

Sasha A Singh

List of Publications by Year in descending order

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70
papers

2,703
citations

257450

24
h-index

197818

49
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78
all docs

78
docs citations

78
times ranked

4401
citing authors

#	ARTICLE	IF	CITATIONS
1	A Novel Spectral Annotation Strategy Streamlines Reporting of Mono-ADP-ribosylated Peptides Derived from Mouse Liver and Spleen in Response to IFN- β . <i>Molecular and Cellular Proteomics</i> , 2022, 21, 100153.	3.8	5
2	Lipoprotein(a) Induces Vesicular Cardiovascular Calcification Revealed With Single-Extracellular Vesicle Analysis. <i>Frontiers in Cardiovascular Medicine</i> , 2022, 9, 778919.	2.4	12
3	Second Heart Field-Derived Cells Contribute to Angiotensin II-Mediated Ascending Aortopathies. <i>Circulation</i> , 2022, 145, 987-1001.	1.6	18
4	Prothymosin Alpha: A Novel Contributor to Estradiol Receptor Alpha-Mediated CD8 ⁺ T-Cell Pathogenic Responses and Recognition of Type 1 Collagen in Rheumatic Heart Valve Disease. <i>Circulation</i> , 2022, 145, 531-548.	1.6	12
5	The RiboMaP Spectral Annotation Method Applied to Various ADP-Ribosylome Studies Including INF- β -Stimulated Human Cells and Mouse Tissues. <i>Frontiers in Cardiovascular Medicine</i> , 2022, 9, 851351.	2.4	3
6	A disease-driver population within interstitial cells of human calcific aortic valves identified via single-cell and proteomic profiling. <i>Cell Reports</i> , 2022, 39, 110685.	6.4	16
7	ApoC-III is a novel inducer of calcification in human aortic valves. <i>Journal of Biological Chemistry</i> , 2021, 296, 100193.	3.4	28
8	CROT (Carnitine O-Octanoyltransferase) Is a Novel Contributing Factor in Vascular Calcification via Promoting Fatty Acid Metabolism and Mitochondrial Dysfunction. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2021, 41, 755-768.	2.4	17
9	Dynamamin-related protein 1 inhibition reduces hepatic PCSK9 secretion. <i>Cardiovascular Research</i> , 2021, 117, 2340-2353.	3.8	16
10	Metabolism of PLTP, CETP, and LCAT on multiple HDL sizes using the Orbitrap Fusion Lumos. <i>JCI Insight</i> , 2021, 6, .	5.0	12
11	Systems Approach to Discovery of Therapeutic Targets for Vein Graft Disease: PPAR α Pivotaly Regulates Metabolism, Activation, and Heterogeneity of Macrophages and Lesion Development. <i>Circulation</i> , 2021, 143, 2454-2470.	1.6	21
12	Elastogenesis Correlates With Pigment Production in Murine Aortic Valve Leaflets. <i>Frontiers in Cardiovascular Medicine</i> , 2021, 8, 678401.	2.4	4
13	Highly Selective PPAR α (Peroxisome Proliferator-Activated Receptor α) Agonist Pemafibrate Inhibits Stent Inflammation and Restenosis Assessed by Multimodality Molecular-Microstructural Imaging. <i>Journal of the American Heart Association</i> , 2021, 10, e020834.	3.7	7
14	Integration of Functional Imaging, Cytometry, and Unbiased Proteomics Reveals New Features of Endothelial-to-Mesenchymal Transition in Ischemic Mitral Valve Regurgitation in Human Patients. <i>Frontiers in Cardiovascular Medicine</i> , 2021, 8, 688396.	2.4	0
15	Computational Screening Strategy for Drug Repurposing Identified Niclosamide as Inhibitor of Vascular Calcification. <i>Frontiers in Cardiovascular Medicine</i> , 2021, 8, 826529.	2.4	5
16	Multiorgan Systems Study Reveals Igfbp7 as a Suppressor of Gluconeogenesis after Gastric Bypass Surgery. <i>Journal of Proteome Research</i> , 2020, 19, 129-143.	3.7	4
17	Annexin A1-dependent tethering promotes extracellular vesicle aggregation revealed with single-extracellular vesicle analysis. <i>Science Advances</i> , 2020, 6, .	10.3	65
18	S100A9-RAGE Axis Accelerates Formation of Macrophage-Mediated Extracellular Vesicle Microcalcification in Diabetes Mellitus. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2020, 40, 1838-1853.	2.4	52

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19	The impact of PARPs and ADP-ribosylation on inflammation and host-pathogen interactions. <i>Genes and Development</i> , 2020, 34, 341-359.	5.9	157
20	Calcific Aortic Valve Disease Is Timely, But Are We Looking Too Late?. <i>JACC Basic To Translational Science</i> , 2020, 5, 1178-1180.	4.1	5
21	Sphingosine 1-phosphate-regulated transcriptomes in heterogenous arterial and lymphatic endothelium of the aorta. <i>ELife</i> , 2020, 9, .	6.0	34
22	Abstract 13401: Prothymosin Alpha (Prot±) Associates With Pathogenesis and Sex Predisposition in Rheumatic Heart Valve Disease. <i>Circulation</i> , 2020, 142, .	1.6	0
23	Effects of Replacing Dietary Monounsaturated Fat With Carbohydrate on HDL (High-Density) Tj ETQq1 1 0.784314 rgBT /Overlock 10 and <i>Vascular Biology</i> , 2019, 39, 2411-2430.	2.4	15
24	Standardization of Human Calcific Aortic Valve Disease in vitro Modeling Reveals Passage-Dependent Calcification. <i>Frontiers in Cardiovascular Medicine</i> , 2019, 6, 49.	2.4	49
25	A Study into the ADP-Ribosylome of IFN- β -Stimulated THP-1 Human Macrophage-like Cells Identifies ARTD8/PARP14 and ARTD9/PARP9 ADP-Ribosylation. <i>Journal of Proteome Research</i> , 2019, 18, 1607-1622.	3.7	21
26	Uremic Toxin Indoxyl Sulfate Promotes Proinflammatory Macrophage Activation Via the Interplay of OATP2B1 and DLL4-Notch Signaling. <i>Circulation</i> , 2019, 139, 78-96.	1.6	126
27	¹⁸ F-Fluoride Signal Amplification Identifies Microcalcifications Associated With Atherosclerotic Plaque Instability in Positron Emission Tomography/Computed Tomography Images. <i>Circulation: Cardiovascular Imaging</i> , 2019, 12, e007835.	2.6	92
28	XINA: A Workflow for the Integration of Multiplexed Proteomics Kinetics Data with Network Analysis. <i>Journal of Proteome Research</i> , 2019, 18, 775-781.	3.7	13
29	Abstract MP41: A Diet High in Carbohydrate and Low in Fat Alters the HDL Proteome and Metabolism of 9 HDL Proteins in Humans. <i>Circulation</i> , 2019, 139, .	1.6	1
30	Spatiotemporal Multi-Omics Mapping Generates a Molecular Atlas of the Aortic Valve and Reveals Networks Driving Disease. <i>Circulation</i> , 2018, 138, 377-393.	1.6	180
31	Context-enriched interactome powered by proteomics helps the identification of novel regulators of macrophage activation. <i>ELife</i> , 2018, 7, .	6.0	11
32	Abstract 608: Integrated Omics and Network Analysis Identify Drivers of Calcific Bicuspid Aortic Valve Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2018, 38, .	2.4	0
33	Abstract 166: Effects of Dietary Unsaturated Fat and Carbohydrate on the HDL Proteome and Metabolism of 9 HDL Proteins Across 6 HDL Size Fractions in Humans. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2018, 38, .	2.4	0
34	Abstract 663: Nuclear RSK1 Induces Pro-inflammatory Activation of Macrophages through STAT1 Phosphorylation at Ser727. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2018, 38, .	2.4	0
35	Abstract 175: Dynamin-Related Protein 1 Regulates Proteostasis and Proprotein Convertase Subtilisin/Kexin Type 9 Secretion. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2018, 38, .	2.4	0
36	Abstract 708: Using Global Proteomics and Network Science to Explore Therapeutic Targets for Abdominal Aortic Aneurysm (AAA). <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2018, 38, .	2.4	0

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37	Abstract 329: Systems-based Target Discovery Reveals PPAR α as a Key Metabolic Regulator for Macrophage Activation in Vein Graft Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2018, 38, .	2.4	0
38	Abstract 228: Multi-omics Mapping Generates a Molecular Atlas of the Aortic Valve and Reveals Networks Driving Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2018, 38, .	2.4	0
39	Unbiased and targeted mass spectrometry for the HDL proteome. <i>Current Opinion in Lipidology</i> , 2017, 28, 68-77.	2.7	15
40	Inside Front Cover: PRM technology enables detection and reliable automated quantification of low tracer incorporation into circulating apolipoproteins for HDL metabolism studies. <i>Proteomics</i> , 2017, 17, 1770014.	2.2	0
41	Automation of PRM-dependent D3 ϵ -Leu tracer enrichment in HDL to study the metabolism of apoA ϵ , LCAT and other apolipoproteins. <i>Proteomics</i> , 2017, 17, 1600085.	2.2	4
42	Current Trends and Future Perspectives of State-of-the-Art Proteomics Technologies Applied to Cardiovascular Disease Research. <i>Circulation Journal</i> , 2016, 80, 1674-1683.	1.6	11
43	PARP9 and PARP14 cross-regulate macrophage activation via STAT1 ADP-ribosylation. <i>Nature Communications</i> , 2016, 7, 12849.	12.8	214
44	Endophenotype Network Models: Common Core of Complex Diseases. <i>Scientific Reports</i> , 2016, 6, 27414.	3.3	72
45	Multiple apolipoprotein kinetics measured in human HDL by high-resolution/accurate mass parallel reaction monitoring. <i>Journal of Lipid Research</i> , 2016, 57, 714-728.	4.2	35
46	Sortilin mediates vascular calcification via its recruitment into extracellular vesicles. <i>Journal of Clinical Investigation</i> , 2016, 126, 1323-1336.	8.2	196
47	Abstract 538: Metabolism of Multiple Apolipoproteins Across HDL Size in Humans. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2016, 36, .	2.4	0
48	New CETP inhibitor K-312 reduces PCSK9 expression: a potential effect on LDL cholesterol metabolism. <i>American Journal of Physiology - Endocrinology and Metabolism</i> , 2015, 309, E177-E190.	3.5	38
49	mIMT-visHTS: A novel method for multiplexing isobaric mass tagged datasets with an accompanying visualization high throughput screening tool for protein profiling. <i>Journal of Proteomics</i> , 2015, 128, 132-140.	2.4	7
50	Mass spectrometry meets the challenge of understanding the complexity of the lipoproteome: recent findings regarding proteins involved in dyslipidemia and cardiovascular disease. <i>Expert Review of Proteomics</i> , 2015, 12, 519-532.	3.0	7
51	Abstract 322: HDL Dynamics in Circulation: Complexity of Protein Distribution and Metabolism Across HDL Size. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2015, 35, .	2.4	0
52	Cystathionine β -lyase Accelerates Osteoclast Differentiation. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2014, 34, 626-634.	2.4	37
53	Co-regulation proteomics reveals substrates and mechanisms of APC/C-dependent degradation. <i>EMBO Journal</i> , 2014, 33, 385-399.	7.8	72
54	Enrichment of calcifying extracellular vesicles using density-based ultracentrifugation protocol. <i>Journal of Extracellular Vesicles</i> , 2014, 3, 25129.	12.2	39

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55	Abstract 18687: PARP9 and PARP14 are Novel Regulators of Macrophage Activation. <i>Circulation</i> , 2014, 130, .	1.6	0
56	FLEXIQinase, a mass spectrometryâ€‘based assay, to unveil multikinase mechanisms. <i>Nature Methods</i> , 2012, 9, 504-508.	19.0	26
57	Tiki1 Is Required for Head Formation via Wnt Cleavage-Oxidation and Inactivation. <i>Cell</i> , 2012, 149, 1565-1577.	28.9	125
58	A Practical Guide to the FLEXIQuant Method. <i>Methods in Molecular Biology</i> , 2012, 893, 295-319.	0.9	19
59	Interaction of survival of motor neuron (SMN) and HuD proteins with mRNA cpg15 rescues motor neuron axonal deficits. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 10337-10342.	7.1	185
60	The Application of High Throughput Mass Spectrometry to the Analysis of Glycoproteins. , 2010, , 103-125.		0
61	A simple and effective method for detecting phosphopeptides for phosphoproteomic analysis. <i>Journal of Proteomics</i> , 2009, 72, 831-835.	2.4	8
62	FLEXIQuant: A Novel Tool for the Absolute Quantification of Proteins, and the Simultaneous Identification and Quantification of Potentially Modified Peptides. <i>Journal of Proteome Research</i> , 2009, 8, 2201-2210.	3.7	109
63	A Phylogenomic Analysis of the Shikimate Dehydrogenases Reveals Broadscale Functional Diversification and Identifies One Functionally Distinct Subclass. <i>Molecular Biology and Evolution</i> , 2008, 25, 2221-2232.	8.9	20
64	The DHQ-dehydroshikimate-SDH-shikimate-NADP(H) Complex: Insights into Metabolite Transfer in the Shikimate Pathway,. <i>Crystal Growth and Design</i> , 2007, 7, 2153-2160.	3.0	16
65	Structure ofArabidopsisDehydroquininate Dehydratase-Shikimate Dehydrogenase and Implications for Metabolic Channeling in the Shikimate Pathwayâ€‘â€‘. <i>Biochemistry</i> , 2006, 45, 7787-7796.	2.5	72
66	Biochemical characterization of prephenate dehydrogenase from the hyperthermophilic bacteriumAquifex aeolicus. <i>Protein Science</i> , 2006, 15, 1417-1432.	7.6	15
67	Crystal Structure of Prephenate Dehydrogenase from Aquifex aeolicus. <i>Journal of Biological Chemistry</i> , 2006, 281, 12919-12928.	3.4	19
68	Crystal Structure of a Novel Shikimate Dehydrogenase from Haemophilus influenzae. <i>Journal of Biological Chemistry</i> , 2005, 280, 17101-17108.	3.4	33
69	Overlapping and non-redundant functions of the Arabidopsis auxin response factors MONOPTEROS and NONPHOTOTROPIC HYPOCOTYL 4. <i>Development (Cambridge)</i> , 2004, 131, 1089-1100.	2.5	302
70	Promise of a Novel Bedside-to-Bench Paradigm: Can Percutaneous Coronary Intervention Proteomics Balloon Into Clinical Practice?. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 0, , .	2.4	0