>	Select and enter the differential heat of adsorption at zero surface coverage. This allows inclusion of all Temkin constants.
<b>Specify monolayer capacity</b> <i>[check box]</i>	Select and enter the monolayer capacity of the sample.
 <p>For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a>.</p>	

## ***VALIDATION REPORT OPTIONS***

This report allows data to be examined by the analysis program to determine if the results are within typical ranges. If the data for any reports selected for validation are determined to be out of range, a warning will display and suggestions are given for corrective action. This information is detailed in the report and plotted on the graph as a unique plot symbol.



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## 9 DIAGNOSTICS

### Unit [n] > Diagnostics

Use to display diagnostic readings, start diagnostic tests, and open saved diagnostic reports. Each test generates a file to the default directory name and path of ...\\...\\Service\\userdiag unless another directory name was specified. These reports can be sent to a Micromeritics Service Representative for examination.

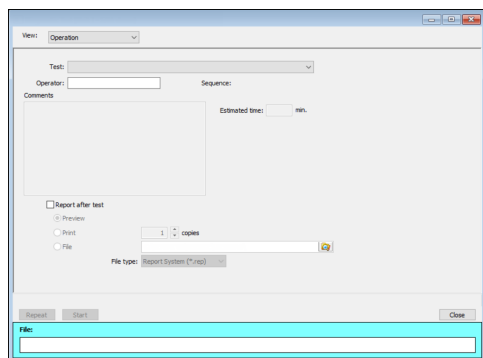
### START DIAGNOSTIC TEST

#### Unit [n] > Diagnostics

Provides a method to start a diagnostic test immediately. Upon completion of the diagnostic test, the file is saved as a .REP file which can be retrieved by going to **Reports > Open Report** and selecting the report file.

Use to clean and verify gas lines when connecting or changing a gas. See [Clean and Verify the Gas Line on page 11 - 5](#) for detailed instructions. Also used to perform diagnostic tests requested by a Micromeritics Service Representative.


This option is typically used and /or requested by a Service Representative. These tests create and save files to the ...\\...\\Service\\userdiag directory. The Micromeritics Service Representative may request an emailed or faxed copy of the file for diagnostic purposes.



### Start Diagnostic Test Fields and Buttons Table

Field or Button	Description
<b>Comments</b> [text box]	Displays comments from the selected diagnostic test.
<b>Estimated time (min.)</b>	Approximate time for test completion.
<b>File</b> [group box]	Shows a status bar of steps complete once the test begins.
<b>Next</b> [button]	Starts the next test.
<b>Operator</b> [text box]	Enter information to identify the person running the service test.

**Start Diagnostic Test Fields and Buttons Table (continued)**

Field or Button	Description
<b>Repeat</b> <i>[button]</i>	Repeats the selected diagnostic test.
<b>Report after test</b> <i>[check box]</i>	Automatically generates reports to the selected destination when the test is complete.
<b>Sequence</b>	Sequence number assigned to the test.
<b>Start</b> <i>[button]</i>	Starts the diagnostic test.
<b>Test</b> <i>[drop-down box]</i>	Select the diagnostic test to be performed.
 For fields and buttons not listed in this table, see <a href="#">Common Fields and Buttons on page 2 - 3</a> .	

## 10 CALIBRATION

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### ***Unit [n] > Calibration***



Service Test Mode is required for this test. See [Service Test Mode on page 11 - 22](#) for instructions on how to enter Service Test Mode.

A calibration file was created specifically for the analyzer and included with the accessories. It is not necessary to recalibrate the system unless it seems out of calibration.

Disabled calibration menu options can be accessed only with the assistance of an authorized Micromeritics Service Representative. Calibrations can be saved to a file and reloaded later.

To review calibration details of the analyzer, go to ***Unit [n] > Unit Configuration***.

Generally it will not be necessary to change the data in the calibration file. However, if a condition occurs during the operational verification that requires changes to the calibration data, changes should be saved in a file. Calibration data files are retained in the analyzer history file and can be reloaded in the event that calibration data becomes corrupt.

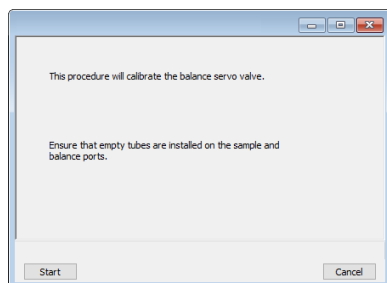
## **BALANCE SERVO VALVE**

### **Unit [n] > Calibration > Balance Servo**

Use to calibrate the servo valve to the manifold pressure transducer. The servo valve should always be recalibrated after a pressure calibration has been performed. The pressure transducer should be calibrated before starting this calibration procedure.



Ensure the pressure transducer has been calibrated before performing this procedure. Go to **Unit [n] > Unit Configuration** and view the calibration information. Contact your Micromeritics Service Representative if calibration dates are not listed.



Click **Start**. The window closes when the calibration is complete. Click **Cancel** to stop the calibration process.



## ***MATCH TRANSDUCERS***

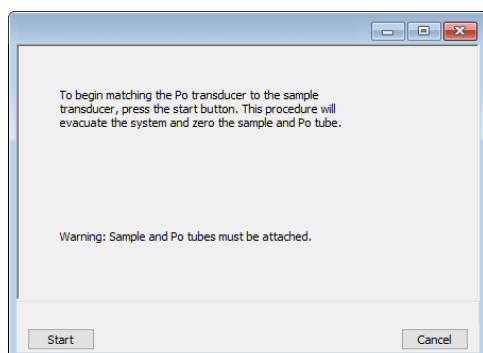
### ***Unit [n] > Calibration > Match Transducers***

This option is disabled for the Gemini 2390a model.

Use to evacuate the system and zero the pressure transducers, then adjust the scale to match them to the manifold transducer near full scale pressure.



A blank sample tube or small plug must be installed on each selected port prior to starting this process.

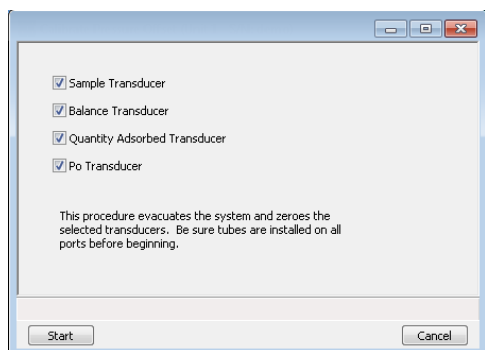


1. Install a blank sample tube or small plug on each port.
2. Click **Start**. The window closes when the operation is complete. Click **OK** when the process is complete.

## ***ZERO PRESSURE***

### ***Unit [n] > Calibration > Zero Pressure***

This procedure evacuates the system and zeroes the pressure transducers. This calibration should only be performed by qualified service personnel. In order to perform this procedure, sample tubes must be attached to each port.



1. Install a blank sample tube or small plug on each applicable port.
2. Ensure that all applicable transducers are selected, then click **Start**. Click **OK** when the process is complete.

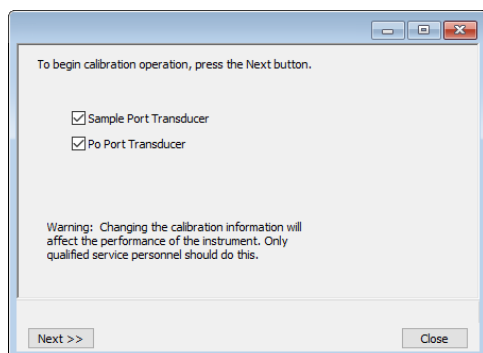
## PRESSURE SCALE CALIBRATION

**Unit [n] > Calibration > Pressure Scale**



Service Test Mode is required for this test. See [Service Test Mode on page 11 - 22](#) for instructions on how to enter Service Test Mode.

Use to adjust the scale of a transducer to match a reference standard.



Changing the calibration information will affect the performance of the instrument. Only qualified personnel should do this.

## QUANTITY ADSORBED

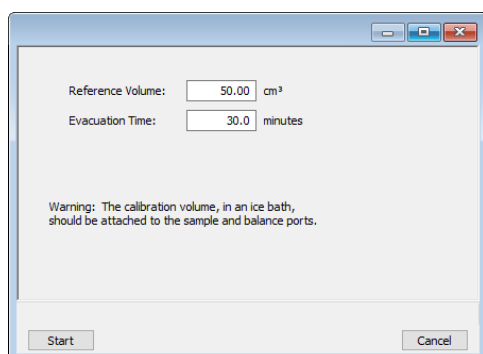
**Unit [n] > Calibration > Quantity Adsorbed**



Service Test Mode is required for this test. See [Service Test Mode on page 11 - 22](#) for instructions on how to enter Service Test Mode.



Ensure the calibration volume (in an ice bath) is attached to the sample and balance ports.



Click **Start**. The window closes when the calibration is complete. Click **Cancel** to stop the calibration process.

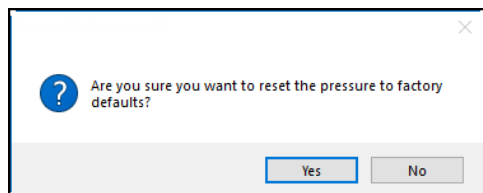
## RESET PRESSURE CALIBRATION

**Unit [n] > Calibration > Reset Pressure Calibration**



Service Test Mode is required for this test. See [Service Test Mode on page 11 - 22](#) for instructions on how to enter Service Test Mode.

This procedure resets the pressure calibration to the factory default settings.



The servo valve should always be recalibrated after a pressure calibration has been performed. The pressure transducer should be calibrated before starting this procedure.

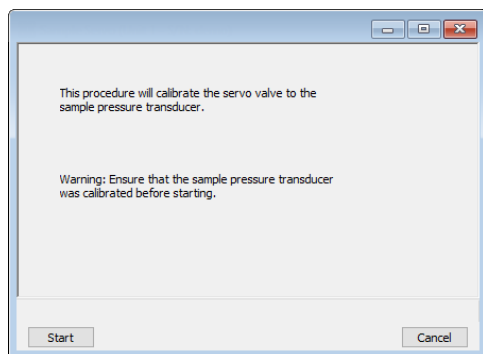
## **SAMPLE SERVO VALVE**

### **Unit [n] > Calibration > Sample Servo**

Use to calibrate the servo valve to the manifold pressure transducer. The servo valve should always be recalibrated after a pressure calibration has been performed. The pressure transducer should be calibrated before starting this calibration procedure.



Ensure the pressure transducer has been calibrated before performing this procedure. Go to **Unit [n] > Unit Configuration** and view the calibration information. Contact your Micromeritics Service Representative if calibration dates are not listed.



Click **Start**. The window closes when the calibration is complete. Click **Cancel** to stop the calibration process.

## ***LOAD CALIBRATION FROM FILE***

### ***Unit [n] > Calibration > Load from File***

Use to load a previously saved calibration file.

It is recommended that the current calibration settings be saved using ***Unit [n] > Calibration > Save to File*** prior to loading another calibration file. When loading a previously saved calibration file, a backup of the current file is created and saved as *[SN]last.cal*. The backup file is overwritten each time a new one is created.



Changing the calibration may affect the analyzer's performance.

## ***SAVE CALIBRATION TO FILE***

### ***Unit [n] > Calibration > Save to File***

Use to save the current calibration settings to a backup file which can later be reloaded using the ***Unit [n] > Calibration > Load from File*** menu option.

The default file naming convention for calibration files can be used or the file name can be changed. The default file name of 0217-2013-04-25.CAL is interpreted as:

<b>0217</b>	is the analyzer serial number
<b>2013-04-25</b>	is the date the calibration file was saved
<b>.CAL</b>	is the file name extension

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## 11 TROUBLESHOOTING AND MAINTENANCE

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The analyzer has been designed to provide efficient and continuous service; however, certain maintenance procedures should be followed to obtain the best results over the longest period of time. When unexpected results occur, some common operational problems not indicated on the window and their respective causes and solutions are provided:

Log in to your [customer portal](#) to access error messages. Parts and accessories can be found online at [www.Micromeritics.com](http://www.Micromeritics.com).

Most operational problems are caused by:

- Leaks (commonly found at the sample tube O-ring at the analysis port)
- Sample weighing errors
- Use of too much analysis bath fluid in the dewar at the start of an analysis
- Entry of incorrect system volume for analysis
- Impure gas supply

When unexpected analysis results occur, check the above first. Some common operational problems not indicated on the window and their respective causes and solutions are provided below:

### **Elevator cannot be raised or lowered.**

*Cause:* Dewar elevator stuck.

*Action:* Check for possible obstruction to elevator movement.

### **Elevator is noisy.**

*Cause:* The elevator screw may need greasing.

*Action:* Contact your Micromeritics Service Representative.

### **Expected results are not within range.**

*Cause A:* Sample improperly degassed.

*Action A:* Verify that the degassing temperature and degas time were set properly.

*Cause B:* Gas cylinder(s) may be almost empty - causing gas impurity.

*Action B:* Replace gas cylinder. See [Replace a Gas Cylinder on page 11 - 8](#).

*Cause C:* Undetermined.

*Action C:* Do the following:

1. Perform an empty tube analysis. See [Perform a Leak Test on page 11](#)

- [12](#) and print the results.
- 2. Perform a reference material analysis. See [Perform a Reference Material Analysis on page 6 - 13](#) . Print the results.
- 3. Contact your Micromeritics Service Representative.

**Specified pressure not reached.**

*Cause A:* Sample or balance tube not properly attached.

*Action A:* Reattach sample and balance tubes. Ensure both are securely attached to ports. Replace O-rings if defective. See [Sample Tube O-ring Replacement on page 11 - 21](#).

*Cause B:* Leak in gas line.

*Action B:* Perform the Adsorptive and Helium Line tests to determine if a leak exists. See [Diagnostics on page 9 - 1](#).

**Unable to reach a satisfactory vacuum.**

*Cause A:* Vacuum pump oil level is low or needs to be changed.

*Action A:* Inspect the oil to see if it is low or needs changing. Log in to your [customer portal](#) to access the Vacuum Pump Guide.

*Cause B:* Vacuum pump is leaking.

*Action B:* Locate and repair the source of the leak or replace pump.

*Cause C:* Centering ring has become too flat and unable to hold a vacuum.

*Action C:* Check the centering ring at the pump intake port. Also check the centering ring at the top of the oil vapor trap. Replace if necessary. Log in to your [customer portal](#) to access the Vacuum Pump Guide.

**Nitrogen or helium drained from tank, or depleted in a short period of time.**

*Cause:* Leak in the gas line connection.

*Action:* Perform the Adsorptive (or Helium) Line test to determine the location of the leak. See [Diagnostics on page 9 - 1](#).

**Unit does not work when powered ON.**

*Cause A:* Power cord not fully inserted at one end.

*Action A:* Insert power plug firmly into power source and analyzer power connector.

*Cause B:* No power at outlet.

*Action B:* Plug in another electrical device to test the outlet. If there is no power, contact an electrician.

*Cause C:* Plug prongs bent so that contact not made at outlet.

*Action C:* Gently move power plug at outlet while watching power indicator light. If indicator light comes on, have electrician replace outlet or plug.

*Cause D:* Power cord damaged.

*Action D:* Have electrician check cord using a test meter. Replace the cord if defective.

*Cause E:* Loose internal connection or broken wire.

*Action E:* Call a Micromeritics Service Representative for repair or replacement information.

### **Valves cannot be operated.**

*Cause:* Cable from computer to the instrument is loose.

*Action:* Reconnect the cable.

### **Vacuum pump is noisy.**

*Cause A:* Sample tube connector is loose.

*Action A:* Tighten fitting. Replace O-ring.

*Cause B:* Sample tube O-ring is worn or cracked.

*Action B:* Replace O-ring. See [Sample Tube O-ring Replacement on page 11 - 21](#).

*Cause C:* Sample tube is cracked.

*Action C:* Replace with new sample tube.

*Cause D:* No sample tube loaded on a selected port.

*Action D:* Install plug or empty sample tube.

*Cause E:* Gas inlet valve open while vacuum valve open.

*Action E:* With manual control enabled, use the instrument schematic to close gas inlet valve.

## ***GUIDELINES FOR CONNECTING GASES***

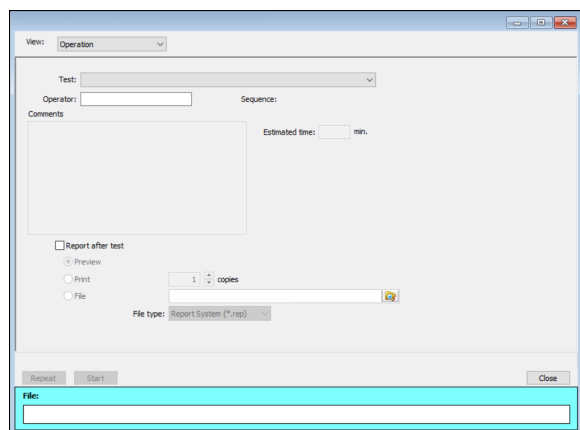
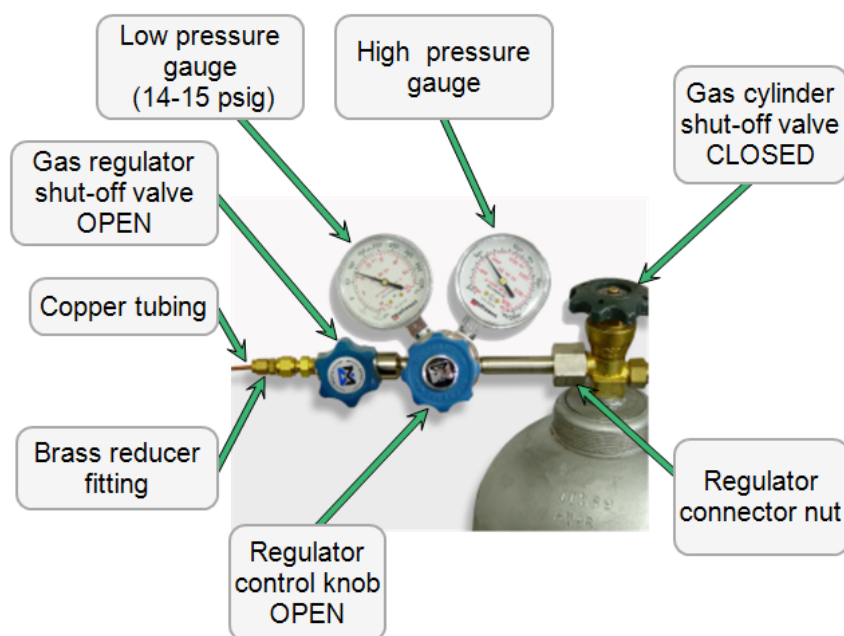
- Place gas cylinders within 6 feet (2 m) of the gas inlets of the analyzer. Using gas line extenders on gas cylinders located in remote areas may degrade gas quality and reduce pressure. Gas lines are typically five to six feet long. Place the cylinders close enough to allow for proper connection at the analyzer inlet.
- Use a retaining strap (or other appropriate tether) to secure the gas cylinder.
- Always use the gas lines provided with the analyzer. It is very important that proper gas lines are used with the analyzer.
  - **Do not use** polymer tubing for the gas line.
  - **Do not use** flexible gas lines. Some flexible lines may appear to be appropriate, such as those with a herringbone covering, but the line may be coated internally with a polymer.
- Long gas lines, such as those used with gas cylinders placed in remote areas, must be evacuated for an extended period of time to remove ambient gases. When possible, avoid placing gas cylinders in remote locations. It is always best to have gas cylinders located near the analyzer.
- Carefully route the gas lines from the cylinder to the analyzer avoiding overlapping or entangling gas lines. This will help avoid confusion when maintenance is required.
- Label the gas line at the analyzer inlet for proper identification and maintenance.
- Replace gas cylinders before gas is depleted. It is best to replace a gas cylinder when the pressure reads approximately 500 psi (3500 kPa) on the high pressure gauge. Contaminants absorbed to the walls of the cylinder will desorb as the pressure decreases.
- Ensure the gas cylinder is closed before connecting to the analyzer.

## CLEAN AND VERIFY THE GAS LINE

### Unit [n] > Diagnostics

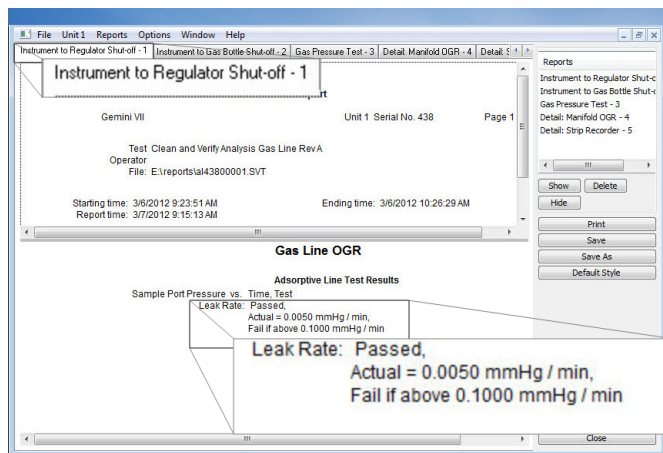
Always clean the gas lines and verify there are no leaks at the connections after a gas cylinder is connected. This test examines the gas line from the analyzer to the gas cylinder, then from the analyzer to the regulator shutoff valve. A report is generated at the completion of the test to verify that it has passed or failed. Causes and corrective action for a failure are provided.

Before beginning, confirm that the state for valves and the low-pressure gauge are as follows:



1. Click the down arrow to the right of the *Test* field and select *Clean and Verify Analysis Gas Line Rev [n]*. This procedure uses the analysis gas as an example. Select *Clean and Verify*

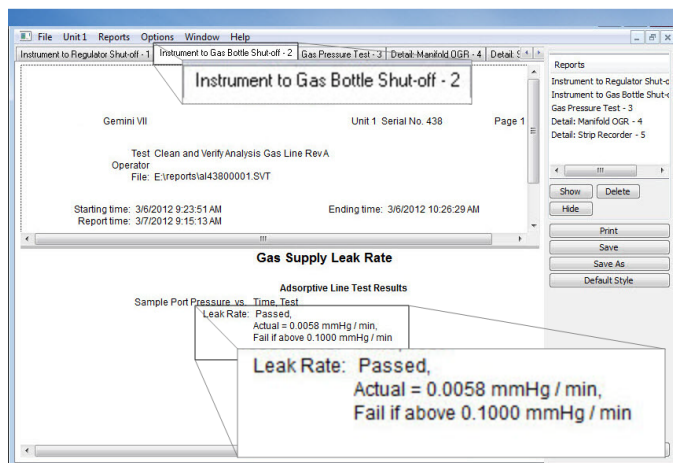
- Helium Gas Line Rev [n]* if the helium cylinder is being verified.
2. Select *Report after test* and select *Preview* as the destination. Resize the window if this option does not display.
  3. Click **Start**. From the *View* drop-down list, select either *Operation*, *Instrument Log*, or *Instrument Schematic*.
  4. A series of prompts display requiring operator response. The length of time a test will run is also indicated.
  5. A popup window indicates the test is complete. Click **OK**.
  6. Click the *Instrument to Regulator Shut-off Valve - 1* tab to display its report.



The *Leak Rate* field should read *Passed* indicating the gas line from the analyzer to the regulator is clean and leak-free.

If the *Leak Rate* field indicates *Failed*, a leak is indicated. Check the connections from the analyzer to the gas regulator valve. Tighten as necessary, then run the test again.

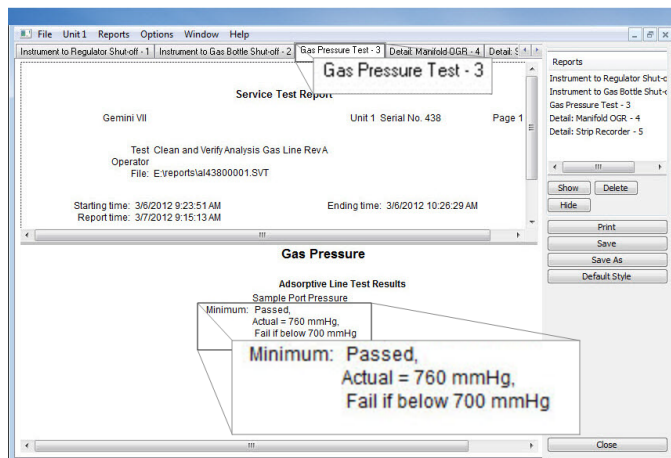
7. Click the *Instrument to Gas Bottle Shut-off Valve - 2* tab to display its report.



The *Leak Rate* field should read *Passed* indicating the gas line from the analyzer to the regulator is clean and leak-free.

If the *Leak Rate* field indicates *Failed*, a leak is indicated. Check the connections from the regulator shut-off valve to the gas cylinder shut-off valve. Tighten as necessary, then run the test again.

8. Click the *Gas Pressure Test - 3* tab to display its report.



On the report, the *Minimum* field should read *Passed* indicating all valves are in the proper state for operation.

If the *Minimum* field indicates *Failed*, one or more valves is not in the proper position. Set the valves as shown at the beginning of this section and ensure the appropriate pressure is displayed on the low-pressure gauge.

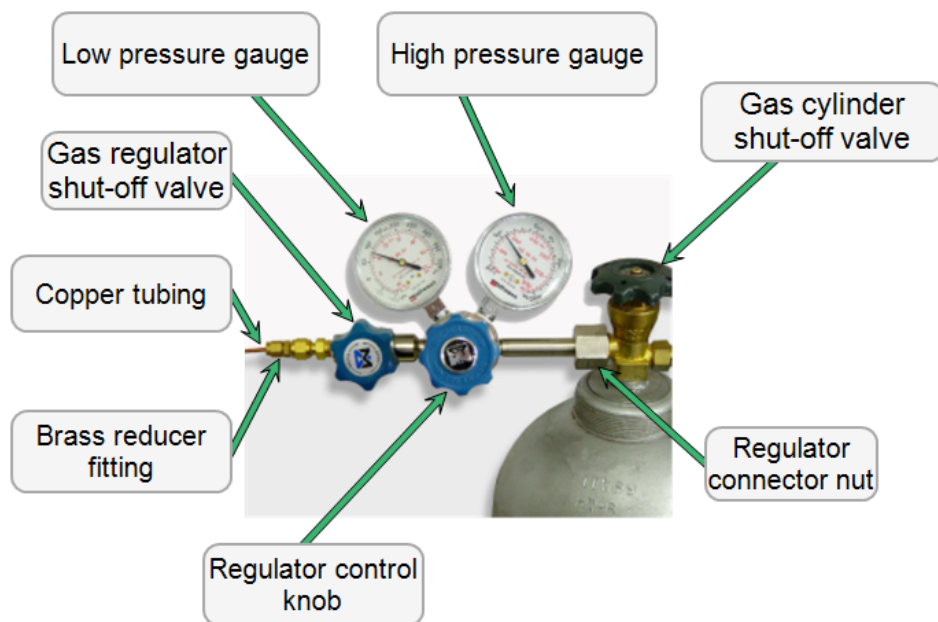
If running the test again, close the gas cylinder valve before starting the test.

9. Click **Close** to close the test report. Click **Close** again to close the test.

## REPLACE A GAS CYLINDER



These instructions apply to working with inert gases only. When working with hazardous gases, follow the safety procedures established by your lab.





***DISCONNECT THE DEPLETED GAS CYLINDER***

1. Close the regulator shut-off valve and gas cylinder shut-off valve by turning the knobs clockwise.
2. Disconnect the gas line from the regulator. Gas will be vented from the line. It is not necessary to disconnect the gas line from the analyzer inlet if the cylinder will be replaced immediately with one of the same type.
3. Open the gas regulator shut-off valve by turning the knob counter-clockwise. Gas will be vented from the regulator.
4. Turn the regulator control knob clockwise to open and vent any remaining gas. Both gauges should read at or near zero. If not, make sure the gas regulator shut-off valve is open.
5. Close the regulator by turning the control knob counter-clockwise.
6. Use an appropriate wrench to loosen the nut at the regulator connector nut then remove the regulator from the cylinder.
7. Replace the protective cap on the depleted cylinder. Disconnect the retaining strap and move the cylinder to an appropriate location.

## CONNECT A GAS CYLINDER

### Regulator Pressure Settings

Analyzer Series	Gauge should indicate
3Flex	15 psig (103 kPag)
AccuPyc	25 psig (172 kPag)
ASAP	15 psig (103 kPag)
AutoChem	15-18 psig (103 - 124 kPag)
AutoPore	50-60psig (345 - 404 kPag)
Gemini	15-18 psig (103 - 124 kPag)
TriStar	15-18 psig (103 - 124 kPag)

Move the replacement cylinder close to the analyzer and tether it into place.

1. Use an appropriate cylinder wrench to remove the protective cap from the replacement gas cylinder. Place the protective cap in a secure location. It will be needed to recap the gas cylinder when it is depleted and replaced.
2. Attach the gas regulator to the gas cylinder connector. Hand tighten the nut, then use an appropriate wrench to tighten an additional 3/4 turn.



Over-tightening the fitting may cause a leak.

3. Check for leaks at the high-pressure side of the regulator and in the connector.
  - a. Turn the regulator control knob fully counter-clockwise.
  - b. Slowly open the gas cylinder shut-off valve, then quickly close it.
  - c. Observe the pressure on the high-pressure gauge for approximately one minute:
    - If the pressure is stable, proceed with the next step.
    - If the pressure decreases, tighten the regulator connector nut until it becomes stable. If the pressure does not remain stable, remove the regulator and clean all contacts at the regulator connection, then reinstall the regulator.
4. Purge the air from the lines.



Purge the regulator before proceeding to prevent contamination of the analysis gas supply.

- a. Open the gas cylinder valve to pressurize the regulator, then close the valve.
- b. Adjust the *Pressure Control* knob to approximately 5 psi.

- c. Turn the regulator *Shut-off* valve counter-clockwise to open. Allow gas to flow until both gauges read approximately zero.
- d. Close the regulator *Shut-off* valve to stop gas flow.
- e. Reconnect the gas line to the regulator.
- f. Use two 7/16 in. (11 mm) wrenches to tighten the gas line connection. Hold one wrench fitting steady and the other to tighten the connector nut.
5. Set the analyzer pressure.
  - a. Turn the *Regulator Control* knob clockwise until the low pressure gauge indicates the appropriate pressure. See [Regulator Pressure Settings on the previous page](#).
  - b. Open the regulator *Shut-off* valve.
  - c. Open the gas cylinder *Shut-off* valve and flow gas for 10 to 30 seconds.
  - d. Close the gas cylinder *Shut-off* valve.
  - e. Close the gas cylinder valve.
6. If the gas line to the instrument inlet was previously disconnected, reconnect it now.

## ENABLE MANUAL CONTROL

### Unit [n] > Enable Manual Control



See also:

[Show Instrument Schematic on page 2 - 19](#)

Use to enable the manual control of certain system valves and elevator components on the analyzer schematic. When this option is enabled, a checkmark appears to the left of **Unit [n] > Enable Manual Control**. If the analyzer schematic is not immediately visible, go to **Unit [n] > Show Instrument Schematic**.

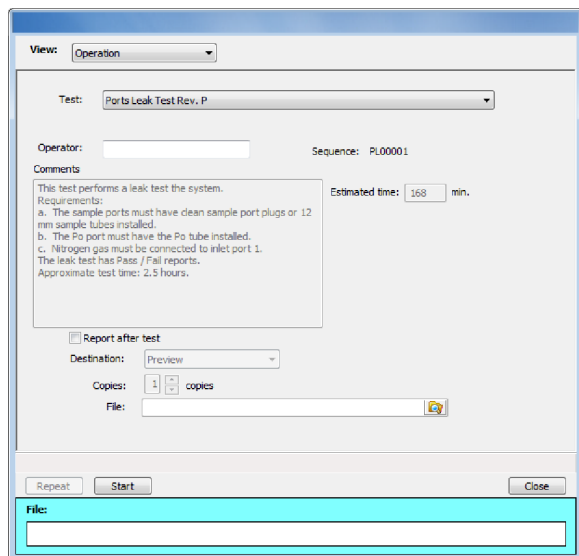
## PERFORM A LEAK TEST

### Unit [n] > Diagnostics

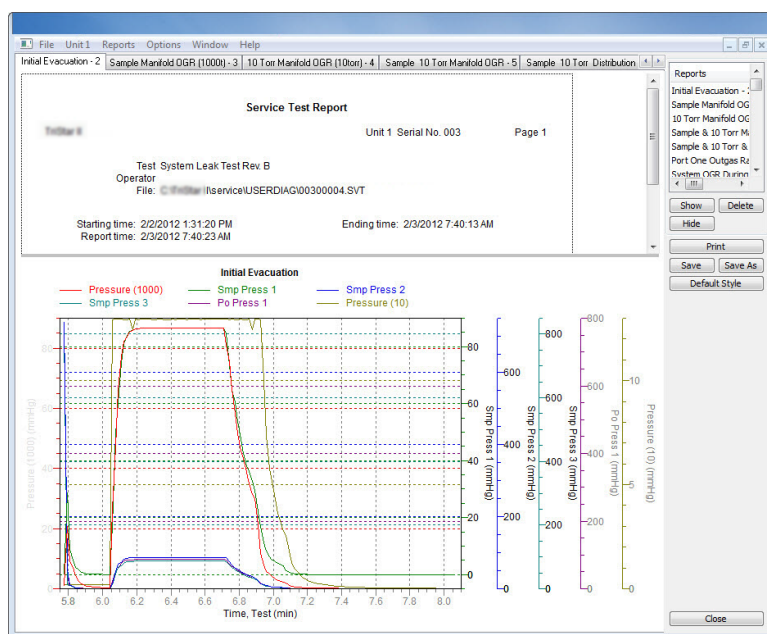
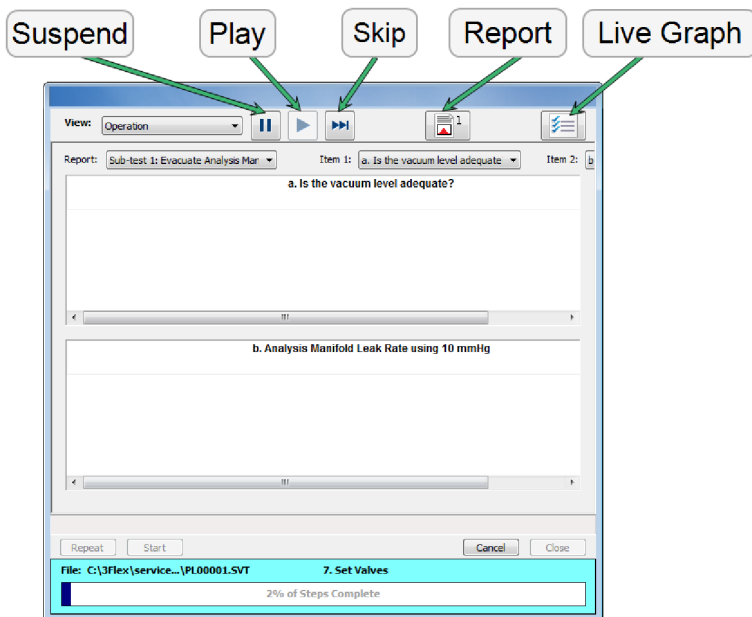
A Micromeritics Service Representative may request that a leak test be performed to determine if there is a system leak and may also require a copy of the report generated by this test.

The test provides:

- Prompts on preparing the analyzer for the test
- Approximate time period of the test
- Prompts in which an operator response is required



1. Select the test to run.
2. Select *Report After Analysis* and choose *Preview* as the destination.
3. Click **Start**.
4. Verify all tests have a *Passed* status by selecting the tabs and looking for the *Passed* status for each test run.
5. Click **Save As** to save the test file results.



## PERFORM REFERENCE MATERIAL ANALYSIS



See:

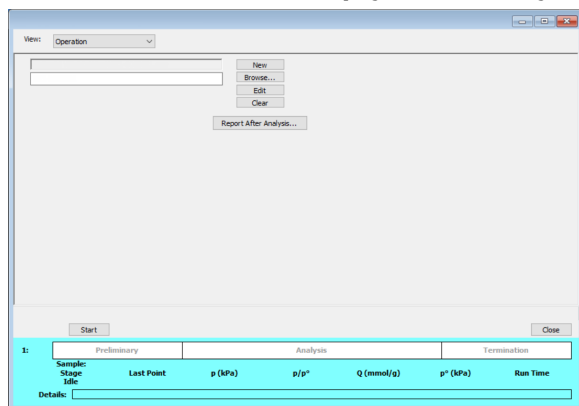
[Perform a Reference Material Analysis on page 6 - 13](#)

## PERFORM AN EMPTY TUBE ANALYSIS FOR DIAGNOSTIC PURPOSES

An empty tube analysis is a means of pinpointing operational problems in the analyzer. Leaks, as well as electronic problems, will result in an unsatisfactory blank analysis. This test should be performed prior to detailed troubleshooting. The data produced during the blank analysis can be helpful in identifying the cause of a problem and should be performed before contacting a Micromeritics Service Representative.

Helium is required for performing an empty tube analysis since a measured free space is required. If helium is unavailable, contact a Micromeritics Service Representative for an alternative method of performing this procedure.

1. Install empty, straight-wall sample tubes (free of cracks or chips) of the same size in the sample and balance ports. Ensure the O-rings are in good condition.
2. Place the dewar support on the elevator, then place a dewar of water at ambient temperature on the dewar support. (A dewar support is not necessary for the larger dewar used with the Gemini VII 2390t. Place the dewar directly on the elevator.)
3. Go to **Unit [n] > Empty Tube Analysis**.



4. Click **Browse** and select a file or to create a new one. Click **Open**.
5. Click **Edit** to make modifications or replace all parameters in the open file using the **Replace All** button. Click **Save**, then click **Close**.
6. Click **Report after analysis** to generate reports to specify the report destination when the analysis is complete. Click **OK**.

7. Click **Start** to begin the analysis. A window displays data as they are collected. A short delay is encountered before the port status changes from the Idle state at the bottom of the screen. Review the analysis results to see if they are within the tolerances shown in the following table. If these results are not achieved, call a Micromeritics Service Representative.

<b>P/P<sub>0</sub></b>	<b>Tolerance in cm<sup>3</sup></b>
<b>0.1</b>	±0.008
<b>0.2</b>	±0.010
<b>0.3</b>	±0.012
<b>0.4</b>	±0.014
<b>0.5</b>	±0.016
<b>0.6</b>	±0.018
<b>0.7</b>	±0.020
<b>0.8</b>	±0.022
<b>0.9</b>	±0.024

## ***PREVENTIVE MAINTENANCE***

Perform the following preventive maintenance procedures to keep the analyzer operating at peak performance. Micromeritics also recommends that preventive maintenance procedures and calibration be performed by a Micromeritics Service Representative every 12 months.

<b>Maintenance Required</b>	<b>Frequency</b>
<b>Alumina in vapor oil traps *</b>	Replace as required.
<b>Analyzer exterior</b>	Clean as required or every 3 to 6 months.
<b>Dewar</b>	Check and clean weekly.
<b>Diaphragm vacuum pump **</b>	If an oil free pump is used, replace diaphragm(s) every 12 months.
<b>Dry forepumps</b>	As required or every 24 months.
<b>Port filters and O-rings</b>	Replace every 3 to 6 months.
<b>Sample tube O-ring</b>	Replace as required or every 3 to 6 months.
<b>Test for leaks</b>	As required or every 12 months.
<b>Vacuum pump exhaust filter</b>	Replace as required or every 12 months.
<b>Vacuum pump fluid *</b>	Inspect and change as required or every 3 to 6 months.

\* Oil sealed vacuum pumps only.

\*\* For oil-free pumps only: After 12 to 18 months of operation, the diaphragm(s) in the pump will wear out and the pump becomes completely inoperable. To prevent instrument downtime due to an inoperable pump, it is recommend that the diaphragm(s) be replaced by a Micromeritics Service Representative every 12 months.



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## CHECK AND CLEAN THE DEWAR



When handling dewars, follow the precautions outlined in [Dewar Precautions on page 6 - 1](#).



Always handle glass dewars with care. Any product incorporating a vacuum is a potential safety hazard and should be treated with caution. If in doubt, contact your safety officer.

Ice and suspended frost particles may accumulate in the bottom of the analysis port dewar. Particles or deposits exceeding 1/4 in. in depth may jam between the bottom of the sample tube and the bottom of the dewar. Accumulations of fine particles impede liquid nitrogen circulation around the bottom of the sample tube. This causes the sample temperature to be slightly higher which, in turn, can cause pore volume measurement errors in those samples exhibiting high isotherm slope above 0.97 relative pressure. Accumulated ice is likely to melt and form a pool of water in the dewar if all liquid nitrogen evaporates. The water must be removed, otherwise it will solidify when liquid nitrogen is added and could press on the bottom of the sample tube causing breakage.

To ensure problems do not develop due to ice accumulation, check the dewar after each use. Clean on a weekly basis.

1. Remove the dewar from the analyzer.
2. Pour out liquid nitrogen into an appropriate cryogenic container. Do not re-use liquid nitrogen.



Do not pour liquid nitrogen directly into a sink. Doing so may cause drain pipes to burst.

3. Rinse the dewar with warm water to melt any remaining ice accumulation which may remain. Dry thoroughly.
4. Replace the dewar.

## ***CLEAN THE ANALYZER***

The exterior casing of the analyzer may be cleaned using a clean cloth dampened with isopropyl alcohol (IPA), a mild detergent, or a 3% hydrogen peroxide solution. Do not use any type of abrasive cleaner.



- Do not allow liquid to penetrate the casing of the analyzer. Doing so could result in damage to the unit.
- Use only a mild detergent in water to clean safety shields. The use of isopropyl alcohol can damage the shield surface.

## ***CLEAN THE GAS DELIVERY TUBES***

If using a FlowPrep or SmartPrep, gas delivery tubes should be wiped with a lint-free cloth after each use to remove any particles of sample that may have adhered to the tube. If a gas delivery tube becomes clogged or damaged, it should be replaced.

1. Remove the gas delivery tube from the flexible tubing.
2. Attach a new gas delivery tube by pressing the flexible tubing over the gas delivery tube. Ensure that the flexible tubing completely covers the ferrule at the end of the gas delivery tube.

## ***LUBRICATE THE ELEVATOR DRIVE ASSEMBLY***

The elevator screw is lubricated before it leaves the factory and should not require lubricating. If the elevator starts to vibrate or becomes noisy when traveling, contact a Micromeritics Service Representative for disposition.

Should lubrication become necessary, apply a light coat of Teflon Magnalube-G grease (Micromeritics part number 004-16163-00) to the elevator screw. Do not grease the guide rods.

## POWER ANALYZER ON AND OFF



DO NOT connect or disconnect cables when the analyzer is powered ON.

If a Smart VacPrep is used, it is recommended that the power to the Smart VacPrep remain ON when the analyzer is powered on. If it does become necessary to power off the Smart VacPrep, exit the analyzer program first. Restart the analyzer program, then power on the Smart VacPrep.

Power ON the equipment in the following order:

1. Computer, monitor, and printer
2. Analyzer
3. External vacuum pump (the pump must warm approximately two hours before performing analyses)
4. Degasser

Power OFF the equipment in the following order:

1. Exit the analysis program. Failure to do so could result in loss of data. If an analysis is in progress when closing the application, the following message is displayed:

**2459 - An Instrument is busy. A delay in restarting this application could result in loss of new data. Continue program exit? Yes / No**

**Yes.** Closes the program. The analysis continues and data continue to be collected. The data will be restored when the application is restarted. Reports queued in the print manager will print. If a power failure occurs and an uninterruptible power supply (UPS) is not attached to the analyzer, the data collected after exiting the analysis program are lost.

**No.** The program remains open and the analysis continues to run.

2. Computer, monitor, and printer
3. Analyzer
4. External pump
5. Smart VacPrep

## SAMPLE PORT FRIT REPLACEMENT



The equipment images in this topic may differ slightly from your equipment, however the instructions are the same unless otherwise noted.

A frit is located in the connecting nut attached to each analysis port. If the frit becomes contaminated, the contaminant may adsorb or desorb during analysis, affecting the results. A contaminated frit on the analysis port may be indicated as a leak or a free space reading much lower than normal.



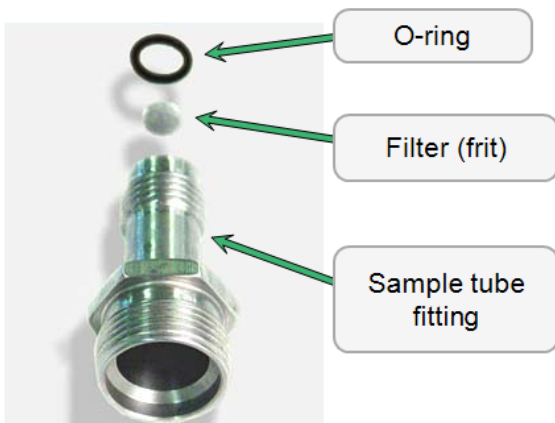
Use the proper size frit for the sample port. The analyzer will not operate properly if an incorrect size is used.

1. Go to **Unit [n] > Enable Manual Control**. Ensure a checkmark displays to the left of the menu item. If the analyzer schematic does not display, go to **Unit [n] > Show Instrument Schematic**.
2. Right click on the valve of the appropriate port. If the valve is open, click **Close** to close the valve.
3. Use a wrench to remove the connecting nut from the sample port. Pry out the filter (frit) and O-ring.



To avoid degassing problems, the frit should be clean and should not be touched with bare hands.

4. Place a new frit into the connecting nut.



5. Replace the filter and O-ring. Carefully reassemble the sample tube fitting and reinstall on the sample port. Tighten by hand, then with a wrench to prevent leaks.

## SAMPLE TUBE O-RING REPLACEMENT

It is important to maintain a vacuum-tight seal near the top of the sample tube stem. If an O-ring becomes worn or cracked, it does not provide a good seal and will need to be replaced.



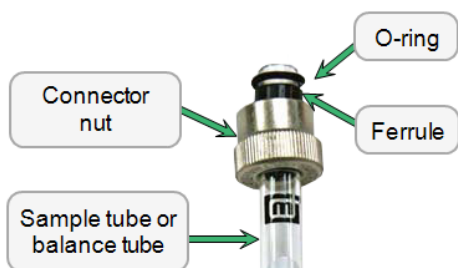
Before removing (or installing) a sample tube, ensure that the port valve is closed. Observe the analyzer schematic to verify valve status.



Do not allow the sample tube connector nut drop onto the bottom of the tube as it may break the tube.

Use the proper size frit for the sample port. The analyzer will not operate properly if an incorrect size is used.

1. Carefully remove the dewar from the analyzer. Take care not to bump the sample tube bulbs with the dewar during this process. Place the dewar aside.
2. Hold the sample tube firmly with one hand, loosen the sample tube connector nut by turning counter-clockwise.
3. Carefully pull the sample tube down until it is free from the port. It may be necessary to grasp the sample tube with both hands.



4. Remove the O-ring from the top of the sample tube and replace it with a new one.



If the O-ring remains inside the sample port, use the O-ring removal tool from the accessory kit (tool part number: 004-54618-00) or use a pair of tweezers or needle-nose pliers to remove it.

5. After the new O-ring is in place, insert the sample tube back into the sample port until it is fully seated.
6. Slide the sample tube connector nut up the tube until it comes in contact with the port fitting (the ferrule and O-ring will move along with the connector nut). Then, turning clockwise, hand tighten the connector nut to the sample connector.

## ***RECOVER FROM A POWER FAILURE***

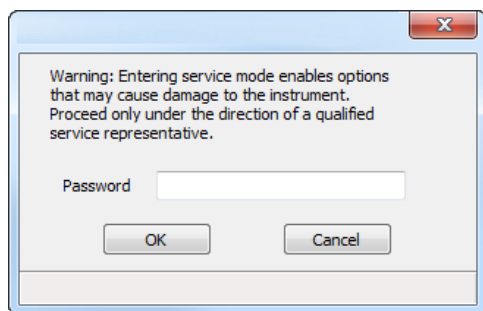
The analyzer saves entered and collected data in case of power failure. File parameters and any other data entered will still be present when power is restored. If an analysis was in progress when the power failure occurred, it will be canceled when the analyzer restarts. Any data collected during the analysis will still be present, but the analysis should be restarted in order to produce complete results.

## ***SERVICE TEST MODE***

### ***Options > Service Test Mode***

Service Test Mode is a password protected option used to perform certain service tests with the assistance of a trained Micromeritics Service Representative. This password is supplied by your Micromeritics Service Representative.

If a menu item is grayed out, it is usually an indication that Service Test Mode is required.



To exit Service Test Mode, go to ***Options > Service Test Mode*** and deselect the *Service Test Mode* option or close the application.

## ***OIL BASED VACUUM PUMP***

Log in to your [customer portal](#) to access the Vacuum Pump Guide.

## A ANALYZE SAMPLES WITH A TOTAL SURFACE AREA OF 1.0M<sup>2</sup> OR LESS

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The unique balanced measurement method used in the Gemini permits small amounts of surface area to be measured with nitrogen gas that otherwise would be measurable only with krypton. Low surface area samples often displace many times more nitrogen than they adsorb, especially if composed of low-density materials of large particle size. The standard, built-in, helium free-space difference measurement and mathematical compensation routine typically removes the effects of more than 99% of this sample displacement, but the small amount remaining uncompensated can still be significant.

A technique for adding a compensating volume with negligible surface area into the balance tube has been developed. It can reduce the size of the initial imbalance to low levels and allow precise measurement of adsorbed gas.

This technique can be performed by the following methods:

- Using glass beads in the balance tube
- Using filler rods in the sample and balance tubes

Some users find that a combination of the two methods — using glass beads and filler rods — produces even better results. This, however, will depend on the type of sample you are analyzing.

Filler rods and glass beads are included in the accessories kit. Either straight-wall or bulb sample tubes may be used.

### USING GLASS BEADS

This method typically produces the best results.

1. Place an appropriate quantity of sample in a clean sample tube.
2. Load a second sample tube of the same size with glass beads that have a total volume approximately the same as the sample volume.
  - a. Determine the volume (v) of the sample in cm<sup>3</sup>:

$$v = \frac{w}{\rho}$$

where

w = mass of sample (g)

ρ = density of sample (g/cm<sup>3</sup>); if density is unknown, refer to your laboratory handbook.

- b. Determine the number (n) of glass beads needed to equal the sample volume:

$$n = \frac{v}{0.014\text{cm}^3}$$

where

0.014cm<sup>3</sup> = approximate volume of one bead

3. Outgas the sample in the sample tube at an appropriate temperature for an appropriate amount of time.
4. Install the sample tube (containing the outgassed sample) onto the analysis port and the sample tube (containing the glass beads) onto the balance port.
5. Specify a one-point measurement ( $P/P_0 = 0.05$  to  $0.1$ ) so that the initial free-space measurement can quickly be determined; then perform the measurement.
6. Using the measured free space absolute value and the following relationship, determine the mass of glass beads to remove from (or add to) the balance tube to reduce the free-space imbalance:

$$\frac{W}{T}$$

where

2.515 g/cm<sup>3</sup> = density of glass beads

3.53 = thermal correction (no units)



Note that the volume of one glass bead is approximately 0.014 cm<sup>3</sup>. Therefore, if the measured free space is less than 0.02 cm<sup>3</sup>, it is unnecessary to correct the free space.

7. Use a beaker of warm water to bring the balance tube to room temperature before removing it from the balance port of the Gemini to remove (or add) glass beads. This prevents condensation of moisture from the laboratory atmosphere onto the cold glass beads.
8. Remove the balance tube:
  - If the measured free space is negative (–), add the calculated mass of glass beads into the balance tube.
  - If the measured free space is positive (+), remove the calculated mass of glass beads from the balance tube.
9. Reinstall the balance tube onto the balance port of the Gemini analyzer.



For subsequent samples of the same material, use the same mass of sample that was used for the initial sample so that the original bead quantity may be left undisturbed on the balance port.

10. Prepare the analysis dewar and place it on the elevator.
11. Close the sample compartment door and start the analysis.

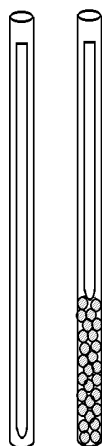


## USING FILLER RODS

1. Clean the sample tube, balance tube, and filler rods; label your sample and balance tubes.
2. Prepare the sample and place it into the sample tube; insert the shorter filler rod into the sample tube.

If using bulbous sample tubes, use the same size filler rod as used in the balance tube.

3. Ensure that the filler rods are equidistant from the top of each tube. If they are not, add or remove sample until they are the same distance from the top of the tube.



Packing of some powders may restrict gas access to the powder and cause slower equilibration and/or lower results.

4. Attach the sample tube to the analysis port and the balance tube to the balance port.
5. Prepare the analysis dewar and place it on the elevator.
6. Close the sample compartment door and start the analysis.

## ***USING GLASS BEADS AND FILLER RODS***

This method may further improve results, depending on the sample material being analyzed.

1. After determining and loading the amount of glass beads to use, insert a filler rod into the balance tube.
2. Insert a filler rod into the sample tube. Ensure that the filler rods are equidistant from the top of each tube. If they are not, add or remove sample until they are the same distance from the top of the tube.
3. If using bulbous sample tubes, use the same size filler rod as used in the balance tube.
4. Attach the sample tube to the analysis port and the balance tube to the balance port.
5. Prepare the analysis dewar and place it on the elevator.
6. Close the sample compartment door and start the analysis.

## ***B DFT MODELS***

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Theories are developed by scientists in an attempt to explain a class of observed behavior. In the experimental physical sciences, theories are often expressed in terms of a model that can be visualized and described mathematically. Early models of physical adsorption were quite simple, both conceptually and mathematically, for very practical reasons — hand computations were required. Today we can explore complex models that describe adsorption systems on the atomic scale of size and sub-picosecond time frame. This is not because scientists are smarter, but because of available tools. The DFT models are created by classical approaches to adsorption as well as models based on modern statistical thermodynamics.

### ***MODELS BASED ON STATISTICAL THERMODYNAMICS***

Included in this group are methods that model the adsorption system in terms of forces acting between individual molecules.

#### ***THEORETICAL BACKGROUND***

Traditional adsorption theories attempt to describe experimental adsorption isotherms with an isotherm equation containing a small number of parameters. At a minimum, these parameters include the extent of the surface, such as the monolayer capacity ( $V_m$ ), and the molar intensity of the gas-surface interaction, such as the Langmuir “K” constant or the BET “C” constant. In some equations, additional parameters take into account the lateral interaction of adsorbed molecules with each other. Other theories, such as the Dubinin-Astakhov approach, also include parameters for the effect of adsorbent porosity.

Instead of this classical kinetic or phenomenological approach, we can use a molecular-based statistical thermodynamic theory that allows us to relate the adsorption isotherm to the microscopic properties of the system: the fluid-fluid and fluid-solid interaction energy parameters, the pore size, the pore geometry, and the temperature.

The following example is given so that you may understand how such a theory is constructed:

A clean sample of a solid material containing slit-shaped pores of a single width is placed in an evacuated space. It is kept at a fixed temperature as a known quantity of pure argon gas is admitted into the space surrounding the sample. The pressure within the space is recorded over time. In this situation, the pressure falls rapidly from its initial value and gradually approaches a steady reading, called the equilibrium pressure. The amount adsorbed corresponds to the quantity of gas effectively removed from the gas phase by the solid surface. A graph that plots amount adsorbed versus equilibrium pressure is called an adsorption isotherm.

Under such conditions, the argon atoms that randomly enter the pore space feel the presence of the solid surface as the action of an external attractive force (the dispersion forces or Van der Waal's forces) and spend more time near the surface. As a result, the space near the surface acquires a greater average density of argon atoms than regions farther removed.

If the equilibrium distribution of the gas atoms near the surface could be described as a function of pressure and the molecular properties of the components of the system, then a model could be constructed for the adsorption isotherm for the system. Modern physical chemistry provides several ways to calculate this distribution. All these methods are based on the fundamental thermodynamic law that such a system adopts a configuration of minimum free energy at equilibrium. Also needed is a description of the pairwise interaction energy between atoms,  $U(s)$ , commonly given by a Lennard-Jones potential:

$$U(s) = 4\epsilon\left(\frac{\sigma}{s}\right)^{12} - \left(\frac{\sigma}{s}\right)^6$$

where

$\epsilon$  = a characteristic energy of the adsorptive,  
 $\sigma$  = the diameter of the adsorptive molecule, and  
 $s$  = the separation distance.

## ***MOLECULAR SIMULATION METHODS***

Two simulation techniques are commonly used to determine the distribution of gas molecules in a system in equilibrium: the molecular dynamics method and the Monte Carlo method. Both of these are used as reference methods because their results are considered exact.

### ***MOLECULAR DYNAMICS METHOD***

In the molecular dynamics method, the position and velocity of individual gas particles are calculated over time at very short intervals. This method takes into account both the forces acting between the gas particles themselves and those acting between the gas particles and the atoms of the simulated surface. As the simulated particles collide with each other and with the surface, the average concentration of particles in the space near the surface is calculated; this calculation yields the amount of gas adsorbed.

This method can be thought of as a way to determine the chronological record of the movement of each particle in the system using time steps of 10-14 seconds. Although the mathematics are simple, the number of calculations required for a system of even a few hundred particles is astronomical and challenges even the fastest computers.

## **MONTE CARLO METHOD**

In the Monte Carlo method, determination of the system equilibrium distribution begins with an assumption (which may be only approximate) about the initial configuration of particles in the system. The system is “equilibrated” through a process of randomly selecting one particle and conditionally moving it a random distance in a random direction.

If the move results in a configuration of *lower total energy*, then the move is completed and another particle is randomly selected to be moved.

If the move results in a configuration of *higher energy*, a probability for that event is calculated, and a random number between zero and one is generated. If the generated number is smaller than the probability of the event, then the move is accepted; otherwise, another particle is selected and the process is repeated. This process continues until the average total energy of the system no longer decreases; at this point, average configuration data are accumulated to yield the mean density distribution of particles in the system.

Monte Carlo simulations require considerably less computation time than molecular dynamic simulations and can yield the same results; however, neither method provides a really practical way to calculate complete isotherms.

## **DENSITY FUNCTIONAL FORMULATION**

*Density functional theory* offers a practical alternative to both molecular dynamic and Monte Carlo simulations. When compared to reference methods based on molecular simulation, this theory provides an accurate method of describing inhomogeneous systems yet requires fewer calculations. Because the density functional theory provides accuracy and a reduced number of calculations, it is the basis embodied in the DFT models.

The system being modeled consists of a single pore represented by two parallel walls separated by a distance  $H$ . The pore is open and immersed in a single component fluid (adsorptive) at a fixed temperature and pressure. Under such conditions, the fluid responds to the walls and reaches an equilibrium distribution. In this condition (by the definition of equilibrium), the chemical potential at every point equals the chemical potential of the bulk fluid. The bulk fluid is a homogenous system of constant density; its chemical potential<sup>1)</sup> is determined by the pressure of the system using well-known equations. The fluid near the walls is not of constant density; its chemical potential is composed of several position-dependent contributions that must total at every point to the same value as the chemical potential of the bulk fluid.

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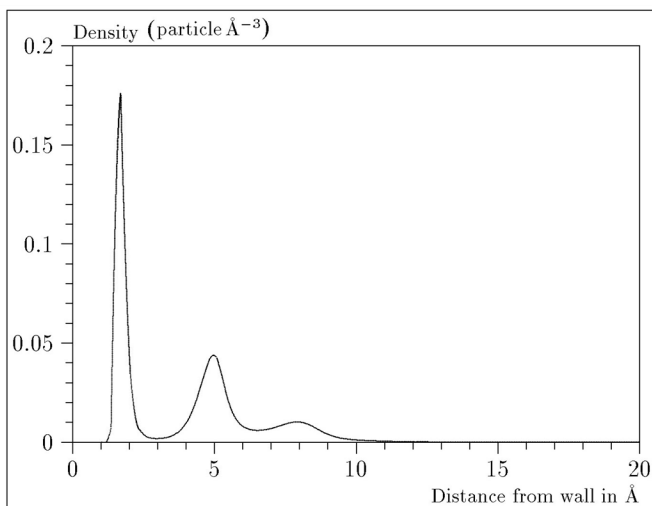
<sup>1)</sup> Chemical potential may be thought of as the energy change felt by a probe particle when it is inserted into the system from a reference point outside the system. It can also be defined as the partial derivative of the grand potential energy with respect to density (or concentration).

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As noted previously, at equilibrium, the whole system has a minimum (Helmholtz) free energy, known thermodynamically as the grand potential energy (GPE). Density functional theory describes the thermodynamic grand potential as a functional of the single-particle density distribution; therefore, calculating the density profile that minimizes the GPE yields the equilibrium density profile. The calculation method requires the solution of a system of complex integral equations that are implicit functions of the density vector. Since analytic solutions are not possible, the problem must be solved using iterative numerical methods. Although calculations using these methods still require supercomputing speed, the calculation of many isotherm pressure points for a wide range of pore sizes is a feasible task. The complete details of the theory and the mathematics can be found in the papers listed under [DFT Model References on page B - 17](#).

The following graphs and accompanying text illustrate the results of using density functional theory to predict the behavior of a model system.

Figure 1 shows the density profile for argon at a carbon surface as calculated by density functional theory for a temperature of 87.3 K and a relative pressure of about 0.5.

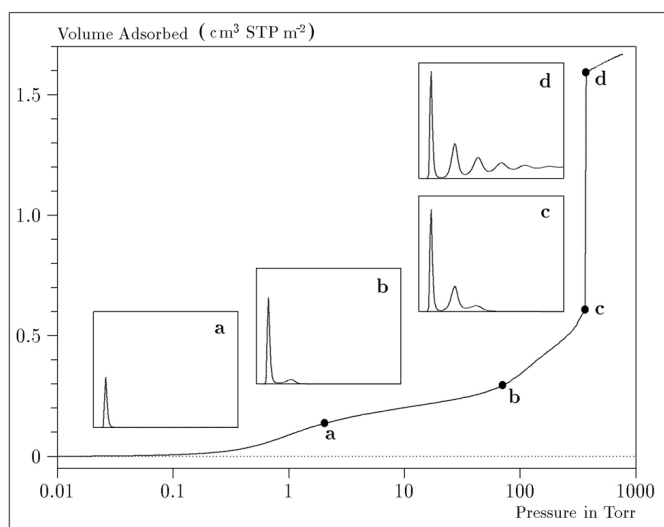


**Figure 1. Density Profile for Argon on Carbon at 87.3 K and a Relative Pressure of 0.5**

This figure represents a cross-section of the region near the surface. Note the layerwise distribution of adsorbate; the first monolayer is sharply defined and a third layer can be distinguished. The area under the profile curve represents the amount adsorbed per unit area at this pressure. The positions of the maxima are separated by a distance determined by the size of the adsorptive atom.

Given the density profile, the amount adsorbed at the stated pressure can be easily calculated as the integral over the profile. Repeating this calculation over a range of pressures yields the adsorption isotherm for the model. If the value of  $H$  is very large, the isotherm obtained corresponds to that of an external, or *free*, surface. If  $H$  is smaller, a range of pressures is reached where two minima exist for the grand potential, showing the presence of two metastable phases having different density distributions but the same chemical potential. The phase with the lower GPE is the stable one. As the pressure is increased, a point is reached where the other phase becomes the stable one. This phase transition reflects condensation of adsorbate in the pore; the pressure at which it occurs is called the *critical pore-filling pressure*. This pressure is analogous to the condensation pressure predicted by the Kelvin equation in the classical model of pore filling.

Figure 2 shows how the profiles change with pressure for a model pore with  $H = 40$  angstroms. The inset shows the density profiles for the corresponding points of the isotherm.



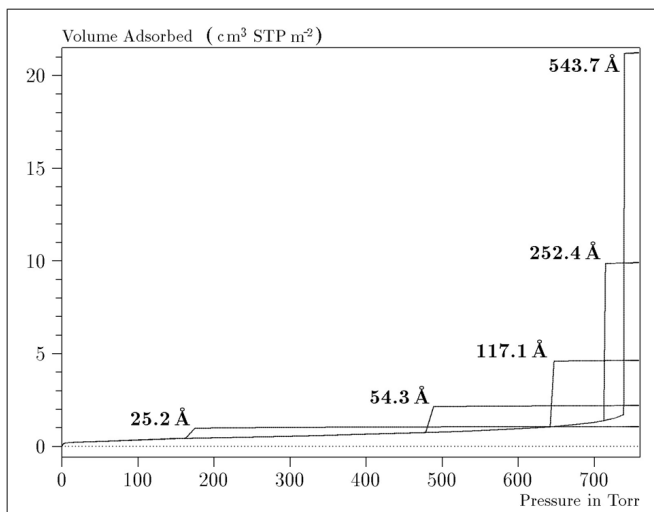
**Figure 2. Model Isotherm for Argon at 87.3 K in a 40 Å Slit in a Carbon Substrate**

The profiles show the density distribution from one wall to the center of the slit; the other half of the distribution is a mirror image of the profile shown.

As the pressure is first increased from zero, almost all the adsorbed atoms occupy a position close to the surface.

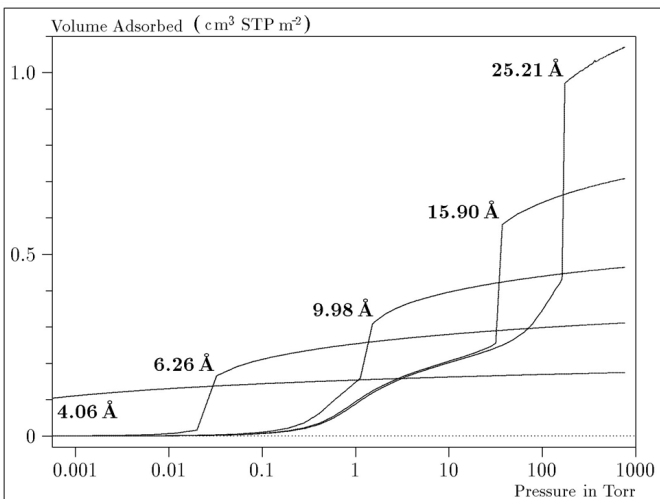
- Inset **a** shows the profile corresponding to point a on the isotherm where the surface is about half covered.
- At point **b**, the first layer is so full that it is more favorable for atoms to start a new layer.
- At point **c**, a third layer is forming. Point **c**, for this size slit, is the critical pore-filling pressure. In inset **c**, the profile shows the density decreasing to near zero (actually the bulk gas density) at 4 or 5 molecular diameters from the surface.
- Inset **d** shows the profile converging on a density similar to that of bulk liquid argon in the center of the pore, indicating a phase transition.

Note that the adsorption isotherms for pores larger than the one shown in the previous graph is identical up to point **c**. The lower branch of the isotherm simply continues to a higher pressure for larger pores. This trend is illustrated in the Figure 3, where isotherms for some larger size pores are shown. It is clear that pore size is uniquely characterized by a corresponding critical pore-filling pressure. At large pore sizes, density functional theory produces results for the critical filling pressures that are in good agreement with those produced by the Kelvin equation.



**Figure 3. Model Isotherms for Some Larger Pore Widths Argon on Carbon at 87.3 K**

Figure 4 shows model isotherms for pores in the micropore size range. Note the logarithmic scale for pressure.



**Figure 4. Model Isotherms in the Micropore Size Range of Pore Width Argon on Carbon at 87.3 K**



Pores of 4 Å width, barely larger than the argon atom (3.38 Å), fill at pressures below 1 millitorr. Pores below 15 Å fill before a monolayer is completed on the surface of the larger pores. In the micropore size range, the pore volume fills more gradually with pressure and the total shape of the isotherm is important in characterizing the pore size.

### **Models Included**

#### **Non-Local Density Functional Theory with Density-Independent Weights**

##### **N2 - DFT Model**

##### **AR - DFT Model**

<b>Geometry:</b>	Slit
<b>Substrate:</b>	Carbon (graphite)
<b>Category:</b>	Porosity
<b>Method:</b>	Nitrogen at 77 K; Argon at 87 K

Using the methods of non-local density functional theory, two sets of isotherms have been calculated to serve as kernel functions for the characterization of porous solids from adsorption data. The model isotherms are stored in binary format files. These models assume a *slit-like pore geometry*. The pore size range from 4.0 to 4000 Å is covered in 91 classes in a geometric progression. The class intervals are rounded to the nearest 0.02 molecular diameters. A model for the free or external surface is included to account for unfilled pores. Each of the 92 model isotherms has been calculated at 181 pressure points from near  $1 \times 10^{-6}$  to near 1.00 relative pressure.

These models are identical to those supplied with the original DOS version of DFT software. Some slight difference from the DOS results may be noted when they are applied to the same data due to improvements in the deconvolution algorithm and better regularization of the current software.

#### **Non-Local Density Functional Theory with Density-Dependent Weights**

##### **N2 - Modified Density Functional**

<b>Geometry:</b>	Free surface
<b>Substrate:</b>	Surface energy
<b>Method:</b>	Nitrogen at 77K

Using the modified Tarazona prescription described by Olivier (see [DFT Model References on page B - 17](#) [items 3 and 4]), model isotherms were calculated for a wide range of adsorptive energies to a relative pressure of 0.6. The model makes no provision for pore filling in the micropore region. If the sample solid contains small mesopores, the isotherm data should be truncated (using the *Select Data Points* window) to a suitably low relative pressure to avoid trying to fit this region; mesopore filling reports as a large area of low energy in the calculated distribution of adsorptive potential.

The surface energy is reported in terms of the effective Lennard-Jones interaction parameter, ie, for the adsorptive / adsorbent pair divided by Boltzmann constant. The units are therefore Kelvin.

**N2 - Cylindrical Pores - Oxide Surface****AR - Cylindrical Pores - Oxide Surface**

<b>Geometry:</b>	Cylinder
<b>Substrate:</b>	Oxide
<b>Category:</b>	Porosity
<b>Method:</b>	Nitrogen at 77K; Argon at 87K

Model isotherms were calculated using a combination of statistical mechanical calculations and experimental observations for macroporous silicas and MCM-41 mesoporous silicas as well as zeolites. The pore-filling pressures were determined as a function of the pore size from adsorption isotherms on MCM-41 materials characterized by X-ray and other techniques. The variation of the pore fluid density with pressure and pore size has been accounted for by density functional theory calculations. The N2 model reports pore sizes ranging from 3.8 to 387 Å and the AR model from 3.8 to over 500 angstroms.

**References:** M. Jaroniec, M. Kruk, J.P. Olivier, and S. Koch, "A New Method for the Accurate Pore Size Analysis of MCM-41 and Other Silica-Based Mesoporous Materials," Proceedings of COPS-V, Heidelberg, Germany (1999).

**N2 – Cylindrical Pores – Pillared Clay Surface (Montmorillonite)**

<b>Geometry:</b>	Cylinder
<b>Substrate:</b>	Crystalline Silicate
<b>Category:</b>	Porosity
<b>Method:</b>	Nitrogen at 77K

Model isotherms were calculated using a combination of statistical thermodynamic Non-Local Density Functional Theory (NLDFT) calculations and experimental isotherms for reference samples of montmorillonite. The construction method for the hybrid models was analogous to that described in the first reference below (Jaroniec et al, 1999). The additional references add additional theoretical details as well as examples of the application of the model to pillared clay catalysts. This model reports pore widths from 3.8 to 387 angstroms.

**References:** Mietec Jaroniec, Michal Kruk, James P. Olivier and Stefan Koch, "A New Method for the Characterization of Mesoporous Silicas," Proceedings of COPS-V, 1999, Studies in Surface Science, Vol 128, *Characterization of porous Solids V*, Unger, et al, Eds, Elsevier, Amsterdam, 2000.

James P. Olivier and Mario L. Occelli, "Surface Area and Microporosity of a Pillared Interlayered Clay (PILC) from a Hybrid Density Functional Theory

(DFT) Method,” *The Journal of Physical Chemistry B*; 2001, 105(3), 623-629.

M. L. Occelli, J. P. Olivier, J. A. Perdigon-Melon, and A. Auroux, “Surface Area, Pore Volume Distribution, and Acidity in Mesoporous Expanded Clay Catalysts from Hybrid Density Functional Theory (DFT) and Adsorption Microcalorimetry Methods,” *Langmuir* 2002, 18, 9816-9823.9b.

James P. Olivier, “The Importance of Surface Heterogeneity in Developing Characterization Methods.” *6<sup>th</sup> International Symposium on the Characterization of Porous Solids*, Studies in Surface Science and Catalysis 144, Elsevier, 2002.

James P. Olivier and Mario L. Occelli, “Surface Area and Microporosity of Pillared Rectorite Catalysts from a Hybrid Density Functional Theory Method,” *Microporous and Mesoporous Materials* 2003, 57, 291-296.

### **C02 - DFT Model**

**Geometry:** Slit  
**Substrate:** Carbon  
**Category:** Porosity  
**Method:** Carbon dioxide at 273 K

Model isotherms were calculated using the non-local prescription of Tarazona, employing molecular parameters derived from the known bulk properties of carbon dioxide.

### **AR - Modified Density Functional Model**

**Geometry:** Free surface  
**Substrate:** Any  
**Category:** Surface energy  
**Method:** Argon at 87K

This model was produced in the same manner as the N2 Modified Density Functional model listed earlier, except applicable to argon adsorbed at 87.3 K.

### **N2 - Tarazona NLDFT, Esf = 30.0K**

**Geometry:** Cylinder  
**Substrate:** Oxide  
**Category:** Porosity  
**Method:** Nitrogen at 77K

Model isotherms were calculated using the prescriptions of Tarazona for density dependent weighting functions and a cylindrical pore geometry. The wall potential used is  $k = 30$  K, typical for a silica or alumina surface.

This model file is particularly useful for sizing zeolites or zeolite containing materials that have substantial micropore volume. The reported pore size range is 3.8 to 387 angstroms.

**References:** P. Tarazona, Phys. Rev. A 31: 2672 (1985).  
Idem, Phys. Rev. A 32: 3148 (1985).  
P. Tarazona, U. M. B. Marconi, and R. Evans, Mol. Phys. 60: 573 (1987).

#### **N2 - Carbon Slit Pores by NLDFT**

#### **Ar - Carbon Slit Pores by NLDFT**

**Geometry:** Slit  
**Substrate:** Carbon  
**Category:** Porosity  
**Method:** Nitrogen at 77K; Argon at 87K

Model isotherms were calculated using the prescriptions of Tarazona for density dependent weighting functions and a slit-like pore geometry. These models are slightly different from N2-DFT and Ar-DFT models that were calculated using NLDFT with density independent weighting functions.

The reported pore size range is from 3.5 to 1000 angstroms.

**References:** P. Tarazona, Phys. Rev. A 31: 2672 (1985).  
Idem, Phys. Rev. A 32: 3148 (1985).  
P. Tarazona, U. M. B. Marconi, and R. Evans, Mol. Phys. 60: 573 (1987).

#### **N2 - Carbon Finite Pores, As=6, 2D-NLDFT**

#### **Ar - Carbon Finite Pores, As=6, 2D-NLDFT**

**Geometry:** Finite Slit  
**Substrate:** Carbon  
**Category:** Porosity  
**Method:** Nitrogen at 77K; Argon at 87K

Model isotherms were calculated using the prescriptions of Tarazona for density dependent weighting functions assuming 2D model of finite slit pores having a diameter-to-width aspect ratio of 6.

This model is particularly useful for microporous carbon materials. The reported pore size range is from 3.5 to 250 angstroms.

**References:** Jacek Jagiello and James P. Olivier. "A simple two-dimensional NLDFT model of gas adsorption in finite carbon pores. Application to pore structure analysis.," The Journal of Physical Chemistry C, 113(45):19382-19385, 2009.

**N2 - Carbon Finite Pores, As=12, 2D-NLDFT**

**Ar - Carbon Finite Pores, As=12, 2D-NLDFT**

**Geometry:** Finite Slit  
**Substrate:** Carbon  
**Category:** Porosity  
**Method:** Nitrogen at 77K; Argon at 87K

Model isotherms were calculated using the same methods and assumptions that were used in the model above except in this model, the aspect ratio is equal to 12.

These two finite pore models may be used as a research tool in conjunction with independent analytical techniques such as high-resolution transmission electron microscopy (HRTEM) and/or X-ray diffraction (XRD) to obtain comprehensive information about the structure of studied carbon material.

**References:** Jacek Jagiello and James P. Olivier. "A simple two-dimensional NLDFT model of gas adsorption in finite carbon pores. Application to pore structure analysis.," The Journal of Physical Chemistry C, 113(45):19382-19385, 2009.

**N2 - Carbon Cylinder, single-wall nanotube by NLDFT**

**Ar - Argon Cylinder, single-wall nanotube by NLDFT**

**Geometry:** Cylinder  
**Substrate:** Carbon  
**Category:** Porosity  
**Method:** Nitrogen at 77 K; Argon at 87 K

Model isotherms were calculated using the prescriptions of Tarazona for density dependent weighting functions and cylindrical pore geometry. The pore wall potential is described by the Lennard-Jones potential of interaction between a gas molecule and the graphitic surface of an infinitely long cylinder.

This model is particularly useful for characterizing carbon single-wall nanotubes. The reported pore size range is from 3.5 to 1000 angstroms.

**References:** P. Tarazona, Phys. Rev. A 31: 2672 (1985).  
Idem, Phys. Rev. A 32: 3148 (1985).  
P. Tarazona, U. M. B. Marconi, and R. Evans, Mol. Phys. 60: 573 (1987).

**N2 - Carbon Cylinder, multi-wall nanotube by NLDFT****Ar - Argon Cylinder, multi-wall nanotube by NLDFT**

<b>Geometry:</b>	Cylinder
<b>Substrate:</b>	Carbon
<b>Category:</b>	Porosity
<b>Method:</b>	Nitrogen at 77 K; Argon at 87 K

Model isotherms were calculated using the prescriptions of Tarazona for density dependent weighting functions and cylindrical pore geometry. The pore wall potential is described by the Lennard-Jones potential of interaction between a gas molecule and multiple concentric graphitic surfaces of infinitely long cylinders.

This model is particularly useful for characterizing carbon multi-wall nanotubes. The reported pore size range is from 3.5 to 1000 angstroms.

**References:** P. Tarazona, Phys. Rev. A 31: 2672 (1985).  
Idem, Phys. Rev. A 32: 3148 (1985).  
P. Tarazona, U. M. B. Marconi, and R. Evans, Mol. Phys. 60: 573 (1987)

**Ar - Zeolites H-Form by NLDFT**

<b>Geometry:</b>	Cylinder
<b>Substrate:</b>	Zeolite
<b>Category:</b>	Porosity
<b>Method:</b>	Argon at 77 K

Model isotherms were calculated using the prescriptions of Tarazona for density dependent weighting functions and cylindrical pore geometry. The pore wall potential is described by the Lennard-Jones potential of interaction between a gas molecule and the oxide surface of an infinitely long cylinder.

This model is particularly useful for characterizing oxides and H<sup>+</sup> and (NH<sub>4</sub>)<sup>+</sup> exchanged zeolites. The reported pore size range is from 3.5 to 300 angstroms.

**Ar - Zeolites Me-Form by NLDFT**

<b>Geometry:</b>	Cylinder
<b>Substrate:</b>	Zeolite
<b>Category:</b>	Porosity
<b>Method:</b>	Argon at 77 K

Model isotherms were calculated using the prescriptions of Tarazona for density dependent weighting functions and cylindrical pore geometry. The pore wall potential is described by the Lennard-Jones potential of interaction between a gas molecule and the oxide surface of an infinitely long cylinder.

This model is similar to the model above, but it more appropriate is for characterizing alkali metal exchanged zeolites. The reported pore size range is from 3.5 to 300 angstroms.

## ***MODELS BASED ON CLASSICAL THEORIES***

Both surface energy distribution and pore size distribution may be evaluated using classical approaches to model kernel functions for use with equation (1) of the DFT Theory. ( Log in to your [customer portal](#) to access the Calculations document.) Be aware that the deconvolution method only provides a fitting mechanism; it does not overcome any inherent shortcomings in the underlying theory.

### ***SURFACE ENERGY***

The use of classical theories to extract adsorptive potential distribution is mostly of historical interest. At a minimum, the equation must contain a parameter dependent on adsorption energy and another dependent on monolayer capacity or surface area. This is sufficient to permit the calculation of the set of model isotherms that is used to create a library model. The Langmuir equation has been used in the past, as have the Hill-de Boer equation and the Fowler-Guggenheim equation. All of these suffer from the fact that they only describe monolayer adsorption, whereas the data may include contributions from multilayer formation.

### ***PORE SIZE***

It is well established that the pore space of a mesoporous solid fills with condensed adsorbate at pressures somewhat below the prevailing saturated vapor pressure of the adsorptive. When combined with a correlating function that relates pore size with a critical condensation pressure, this knowledge can be used to characterize the mesopore size distribution of the adsorbent. The correlating function most commonly used is the Kelvin equation. Refinements make allowance for the reduction of the physical pore size by the thickness of the adsorbed film existing at the critical condensation pressure. Still further refinements adjust the film thickness for the curvature of the pore wall.

The commonly used practical methods of extracting mesopore distribution from isotherm data using Kelvin-based theories, such as the BJH method, were for the most part developed decades ago and were designed for hand computation using relatively few experimental points. In general, these methods visualize the incremental decomposition of an experimental isotherm, starting at the highest relative pressure or pore size. At each step, the quantity of adsorptive involved is divided between pore emptying and film thinning processes and exactly is accounted for. This computational algorithm frequently leads to inconsistencies when carried to small mesopore sizes. If the thickness curve used is too steep, it finally will predict a larger increment of adsorptive for a given pressure increment than is actually observed; since a negative pore volume is non-physical, the algorithm must stop. Conversely, if the thickness curve used underestimates film thinning, accumulated error results in the calculation of an overly large volume of (possibly nonexistent) small pores.



The use of equation (1) represents an improvement over the traditional algorithm. Kernel functions corresponding to various classical Kelvin-based methods have been calculated for differing geometries and included in the list of models.

## ***MODELS INCLUDED***

### **Kelvin Equation with Halsey Thickness Curve**

#### **N2 - Halsey Thickness Curve**

<b>Geometry:</b>	Slit
<b>Substrate:</b>	Average
<b>Category:</b>	Porosity
<b>Method:</b>	Nitrogen 77 K

The kernel function is calculated using the Halsey equation with standard parameters:

$$t = 3.54 \left( \frac{-5.00}{\ln(P/P_0)} \right)^{1/3}$$

The nitrogen properties used in the Kelvin equation are:

<b>Surface tension =</b>	8.88 dynes cm <sup>-1</sup>
<b>Molar density =</b>	0.02887 g cm <sup>-3</sup>

#### **N2 - Halsey Thickness Curve**

<b>Geometry:</b>	Cylinder
<b>Substrate:</b>	Average
<b>Category:</b>	Porosity
<b>Method:</b>	Nitrogen 77 K

The calculation is the same as above except that cylindrical geometry is assumed.

<b>Reference:</b>	G. Halsey, J. Chem. Phys 16, 931 (1948).
-------------------	--

### **Kelvin Equation with Harkins and Jura Thickness Curve**

#### **N2 - Harkins and Jura Thickness Curve**

<b>Geometry:</b>	Slit
<b>Substrate:</b>	Average
<b>Category:</b>	Porosity

**Method:** Nitrogen 77 K

The kernel function is calculated using the Harkins and Jura equation with standard parameters:

$$t = 3.54 \left( \frac{13.99}{0.034 - \log(P/P_0)} \right)^{1/2}$$

The nitrogen properties used in the Kelvin equation are:

**Surface tension =** 8.88 dynes cm<sup>-1</sup>

**Molar density =** 0.02887 g cm<sup>-3</sup>

## **N2 - Harkins and Jura Thickness Curve**

**Geometry:** Cylinder

**Substrate:** Average

**Category:** Porosity

**Method:** Nitrogen 77 K

The calculation is the same as above except that cylindrical geometry is assumed.

**References:** W. D. Harkins and G. Jura, J.A.C.S. 66, 1366 (1944).  
J. H. DeBoer et al., J. Colloid and Interface Sci. 21, 405 (1966).

## **Kelvin Equation with Broekhoff-de Boer Thickness Curve**

### **N2 - Broekhoff-de Boer Model**

**Geometry:** Cylinder

**Substrate:** Average

**Category:** Porosity

**Method:** Nitrogen 77 K

The kernel function is calculated using the Broekhoff-de Boer equation with standard parameters:

$$\log\left(p/p^0\right) = \frac{-16.11}{t^2} + 0.1682^{-0.1137t}$$

The nitrogen properties used in the Kelvin equation are:

**Surface tension =** 8.88 dynes cm<sup>-1</sup>

**Molar density =** 0.02887g cm<sup>-3</sup>

### **N2 - Broekhoff-de Boer Model**

**Geometry:** Cylinder  
**Substrate:** Average  
**Category:** Porosity  
**Method:** Nitrogen 77 K

The calculation is similar to the above except that cylindrical geometry is assumed, and the film thickness depends on pore size (see reference).

**References:** Specifically, equations 20 and 21 in: J.C.P. Broekhoff and J.H. de Boer, "The Surface Area in Intermediate Pores," Proceedings of the International Symposium on Surface Area Determination, D.H. Everett, R.H. Ottwill, eds., U.K. (1969).

## ***DFT MODEL REFERENCES***

The papers listed below provide additional information on DFT models:

1. "Determination of Pore Size Distribution from Density Functional Theoretic Models of Adsorption and Condensation within Porous Solids," J.P. Olivier and W.B. Conklin, Micromeritics Instrument Corp; presented at the International Symposium on the Effects of Surface Heterogeneity in Adsorption and Catalysts on Solids, Kazimierz Dolny, Poland (July 1992).
2. "Classification of Adsorption Behavior: Simple Fluids in Pores of Slit-shaped Geometry," Perla B. Balbuena and Keith E. Gubbins, *Fluid Phase Equilibria*, 76, 21-35, Elsevier Science Publishers, B.V., Amsterdam (1992).
3. "Modeling Physical Adsorption on Porous and Nonporous solids Using Density Functional Theory," J.P. Olivier, *Journal of Porous Materials*, 3, 9-17 (1995).
4. "The Determination of Surface Energetic Heterogeneity Using Model Isotherms Calculated by Density Functional Theory," J.P. Olivier; presented at the Fifth International Conference on the Fundamentals of Adsorption, Pacific Grove, CA (1995).

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## C EXPORTED DATA EXAMPLE

### Sample Information

#### Sample Information Parameters

Sample: SiAl Reference Material 004/16821/00 - Full  
 Operator: SV  
 Submitter: Micromeritics  
 Mass type: Entered  
 Mass: 0.3054 g  
 Density: 1.000 g/cm<sup>3</sup>  
 Type of data: Automatically collected  
 Instrument type: 2390  
 Original instrument type: 2390  
 Comments:

### Sample Tube

#### Sample Tube Options

Sample Tube: Sample Tube  
 Warm free space: 1.0000 cm<sup>3</sup>  
 Cold free space: 1.0000 cm<sup>3</sup>  
 Non-ideality factor: 0.0000620  
 Use isothermal jacket: No  
 Use filler rod: No  
 Vacuum seal type: None

### Degas Conditions

#### Degas Conditions Options

Description: Degas Conditions

#### Evacuation Phase

Temperature Ramp Rate: 1.0 °C/min  
 Target Temperature: 30 °C  
 Evacuation Rate: 0.67 kPa/s  
 Unrest. Evacuation from: 0.67 kPa  
 Vacuum Setpoint: 1.333224e-003 kPa  
 Evacuation Time: 0 min

#### Heating Phase

Sample Prep: Stage	Temperature (°C)	Ramp Rate (°C/min)	Time (min)
1	90	10	60
2	350	10	240

#### Evacuation and Heating Phases

Hold Pressure: 13.3 kPa

Backfill Sample Tube: Automatic

## Analysis Conditions

### Analysis Conditions Options

Description: BET

#### Pressure Table

Relative Pressure (p/p°)	Rel. Pressure Increment (p/p°)
0.050000000	
0.070000000	
0.090000000	
0.110000000	
0.135000000	
0.150000000	
0.175000000	
0.200000000	
0.225000000	
0.250000000	
0.275000000	
0.300000000	
0.320000000	
0.340000000	
0.380000000	
0.420000000	
0.460000000	
0.500000000	
0.540000000	
0.580000000	
0.620000000	
0.660000000	
0.700000000	
0.740000000	
0.780000000	
0.820000000	
0.860000000	
0.890000000	
0.900000000	
0.910000000	
0.920000000	
0.930000000	
0.940000000	
0.950000000	
0.955000000	
0.960000000	

#### Preparation

Evacuation rate: 133.32 kPa/min  
Evacuation time: 20.00 min

#### Free Space

Measured before analysis

#### p° and Temperature

p° type: Measured in Psat tube for each point  
Temperature type: Entered  
Temperature: 77.300 K

#### Analysis Method

Analysis Mode: Equilibration  
Equilibration time: 10 s

## Adsorptive Properties

### Adsorptive Properties Options

Adsorptive: Nitrogen (N<sub>2</sub>)  
 Non-condensing adsorptive: No  
 Maximum manifold pressure: 139.989 kPa  
 Molecular cross-sectional area: 0.162 nm<sup>2</sup>  
 Non-ideality factor: 0.0000620  
 Density conversion factor: 0.0015468  
 Adsorbate molecular weight: 28.01  
 Dosing method: Normal

### Psat vs. Temperature Table

	Saturation Pressure (kPa)	Temperature (K)
1	600.193	75.400
2	634.512	75.850
3	674.383	76.350
4	720.420	76.900
5	742.119	77.150
6	759.833	77.350
7	777.867	77.550
8	805.525	77.850
9	853.268	78.350
10	903.122	78.850

## Report Options

### Report Options

Description: Sil-Alum Reference Material 004/16821/00  
 Show report title: Yes  
 Report title: Micromeritics Instrument Corporation  
 Show graphic: Yes  
 Graphic file: miclogo.emf  
 Graphic width: 2.000 inches  
 Graphic height: 0.250 inches

### Surface Area

Single-point BET: Yes  
 Multi-point BET: Yes  
 Langmuir: No  
 t-Plot Micropore: Yes  
 t-Plot External: Yes  
 BJH Cumulative Adsorption: Yes  
 BJH Cumulative Desorption: No  
 D-H Cum. Adsorption: No  
 D-H Cum. Desorption: No

### Pore Volume

Adsorption Total: Yes  
 Relative Pressure: 0.997721194 p/p°  
 Desorption Total: No  
 t-Plot Micropore: Yes  
 BJH Cum. Adsorption: Yes  
 BJH Cum. Desorption: No  
 D-H Cum. Adsorption: No  
 D-H Cum. Desorption: No

**Pore Size**

Average pore diameter: Yes  
BJH adsorption avg.: Yes  
BJH desorption avg.: Yes  
D-H adsorption avg.: No  
D-H desorption avg.: No

**Other**

Freundlich: No  
Temkin: No  
Alpha-S method: No  
DFT Pore Size: No  
DFT Surface Energy: No  
Nanoparticle Size: No

**Horvath-Kawazoe**

Maximum pore volume: No  
Median pore width: No

**Dubinin-Radushkevich**

Micropore surface area: No  
Monolayer capacity: No

**Dubinin-Astakhov**

Micropore surface area: No  
Limiting micropore volume: No

**MP-Method**

Cumulative surface area: No  
Cumulative pore volume: No  
Average pore hydraulic radius: No

**Pass/Fail**

Pass/Fail 1: S A:Multi-point BET:  
High Range Value: 222.0000 m<sup>2</sup>/g  
High Range Message: 004/16821/00 Lot A-501-48 Upper Limit  
Low Range Value: 210.0000 m<sup>2</sup>/g  
Low Range Message: 004/16821/00 Lot a-501-48 Lower Limit  
Pass/Fail 2: S A:Single-point BET:  
High Range Value: 217.0000 m<sup>2</sup>/g  
High Range Message: 004/16821/00 Lot A-501-48 Upper Limit  
Low Range Value: 205.0000 m<sup>2</sup>/g  
Low Range Message: 004/16821/00 Lot A-501-48 Lower Limit  
Pass/Fail 3: P V:Adsorption Total:  
High Range Value: 0.680000 cm<sup>3</sup>/g  
High Range Message: 004/16821/00 Lot A-501-48 Upper Limit  
Low Range Value: 0.520000 cm<sup>3</sup>/g  
Low Range Message: 004/16821/00 Lot A-501-48 Lower Limit  
Pass/Fail 4: P S:Average pore diameter:  
High Range Value: 129.000 Å  
High Range Message: 004/16821/00 Lot A-501-48 Upper Limit



**Pass/Fail**

Low Range Value: 99.000 Å  
Low Range Message: 004/16821/00 Lot A-501-48 Lower Limit

Isotherm: Yes

**Isotherm Reports**

Elapsed time: Yes  
Time between points: No  
Plot adsorption branch: Yes  
Plot desorption branch: Yes  
Quantity adsorbed: Per Gram

Tabular report selected: Yes  
Linear plot selected: Yes  
Logarithmic plot selected: No  
Linear absolute plot selected: No  
Pressure composition plot selected: No

**Isotherm Linear Plot axis data**

Plot curve: Yes  
Plot points: Yes  
Overlay samples: No  
Autoscale X axis: Yes  
Autoscale Y axis: Yes

BET: Yes

**BET Reports**

Tabular report selected: Yes  
Transform plot selected: Yes  
Overlay Samples: No  
Autoscale X axis: Yes  
Autoscale Y axis: Yes

Isotherm plot selected: No  
Overlay Samples: No  
Autoscale X axis: Yes  
Autoscale Y axis: Yes

Fit pressure range: 0.000000000 to 1.000000000 p/p<sup>o</sup>

Langmuir: No

Freundlich: No

Temkin: No

t-Plot: Yes

**t-Plot Reports**

Tabular report selected: Yes  
Plot report selected: Yes  
Minimum thickness: 3.5000 Å  
Maximum thickness: 5.0000 Å  
Surface Area Correction Factor: 1.000  
Harkins and Jura Thickness Equation  
Numerator: 13.9900  
Denominator: 0.0340  
Exponent: 0.500  
Surface Area Type: BET

Alpha-S Method: No

f-Ratio Method: No

BJH Adsorption: Yes

#### BJH Adsorption Reports

Halsey Thickness Equation

Numerator: -5.000

Multiplier: 3.540

#### BJH Adsorption Reports

Exponent: 0.333

Minimum Pore Width: 17.000 Å

Maximum Pore Width: 3,000.000 Å

Fraction of pores open at both ends: 0.00

Smooth differential values: Yes

Cumulative reports: Larger

Adsorbate Property

Adsorptive Factor (Å)

N2 9.53000

AR 10.44000

BJH Tabular Report Yes

Cumulative Pore Volume Yes

dV/dw Pore Volume No

dV/dlog(w) Pore Volume Yes

Cumulative Pore Area Yes

dA/dw Pore Area No

dA/dlog(w) Pore Area Yes

#### BJH Tabular Report

Tabular data defined by: Collected points

Tabular Report Columns

Column 1: w Interval

Column 2: Average w

Column 3: Incremental Pore Volume

Column 4: Cumulative Pore Volume

Column 5: Incremental Pore Area

Column 6: Cumulative Pore Area

#### Cumulative Pore Volume axis data

Overlay samples: No

Autoscale X axis: Yes

Autoscale Y axis: Yes

#### Cumulative Pore Volume Overlay

Overlay plot: None

#### dV/dlog(w) Pore Volume axis data

Overlay samples: No

Autoscale X axis: Yes

Autoscale Y axis: Yes

**dV/dlog(w) Pore Volume Overlay**

Overlay plot: None

**Cumulative Pore Area axis data**

Overlay samples: No

Autoscale X axis: Yes

Autoscale Y axis: Yes

**Cumulative Pore Area Overlay**

Overlay plot: None

**dA/dlog(w) Pore Area axis data**

Overlay samples: No

Autoscale X axis: Yes

Autoscale Y axis: Yes

**dA/dlog(w) Pore Area Overlay**

Overlay plot: None

BJH Desorption: No

Dollimore-Heal Adsorption: No

Dollimore-Heal Desorption: No

Horvath-Kawazoe: No

DFT Pore Size: No

DFT Surface Energy: No

Dubinin: No

MP-Method: No

Options: Yes

Sample Log: No

Validation: No

Manufacturing: No

## Collected Data

### Collected Data Options

Start time: 1/5/2009 9:21:04 AM

End time: 1/6/2009 6:50:21 AM

Type of instrument: v

Unit number: 1

Serial number: 107

Analysis Conditions modified: No

### p<sup>s</sup> and Temperature

p<sup>s</sup> type: Measured in Psat tube for each pointp<sup>s</sup>: 99.8223 kPa

Temperature type: Entered

Temperature: 77.300 K

### Psat measurements

	Run time (min)	Pressure (kPa)
1	63	100.4412
2	72	100.3459
3	78	100.3833
4	84	100.3445
5	89	100.3576
6	95	100.3401
7	101	100.3560
8	107	100.3373
9	112	100.3623
10	118	100.3621
11	125	100.3535
12	131	100.3202
13	137	100.2948
14	144	100.2936
15	150	100.2911
16	159	100.2716
17	168	100.2746
18	178	100.2715
19	190	100.2008
20	203	100.1834
21	217	100.1994
22	236	100.1182
23	260	100.0388
24	292	100.0147
25	337	99.9394
26	405	99.8515
27	470	99.8816
28	531	99.7750
29	558	99.8090
30	567	99.7827
31	575	99.7786
32	582	99.7977
33	589	99.8023
34	595	99.7830
35	601	99.8200
36	608	99.8174
37	611	99.7740

## Sample log

Sample log		
Date	Time	Log Message
1/5/2009	9:21:04 AM	Analysis Started - Serial #: 107.
1/6/2009	6:50:21 AM	Analysis Done - Serial #: 107.

## ***D O-RING COMPATIBILITY***

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Viton or Buna-N may also be suitable for analyses similar to the reference material example file. The ability to re-use Buna-N or Viton O-rings may be limited, while the re-use of Kalrez O-rings should be more broad. Frequency of use and, potentially, several other factors affect the duration of O-ring use, so rigid rules cannot be specified for these materials. Leak rate and ultimate vacuum levels may be used as indicators for O-ring performance.

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## ***E WORKSHEETS***

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Worksheets in this section may be copied as needed.

*[Sample Data Worksheet for Gas Adsorption on the next page](#)*

Sample tube identification:

Degas Information	
Degas apparatus	
Temperature (°C)	
Time (hours)	
Actual time started	
Actual time finished	

Degas Notes:



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