Comparative Shotgun Proteomics Using Spectral Count Data and Quasi-Likelihood Modeling *J. Proteome Res.* **2010,** *9,* 4295–4305. Ming Li, William Gray, Haixia Zhang, Christine H. Chung, Dean Billheimer, Wendell G. Yarbrough, Daniel C. Liebler, Yu Shyr, and Robbert J. C. Slebos*

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Supporting Information Available: IDPicker report of CPTAC yeast spike data, filtered CPTAC data set used for analysis, and detailed methods description were not linked to the original document but were cited in the text. These files are now available. This material is available free of charge via the Internet at http://pubs.acs.org.

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