## Sobia Idrees

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11658180/publications.pdf

Version: 2024-02-01

		1163117	1058476	
14	235	8	14	
papers	citations	h-index	g-index	
15	15	15	368	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	A brief review on dengue molecular virology, diagnosis, treatment and prevalence in Pakistan. Genetic Vaccines and Therapy, 2012, 10, 6.	1.5	54
2	RNAi: antiviral therapy against dengue virus. Asian Pacific Journal of Tropical Biomedicine, 2013, 3, 232-236.	1.2	44
3	Medicinal plants against hepatitis C virus. World Journal of Gastroenterology, 2014, 20, 2941.	3.3	27
4	Computer Aided Screening of Phytochemicals from Garcinia against the Dengue NS2B/NS3 Protease. Bioinformation, 2014, 10, 115-118.	0.5	23
5	Structural analysis and epitope prediction of HCV E1 protein isolated in Pakistan: an in-silico approach. Virology Journal, 2013, 10, 113.	3.4	18
6	Missense mutation in SLC4All in two Pakistani families affected with congenital hereditary endothelial dystrophy (CHED2). Australasian journal of optometry, The, 2016, 99, 73-77.	1.3	16
7	Discovery and design of cyclic peptides as dengue virus inhibitors through structure-based molecular docking. Asian Pacific Journal of Tropical Medicine, 2014, 7, 513-516.	0.8	11
8	SLiM-Enrich: computational assessment of protein–protein interaction data as a source of domain-motif interactions. PeerJ, 2018, 6, e5858.	2.0	11
9	In silico analysis of five missense mutations in CYP1B1 gene in Pakistani families affected with primary congenital glaucoma. International Ophthalmology, 2018, 38, 807-814.	1.4	8
10	Molecular screening of phytochemicals from Amelanchier Alnifolia against HCV NS3 protease/helicase using computational docking techniques. Bioinformation, 2013, 9, 978-982.	0.5	8
11	HCV Envelope protein 2 sequence comparison of Pakistani isolate and In-silico prediction of conserved epitopes for vaccine development. Journal of Translational Medicine, 2013, 11, 105.	4.4	5
12	Development of global consensus sequence of HCV glycoproteins involved in viral entry. Theoretical Biology and Medical Modelling, 2013, 10, 24.	2.1	5
13	Global Consensus Sequence Development and Analysis of Dengue NS3 Conserved Domains. BioResearch Open Access, 2013, 2, 392-396.	2.6	3
14	Gene Expression Profiling of Immune Responsive and Fibrosis Genes in Hepatitis C Virus Infected Patients. Viral Immunology, 2014, 27, 250-254.	1.3	2