

SCIEX WIFF

.wiff, multi-sample #n, MRM-centric extract, Clearcore2, method from WIFF.

How Quanto reads SCIEX .wiff files and builds MRM methods from the WIFF.

Raw data

Item	Notes
File	.wiff (and companion files as required by the vendor reader)
Multi-sample	Samples inside a WIFF are addressed as #n (sample index)
Extract style	MRM-centric — transitions from the WIFF

Reader add-in: SciexWiff using **Clearcore2** / SCIEX vendor APIs. Support DLLs live under:

C:\Program Files\Quanto\AddIns\VendorSupportFiles\Sciex\

Method import

Source	What you point at	What comes in
SCIEX method from data	.wiff file	MRM compounds / transitions (SciexWiffMethod)

Standalone .msm is **not** supported with the current minimal SCIEX DLL set — use method-from-WIFF.

Import: Method → Load / Import — Replace or Append ([Method design](#)).

Empty or odd chromatograms

1. Confirm the Quant MRM matches a transition in the WIFF.
2. Confirm the correct multi-sample #n was added.
3. Confirm VendorSupportFiles\Sciex is present for this build.

See [Troubleshooting](#).

Related

- [Compatibility](#)
- [Reports, import, and export](#)
- [Creating add-ins](#)