

NOVA 800 User Instruction Manual



Zaera Research Group
prepared on August 21, 2023

Table of Contents

1	Introduction	Safety Sign Overview Summary & Keynotes FACES (https://faces.ccrcc.uga.edu)
2	Sample Loading	Sample Amounts Login Kaomi Instrument Connection Manage Sample IDs/weight
3	Sample Degassing	Sample Loading Start Degassing
4	Sample Cell Setting	p ₀ cell/ CLS
5	Start Analysis	Profile Selection New Profile LN ₂ Refill
6	After Analysis	Sample Removal Cleaning Cells and Rods Data Copy Report any Issue
7	Step by Step Guide	

1. Introduction

Safety Signs

	<i>Wear protective gloves.</i>
	<i>Wear safety goggles.</i>
	<i>Use protective clothing.</i>

Overview

Instrument	Analysis stations	Degas stations	Design features	Analysis results
NOVA 800	4	4	<ul style="list-style-type: none">• Long sample cells and two-liter Dewar (40+ hours) enable the most detailed pore size analyses• Flexible software for complete surface area, pore size and pore volume analyses, and data reduction• Moderate sample throughput with 2 analysis Stations• Highest sample throughput with 4 analysis stations	<ul style="list-style-type: none">• Surface area: BET, NSA, STSA, Langmuir• Pore size: BJH, DFT• Pore volume

Abbreviations: *BET - Brunauer-Emmett-Teller; NSA - Nitrogen Surface Area, STSA: Statistical Thickness Surface Area., BJH - Barrett, Joyner & Halenda; DFT - Density Functional Theory.*

Summary & Keynotes

- All users should **make reservations ahead, with the FACES system**. You should consider drying time for sample cells after washing.
- Maximum heating temperature for degassing: 425 °C.

(Where degassing temp. \geq 350 °C, **quartz sample cell** should be used instead of standard borosilicate sample cell)

- If **any types of issues** happen, please let the manager know about it.

(Issues including broken sample cells/rods, unexpected abort, problem with pump etc.)

- If you notice **LN₂** is running out after your analysis, please let the manager know about it for the next user.

2 & 3. Sample Loading & Degassing

The recommended amount of powder samples for the analysis can be found as follows:

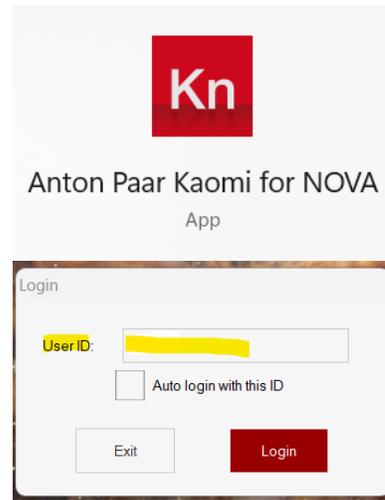
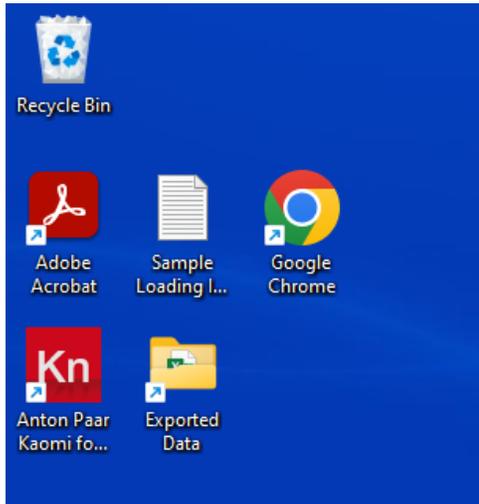
(1) Total surface area (m²) in the cell: 2 – 5 m²

e.g., Sample A has 100 m²/g as BET surface area.

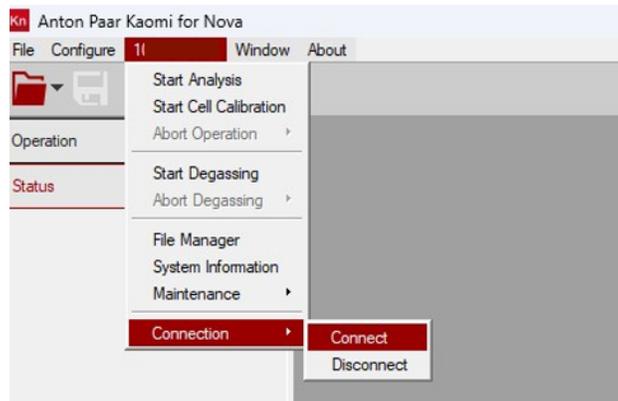
BET surface area (m ² /g)	Loaded mass (g)	Total surface area in the cell (m ²)	Result
100	0.005	100 * 0.005 = 0.5	<< 2, not promising
	0.1	100 * 0.1 = 10	>> 5, not promising
	0.05	100 * 0.05 = 5	Just right

(2) Minimum of sample mass in the cell: 25 mg (after degassing)

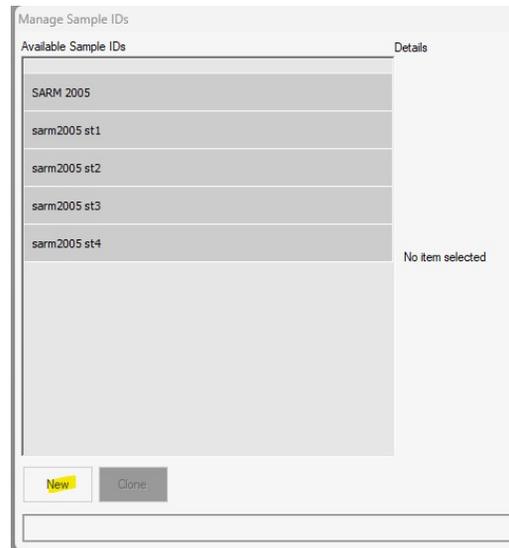
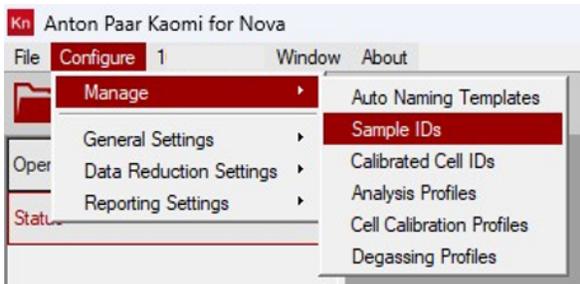
You should consider and expect sample loss during (a) transferring after weighing out and (b) degassing.



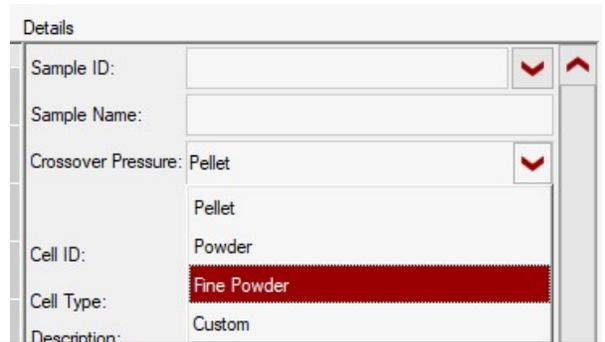
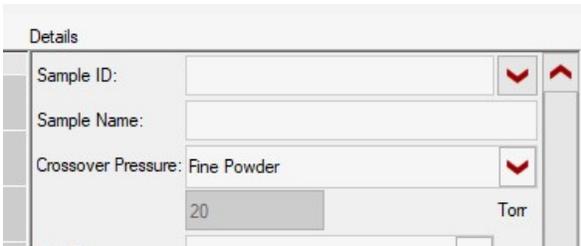
- Run 'Anton Paar Kaomi for NOVA' program, insert your UCR NetID at 'User ID' and click Login button.



- Connect with instrument. (S/N (1*****)) > Connection > Connect)



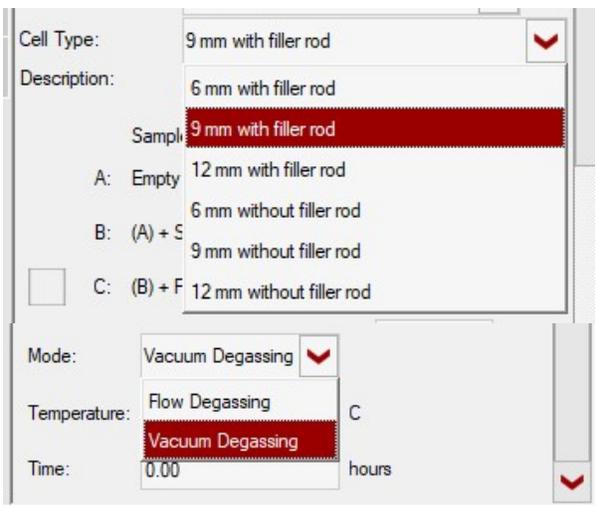
c. Make **new sample IDs** via Kaomi program. (Configure > Manage > Sample IDs > New)



Sample ID & Name; whichever you can remember and distinguish between samples after analysis

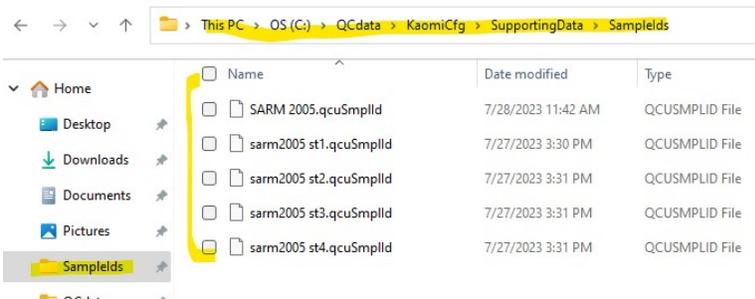
Crossover Pressure;

Powder	Fine Powder (Recommended)	Custom
50 Torr	20 Torr	e.g., 10 Torr



Cell type: 9 mm with filler rod (fixed)
(only available ones in our lab)

Mode: Vacuum Degassing (fixed)
(Inform operator when 'Flow Degassing' is required for your samples)



You can delete your Sample IDs at location of SampleIDs folder (C: > QCdata>KaomiCfg>SupportingData>SampleIDs).

c.1. Weigh out and insert the values for each part as follows:

Details

Sample Weight Calculator

A: Empty Cell (g):

B: (A) + Sample (g):

C: (B) + Filler Rod (g):

D: (B) after Degassing (g):

Calculated Sample Weight (g):

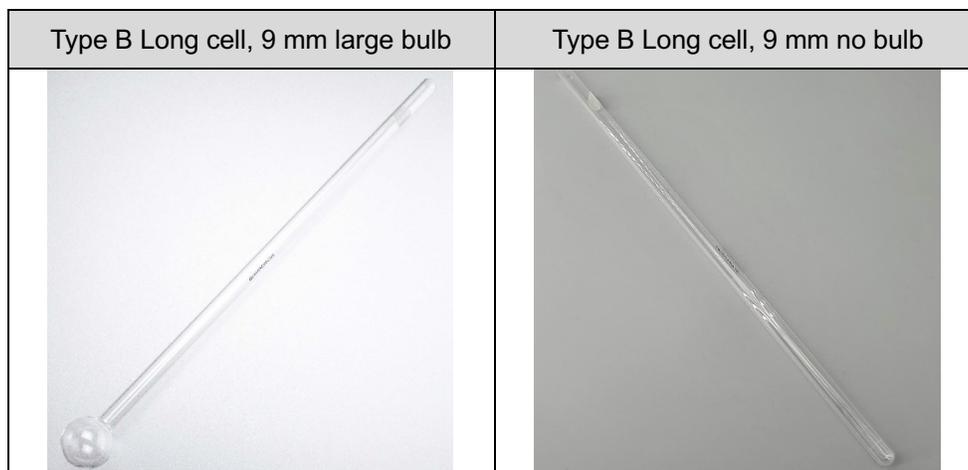
Enter Sample Weight (g):

A. Empty Cell (g)	Measure the weight of the empty cell w/o the rod
B. (A) + sample (g)	Measure the weight of (empty cell + powder sample) w/o the rod
D. (B) after degassing	Measure the weight of (empty cell + powder sample) w/o the rod after degassing

Do not use the rod during degassing step to avoid sample loss.

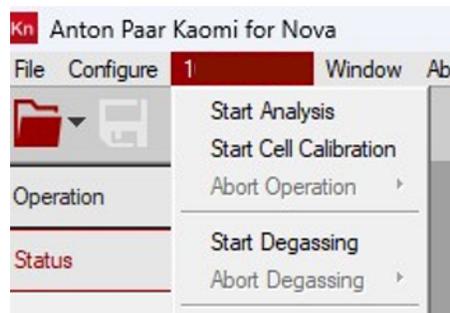
c.2. Reference the video on our lab website for sample transfer from weighing paper to the sample cells.

c.3. Two types of sample cells

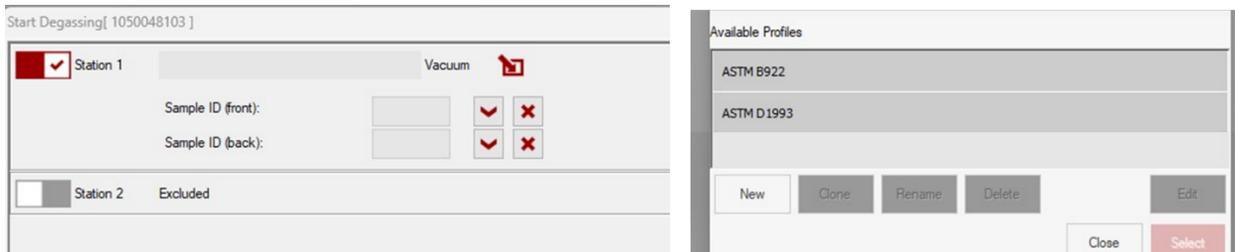




d. Load the sample cell on the degas station (②).

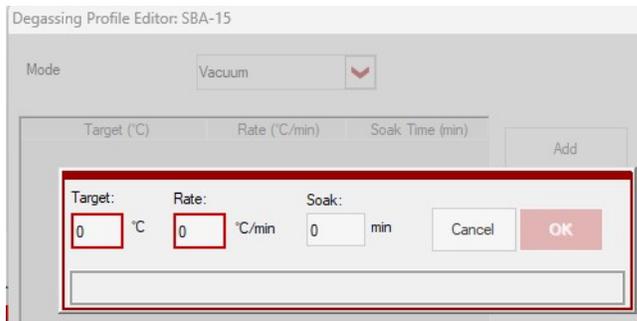


e. Start Degassing (S/N > Start Degassing).

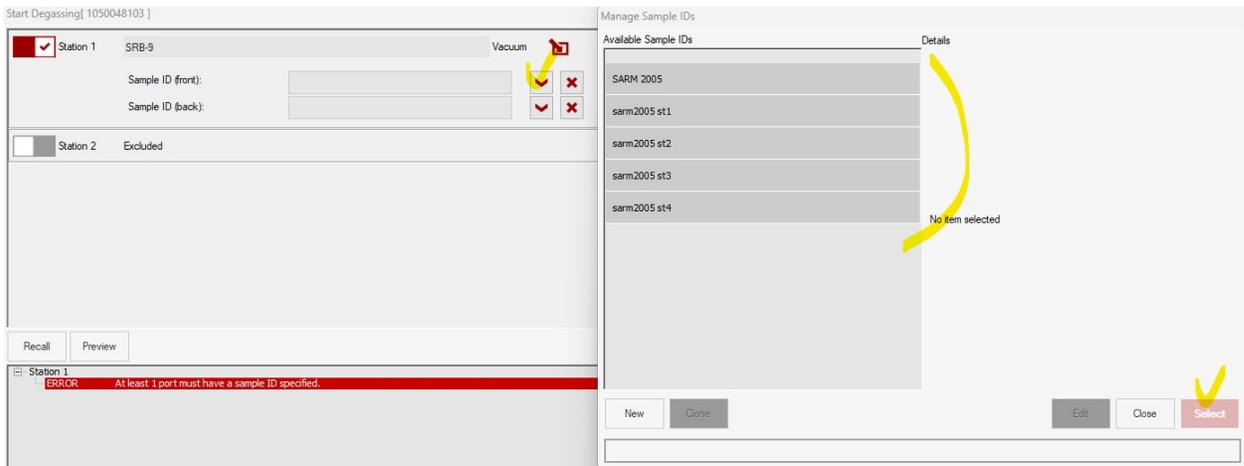


e.1. Enable Station 1 (or 2) by selecting its on/off control () (make sure the check box is checked).

e.2. Select the degas profile. If needed, click 'New' and make new degassing method.



Max. heating temp. : 425 °C



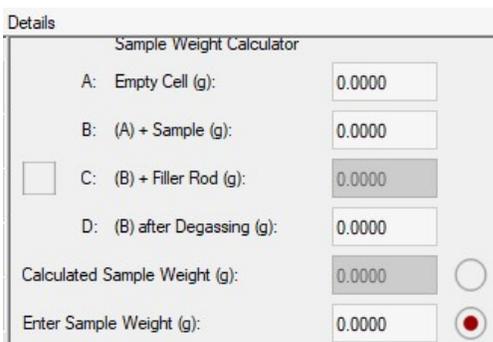
e.4. Open the Sample ID drop-down list.

e.5. Click 'Start' to begin the degassing process. Keep track of progress by selecting Status from the Kaomi for Nova Control Center/ Sidebar, or the Log View icon on the touchscreen.

e.6. Always use **Software method** to run degassing, if you are not familiar with Touchscreen method.

f. After degassing is complete, the sample cell is cooled down, and backfill is complete, remove the sample cell from the degassing station.

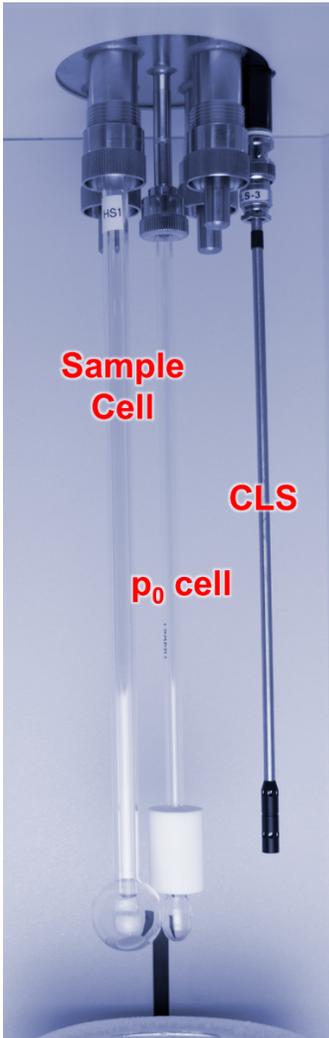
g. Re-weigh the sample cell to determine the net weight for the analysis.



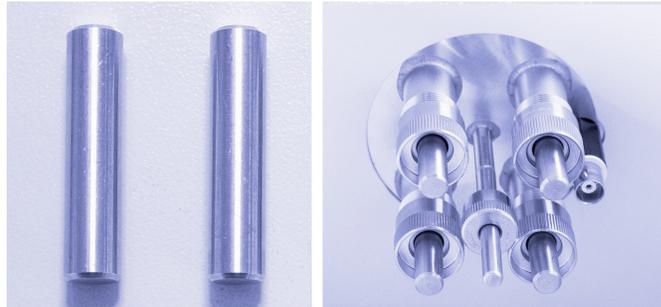
D. (B) after degassing	Measure the weight of (empty cell + powder sample) w/o the rod after degassing
------------------------	--

h. Update the Sample ID.

4 & 5. Sample Cell Setting and Start Analysis

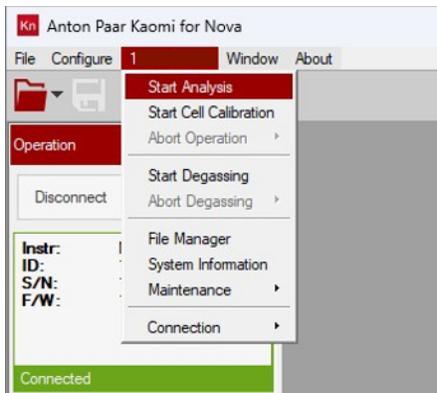


- Fill the Dewar with LN₂ to the level indicator and allow the cryogen to settle. If needed, add more cryogen to ensure it is at the level indicator.
- To set up and run an analysis, ensure the **p₀ cell** and **CLS** are installed and attach the sample cell you have been working with to Analysis Station 1 as shown in left hand side figure.
- Inform the manager, if there is any issue with compartments (e.g., O-ring and/or fittings)
- Ensure all unused stations are sealed with dowel pins.

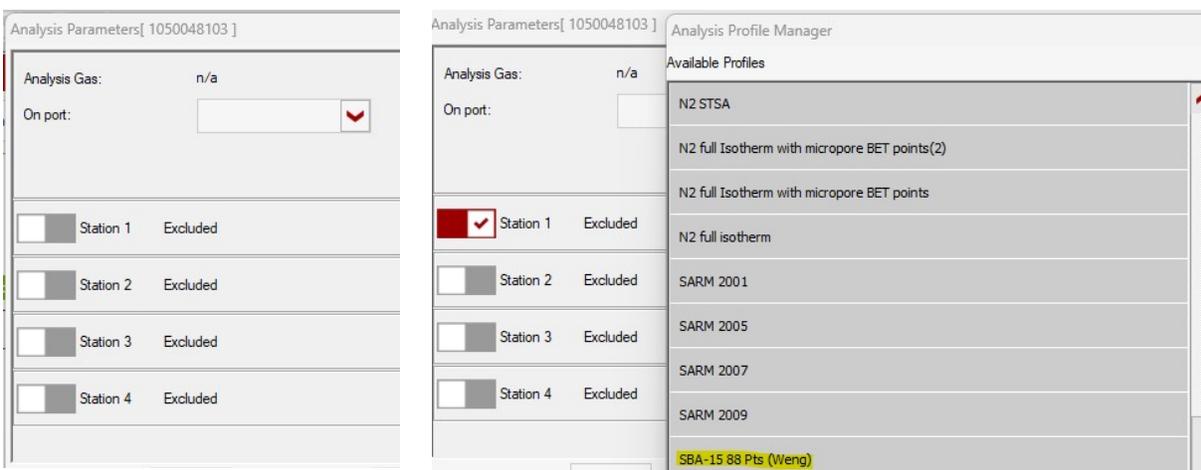


(left: image of 'dowel pins', right: as inserted in analysis stations)

- Note: Make sure to use only one O-ring when installing the sample cell into the fitting (using two or more O-rings may cause erroneous results).



a. Start Analysis (S/N > Start Analysis).



- b. Select on using Station and corresponding analysis profile.
 - b.1. If needed, make a new profile for your own sample.
- c. Fill in the remaining fields, such as:
 - c.1. On port: set to nitrogen.
 - c.2. Sample ID as same as used for degassing
 - c.3. Net sample weight (calculated earlier) in the Sample Weight entry field.
 - c.4. File name for the results file (your choice if not using the auto-naming templates feature)
- d. Click Start to begin the analysis process.
- e. Keep track of progress by selecting Status from the Kaomi for Nova Control Center/Sidebar, or the Log View icon on the touchscreen.

※ **Because of the limited LN₂ volume in the Dewar, you should consider how long the analysis will take your sample. (It can stay up to 40 h once it is refilled.)**

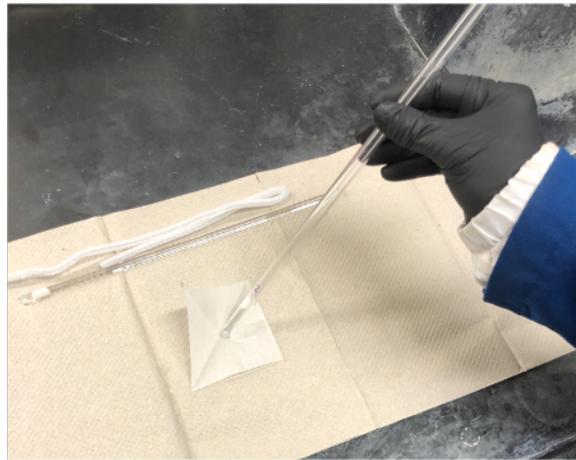
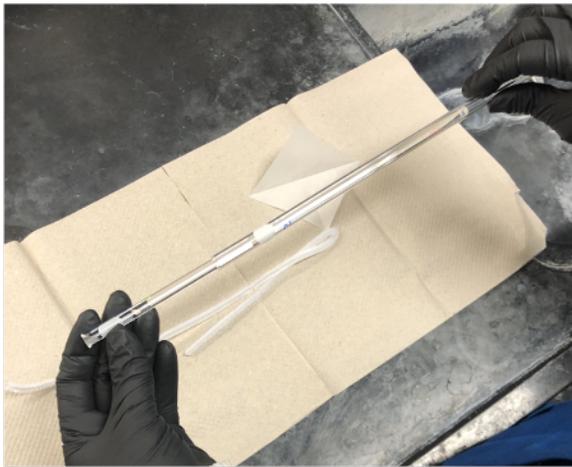
e.g., *Student A* planned to analyze 4 samples, but it turns out that 2 samples would already take 30 hours. → Analyze 2 samples first (Station 1 and 2 are on first), refill LN₂ after two samples and analyze the rest of the samples.

6. After Analysis

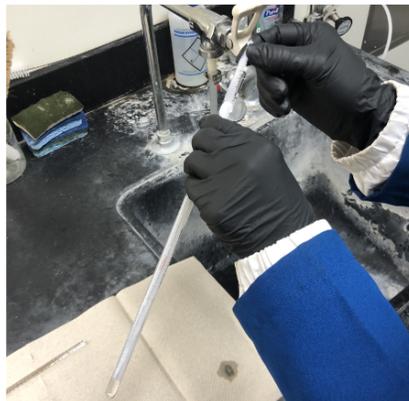
- a. Remove the sample cells from the Instrument. Make sure all unused stations are sealed with dowel pins.
- b. Clean up the used cell.
 - b.1. Prepare following tools: pipe brush and container for collected powder (vial or waste)



- b.2. Separate the rod carefully and collect used powder samples.



- b.3. Clean up the cell with deionized water by using a pipe brush. Repeat 3-4 times if needed.



b.4. Clean up the cell with acetone by using a pipe brush. Repeat 1-2 times if needed.



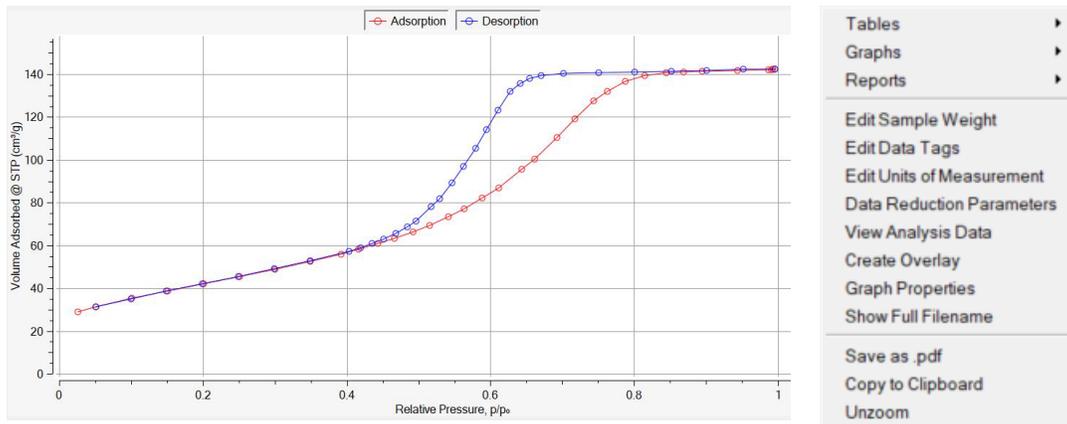
b.5. Clean up the rod in a such way. Wipe down with Kimwipes.



b.6. Dry the cleaned ones inside of the oven upside-down.

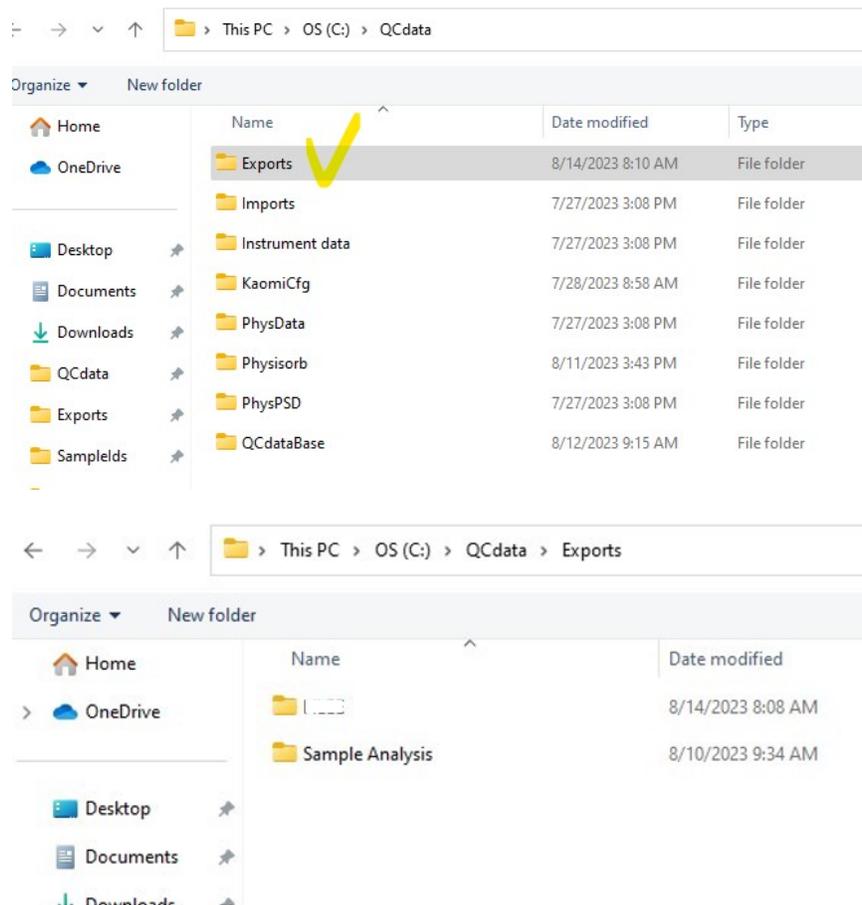


c. Inform the manager, if any types of replacement is needed.

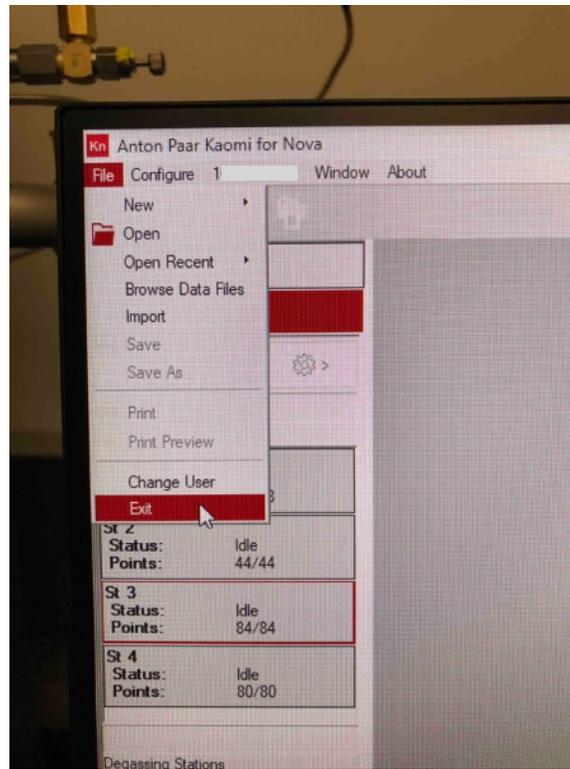


d. Right-click the displayed figure displayed in Kaomi software for reporting options.

e. Save your data in your designated folder only. If you don't have your own folder, please make one in This PC > OS (C:) > QCdata > [your name] as you see below,



- f. Close the results windows
- g. Click the File menu and select Exit to finish the program.
- h. Clean the bench for the next user.



Step by Step Guide

1. Take a clean cell from the oven
2. Cool the cell down to room temperature
3. Run [Anton Paar Kaomi for NOVA]
4. Type your UCR NetID in [User ID]
5. Click [Login] button
6. Select [1*****] > [Connection] > [Connect]
7. Select [Configure] > [Manage] > [Sample IDs] > [New]
8. Fill Sample ID and Sample Name
9. Select [Fine Powder] on Crossover Pressure
10. Choose [9 mm with filler rod]
11. Click [Vacuum Degassing] on Mode
12. Weigh the empty cell
13. Type the weight of empty cell (Line A)
14. Fill the cell with your sample
15. Weigh the sample cell
16. Type the weight of the sample cell (Line B)
17. Load the sample cell onto the degas station
18. Select [1*****] > [Start Degassing]
19. Enable [Station 1]
20. Select the degas profile, [SBA-15]
21. Open the Sample ID drop-down list
22. Choose the right Sample ID
23. Click [Select] button
24. Click [Start] button
25. Wait until the sample cell is fully cooled down after degassing is complete
26. Remove the sample cell from the degassing station
27. Re-weigh the sample cell
28. Type the weight of the degassed sample cell (Line D)
29. Fill the Dewar (use cryogenic gloves) with LN2 to the level indicator
30. Allow LN2 to be settled.
31. Ensure P0 cell and CLS are installed
32. Put a clean rod into the sample cell
33. Attach the sample cell to the analysis station (1 to 4)
34. Ensure any unused station is sealed with dowel pins
35. Select [1*****] > [Start Analysis]
36. Check all the stations you attached your sample cells
37. Select SBA-15 analysis profile
38. Set Nitrogen to on port

39. Type the net sample weight in Sample Weight entry field
40. Name your data file if not using the auto-naming templates feature
41. Click [Start] button
42. Wait until all the analysis is done
43. Mouse right-click on the result figure for reporting options
44. Select [Tables] for numeric data
45. Select [Graphs] for image data
46. Close the results window
47. Click [File] > [Exit]
48. Remove the sample cells from the analysis station
49. Put the dowel pins back to the ports
50. Have a pipe brush and a vial
51. Separate the rods from the sample cells
52. Collect your powder samples from the cells and place them in the vials
53. Rinse the cells and rods with deionized water by using a pipe brush (3-4 times)
54. Clean up the cells and rods with acetone by using a pipe brush (1-2 times)
55. Wipe the cells and rods with Kimwipes
56. Put the cleaned cells and rods (upside-down) back into the oven