

## Publication List

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- ▶ Adam Rauch, Matthew Bellew, Jimmy Eng, Matthew Fitzgibbon, Ted Holzman, Peter Hussey, Mark Igra, Brendan Maclean, Chen Wei Lin, Andrea Detter, Ruihua Fang, Vitor Faca, Phil Gafken, Heidi Zhang, Jeffrey Whitaker, David States, Sam Hanash, Amanda Paulovich, and Martin W. McIntosh. 2006. Computational Proteomics Analysis System (CPAS): An Extensible, Open-Source Analytic System for Evaluating and Publishing Proteomic Data and High Throughput Biological Experiments. *Journal of Proteome Research*, **5**, 112-121.  
<http://pubs.acs.org/cgi-bin/abstract.cgi/jprobs/2006/5/i01/abs/pr0503533.htm>
- ▶ Brendan MacLean Jimmy K. Eng, Ronald C. Beavis and Martin McIntosh. 2006 General framework for developing and evaluating database scoring algorithms using the TANDEM search engine. *Bioinformatics* **22(22)**:2830-2832.  
<http://bioinformatics.oxfordjournals.org/cgi/content/full/22/22/2830>
- ▶ Matthew Bellew, Marc Coram, Matthew Fitzgibbon, Mark Igra, Tim Randolph, Pei Wang, Damon May, Jimmy Eng, Ruihua Fang, ChenWei Lin, Jinzhi Chen, David Goodlett, Jeffrey Whiteaker, Amanda Paulovich and Martin McIntosh. 2006. A suite of algorithms for the comprehensive analysis of complex protein mixtures using high-resolution LC-MS. *Bioinformatics* **22(15)**:1902-1909.  
<http://bioinformatics.oxfordjournals.org/cgi/content/abstract/22/15/1902>
- ▶ Andrew R Jones, Michael Miller, Ruedi Aebersold, Rolf Apweiler, Catherine A Ball, Alvis Brazma, James DeGreef, Nigel Hardy, Henning Hermjakob, Simon J Hubbard, Peter Hussey, Mark Igra, Helen Jenkins, Randall K Julian, Jr, Kent Laursen, Stephen G Oliver, Norman W Paton, Susanna-Assunta Sansone, Ugis Sarkans, Christian J Stoeckert, Jr, Chris F Taylor, Patricia L Whetzel, Joseph A White, Paul Spellman & Angel Pizarro. 2007. The Functional Genomics Experiment model (FuGE): an extensible framework for standards in functional genomics. *Nature Biotechnology* **25**, 1127 – 1133.  
<http://www.nature.com/nbt/journal/v25/n10/pubmed/nbt1347.html>