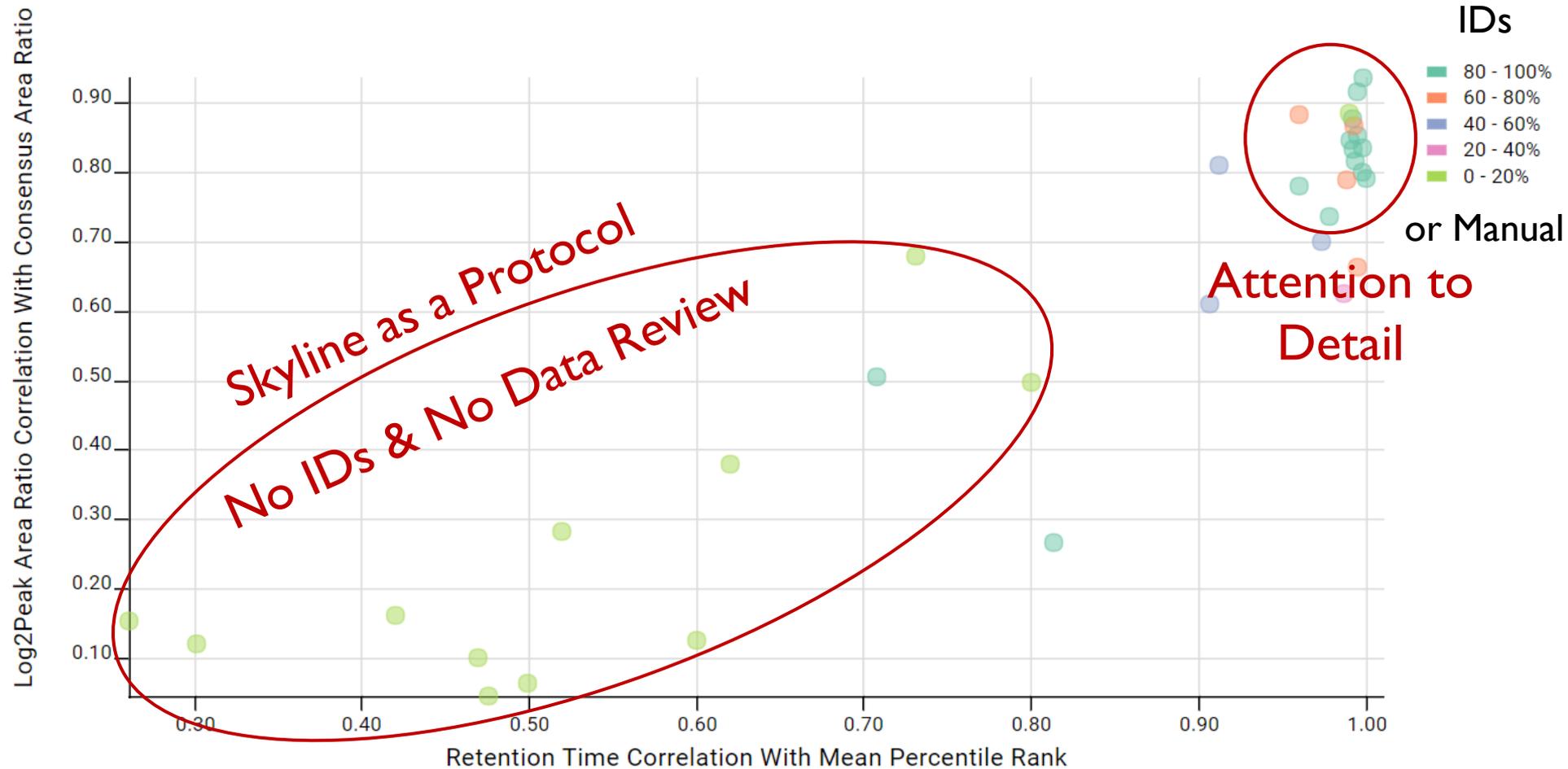


ABRF sPRG 2013 Study

▶ A cautionary tale

Results - Log2Peak Area Ratio Correlation With Consensus Area Ratio



Keys to Success with MS1 in Skyline

- ▶ Make sure you have ID annotations
 - ▶ Diagnose with Spectral Library Explorer
 - ▶ <http://tinyurl.com/Skyline-missing-ids>
 - ▶ Use File > Import > Peptide Search
- ▶ Review RT alignment in alignment viewer
- ▶ **Got HYPOTHESIS??**
- ▶ Review and manually adjust <5% of peaks (targeted)

- ▶ Use tutorial and webinars

