

A Mass Spectrometric-Derived Cell Surface Protein Atlas

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Citation Report

#	ARTICLE	IF	CITATIONS
1	A Combined Omics Approach to Generate the Surface Atlas of Human Naive CD4+ T Cells during Early T-Cell Receptor Activation. <i>Molecular and Cellular Proteomics</i> , 2015, 14, 2085-2102.	2.5	40
2	Bioinformatics Analysis of the Human Surfaceome Reveals New Targets for a Variety of Tumor Types. <i>International Journal of Genomics</i> , 2016, 2016, 1-7.	0.8	13
3	Identification of glycoproteins associated with HIV latently infected cells using quantitative glycoproteomics. <i>Proteomics</i> , 2016, 16, 1872-1880.	1.3	10
4	N-glycosylation proteome enrichment analysis in kidney reveals differences between diabetic mouse models. <i>Clinical Proteomics</i> , 2016, 13, 22.	1.1	13
5	Mapping the Cell-Surface N-Glycoproteome of Human Hepatocytes Reveals Markers for Selecting a Homogeneous Population of iPSC-Derived Hepatocytes. <i>Stem Cell Reports</i> , 2016, 7, 543-556.	2.3	44
6	Highlights of the Biology and Disease-driven Human Proteome Project, 2015-2016. <i>Journal of Proteome Research</i> , 2016, 15, 3979-3987.	1.8	21
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15	Identification of Extracellular Segments by Mass Spectrometry Improves Topology Prediction of Transmembrane Proteins. <i>Scientific Reports</i> , 2017, 7, 42610.	1.6	15
16	Single-cell RNA-seq reveals new types of human blood dendritic cells, monocytes, and progenitors. <i>Science</i> , 2017, 356, .	6.0	1,846
17	Global analysis of glycoproteins identifies markers of endotoxin tolerant monocytes and GPR84 as a modulator of TNF α expression. <i>Scientific Reports</i> , 2017, 7, 838.	1.6	39
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19	Using hyperLOPIT to perform high-resolution mapping of the spatial proteome. <i>Nature Protocols</i> , 2017, 12, 1110-1135.	5.5	113
20	Aminopeptidase N is not required for porcine epidemic diarrhea virus cell entry. <i>Virus Research</i> , 2017, 235, 6-13.	1.1	74
21	Monitoring Cell-surface N-Glycoproteome Dynamics by Quantitative Proteomics Reveals Mechanistic Insights into Macrophage Differentiation. <i>Molecular and Cellular Proteomics</i> , 2017, 16, 770-785.	2.5	41
22	<i>N</i>-glycoprotein surfaceome of human induced pluripotent stem cell derived hepatic endoderm. <i>Proteomics</i> , 2017, 17, 1600397.	1.3	19
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