

## Protein Quantitation MicroApp Software

Version 1.0

### Introduction

This document describes version 1.0 of the Protein Quantitation MicroApp software – a plug-in to the PeakView™ 1.1.1 software. It is an application for selecting peptides and proteins, extracting the MS peak areas, and exporting the data to MarkerView™ software. The software has been tested on Windows XP.

### Where to Get Help

#### Other Documentation

- A user guide that provides an introduction to all of the main features of the application.

#### Support

For support, contact AB SCIEX.

- Email: [support@absciex.com](mailto:support@absciex.com)
- Web: [www.absciex.com](http://www.absciex.com)

### Requirements

- PeakView™ 1.1.1 software must be installed
- MarkerView™ 1.2.1 software must be used to view the exported data.
- If you have previously installed a version of the application, then you must uninstall it using Add or Remove Programs in Control Panel before you can install this version of the application.

### Installation Instructions

- To install the software, double-click the .msi installer downloaded from the AB SCIEX Web site and then follow the instructions.

### Known Issues and Limitations

- When saving the MarkerView software files for peptides and proteins, the "Save Window" may disappear behind existing windows. Use Alt-Tab to navigate to the correct window.
- Do not process data over the network. Always put the data on your local drive.
- Occasionally, small differences in peak areas between the tables in MarkerView software and XICs generated manually in PeakView software might be observed. This is due to slight differences in the baseline calculations during peak integration and should not affect the quality of the results.

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