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List of Publications by Year in descending order

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Version: 2024-02-01

31
papers

14,834
citations

394286

19
h-index

526166

27
g-index

33
all docs

33
docs citations

33
times ranked

27303
citing authors

#	ARTICLE	IF	CITATIONS
1	ML-AdVInfect: A Machine-Learning Based Adenoviral Infection Predictor. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 647424.	1.6	10
2	Comparison of protein expression profile of limb regeneration between neotenic and metamorphic axolotl. <i>Biochemical and Biophysical Research Communications</i> , 2020, 522, 428-434.	1.0	19
3	Preclinical Molecular Signatures of Spinal Cord Functional Restoration: Optimizing the Metamorphic Axolotl (<i>Ambystoma mexicanum</i>) Model in Regenerative Medicine. <i>OMICS A Journal of Integrative Biology</i> , 2020, 24, 370-378.	1.0	8
4	In Silico Analysis of a Highly Mutated Gene in Cancer Provides Insight into Abnormal mRNA Splicing: Splicing Factor 3B Subunit 1K700E Mutant. <i>Biomolecules</i> , 2020, 10, 680.	1.8	6
5	Taxonomic diversity-based domain interaction prediction. <i>Pamukkale University Journal of Engineering Sciences</i> , 2019, 25, 215-222.	0.2	0
6	In silico Analysis of Vaccination Adverse Events. <i>Journal of Allergy and Clinical Immunology</i> , 2015, 135, AB104.	1.5	0
7	UniRef clusters: a comprehensive and scalable alternative for improving sequence similarity searches. <i>Bioinformatics</i> , 2015, 31, 926-932.	1.8	1,223
8	UniProt: a hub for protein information. <i>Nucleic Acids Research</i> , 2015, 43, D204-D212.	6.5	4,370
9	In silico analysis of autoimmune diseases and genetic relationships to vaccination against infectious diseases. <i>BMC Immunology</i> , 2014, 15, 61.	0.9	21
10	Activities at the Universal Protein Resource (UniProt). <i>Nucleic Acids Research</i> , 2014, 42, D191-D198.	6.5	1,162
11	A Novel Knowledge Representation Framework for the Statistical Validation of Quantitative Imaging Biomarkers. <i>Journal of Digital Imaging</i> , 2013, 26, 614-629.	1.6	6
12	Quantitative Imaging Biomarker Ontology (QIBO) for Knowledge Representation of Biomedical Imaging Biomarkers. <i>Journal of Digital Imaging</i> , 2013, 26, 630-641.	1.6	12
13	A fast Peptide Match service for UniProt Knowledgebase. <i>Bioinformatics</i> , 2013, 29, 2808-2809.	1.8	101
14	In silico analysis of autoimmune diseases and genetic relationships to vaccination against infectious diseases. , 2013, , .		0
15	Reorganizing the protein space at the Universal Protein Resource (UniProt). <i>Nucleic Acids Research</i> , 2012, 40, D71-D75.	6.5	1,196
16	Update on activities at the Universal Protein Resource (UniProt) in 2013. <i>Nucleic Acids Research</i> , 2012, 41, D43-D47.	6.5	620
17	Life sciences domain analysis model. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2012, 19, 1095-1102.	2.2	6
18	Informatics and data quality at collaborative multicenter Breast and Colon Cancer Family Registries. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2012, 19, e125-e128.	2.2	10

#	ARTICLE	IF	CITATIONS
19	Ongoing and future developments at the Universal Protein Resource. <i>Nucleic Acids Research</i> , 2011, 39, D214-D219.	6.5	649
20	A comprehensive protein-centric ID mapping service for molecular data integration. <i>Bioinformatics</i> , 2011, 27, 1190-1191.	1.8	70
21	The Universal Protein Resource (UniProt) 2009. <i>Nucleic Acids Research</i> , 2009, 37, D169-D174.	6.5	548
22	Infrastructure for the life sciences: design and implementation of the UniProt website. <i>BMC Bioinformatics</i> , 2009, 10, 136.	1.2	405
23	Orchestrating caGrid Services in Taverna. , 2008, , .		3
24	UniRef: comprehensive and non-redundant UniProt reference clusters. <i>Bioinformatics</i> , 2007, 23, 1282-1288.	1.8	1,144
25	The Universal Protein Resource (UniProt). <i>Nucleic Acids Research</i> , 2007, 35, D193-D197.	6.5	488
26	The Universal Protein Resource (UniProt). <i>Nucleic Acids Research</i> , 2007, 36, D190-D195.	6.5	852
27	The Universal Protein Resource (UniProt): an expanding universe of protein information. <i>Nucleic Acids Research</i> , 2006, 34, D187-D191.	6.5	961
28	PIRSF: family classification system at the Protein Information Resource. <i>Nucleic Acids Research</i> , 2004, 32, 112D-114.	6.5	193
29	The Protein Information Resource. <i>Nucleic Acids Research</i> , 2003, 31, 345-347.	6.5	385
30	The Protein Information Resource: an integrated public resource of functional annotation of proteins. <i>Nucleic Acids Research</i> , 2002, 30, 35-37.	6.5	186
31	A probabilistic method for identifying start codons in bacterial genomes. <i>Bioinformatics</i> , 2001, 17, 1123-1130.	1.8	180