## Rahul S Samant

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Alternative systems for misfolded protein clearance: life beyond the proteasome. FEBS Journal, 2021, 288, 4464-4487.	4.7	49
2	Native mass spectrometry analyses of chaperonin complex TRiC/CCT reveal subunit N-terminal processing and re-association patterns. Scientific Reports, 2021, 11, 13084.	3.3	7
3	Dosage compensation plans: protein aggregation provides additional insurance against aneuploidy. Genes and Development, 2019, 33, 1027-1030.	5.9	3
4	Methods for measuring misfolded protein clearance in the budding yeast Saccharomyces cerevisiae. Methods in Enzymology, 2019, 619, 27-45.	1.0	7
5	Distinct proteostasis circuits cooperate in nuclear and cytoplasmic protein quality control. Nature, 2018, 563, 407-411.	27.8	141
6	Mechanisms and Functions of Spatial Protein Quality Control. Annual Review of Biochemistry, 2017, 86, 97-122.	11.1	225
7	Abstract 3025: Discovery of preclinical development candidate inhibitors of the mediator complex-associated kinases CDK8 and CDK19 and evaluation of their therapeutic potential. , 2016, , .		1
8	Abstract 4549: Proteomic analysis of ubiquitination identifies the interplay between HSP90 inhibition and CUL5 in the control of autophagy. , 2016, , .		0
9	Abstract 3869: Phosphoproteomic-based identification of CDK8/CDK19 substrates in colorectal cancer. , 2016, , .		0
10	A selective chemical probe for exploring the role of CDK8 and CDK19 in human disease. Nature Chemical Biology, 2015, 11, 973-980.	8.0	114
11	E3 ubiquitin ligase Cullin-5 modulates multiple molecular and cellular responses to heat shock protein 90 inhibition in human cancer cells. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 6834-6839.	7.1	47
12	ATP-competitive inhibitors block protein kinase recruitment to the Hsp90-Cdc37 system. Nature Chemical Biology, 2013, 9, 307-312.	8.0	132
13	The expanding proteome of the molecular chaperone HSP90. Cell Cycle, 2012, 11, 1301-1308.	2.6	37
14	Choose your protein partners. Nature, 2012, 490, 351-352.	27.8	10