

Judith A Blake

List of Publications by Year in descending order

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101
papers

47,361
citations

41344

49
h-index

38395

95
g-index

117
all docs

117
docs citations

117
times ranked

68716
citing authors

#	ARTICLE	IF	CITATIONS
1	Mouse Genome Informatics (MGI): latest news from MGD and GXD. <i>Mammalian Genome</i> , 2022, 33, 4-18.	2.2	30
2	Harmonizing model organism data in the Alliance of Genome Resources. <i>Genetics</i> , 2022, 220, .	2.9	52
3	Reactome and the Gene Ontology: digital convergence of data resources. <i>Bioinformatics</i> , 2021, 37, 3343-3348.	4.1	19
4	Mouse Genome Database (MGD): Knowledgebase for mouseâ€“human comparative biology. <i>Nucleic Acids Research</i> , 2021, 49, D981-D987.	14.5	179
5	The Gene Ontology resource: enriching a GOld mine. <i>Nucleic Acids Research</i> , 2021, 49, D325-D334.	14.5	2,416
6	Alliance of Genome Resources Portal: unified model organism research platform. <i>Nucleic Acids Research</i> , 2020, 48, D650-D658.	14.5	145
7	Investigation of COVID-19 comorbidities reveals genes and pathways coincident with the SARS-CoV-2 viral disease. <i>Scientific Reports</i> , 2020, 10, 20848.	3.3	32
8	Cisplatin-resistant triple-negative breast cancer subtypes: multiple mechanisms of resistance. <i>BMC Cancer</i> , 2019, 19, 1039.	2.6	77
9	An effective biomedical document classification scheme in support of biocuration: addressing class imbalance. <i>Database: the Journal of Biological Databases and Curation</i> , 2019, 2019, .	3.0	15
10	Curating gene sets: challenges and opportunities for integrative analysis. <i>Database: the Journal of Biological Databases and Curation</i> , 2019, 2019, .	3.0	5
11	RNAcentral: a hub of information for non-coding RNA sequences. <i>Nucleic Acids Research</i> , 2019, 47, D221-D229.	14.5	153
12	Mouse Genome Database (MGD) 2019. <i>Nucleic Acids Research</i> , 2019, 47, D801-D806.	14.5	625
13	Improving Interpretation of Cardiac Phenotypes and Enhancing Discovery With Expanded Knowledge in the Gene Ontology. <i>Circulation Genomic and Precision Medicine</i> , 2018, 11, e001813.	3.6	24
14	Model organism data evolving in support of translational medicine. <i>Lab Animal</i> , 2018, 47, 277-289.	0.4	35
15	Mouse Genome Database (MGD)-2018: knowledgebase for the laboratory mouse. <i>Nucleic Acids Research</i> , 2018, 46, D836-D842.	14.5	241
16	Protein Ontology (PRO): enhancing and scaling up the representation of protein entities. <i>Nucleic Acids Research</i> , 2017, 45, D339-D346.	14.5	73
17	Mouse Genome Database (MGD)-2017: community knowledge resource for the laboratory mouse. <i>Nucleic Acids Research</i> , 2017, 45, D723-D729.	14.5	255
18	Mouse Genome Informatics (MGI): Resources for Mining Mouse Genetic, Genomic, and Biological Data in Support of Primary and Translational Research. <i>Methods in Molecular Biology</i> , 2017, 1488, 47-73.	0.9	76

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19	The Non-Coding RNA Ontology (NCRO): a comprehensive resource for the unification of non-coding RNA biology. <i>Journal of Biomedical Semantics</i> , 2016, 7, 24.	1.6	10
20	The Cell Ontology 2016: enhanced content, modularization, and ontology interoperability. <i>Journal of Biomedical Semantics</i> , 2016, 7, 44.	1.6	201
21	The development of non-coding RNA ontology. <i>International Journal of Data Mining and Bioinformatics</i> , 2016, 15, 214.	0.1	9
22	Gene regulation knowledge commons: community action takes care of DNA binding transcription factors. <i>Database: the Journal of Biological Databases and Curation</i> , 2016, 2016, baw088.	3.0	12
23	Modeling biochemical pathways in the gene ontology. <i>Database: the Journal of Biological Databases and Curation</i> , 2016, 2016, baw126.	3.0	11
24	OmniSearch: a semantic search system based on the Ontology for MicroRNA Target (OMIT) for microRNA-target gene interaction data. <i>Journal of Biomedical Semantics</i> , 2016, 7, 25.	1.6	27
25	Mouse genome database 2016. <i>Nucleic Acids Research</i> , 2016, 44, D840-D847.	14.5	80
26	Mouse Genome Database: From sequence to phenotypes and disease models. <i>Genesis</i> , 2015, 53, 458-473.	1.6	13
27	The Mouse Genome Database (MGD): facilitating mouse as a model for human biology and disease. <i>Nucleic Acids Research</i> , 2015, 43, D726-D736.	14.5	335
28	A semantic approach for knowledge capture of MicroRNA-Target gene interactions. , 2015, , .		10
29	A domain ontology for the Non-Coding RNA field. , 2015, , .		0
30	Finding Our Way through Phenotypes. <i>PLoS Biology</i> , 2015, 13, e1002033.	5.6	178
31	Application of comparative biology in GO functional annotation: the mouse model. <i>Mammalian Genome</i> , 2015, 26, 574-583.	2.2	11
32	Orthology for comparative genomics in the mouse genome database. <i>Mammalian Genome</i> , 2015, 26, 305-313.	2.2	9
33	Mouse Genome Informatics (MGI): reflecting on 25 years. <i>Mammalian Genome</i> , 2015, 26, 272-284.	2.2	34
34	Representing Kidney Development Using the Gene Ontology. <i>PLoS ONE</i> , 2014, 9, e99864.	2.5	17
35	OMIT: Dynamic, Semi-Automated Ontology Development for the microRNA Domain. <i>PLoS ONE</i> , 2014, 9, e100855.	2.5	18
36	Methodology for the inference of gene function from phenotype data. <i>BMC Bioinformatics</i> , 2014, 15, 405.	2.6	5

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37	Protein Ontology: a controlled structured network of protein entities. Nucleic Acids Research, 2014, 42, D415-D421.	14.5	63
38	Standardized description of scientific evidence using the Evidence Ontology (ECO). Database: the Journal of Biological Databases and Curation, 2014, 2014, bau075-bau075.	3.0	95
39	A promoter-level mammalian expression atlas. Nature, 2014, 507, 462-470.	27.8	1,838
40	DFLAT: functional annotation for human development. BMC Bioinformatics, 2014, 15, 45.	2.6	27
41	A method for increasing expressivity of Gene Ontology annotations using a compositional approach. BMC Bioinformatics, 2014, 15, 155.	2.6	78
42	The Mouse Genome Database: integration of and access to knowledge about the laboratory mouse. Nucleic Acids Research, 2014, 42, D810-D817.	14.5	196
43	Ontology based molecular signatures for immune cell types via gene expression analysis. BMC Bioinformatics, 2013, 14, 263.	2.6	13
44	Gene Ontology annotation of sequence-specific DNA binding transcription factors: setting the stage for a large-scale curation effort. Database: the Journal of Biological Databases and Curation, 2013, 2013, bat062-bat062.	3.0	33
45	Ten Quick Tips for Using the Gene Ontology. PLoS Computational Biology, 2013, 9, e1003343.	3.2	45
46	The Mouse Genome Database: Genotypes, Phenotypes, and Models of Human Disease. Nucleic Acids Research, 2013, 41, D885-D891.	14.5	61
47	On the Use of Gene Ontology Annotations to Assess Functional Similarity among Orthologs and Paralogs: A Short Report. PLoS Computational Biology, 2012, 8, e1002386.	3.2	91
48	Manual Gene Ontology annotation workflow at the Mouse Genome Informatics Database. Database: the Journal of Biological Databases and Curation, 2012, 2012, bas045-bas045.	3.0	19
49	A Resource of Quantitative Functional Annotation for Homo sapiens Genes. G3: Genes, Genomes, Genetics, 2012, 2, 223-233.	1.8	6
50	An ontology-based MicroRNA knowledge sharing and acquisition framework. , 2012, , .		6
51	The Mouse Genome Database (MGD): comprehensive resource for genetics and genomics of the laboratory mouse. Nucleic Acids Research, 2012, 40, D881-D886.	14.5	233
52	Disease model curation improvements at Mouse Genome Informatics. Database: the Journal of Biological Databases and Curation, 2012, 2012, bar063-bar063.	3.0	10
53	Providing the Missing Link: the Exposure Science Ontology ExO. Environmental Science & Technology, 2012, 46, 3046-3053.	10.0	57
54	Concept annotation in the CRAFT corpus. BMC Bioinformatics, 2012, 13, 161.	2.6	188

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55	The Mouse Genome Database (MGD): premier model organism resource for mammalian genomics and genetics. <i>Nucleic Acids Research</i> , 2011, 39, D842-D848.	14.5	228
56	Hematopoietic cell types: Prototype for a revised cell ontology. <i>Journal of Biomedical Informatics</i> , 2011, 44, 75-79.	4.3	35
57	Autism candidate genes via mouse phenomics. <i>Journal of Biomedical Informatics</i> , 2011, 44, S5-S11.	4.3	16
58	The representation of protein complexes in the Protein Ontology (PRO). <i>BMC Bioinformatics</i> , 2011, 12, 371.	2.6	14
59	Logical Development of the Cell Ontology. <i>BMC Bioinformatics</i> , 2011, 12, 6.	2.6	117
60	The Protein Ontology: a structured representation of protein forms and complexes. <i>Nucleic Acids Research</i> , 2011, 39, D539-D545.	14.5	102
61	Ontology engineering. <i>Nature Biotechnology</i> , 2010, 28, 128-130.	17.5	113
62	The Mouse Genome Database: enhancements and updates. <i>Nucleic Acids Research</i> , 2010, 38, D586-D592.	14.5	78
63	The Mouse Genome Database genotypes::phenotypes. <i>Nucleic Acids Research</i> , 2009, 37, D712-D719.	14.5	101
64	The Gene Ontology's Reference Genome Project: A Unified Framework for Functional Annotation across Species. <i>PLoS Computational Biology</i> , 2009, 5, e1000431.	3.2	148
65	Using ontology visualization to facilitate access to knowledge about human disease genes. <i>Applied Ontology</i> , 2009, 4, 35-49.	2.0	3
66	TGF-beta signaling proteins and the Protein Ontology. <i>BMC Bioinformatics</i> , 2009, 10, S3.	2.6	9
67	Access to immunology through the Gene Ontology. <i>Immunology</i> , 2008, 125, 154-160.	4.4	33
68	Gene Ontology annotations: what they mean and where they come from. <i>BMC Bioinformatics</i> , 2008, 9, S2.	2.6	124
69	The Gene Ontology (GO) Project: Structured Vocabularies for Molecular Biology and Their Application to Genome and Expression Analysis. <i>Current Protocols in Bioinformatics</i> , 2008, 23, Unit 7.2.	25.8	94
70	A critical assessment of <i>Mus musculus</i> gene function prediction using integrated genomic evidence. <i>Genome Biology</i> , 2008, 9, S2.	9.6	214
71	An en masse phenotype and function prediction system for <i>Mus musculus</i> . <i>Genome Biology</i> , 2008, 9, S8.	9.6	20
72	Using bio-ontologies as data annotation, integration & analytical tools at the Mouse Genome Informatics resource. , 2008, , .		1

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73	Ontology development for biological systems: immunology. <i>Bioinformatics</i> , 2007, 23, 913-915.	4.1	49
74	The mouse genome database (MGD): new features facilitating a model system. <i>Nucleic Acids Research</i> , 2007, 35, D630-D637.	14.5	100
75	Mouse Genome Informatics (MGI) Resources for Pathology and Toxicology. <i>Toxicologic Pathology</i> , 2007, 35, 456-457.	1.8	27
76	The Mouse Genome Database (MGD): mouse biology and model systems. <i>Nucleic Acids Research</i> , 2007, 36, D724-D728.	14.5	365
77	Beyond the data deluge: Data integration and bio-ontologies. <i>Journal of Biomedical Informatics</i> , 2006, 39, 314-320.	4.3	128
78	The Mouse Genome Database (MGD): updates and enhancements. <i>Nucleic Acids Research</i> , 2006, 34, D562-D567.	14.5	72
79	A procedure for assessing GO annotation consistency. <i>Bioinformatics</i> , 2005, 21, i136-i143.	4.1	40
80	Ontological visualization of protein-protein interactions. <i>BMC Bioinformatics</i> , 2005, 6, 29.	2.6	11
81	The Mouse Genome Database (MGD): integrating biology with the genome. <i>Nucleic Acids Research</i> , 2004, 32, 476D-481.	14.5	66
82	Systems biology of the 2-cell mouse embryo. <i>Cytogenetic and Genome Research</i> , 2004, 105, 240-250.	1.1	128
83	The mouse Gene Expression Database (GXD): updates and enhancements. <i>Nucleic Acids Research</i> , 2004, 32, 568D-571.	14.5	61
84	Bio-ontologies—fast and furious. <i>Nature Biotechnology</i> , 2004, 22, 773-774.	17.5	67
85	A short study on the success of the Gene Ontology. <i>Web Semantics</i> , 2004, 1, 235-240.	2.9	61
86	The Mouse Genome Database (MGD): from genes to mice—a community resource for mouse biology. <i>Nucleic Acids Research</i> , 2004, 33, D471-D475.	14.5	217
87	MGD: the Mouse Genome Database. <i>Nucleic Acids Research</i> , 2003, 31, 193-195.	14.5	212
88	Human Disease Genes and Their Cloned Mouse Orthologs: Exploration of the FANTOM2 cDNA Sequence Data Set. <i>Genome Research</i> , 2003, 13, 1496-1500.	5.5	7
89	Connecting Sequence and Biology in the Laboratory Mouse. <i>Genome Research</i> , 2003, 13, 1505-1519.	5.5	18
90	The Gene Ontology (GO) Project: Structured Vocabularies for Molecular Biology and Their Application to Genome and Expression Analysis. <i>Current Protocols in Bioinformatics</i> , 2003, 00, Unit 7.2.	25.8	23

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91	The Mouse Genome Database (MGD): the model organism database for the laboratory mouse. <i>Nucleic Acids Research</i> , 2002, 30, 113-115.	14.5	135
92	Extension and Integration of the Gene Ontology (GO): Combining GO Vocabularies With External Vocabularies. <i>Genome Research</i> , 2002, 12, 1982-1991.	5.5	81
93	Rules and Guidelines for Mouse Gene, Allele, and Mutation Nomenclature: A Condensed Version. <i>Genomics</i> , 2002, 79, 471-474.	2.9	29
94	Corralling conditional mutations: A unified resource for mouse phenotypes. <i>Genesis</i> , 2002, 32, 63-65.	1.6	13
95	The Mouse Genome Database and The Gene Expression Database: Genotype to Phenotype. , 2002, , 119-128.		0
96	PROGRAM DESCRIPTION. <i>Genomics</i> , 2001, 74, 121-128.	2.9	47
97	Creating the Gene Ontology Resource: Design and Implementation. <i>Genome Research</i> , 2001, 11, 1425-1433.	5.5	881
98	Gene Ontology: tool for the unification of biology. <i>Nature Genetics</i> , 2000, 25, 25-29.	21.4	34,499
99	Informatics for Mouse Genetics and Genome Mapping. <i>Methods</i> , 1998, 14, 179-190.	3.8	16
100	Mouse and Rat Genome Informatics. , 0, , 119-142.		0
101	A Short Study on the Success of the Gene Ontology. <i>SSRN Electronic Journal</i> , 0, , .	0.4	3