

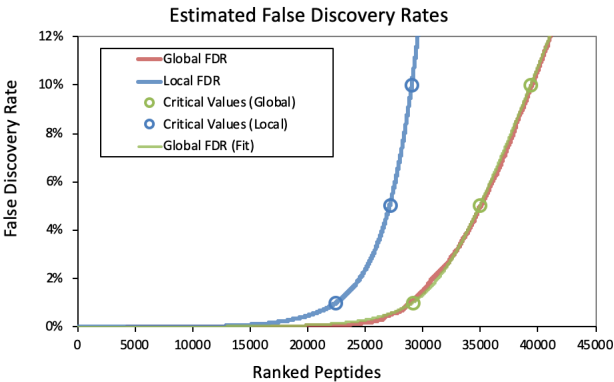
Figure S3. FDR analysis of combined IDA library of retina at peptide level generated by Protein Pilot software

Distinct Peptide Level FDR Analysis

Peptides Identified at Critical False Discovery Rates

Number of Peptides Identified			
Critical FDR	Local FDR	Global FDR	Global FDR from Fit
1.0%	22431	28853	29066
5.0%	27096	35013	34921
10.0%	29021	41853	39356

* It is recommended you use numbers in bold and avoid using numbers in italics.



Correspondence between FDR Levels and ProteinPilot Reported Confidences

Corresponding ProteinPilot Confidence			
Critical FDR	Local FDR	Global FDR	Global FDR from Fit
1.0%	99.3%	90.3%	89.3%
5.0%	95.4%	47.3%	47.8%
10.0%	89.5%	29.1%	29.5%

* It is recommended you use numbers in bold and avoid using numbers in italics.

