Sabyasachi Das

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6758040/publications.pdf

Version: 2024-02-01

331538 377752 2,228 38 21 34 h-index citations g-index papers 39 39 39 3236 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Sequencing of the sea lamprey (Petromyzon marinus) genome provides insights into vertebrate evolution. Nature Genetics, 2013, 45, 415-421.	9.4	588
2	Molecular signature of hypersaline adaptation: insights from genome and proteome composition of halophilic prokaryotes. Genome Biology, 2008, 9, R70.	13.9	282
3	Evolutionary implications of a third lymphocyte lineage in lampreys. Nature, 2013, 501, 435-438.	13.7	180
4	The Evolution of Adaptive Immunity in Vertebrates. Advances in Immunology, 2011, 109, 125-157.	1.1	158
5	MicroRNA 125b inhibition of B cell differentiation in germinal centers. International Immunology, 2010, 22, 583-592.	1.8	141
6	Evolution of Alternative Adaptive Immune Systems in Vertebrates. Annual Review of Immunology, 2018, 36, 19-42.	9.5	92
7	Analysis of Nanoarchaeum equitans genome and proteome composition: indications for hyperthermophilic and parasitic adaptation. BMC Genomics, 2006, 7, 186.	1.2	67
8	Synonymous codon usage in adenoviruses: Influence of mutation, selection and protein hydropathy. Virus Research, 2006, 117, 227-236.	1.1	66
9	Characterization of Lamprey IL-17 Family Members and Their Receptors. Journal of Immunology, 2015, 195, 5440-5451.	0.4	56
10	Evolutionary redefinition of immunoglobulin light chain isotypes in tetrapods using molecular markers. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 16647-16652.	3.3	54
11	Evolutionary dynamics of the immunoglobulin heavy chain variable region genes in vertebrates. Immunogenetics, 2008, 60, 47-55.	1.2	53
12	Definition of a third <i>VLR</i> gene in hagfish. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 15013-15018.	3.3	53
13	Organization of lamprey (i) variable lymphocyte receptor C (li) locus and repertoire development. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 6043-6048.	3.3	49
14	Characterization of Lamprey BAFF-like Gene: Evolutionary Implications. Journal of Immunology, 2016, 197, 2695-2703.	0.4	33
15	Distinct, ecotype-specific genome and proteome signatures in the marine cyanobacteria Prochlorococcus. BMC Genomics, 2010, 11, 103.	1.2	31
16	Codon and Amino Acid Usage in Two Major Human Pathogens of Genus Bartonella Optimization Between Replicational-Transcriptional Selection, Translational Control and Cost Minimization. DNA Research, 2005, 12, 91-102.	1.5	30
17	Analysis of the Immunoglobulin Light Chain Genes in Zebra Finch: Evolutionary Implications. Molecular Biology and Evolution, 2010, 27, 113-120.	3.5	30
18	Evolutionary Genomics of Immunoglobulin-Encoding Loci in Vertebrates. Current Genomics, 2012, 13, 95-102.	0.7	29

#	Article	IF	CITATIONS
19	Comparative Genomics and Evolution of the Alpha-Defensin Multigene Family in Primates. Molecular Biology and Evolution, 2010, 27, 2333-2343.	3.5	28
20	Evolution of two prototypic T cell lineages. Cellular Immunology, 2015, 296, 87-94.	1.4	25
21	Evolutionary Origin and Genomic Organization of Micro-RNA Genes in Immunoglobulin Lambda Variable Region Gene Family. Molecular Biology and Evolution, 2009, 26, 1179-1189.	3.5	22
22	Diazepam Accelerates GABAAR Synaptic Exchange and Alters Intracellular Trafficking. Frontiers in Cellular Neuroscience, 2019, 13, 163.	1.8	22
23	Genomic organization and evolution of immunoglobulin kappa gene enhancers and kappa deleting element in mammals. Molecular Immunology, 2009, 46, 3171-3177.	1.0	20
24	Compositional variation in bacterial genes and proteins with potential expression level. FEBS Letters, 2005, 579, 5205-5210.	1.3	18
25	Genomic donor cassette sharing during <i>VLRA</i> and <i>VLRC</i> assembly in jawless vertebrates. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 14828-14833.	3.3	18
26	Evolutionary Constraints on Codon and Amino Acid Usage in Two Strains of Human Pathogenic Actinobacteria Tropheryma whipplei. Journal of Molecular Evolution, 2006, 62, 645-658.	0.8	17
27	Comparative Analyses of Codon and Amino Acid Usage in Symbiotic Island and Core Genome in Nitrogen-Fixing Symbiotic BacteriumBradyrhizobium japonicum. Journal of Biomolecular Structure and Dynamics, 2005, 23, 221-232.	2.0	14
28	Artemisinin combination therapy fails even in the absence of Plasmodium falciparum kelch13 gene polymorphism in Central India. Scientific Reports, 2021, 11, 9946.	1.6	14
29	Immune Related Genes Underpin the Evolution of Adaptive Immunity in Jawless Vertebrates. Current Genomics, 2012, 13, 86-94.	0.7	11
30	Comparative Genomics and Evolution of Immunoglobulin-Encoding Loci in Tetrapods. Advances in Immunology, 2011, 111, 143-178.	1.1	7
31	Editorial [Hot Topic: Comparative Genomics and Genome Evolution (Guest Editors: Sabyasachi Das and) Tj ETQq1	1.0.78431 0.7	14 rgBT /O
32	Evolution of variable lymphocyte receptor B antibody loci in jawless vertebrates. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118 , .	3.3	6
33	Ancient BCMA-like Genes Herald B Cell Regulation in Lampreys. Journal of Immunology, 2019, 203, 2909-2916.	0.4	3
34	A novel nano-anti-malarial induces redox damage and elicits cytokine response to the parasite. Cytokine, 2021, 144, 155555.	1.4	2
35	B Cells and Antibodies in Jawless Vertebrates. , 2015, , 121-132.		1
36	DDE Transposon as Public Goods. , 2020, , 337-357.		1

#	Article	IF	CITATIONS
37	TRENDS IN CODON AND AMINO ACID USAGE IN HUMAN PATHOGEN <i>TROPHERYMA WHIPPLEI,</i> ONLY KNOWN ACTINOBACTERIA WITH REDUCED GENOME., 2005,,.		O
38	CONSEQUENCES OF MUTATION, SELECTION AND PHYSICO-CHEMICAL PROPERTIES OF ENCODED PROTEINS ON SYNONYMOUS CODON USAGE IN ADENOVIRUSES. , 2005, , .		0