ZT Scan DIA Best Practices



Introduction

Some experimental conditions and samples that are used with ZT Scan DIA methods might cause the data files to be large.

Note:

- We recommend that different computers be used for acquisition and data processing.
- We do not recommend that data be processed or reviewed during acquisition.

Acquisition

The SCIEX Workstation+ has 2 TB of disk space. This disk space might be filled in 48 hours when ZT Scan is used.

Recommendations for Local Acquisition

- Acquire a single sample with ZT Scan to see the file size, and then calculate the quantity of disk space that will be required for the full batch.
- Find the disk space that is available on the computer. If the available space is not sufficient, then move data to a data archive location to make space available.

Note: Microsoft Windows and the installed applications use approximately 0.5 TB of disk space. To calculate the quantity that is available for acquisition data, subtract this quantity from the total disk space available.

 Make backup copies of the data regularly, and then delete the data from the acquisition computer.

Note: Move the data to an external hard drive, a network-attached storage (NAS) device with more than 20 TB, or a local server.

Recommendations for Network Acquisition

- Make sure that the system configuration meets these minimum requirements:
 - · Port speed: 1 GB/s
 - Network speed and latency: GB switch network with a sustained latency of less than 5 ms
 - Destination storage system: Write speeds of 140 MB/s and read speeds of 50 MB/s

Destination storage size: 60 TB

Note: If the cloud is used for long-term storage, then we recommend the use of local (on-premises) storage before the files are uploaded.

Note: The bursting transfer from the acquisition computer to the network (destination) storage location might increase to a maximum of approximately 135 MB/s.

- Make sure that the network location is configured correctly. Refer to the document: SCIEX OS Software Laboratory Director Guide.
 - For information about the network security requirements for network acquisition, refer to the document: SCIEX OS Software Laboratory Director Guide. This document is in the Install\ProductSpecificDocumentation folder in the installation package for the SCIEX OS software.
 - SCIEX Professional Services can supply consulting support for installation, configuration, and testing of network acquisition.
- During initial system configuration, submit three test samples to the queue to make sure that acquisition and transfer of one sample completes before acquisition of the next sample starts.

Processing

Recommendations for Proteomics Processing

To process proteomics data that was acquired with ZT Scan DIA, use the PEAKS Studio software from Bioinformatics Solutions Inc (BSI) or the DIA-NN software.

PEAKS Studio Software

Supply a dedicated processing computer. For the recommended computer configuration, go to BSI. An example configuration for the PEAKS Studio software 12.5 is shown in the figure that follows. For more information, contact support@bioinfor.com.

Note: At this time, SCIEX does not sell a computer configuration that meets the requirements of the PEAKS Studio software.

Figure 1 PEAKS Studio Configuration

Configuration

PEAKS Studio 13 is recommended to be installed on 64-bit Windows operating systems with Windows 10 or later.

Recommended Configurations:

Desktop License: 30+ threads processors and 64 GB+ of RAM with compatible GPU (described below)

Workstation License: 60+ threads processors and 128 GB+ of RAM with compatible GPU (described below) e.g. Intel Core i7/i9/Xeon or AMD Ryzen 7/9/Threadripper processors

For running DeepNovo, it is required that the machine is equipped with an NVIDIA GPU with CUDA compute capability ≥ 8 and at least 8GB of dedicated memory.

For running DIA Database Search, it is recommended that the machine is equipped with an NVIDIA GPU with CUDA compute capability ≥ 5 and at least 8GB of dedicated memory.

The GPU must be updated to CUDA version 12.3 or later.

DIA-NN Software

Supply a dedicated processing computer. For computer recommendations, go to https://github.com/vdemichev/DiaNN/discussions/1292. The computer for the DIA-NN software must meet these minimum requirements:

- CPU: AMD 7/9/Threadripper (for example, Dell T7960)
- RAM: 128 GB +

There are no specifications for GPU capability, but we recommend that the computer meet the GPU requirements for the BSI Workstation.

Recommendations for Metabolomics Processing

To process metabolomics data that was acquired with ZT Scan DIA, use the Analytics workspace in the SCIEX OS software or the MS-DIAL software

SCIEX OS Software

- Supply a dedicated processing computer with a minimum of 64 GB RAM (128 GB recommended).
- Make sure that sufficient disk space is available on the C: drive. If required, then delete files that are not used.
- If out of memory exceptions occur, then increase the **Maximum size (MB)** for the virtual memory to the maximum available.

Note: Windows Administrator rights are required to do this task.

Tip! To increase the size of virtual memory, do this:

1. Push Windows Key+R.

The Run dialog opens.

- 2. Type sysdm.cpl, and then push Enter.
- 3. Go to the Advanced tab.
- 4. In the Performance section, click **Settings**.
- 5. Go to the Advanced tab.
- 6. In the Virtual memory section, click **Change**.
- 7. Make sure that the **Automatically manage paging file size for all drives** check box is not selected.
- 8. Select the drive where the virtual memory is kept. This is usually the C: drive.
- 9. Select Custom size, and then set the Maximum size (MB).
- 10. Click **OK** until all of the dialogs are closed.
- 11. If required, then start the computer again.

MS-DIAL Software

For the requirements for the MS-DIAL software, go to MS-DIAL.

This document is provided to customers who have purchased SCIEX equipment to use in the operation of such SCIEX equipment. This document is copyright protected and any reproduction of this document or any part of this document is strictly prohibited, except as SCIEX may authorize in writing.

Software that may be described in this document is furnished under a license agreement. It is against the law to copy, modify, or distribute the software on any medium, except as specifically allowed in the license agreement. Furthermore, the license agreement may prohibit the software from being disassembled, reverse engineered, or decompiled for any purpose. Warranties are as stated therein.

Portions of this document may make reference to other manufacturers and/or their products, which may contain parts whose names are registered as trademarks and/or function as trademarks of their respective owners. Any such use is intended only to designate such products as those manufacturers' products and does not imply any right and/or license to use or permit others to use such manufacturers' and/or their product names as trademarks.

SCIEX warranties are limited to those express warranties provided at the time of sale or license of its products and are the sole and exclusive representations, warranties, and obligations of SCIEX. SCIEX makes no other warranty of any kind whatsoever, expressed or implied, including without limitation, warranties of merchantability or fitness for a particular purpose, whether arising from a statute or otherwise in law or from a course of dealing or usage of trade, all of which are expressly disclaimed, and assumes no responsibility or contingent liability, including indirect or consequential damages, for any use by the purchaser or for any adverse circumstances arising therefrom.

(GEN-IDV-09-10816-E)

For Research Use Only. Not for use in Diagnostic Procedures.

Trademarks and/or registered trademarks mentioned herein, including associated logos, are the property of AB Sciex Pte. Ltd., or their respective owners, in the United States and/or certain other countries (see sciex.com/trademarks).

AB Sciex[™] is being used under license.

© 2025 DH Tech. Dev. Pte. Ltd.

CE

