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