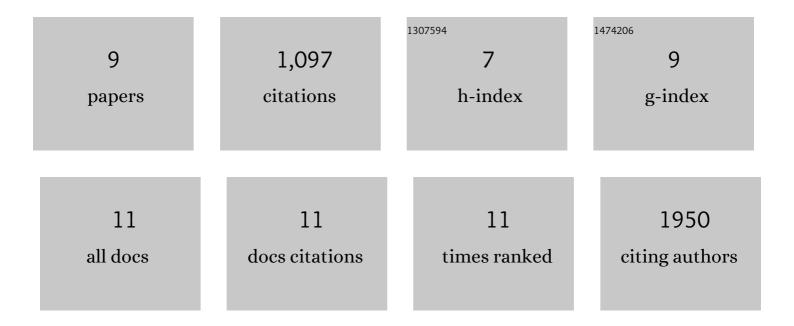
Hasmik Keshishian

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9728155/publications.pdf Version: 2024-02-01



HASMIK KESHISHIAN

#	Article	IF	CITATIONS
1	TAILS Identifies Candidate Substrates and Biomarkers of ADAMTS7, a Therapeutic Protease Target in Coronary Artery Disease. Molecular and Cellular Proteomics, 2022, 21, 100223.	3.8	7
2	A highly multiplexed quantitative phosphosite assay for biology and preclinical studies. Molecular Systems Biology, 2021, 17, e10156.	7.2	12
3	Patterns of substrate affinity, competition, and degradation kinetics underlie biological activity of thalidomide analogs. Blood, 2019, 134, 160-170.	1.4	41
4	Quantitative, multiplexed workflow for deep analysis of human blood plasma and biomarker discovery by mass spectrometry. Nature Protocols, 2017, 12, 1683-1701.	12.0	139
5	Large-Scale Interlaboratory Study to Develop, Analytically Validate and Apply Highly Multiplexed, Quantitative Peptide Assays to Measure Cancer-Relevant Proteins in Plasma. Molecular and Cellular Proteomics, 2015, 14, 2357-2374.	3.8	153
6	Multiplexed, Quantitative Workflow for Sensitive Biomarker Discovery in Plasma Yields Novel Candidates for Early Myocardial Injury. Molecular and Cellular Proteomics, 2015, 14, 2375-2393.	3.8	175
7	Simplified and Efficient Quantification of Low-abundance Proteins at Very High Multiplex via Targeted Mass Spectrometry. Molecular and Cellular Proteomics, 2014, 13, 1137-1149.	3.8	63
8	Developing Multiplexed Assays for Troponin I and Interleukin-33 in Plasma by Peptide Immunoaffinity Enrichment and Targeted Mass Spectrometry. Clinical Chemistry, 2009, 55, 1108-1117.	3.2	243
9	Quantification of Cardiovascular Biomarkers in Patient Plasma by Targeted Mass Spectrometry and Stable Isotope Dilution. Molecular and Cellular Proteomics, 2009, 8, 2339-2349.	3.8	263