



Software Version: 1.9.0.4157

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1. Changes Since Last Release

1.1 Features

- Add Axis Label Size option to all plots

1.2 Updates

- Mismatch corrected between dendrogram and network plots
- Premix Binning mode: The "Epitope Tool" modal is displayed after clicking on the "Undo All" button on the Processing > Referencing/Blank pages
- The "Measurement" feature is present on the Data > Batch Analysis page and Binning page in Premix Binning mode
- Classical / Premix Binning mode: The sensorgram panel is empty on the Binning page after Batch Analysis fitting
- Classical and Premix Binning modes: The "Id" is different in the "Reports" tab in the Binning page each time the application opens
- An unstable error appears when using the "Exclude (remove) all except selected" option
- An unstable error appears when using the "Exclude all except selected ligands/analytes" option
- Array View page: Background color for excluded items isn't displayed on the Analyte Overlay tab and doesn't save to .ebp file
- Classical / Premix Binning mode: The Binning tab has different selections than the state that was saved
- The app stopped working while checking "Is Control" checkbox
- Processing > Y-Align page: Values aren't saved in input fields
- Chords sometimes disappear when moving nodes
- Updating Windows Regional Settings causes errors when opening/saving .ebp files
- Error opening classical binning file with spaces at end of sample names
- Error opening merged premix binning file

1.3 Known Issues

- Coloring of peptide mapping plots can fail to separate distinct peptides

2. Installation/Upgrade Instructions

- i. Run the Epitope installer executable file
- ii. Accept User Account Control permissions
- iii. Complete standard installation steps as instructed
- iv. Open application from Windows menu or desktop shortcut

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