

ChemoView™ 2.0.4 Software Release Notes



Intended Use

The ChemoView™ software is intended for use in the processing and reporting of data generated by flow injection analysis on a SCIEX liquid chromatography-tandem mass spectrometry (LC-MS/MS) system. Supported scan types include MRM, precursor ion scan, and neutral loss scan. The ChemoView™ software is for Research Use Only.

Changes from ChemoView™ 2.0.3 Software

Features

This version adds support for:

- The Windows 7, 64-bit operating system and the Windows 10, 64-bit operating system. Cybersecurity support is included for both the Windows 7, 64-bit operating system and the Windows 10, 64-bit operating system.

Note: The Windows XP operating system and the Windows 7, 32-bit operating system are no longer supported.

- Office 2010, 2013, and 2016, 64-bit versions.

Note: Office 2007 and the 32-bit versions of Office 2010 and 2013 are no longer supported.

Fixes

This version of the software includes the following fixes for the report templates:

- The MRM, neutral loss, and no peaks graph types have been restored. Users can select any of these graph types in a customized template.
- If units are not applicable when the *Test_Units* tag is used, then the word **Ratio** is shown in the report rather than the word **No**.
- The lines in the graphs are now shown in black instead of blue.
- The font size of the peak labels on the graphs has been increased to improve legibility.

Requirements

- A Pentium IV 2.53-GHz system with a minimum of 512 MB of RAM is recommended.
- An English version of the Windows 7, 64-bit operating system or the Windows 10, 64-bit operating system must be installed. Cybersecurity support is included for both operating systems.

Note: Because of the new operating system requirements, previous versions of the ChemoView™ software cannot be upgraded to this version. Version 2.0.4 is only available as a new installation.

- A 64-bit version of Office 2010, 2013, or 2016 must be installed.
- On computers configured with the Windows 7, 64-bit operating system, the Analyst® software, version 1.6.2 or higher, must be installed. On computers configured with the Windows 10, 64-bit operating system, the Analyst® software, version 1.7, must be installed.

Supported Equipment

- All mass spectrometers supported by the Analyst® 1.6.2 or higher software are supported by the ChemoView™ software.
- Dell OptiPlex 9010
- Dell OptiPlex XE2

Scan Types

The ChemoView™ software supports the following scan types:

- Precursor Ion (Prec)
- Neutral Loss (NL)
- Multiple Reaction Monitoring (MRM)

File Compatibility

The ChemoView™ software supports data files created using the Analyst® software, version 1.6 or higher.

Install the Software

Refer to the *ChemoView™ 2.0.4 Software Installation Guide*.

Known Issues and Limitations

Everyone should read this section of the Release Notes. This section applies specifically to issues that are known in the ChemoView™ 2.0.4 software.

- If a single alphanumeric character is added to an existing test, analyte, or processing method name, the ChemoView™ software might not recognize the change and prevent the user from continuing through the workflow. It is recommended that at least two characters be added to an existing name to enable the software to recognize the change.
- Occasionally, when generating a summary report in Microsoft Excel 2010, the report might fail and indicate that the failure was caused by a Visual Basic issue. To resolve this issue:
 1. Click **File > Options**.
 2. On the Excel Options dialog, click **Trust Center**.
 3. In the Microsoft Excel Trust Center section, click **Trust Center Settings**.
 4. Click **Macro Settings**, select the **Trust access to the VBA project object model** check box in the Developer Macro Settings section, and then click **OK**.
 5. Click **OK**.

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Documentation

This version of the document supercedes all previous versions of this document.

To view this document electronically, Adobe Acrobat Reader is required. To download the latest version, go to <https://get.adobe.com/reader>.

For the latest versions of the documentation, visit the SCIEX website at sciex.com.

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Other Documentation

- *Help*

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