1 vial

3050 Spruce Street, Saint Louis, MO 63103 USA Tel: (800) 521-8956 (314) 771-5765 Fax: (800) 325-5052 (314) 771-5757 email: techservice@sial.com sigma-aldrich.com

Universal Proteomics Standard (UPS) Set

Catalog Number **UPS1** Storage Temperature –20 °C

Product Description

The Universal Proteomics Standard (UPS) Set is comprised of one vial containing 48 human source or human sequence recombinant proteins (Catalog Number U6133), and one vial (20 µg) of Proteomics Grade Trypsin (Catalog Number T6567).

There are 5 pmoles of each HPLC purified protein in the vial. Each protein has been quantitated by amino acid analysis (AAA). The proteins have been selected to limit heterogeneous post-translational modifications (PTMs).

This set can be used to standardize and/or evaluate mass spectrometric (e.g., LC-MS/MS, MALDI-TOF-MS, etc.) and electrophoretic analysis conditions prior to the analysis of complex protein samples. Moreover, the set may be used to bracket precious experimental datasets between runs of a known complex standard sample. thereby, confirming the robustness of the analysis method and stability of the instrument employed. Additionally, laboratories generating or comparing mass spectrometric data derived from poorly defined samples may use the standard as an external reference to assist with the evaluation of results and experimental methodology. In this regard the running of the standard may facilitate the comparison of mass spectrometric or other proteomic data that is generated in different laboratories using a wide range of varying workflows, analytical techniques, and instrumentation. Lastly, this set will potentially help identify limitations of proteomics analysis systems and search algorithms.

This protein mixture was extensively evaluated and reported on under the direction of the Association of Biomolecular Resource Facilities (ABRF) Proteomics Standards Research Group (sPRG) during a comprehensive 2005/2006 study. The findings of the study were presented at the ABRF 2006 and US HUPO 2006 conferences. 1,2

Components

Universal Proteomics Standard 5 pmoles each of 48 human proteins, dried in a 2 ml vial Catalog Number U6133

Proteomics Grade Trypsin 20 μg lyophilized enzyme Catalog Number T6567

Precautions and Disclaimer

This product is for R&D use only, not for drug, household, or other uses. Please consult the Material Safety Data Sheet for information regarding hazards and safe handling practices.

Preparation Instructions

The preparation procedure should be compatible with the analysis to be run. For peptide analysis, it is suggested that proteins be resuspended in an appropriate denaturant prior to reduction, alkylation, and tryptic digestion.

Storage/Stability

The set ships in wet ice and storage at $-20~^{\circ}$ C is recommended. After reconstitution and/or digestion, the product should be dispensed into microcentrifuge tubes in single use aliquots and frozen.

References

- Arnott, D.P., et al., sPRG2006 Study: A Proteomics Standard. Presented at the ABRF 2006 Conference, Long Beach, CA, February 11-14, 2006
- Kowalak, J.A., et al., ABRF-sPRG2006 Study: Prototype Proteomics Standard. Presented at the US HUPO 2nd Annual Conference, Boston, MA, March 11-15, 2006.
- UniProt (Universal Protein Resource), © 2006 by UniProt Consortium. http://www.pir.uniprot.org/index.shtml

KR, JAB, GBIS, MAM 01/08-1

UniProt Accession	UniProt Protein Name [Synonym]	MW (Da) (calculated)	Source or recombinant	Host	Tag	Potential PTMs *
Number ³	Alaba I a dalla consis	` ,	N 4:11 -			Observatelies
P00709	Alpha-lactalbumin	14,070	Milk			Glycosylation
P08758 P01008	Annexin A5 Antithrombin-III	35,782	Placenta			Acetylation
P01008 P61769		49,033 11,729	Plasma Urine			Glycosylation
P51769 P55957	Beta-2-microglobulin			E. coli		
P00915	BH3 interacting domain death agonist [BID] Carbonic anhydrase 1	21,978 28,738	Recombinant Erythrocytes	E. COII		Acetylation
P00913 P00918	Carbonic anhydrase 1 Carbonic anhydrase 2	29,095	Erythrocytes			Acetylation
P04040	Carbonic annydrase 2 Catalase	59,583	Erythrocytes			Acetylation
P07339	Cathepsin D	26,624	Liver			Glycosylation
P08311	Cathepsin G	26,751	Sputum			Glycosylation
P01031	Complement C5 [Complement C5a]	8,266	Recombinant	E. coli		Olycosylation
P02741	C-reactive protein	23,030	Plasma	L. 00#		
P06732	Creatine kinase M-type [CK-MM]	43,070	Heart			
P00167	Cytochrome b ₅	16,021	Recombinant	E. coli	6-His	
P99999	Cytochrome c [Apocytochrome c]	11,608	Recombinant	E. coli	0 1 110	
P01133	Epidermal growth factor	6,211	Recombinant	E. coli		
P05413	Fatty acid-binding protein	14,716	Plasma			Acetylation, Phosphorylation
P06396	Gelsolin	82,954	Plasma			Phosphorylation
P08263	Glutathione S-transferase A1 [GST A1-1]	25,482	Recombinant	E. coli		1 Hoophorylation
P09211	Glutathione S-transferase P [GST]	23,220	Placenta			
P01112	GTPase HRas [Ras protein]	21,292	Recombinant	E. coli		
P69905	Hemoglobin alpha chain	15,127	Erythrocytes			
P68871	Hemoglobin beta chain	15,867	Erythrocytes			Acetylation, Nitrosylation, Glycosylation
P12081	Histidyl-tRNA synthetase [Jo-1]	58,223	Recombinant	E. coli		
P01344	Insulin-like growth factor II	7,464	Recombinant	E. coli		
P10145	Interleukin-8	8,381	Recombinant	E. coli		
P02788	Lactotransferrin	78,289	Milk			Glycosylation
P41159	Leptin	16,024	Recombinant	E. coli		
P61626	Lysozyme C	14,692	Milk			
P10636	Microtubule-associated protein tau [Tau protein]	46,810	Recombinant	E. coli	6-His	
P02144	Myoglobin	17,051	Heart			
P15559	NAD(P)H dehydrogenase [quinone] 1 [DT Diaphorase]	30,984	Recombinant	E. coli		
Q15843	Neddylin [Nedd8]	9,071	Recombinant	E. coli		
P62937	Peptidyl-prolyl cis-trans isomerase A [Cyclophilin A]	17,947	Recombinant	E. coli		
Q06830	Peroxiredoxin 1	22,106	Recombinant	E. coli		
P01127	Platelet-derived growth factor B chain	12,286	Recombinant	E. coli		
P02753	Retinol-binding protein	21,065	Urine			
P16083	Ribosyldihydronicotinamide dehydrogenase (quinone) [Quinone oxidoreductase 2 or NQO2]	25,817	Recombinant	E. coli		
P02787	Serotransferrin [Apotransferrin]	75,143	Plasma			Glycosylation
P02768	Serum albumin	66,393	Recombinant	Pichia pastoris		
P63165	Small ubiquitin-related modifier 1 [SUMO-1]	37,420	Recombinant	E. coli	GST	
P00441	Superoxide dismutase [Cu-Zn]	15,800	Erythrocytes			Acetylation
P10599	Thioredoxin	12,424	Recombinant	E. coli	6-His	<u> </u>
P01375	Tumor necrosis factor [TNF-alpha]	17,350	Recombinant	E. coli		
P62988	Ubiquitin	9,387	Recombinant	E. coli	6-His	
P63279	Ubiquitin-conjugating enzyme E2 I [UbcH9]	17,995	Recombinant	E. coli		
O00762	Ubiquitin-conjugating enzyme E2 C [UbcH10]	20,473	Recombinant	E. coli	6-His	
P51965	Ubiquitin-conjugating enzyme E2 E1 [UbcH6]	22,222	Recombinant	E. coli	6-His	

^{*} As reported in UniProt. Potential PTMs have not been verified by Sigma.