

Order	Authors	Title	MS NO.
1	Gilbert S. Omenn, David J. States, Marcin Adamski, Thomas W. Blackwell, Rajasree Menon, Henning Hermjakob, Rolf Apweiler, Brian B. Haab, Richard J. Simpson, James S. Eddes, Eugene A. Kapp, Robert L. Moritz, Daniel W. Chan, Alex J. Rai, Arie Admon, Ruedi Aebersold, Jimmy Eng, William S. Hancock, Stanley A. Hefta, Helmut Meyer, Young-Ki Paik, Jong-Shin Yoo, Peipei Ping, Joel Pounds, Joshua Adkins, Xiaohong Qian, Rong Wang, Valerie Wasinger, Chi Yue Wu, Xiaohang Zhao, Rong Zeng, Alexander Archakov, Akira Tsugita, Ilan Beer, Akhilesh Pandey, Michael Pisano, Philip Andrews, Harald Tammen, David W. Speicher, Samir M. Hanash.	Overview of the HUPO Plasma Proteome Project: Results from the pilot phase with 35 collaborating laboratories and multiple analytical groups, generating a core dataset of 3020 proteins and a publicly-available database	2005-00358
2	Marcin Adamski, Thomas Blackwell, Rajasree Menon, Lennart Martens, Henning Hermjakob, Chris Taylor, Gilbert S. Omenn, David J. States	Data Management and Preliminary Data Analysis in the Pilot Phase of the HUPO Plasma Proteome Project	2005-00186
3	Alex J. Rai, Craig A. Gelfand, Bruce C. Haywood, David Warunek, Catherine Skobe, Mark D. Schuchard, Richard J. Mehig, Steven L. Cockrill, Graham B. I. Scott, Harald Tammen, Peter Schulz-Knappe, David W. Speicher, Frank Vitzthum, Brian B. Haab, Gerard Siest, Daniel W. Chan	Human Proteome Organization--Plasma Proteome Project Specimen Collection and Handling: Towards the Standardization of Parameters for Plasma Proteome Samples	2004-00537.R1
4	Brian B. Haab, Bernhard H. Geierstanger, George Michailidis, Frank Vitzthum, Sara Forrester, Ryan Okon, Petri Saviranta, Achim Brinker, Martin Sorette, Lorah Perlee, Shubha Suresh, Garry Drwal, Joshua N. Adkins, Gilbert S. Omenn	Immunoassay and antibody microarray analysis of the HUPO PPP reference specimens: systematic variation between sample types and calibration of mass spectrometry data	2004-00470.R1
5	Lynn A. Echan, Hsin-Yao Tang, Nadeem Ali-Khan, KiBeom Lee, David W. Speicher	Depletion of multiple high abundance proteins improves protein profiling capacities of human serum and plasma	2004-00518.R1
6	Nina Zolotarjova, James Martosella, Gordon Nicol, Jerome Bailey, Barry E. Boyes and William C. Barrett*	Differences among techniques for high-abundant protein depletion	2004-02021.R1
7	Lei Huang, Gulia Harvie, Jerald S. Feitelson, David A. Herold, Wei-Wei Zhang, Xiangming Fang	Immunoaffinity Separation of Plasma Proteins by IgY Microbeads: Meeting the Needs of Proteomic Sample Preparation	2004-00420.R1
8	Hsin-Yao Tang, Nadeem Ali-Khan, Lynn A. Echan, Natasha Levenkova, John J. Rux and David W. Speicher	A novel 4-dimensional strategy combining protein and peptide separation methods enables detection of low abundance proteins in human plasma and serum proteomes	2004-01099
9	David E. Misek, Rork Kuick, Hong Wang, Vladimir Galchev, Bin Deng, Rong Zhao, John Tra, Michael R. Pisano, Ravi Amunugama, David Allen, Angela K. Walker, John R. Strahler, Philip Andrews, Gilbert S. Omenn, Samir M. Hanash	A Wide Range of Protein Isoforms in Serum and Plasma Uncovered by a Quantitative Intact Protein Analysis System (IPAS)	2005-00103.R1

10	Ziping Yang, William S. Hancock, Tori Richmond-Chew, Leo Bonilla	A Study of Glycoproteins in Human Serum and Plasma Reference Standards (HUPO) using Multi-Lectin Affinity Chromatography Coupled with RPLC-MS/MS	2004-00411.R1
11	Eilon Barnea, Raya Sorkin, Tamar Ziv, Ilan Beer, Arie Admon	Evaluation of Pre-Fractionation Methods as a Preparatory Step for Multidimensional Based Chromatography of Serum Proteins	2004-00412.R1
12	Jin Young Kim, Jeong Hwa Lee, Gun Wook Park ¹ , Kun Cho, Kyung Hoon Kwon, Young Mok Park, Sang Yun Cho, Young Ki Paik, and Jong Shin Yoo	Utility of electrophoretically derived protein mass estimates as additional constraints in proteome analysis of human serum based on MS/MS analysis	2004-00413.R1
13	Sang Yun Cho, Eun-Young Lee, Young-Wook Chun, Joon Seok Lee, Hye-Young Kim, Jae Myun Park, Min-Seok Kwon, Young-Kew Park, Hyoung-Joo Lee, Min-Jung Kang, Jin Young Kim, Jong-Shin Yoo, Sung Jin Park, Jin Won Cho, Young-Ki Paik	Efficient prefractionation of the low-abundance proteins in human plasma and construction of two-dimensional map	2004-00497.R1
14	Robert L. Moritz, Andrew B. Clippingdale, Eugene A. Kapp, James S. Eddes, Hong Ji, Sam Gilbert, Lisa M. Connolly, Richard J. Simpson	Application of Two-Dimensional Free-Flow Electrophoresis / RP-HPLC for Proteomic Analysis of Human Plasma Depleted of Multi High-Abundance Proteins	2005-00096.R1
15	Harald Tammen, Imke Schulte, Rudiger Hess, Christoph Menzel, Markus Kellmann, Thomas Mohring, and Peter Schulz-Knappe	Peptidomic Analysis of Human Blood Specimens: Comparison between Plasma Specimens and Serum by Differential Peptide Display	2004-00419.R1
16	Xiaohai Li, Yan Gong, Ying Wang, Songfeng Wu, Yun Cai, Ping He, Zhuang Lu, Wantao Ying, Yangjun Zhang, Liyan Jiao, Hongzhi He, Zisen Zhang, Fuchu He, Xiaohang Zhao, Xiaohong Qian	Comparison of alternative analytical techniques for the characterization of the human serum proteome in HUPO PPP	2004-00425.R1
17	Ping He, Hong-Zhi He, Jie Dai, Ying Wang, Quan-Hu Sheng, Lan-Ping Zhou, Zi-Sen Zhang, Yu-Lin Sun, Fang Liu, Kun Wang, Jin-Sheng Zhang, Hui-Xin Wang, Zhen-Mei Song, Hai-Rong Zhang, Rong Zeng, Xiaohang Zhao	The Human Plasma Proteome: Analysis of Chinese Serum Using Shotgun Strategy	2004-00422.R1
18	Joshua N. Adkins, Matthew E. Monroe, Kenneth J. Auberry, Yufeng Shen, Jon M. Jacobs, David G. Camp II, Frank Vitzthum, Karin D. Rodland, Richard D. Smith, and Joel G. Pounds	A Proteomic Study of HUPO's Plasma Proteome Project Pilot Samples using an Accurate Mass and Time Tag Strategy	2004-00633.R2
19	Alex J. Rai, Paul M. Stemmer, Zhen Zhang, Bao-ling Adam, William T. Morgan, Rebecca E. Caffrey, Vladimir Podust, Manisha Patel, Lih-yin Lim, Natalia V. Shipulina, Daniel W. Chan, O. John Semmes, Hon-chiu Eastwood Leung	Analysis of HUPO-PPP Reference Specimens using SELDI-TOF Mass Spectrometry: Multi-Institution Correlation of Spectra and Identification of Biomarkers	2004-00606.R1
20	Eugene A Kapp, Frédéric Schütz, Lisa M. Connolly, John A. Chakel, Jose E. Meza, Christine A. Miller, David Fenyo, Jimmy K. Eng, Joshua N. Adkins, Gilbert S. Omenn, Richard J. Simpson	An evaluation, comparison and accurate benchmarking of several publicly-available MS/MS search algorithms: Sensitivity and Specificity analysis	2005-00126
21	Ilan Beer, Eilon Barnea, Arie Admon	Centralized Data Analysis of a Large Inter-Laboratory Proteomics Project – a Feasibility Study	2004-00457.R1

22	Eric W. Deutsch, Jimmy K. Eng, Hui Zhang, Nichole L. King, Alexey I. Nesvizhskii, Biaoyang Lin, Hookeun Lee, Eugene C. Yi, Reto Ossola, Ruedi Aebersold	Human Plasma PeptideAtlas	2005-00160
23	Lennart Martens, Alexey I. Nesvizhskii, Henning Hermjakob, Marcin Adamski, Gilbert S. Omenn, Joël Vandekerckhove and Kris Gevaert	Do we want our data raw? Including binary mass spectrometry data in public proteomics data repositories.	2004-00376.R1
24	Peipei Ping, Thomas M. Vondriska, Chad J. Creighton, TKB Gandhi, Ziping Yang, Rajashree Menon, Min-Seok Kwon, Garry Drwal, Markus Kellman, Suraj Peri, Shubha Suresh, Mads Gronborg, Henrik Molina, Raghothama Chaerkady, B. Rekha, Babylakshmi Muthusamy, Arun S. Shet, Robert E. Gerszten, Haifeng Wu, Mark Raftery, Valerie Wasinger, Peter Schulze-Knappe, Samir M. Hanash, Young Ki-Paik, William S. Hancock, David J. States, Gilbert S. Omenn, Akhilesh Pandey	A Functional Annotation of Subproteomes in Human Plasma	2005-00140
25	Beniam T. Berhane, Chenggong Zong, David A. Liem, Aaron Huang, Steven Le, Ricky D. Edmondson, Richard C. Jones, Julian P. Whitelegge, Peipei Ping, Thomas M. Vondriska.	Cardiovascular-related proteins identified in human plasma by the HUPO Plasma Proteome Project pilot phase	2004-01084.R1
26	Babylakshmi Muthusamy, G. Hanumanthu, R. Reshmi, S. Sriranjini, Shubha Suresh, B. Rekha, D. Srinivas, L. Karthick1, B. M. Vrushabendra1, Salil Sharma, Goparani Mishra, B. P. Rashmi, Kadekar Sandeep, Pritam Chatterjee, K. S. Mangala, H. N. Shivashankar, K. N. Chandrika, Nandan Deshpande, M. Suresh, N. Kannabiran, Vidya Niranjana, Anuradha Nalli, T. S. Keshava Prasad, K. S. Arun, Raghunath Reddy, Sreenath Chandran, Trafina Jadhav, D. Julie, M. Mahesh, S. Lynate John, Kshitish Palvankar, D. Sudhir, P. Bala, M. K. Jithesh, N. S. Rashmi, G. Vishnupriya, Kaushik Dhar, S. Reshma, Raghothama Chaerkady, C. Krishna Prasad, T. K. B. Gandhi, H. C. Harsha, S. Sujatha Mohan, Krishna S. Deshpande, Malabika Sarker, Akhilesh Pandey	Plasma Proteome Database as a resource for proteomics research	2004-00588.R2
27	Lennart Martens, Henning Hermjakob, Philip Jones, Marcin Adamski, Chris Taylor, David States, Kris Gevaert, Joël Vandekerckhove, Rolf Apweiler	PRIDE: The PRoteomics IDentifications database	2004-00647.R1
28	Valerie Wasinger	2D-LC-MS/MS analysis of Gradiflow fractionated native plasma (HUPO PPI)	00297-2004.R2