

Gerald M Rubin

List of Publications by Year in descending order

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Version: 2024-02-01

234
papers

101,100
citations

620

123
h-index

1080

232
g-index

321
all docs

321
docs citations

321
times ranked

93724
citing authors

#	ARTICLE	IF	CITATIONS
1	Connectomic reconstruction predicts visual features used for navigation. <i>Nature</i> , 2024, 634, 181-190.	36.2	0
2	Toward a navigation framework for fetoscopy. <i>International Journal of Computer Assisted Radiology and Surgery</i> , 2023, 18, 2349-2356.	2.9	1
3	A functionally ordered visual feature map in the <i>Drosophila</i> brain. <i>Neuron</i> , 2022, 110, 1700-1711.e6.	8.0	56
4	Meta-analysis of changes in the levels of catecholamines and blood pressure with continuous positive airway pressure therapy in obstructive sleep apnea. <i>Journal of Clinical Hypertension</i> , 2021, 23, 12-20.	2.2	12
5	Performance and Stability of Pearl Millet Varieties for Grain Yield and Micronutrients in Arid and Semi-Arid Regions of India. <i>Frontiers in Plant Science</i> , 2021, 12, 670201.	3.8	23
6	Information flow, cell types and stereotypy in a full olfactory connectome. <i>ELife</i> , 2021, 10, .	5.9	112
7	A connectome of the <i>Drosophila</i> central complex reveals network motifs suitable for flexible navigation and context-dependent action selection. <i>ELife</i> , 2021, 10, .	5.9	200
8	Synaptic targets of photoreceptors specialized to detect color and skylight polarization in <i>Drosophila</i> . <i>ELife</i> , 2021, 10, .	5.9	41
9	Input Connectivity Reveals Additional Heterogeneity of Dopaminergic Reinforcement in <i>Drosophila</i> . <i>Current Biology</i> , 2020, 30, 3200-3211.e8.	4.0	66
10	Complete Connectomic Reconstruction of Olfactory Projection Neurons in the Fly Brain. <i>Current Biology</i> , 2020, 30, 3183-3199.e6.	4.0	146
11	The Neuroanatomical Ultrastructure and Function of a Biological Ring Attractor. <i>Neuron</i> , 2020, 108, 145-163.e10.	8.0	103
12	The Mind of a Mouse. <i>Cell</i> , 2020, 182, 1372-1376.	27.8	142
13	Toward nanoscale localization of memory engrams in <i>Drosophila</i> . <i>Journal of Neurogenetics</i> , 2020, 34, 151-155.	1.4	13
14	Inflammation, reward circuitry and symptoms of anhedonia and PTSD in trauma-exposed women. <i>Social Cognitive and Affective Neuroscience</i> , 2020, 15, 1046-1055.	3.3	45
15	A genetic, genomic, and computational resource for exploring neural circuit function. <i>ELife</i> , 2020, 9, .	5.9	190
16	A connectome and analysis of the adult <i>Drosophila</i> central brain. <i>ELife</i> , 2020, 9, .	5.9	712
17	Spatial readout of visual looming in the central brain of <i>Drosophila</i> . <i>ELife</i> , 2020, 9, .	5.9	43
18	Cell types and neuronal circuitry underlying female aggression in <i>Drosophila</i> . <i>ELife</i> , 2020, 9, .	5.9	74

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19	The connectome of the adult <i>Drosophila</i> mushroom body provides insights into function. <i>ELife</i> , 2020, 9, .	5.9	284
20	Cortical column and whole-brain imaging with molecular contrast and nanoscale resolution. <i>Science</i> , 2019, 363, .	20.9	300
21	Neurogenetic dissection of the <i>Drosophila</i> lateral horn reveals major outputs, diverse behavioural functions, and interactions with the mushroom body. <i>ELife</i> , 2019, 8, .	5.9	140
22	Looking back and looking forward at <i>Janelia</i> . <i>ELife</i> , 2019, 8, .	5.9	4
23	Nitric oxide acts as a cotransmitter in a subset of dopaminergic neurons to diversify memory dynamics. <i>ELife</i> , 2019, 8, .	5.9	108
24	Genetic Reagents for Making Split-GAL4 Lines in <i>Drosophila</i> . <i>Genetics</i> , 2018, 209, 31-35.	2.9	180
25	Neuroarchitecture of the <i>Drosophila</i> central complex: A catalog of nodulus and asymmetrical body neurons and a revision of the protocerebral bridge catalog. <i>Journal of Comparative Neurology</i> , 2018, 526, 2585-2611.	2.0	134
26	Communication from Learned to Innate Olfactory Processing Centers Is Required for Memory Retrieval in <i>Drosophila</i> . <i>Neuron</i> , 2018, 100, 651-668.e8.	8.0	89
27	The glia of the adult <i>Drosophila</i> nervous system. <i>Glia</i> , 2017, 65, 606-638.	5.3	240
28	Moonwalker Descending Neurons Mediate Visually Evoked Retreat in <i>Drosophila</i> . <i>Current Biology</i> , 2017, 27, 766-771.	4.0	68
29	Representations of Novelty and Familiarity in a Mushroom Body Compartment. <i>Cell</i> , 2017, 169, 956-969.e17.	27.8	122
30	The Emergence of Directional Selectivity in the Visual Motion Pathway of <i>Drosophila</i> . <i>Neuron</i> , 2017, 94, 168-182.e10.	8.0	153
31	A Circuit Node that Integrates Convergent Input from Neuromodulatory and Social Behavior-Promoting Neurons to Control Aggression in <i>Drosophila</i> . <i>Neuron</i> , 2017, 95, 1112-1128.e7.	8.0	82
32	Mapping the Neural Substrates of Behavior. <i>Cell</i> , 2017, 170, 393-406.e28.	27.8	203
33	Ultra-selective looming detection from radial motion opponency. <i>Nature</i> , 2017, 551, 237-241.	36.2	137
34	The comprehensive connectome of a neural substrate for α - <i>ON</i> TM motion detection in <i>Drosophila</i> . <i>ELife</i> , 2017, 6, .	5.9	173
35	TO007 COMPLEMENT MODULATION OF PERICYTE-TO-MYOFIBROBLAST TRANS-DIFFERENTIATION (PMT) AND MICROVASCULAR RAREFACTION IN RENAL ISCHEMIA/REPERFUSION (I/R). <i>Nephrology Dialysis Transplantation</i> , 2016, 31, i63-i63.	0.8	0
36	Direct neural pathways convey distinct visual information to <i>Drosophila</i> mushroom bodies. <i>ELife</i> , 2016, 5, .	5.9	134

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37	Dopaminergic neurons write and update memories with cell-type-specific rules. <i>ELife</i> , 2016, 5, .	5.9	262
38	FlyBook: A Preface. <i>Genetics</i> , 2015, 201, 343-343.	2.9	1
39	Control of Sleep by Dopaminergic Inputs to the <i>Drosophila</i> Mushroom Body. <i>Frontiers in Neural Circuits</i> , 2015, 9, 73.	3.0	82
40	P1 interneurons promote a persistent internal state that enhances inter-male aggression in <i>Drosophila</i> . <i>ELife</i> , 2015, 4, .	5.9	181
41	Distinct dopamine neurons mediate reward signals for short- and long-term memories. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 578-583.	7.6	227
42	A Dopamine-Modulated Neural Circuit Regulating Aversive Taste Memory in <i>Drosophila</i> . <i>Current Biology</i> , 2015, 25, 1535-1541.	4.0	89
43	Optimized tools for multicolor stochastic labeling reveal diverse stereotyped cell arrangements in the fly visual system. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E2967-76.	7.6	516
44	Heterosynaptic Plasticity Underlies Aversive Olfactory Learning in <i>Drosophila</i> . <i>Neuron</i> , 2015, 88, 985-998.	8.0	318
45	The Release 6 reference sequence of the <i>Drosophila melanogaster</i> genome. <i>Genome Research</i> , 2015, 25, 445-458.	5.6	403
46	Neural Circuit to Integrate Opposing Motions in the Visual Field. <i>Cell</i> , 2015, 162, 351-362.	27.8	118
47	High-performance probes for light and electron microscopy. <i>Nature Methods</i> , 2015, 12, 568-576.	19.6	244
48	Neuroarchitecture and neuroanatomy of the <i>Drosophila</i> central complex: A GAL4-based dissection of protocerebral bridge neurons and circuits. <i>Journal of Comparative Neurology</i> , 2015, 523, 997-1037.	2.0	282
49	Perspectives on numeracy: reflections from international assessments. <i>ZDM - International Journal on Mathematics Education</i> , 2015, 47, 691-706.	2.2	20
50	Plasticity-driven individualization of olfactory coding in mushroom body output neurons. <i>Nature</i> , 2015, 526, 258-262.	36.2	146
51	Propagation of Homeostatic Sleep Signals by Segregated Synaptic Microcircuits of the <i>Drosophila</i> Mushroom Body. <i>Current Biology</i> , 2015, 25, 2915-2927.	4.0	140
52	A Higher Brain Circuit for Immediate Integration of Conflicting Sensory Information in <i>Drosophila</i> . <i>Current Biology</i> , 2015, 25, 2203-2214.	4.0	146
53	Reward signal in a recurrent circuit drives appetitive long-term memory formation. <i>ELife</i> , 2015, 4, e10719.	5.9	144
54	Wide-Field Feedback Neurons Dynamically Tune Early Visual Processing. <i>Neuron</i> , 2014, 82, 887-895.	8.0	59

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55	Shared mushroom body circuits underlie visual and olfactory memories in <i>Drosophila</i> . <i>ELife</i> , 2014, 3, e02395.	5.9	173
56	A visual motion detection circuit suggested by <i>Drosophila</i> connectomics. <i>Nature</i> , 2013, 500, 175-181.	36.2	648
57	A directional tuning map of <i>Drosophila</i> elementary motion detectors. <i>Nature</i> , 2013, 500, 212-216.	36.2	336
58	Contributions of the 12 Neuron Classes in the Fly Lamina to Motion Vision. <i>Neuron</i> , 2013, 79, 128-140.	8.0	206
59	The effort to make mosaic analysis a household tool. <i>Development (Cambridge)</i> , 2012, 139, 4501-4503.	2.6	27
60	Using translational enhancers to increase transgene expression in <i>Drosophila</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 6626-6631.	7.6	382
61	A Resource for Manipulating Gene Expression and Analyzing cis-Regulatory Modules in the <i>Drosophila</i> CNS. <i>Cell Reports</i> , 2012, 2, 1002-1013.	6.3	117
62	A Survey of 6,300 Genomic Fragments for cis-Regulatory Activity in the Imaginal Discs of <i>Drosophila melanogaster</i> . <i>Cell Reports</i> , 2012, 2, 1014-1024.	6.3	119
63	A GAL4-Driver Line Resource for <i>Drosophila</i> Neurobiology. <i>Cell Reports</i> , 2012, 2, 991-1001.	6.3	1,361
64	A subset of dopamine neurons signals reward for odour memory in <i>Drosophila</i> . <i>Nature</i> , 2012, 488, 512-516.	36.2	564
65	Mushroom body efferent neurons responsible for aversive olfactory memory retrieval in <i>Drosophila</i> . <i>Nature Neuroscience</i> , 2011, 14, 903-910.	14.5	255
66	Multiple new site-specific recombinases for use in manipulating animal genomes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 14198-14203.	7.6	156
67	Refinement of Tools for Targeted Gene Expression in <i>Drosophila</i> . <i>Genetics</i> , 2010, 186, 735-755.	2.9	1,065
68	Quick Preparation of Genomic DNA from <i>Drosophila</i> . <i>Cold Spring Harbor Protocols</i> , 2009, 2009, pdb.prot5198.	0.3	34
69	Recovery of DNA Sequences Flanking P-Element Insertions in <i>Drosophila</i> : Inverse PCR and Plasmid Rescue. <i>Cold Spring Harbor Protocols</i> , 2009, 2009, pdb.prot5199.	0.3	28
70	Tools for neuroanatomy and neurogenetics in <i>Drosophila</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 9715-9720.	7.6	947
71	Biological Annotation of the <i>Drosophila</i> Genome Sequence. <i>Novartis Foundation Symposium</i> , 2008, , 79-83.	0.0	9
72	Global analysis of patterns of gene expression during <i>Drosophila</i> embryogenesis. <i>Genome Biology</i> , 2007, 8, R145.	7.3	398

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73	Global analyses of mRNA translational control during early <i>Drosophila</i> embryogenesis. <i>Genome Biology</i> , 2007, 8, R63.	7.3	74
74	Comparative Analysis of Spatial Patterns of Gene Expression in <i>Drosophila</i> Imaginal Discs. <i>Lecture Notes in Computer Science</i> , 2007, , 533-547.	1.0	9
75	Janelia Farm: An Experiment in Scientific Culture. <i>Cell</i> , 2006, 125, 209-212.	27.8	19
76	Large-Scale Trends in the Evolution of Gene Structures within 11 Animal Genomes. <i>PLoS Computational Biology</i> , 2006, 2, e15.	3.1	70
77	Pervasive regulation of <i>Drosophila</i> Notch target genes by GY-box-, Brd-box-, and K-box-class microRNAs. <i>Genes and Development</i> , 2005, 19, 1067-1080.	5.9	261
78	The ubiquitin ligase <i>Drosophila</i> Mind bomb promotes Notch signaling by regulating the localization and activity of Serrate and Delta. <i>Development (Cambridge)</i> , 2005, 132, 2319-2332.	2.6	145
79	Identification of putative noncoding polyadenylated transcripts in <i>Drosophila melanogaster</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 5495-5500.	7.6	112
80	A computational and experimental approach to validating annotations and gene predictions in the <i>Drosophila melanogaster</i> genome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 1566-1571.	7.6	32
81	Cathepsin D-deficient <i>Drosophila</i> recapitulate the key features of neuronal ceroid lipofuscinoses. <i>Neurobiology of Disease</i> , 2005, 19, 194-199.	4.5	68
82	<i>Drosophila</i> microRNAs exhibit diverse spatial expression patterns during embryonic development. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 18017-18022.	7.6	256
83	Complementary miRNA pairs suggest a regulatory role for miRNA:miRNA duplexes. <i>Rna</i> , 2004, 10, 171-175.	3.6	83
84	<i>Drosophila melanogaster</i> MNK/Chk2 and p53 Regulate Multiple DNA Repair and Apoptotic Pathways following DNA Damage. <i>Molecular and Cellular Biology</i> , 2004, 24, 1219-1231.	2.5	288
85	Nurturing interdisciplinary research. <i>Nature Structural and Molecular Biology</i> , 2004, 11, 1166-1169.	8.1	45
86	The BDGP Gene Disruption Project. <i>Genetics</i> , 2004, 167, 761-781.	2.9	782
87	Computational identification of developmental enhancers: conservation and function of transcription factor binding-site clusters in <i>Drosophila melanogaster</i> and <i>Drosophila pseudoobscura</i> . <i>Genome Biology</i> , 2004, 5, R61.	7.3	184
88	THE <i>DROSOPHILA MELANOGASTER</i> GENOME. <i>Annual Review of Genomics and Human Genetics</i> , 2003, 4, 89-117.	6.3	117
89	Y chromosome and other heterochromatic sequences of the <i>Drosophila melanogaster</i> genome: how far can we go?. <i>Genetica</i> , 2003, 117, 227-237.	1.2	44
90	Quantitative Analysis of Bristle Number in <i>Drosophila</i> Mutants Identifies Genes Involved in Neural Development. <i>Current Biology</i> , 2003, 13, 1388-1396.	4.0	115

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91	Drosophila Matrix Metalloproteinases Are Required for Tissue Remodeling, but Not Embryonic Development. <i>Developmental Cell</i> , 2003, 4, 95-106.	7.0	229
92	Computational identification of Drosophila microRNA genes. <i>Genome Biology</i> , 2003, 4, R42.	7.3	629
93	The Drosophila synaptotagmin-like protein bitesize is required for growth and has mRNA localization sequences within its open reading frame. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 13368-13373.	7.6	42
94	ARGONAUTE1 is required for efficient RNA interference in Drosophila embryos. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 6889-6894.	7.6	166
95	Exploiting transcription factor binding site clustering to identify cis-regulatory modules involved in pattern formation in the Drosophila genome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 757-762.	7.6	545
96	The Drosophila Gene Collection: Identification of Putative Full-Length cDNAs for 70% of D. melanogaster Genes. <i>Genome Research</i> , 2002, 12, 1294-1300.	5.6	181
97	Biological and computational annotation of the Drosophila Genome Sequence. , 2002, , .		0
98	Targeted mutagenesis by homologous recombination in D. melanogaster. <i>Genes and Development</i> , 2002, 16, 1568-1581.	5.9	300
99	An expectation maximization algorithm for training hidden substitution models 1 Edited by F. Cohen. <i>Journal of Molecular Biology</i> , 2002, 317, 753-764.	4.3	69
100	The transposable elements of the Drosophila melanogaster euchromatin: a genomics perspective. <i>Genome Biology</i> , 2002, 3, RESEARCH0084.	9.2	483
101	Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 16899-16903.	7.6	1,628
102	Finishing a whole-genome shotgun: release 3 of the Drosophila melanogaster euchromatic genome sequence. <i>Genome Biology</i> , 2002, 3, research0079.1.	7.3	321
103	Annotation of the Drosophila melanogaster euchromatic genome: a systematic review. <i>Genome Biology</i> , 2002, 3, research0083.1.	7.3	310
104	Heterochromatic sequences in a Drosophila whole-genome shotgun assembly. <i>Genome Biology</i> , 2002, 3, research0085.1.	7.3	239
105	Computational analysis of core promoters in the Drosophila genome. <i>Genome Biology</i> , 2002, 3, research0087.1.	7.3	382
106	A Drosophila full-length cDNA resource. <i>Genome Biology</i> , 2002, 3, research0080.1.	7.3	168
107	Assessing the impact of comparative genomic sequence data on the functional annotation of the Drosophila genome. <i>Genome Biology</i> , 2002, 3, research0086.1.	7.3	122
108	Systematic determination of patterns of gene expression during Drosophila embryogenesis. <i>Genome Biology</i> , 2002, 3, research0088.1.	7.3	619

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109	Evidence for large domains of similarly expressed genes in the <i>Drosophila</i> genome. <i>Journal of Biology</i> , 2002, 1, 5.	2.5	427
110	The Toll and Imd pathways are the major regulators of the immune response in <i>Drosophila</i> . <i>EMBO Journal</i> , 2002, 21, 2568-2579.	8.2	774
111	The Ca ²⁺ -Calmodulin-Activated Protein Phosphatase Calcineurin Negatively Regulates Egf Receptor Signaling in <i>Drosophila</i> Development. <i>Genetics</i> , 2002, 161, 183-193.	2.9	31
112	neuralized Functions Cell-Autonomously to Regulate a Subset of Notch-Dependent Processes during Adult <i>Drosophila</i> Development. <i>Developmental Biology</i> , 2001, 231, 217-233.	2.1	85
113	<i>Drosophila</i> Neuralized Is a Ubiquitin Ligase that Promotes the Internalization and Degradation of Delta. <i>Developmental Cell</i> , 2001, 1, 783-794.	7.0	307
114	<i>Drosophila</i> Fragile X-Related Gene Regulates the MAP1B Homolog Futsch to Control Synaptic Structure and Function. <i>Cell</i> , 2001, 107, 591-603.	27.8	611
115	Comparing species. <i>Nature</i> , 2001, 409, 820-821.	36.2	79
116	Genome-wide analysis of the <i>Drosophila</i> immune response by using oligonucleotide microarrays. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 12590-12595.	7.6	662
117	Gene Ontology: tool for the unification of biology. <i>Nature Genetics</i> , 2000, 25, 25-29.	20.4	35,841
118	The Genome Sequence of <i>Drosophila melanogaster</i> . <i>Science</i> , 2000, 287, 2185-2195.	20.9	5,617
119	Comparative Genomics of the Eukaryotes. <i>Science</i> , 2000, 287, 2204-2215.	20.9	1,583
120	<i>Drosophila</i> p53 Binds a Damage Response Element at the reaper Locus. <i>Cell</i> , 2000, 101, 103-113.	27.8	435
121	A Whole-Genome Assembly of <i>Drosophila</i> . <i>Science</i> , 2000, 287, 2196-2204.	20.9	1,464
122	A Brief History of <i>Drosophila</i> 's Contributions to Genome Research. <i>Science</i> , 2000, 287, 2216-2218.	20.9	220
123	A <i>Drosophila</i> Complementary DNA Resource. <i>Science</i> , 2000, 287, 2222-2224.	20.9	337
124	Insertion site preferences of the P transposable element in <i>Drosophila melanogaster</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 3347-3351.	7.6	137
125	A Genetic Screen for Novel Components of the Ras/Mitogen-Activated Protein Kinase Signaling Pathway That Interact With the yan Gene of <i>Drosophila</i> Identifies split ends, a New RNA Recognition Motif-Containing Protein. <i>Genetics</i> , 2000, 154, 695-712.	2.9	136
126	A Misexpression Screen Identifies Genes That Can Modulate RAS1 Pathway Signaling in <i>Drosophila melanogaster</i> . <i>Genetics</i> , 2000, 156, 1219-1230.	2.9	102

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127	A Genetic Screen for Modifiers of a Kinase Suppressor of Ras-Dependent Rough Eye Phenotype in <i>Drosophila</i> . <i>Genetics</i> , 2000, 156, 1231-1242.	2.9	82
128	<i>mus304</i> encodes a novel DNA damage checkpoint protein required during <i>Drosophila</i> development. <i>Genes and Development</i> , 2000, 14, 666-678.	5.9	111
129	<i>Drosophila</i> and human RecQ5 exist in different isoforms generated by alternative splicing. <i>Nucleic Acids Research</i> , 1999, 27, 3762-3769.	14.0	61
130	Synaptic function modulated by changes in the ratio of synaptotagmin I and IV. <i>Nature</i> , 1999, 400, 757-760.	36.2	150
131	PTP-ER, a Novel Tyrosine Phosphatase, Functions Downstream of Ras1 to Downregulate MAP Kinase during <i>Drosophila</i> Eye Development. <i>Molecular Cell</i> , 1999, 3, 741-750.	9.6	71
132	<i>gigas</i> , a <i>Drosophila</i> Homolog of Tuberous Sclerosis Gene Product-2, Regulates the Cell Cycle. <i>Cell</i> , 1999, 96, 529-539.	27.8	254
133	Identification of Constitutive and Ras-Inducible Phosphorylation Sites of KSR: Implications for 14-3-3 Binding, Mitogen-Activated Protein Kinase Binding, and KSR Overexpression. <i>Molecular and Cellular Biology</i> , 1999, 19, 229-240.	2.5	195
134	The Berkeley <i>Drosophila</i> Genome Project Gene Disruption Project: Single P-Element Insertions Mutating 25% of Vital <i>Drosophila</i> Genes. <i>Genetics</i> , 1999, 153, 135-177.	2.9	734
135	The <i>Drosophila</i> genome project: a progress report. <i>Trends in Genetics</i> , 1998, 14, 340-343.	6.9	26
136	CNK, a RAF-Binding Multidomain Protein Required for RAS Signaling. <i>Cell</i> , 1998, 95, 343-353.	27.8	166
137	BioViews: Java-Based Tools for Genomic Data Visualization. <i>Genome Research</i> , 1998, 8, 291-305.	5.6	28
138	A high throughput screen to identify secreted and transmembrane proteins involved in <i>Drosophila</i> embryogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1998, 95, 9973-9978.	7.6	108
139	A Computer Program for Aligning a cDNA Sequence with a Genomic DNA Sequence. <i>Genome Research</i> , 1998, 8, 967-974.	5.6	683
140	The development of the <i>Drosophila</i> visual system. , 1998, , 474-508.		9
141	A Genetic Screen to Identify Components of the sina Signaling Pathway in <i>Drosophila</i> Eye Development. <i>Genetics</i> , 1998, 148, 277-286.	2.9	90
142	A Genetic Screen to Identify Components of the sina Signaling Pathway in <i>Drosophila</i> Eye Development. <i>Genetics</i> , 1998, 148, 277-286.	2.9	74
143	P element insertion-dependent gene activation in the <i>Drosophila</i> eye. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1997, 94, 5195-5200.	7.6	101
144	Kuzbanian Controls Proteolytic Processing of Notch and Mediates Lateral Inhibition during <i>Drosophila</i> and Vertebrate Neurogenesis. <i>Cell</i> , 1997, 90, 271-280.	27.8	494

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145	PHYL Acts to Down-Regulate TTK88, a Transcriptional Repressor of Neuronal Cell Fates, by a SINA-Dependent Mechanism. <i>Cell</i> , 1997, 90, 459-467.	27.8	223
146	misshapen encodes a protein kinase involved in cell shape control in <i>Drosophila</i> . <i>Gene</i> , 1997, 186, 119-125.	2.3	49
147	KSR stimulates Raf-1 activity in a kinase-independent manner. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1997, 94, 12792-12796.	7.6	165
148	Targets of glass regulation in the <i>Drosophila</i> eye disc. <i>Mechanisms of Development</i> , 1996, 56, 17-24.	1.7	20
149	Pk92b: a <i>drosophila melanogaster</i> protein kinase that belongs to the mekk family. <i>Gene</i> , 1996, 169, 283-284.	2.3	10
150	The Role of the Genome Project in Determining Gene Function: Insights from Model Organisms. <i>Cell</i> , 1996, 86, 521-529.	27.8	453
151	A <i>Drosophila</i> gene regulated by rough and glass shows similarity to ena and VASP. <i>Gene</i> , 1996, 183, 103-108.	2.3	17
152	A Screen for Genes That Function Downstream of Ras1 During <i>Drosophila</i> Eye Development. <i>Genetics</i> , 1996, 143, 315-329.	2.9	254
153	Yan functions as a general inhibitor of differentiation and is negatively regulated by activation of the Ras1/MAPK pathway. <i>Cell</i> , 1995, 81, 857-866.	27.8	333
154	<i>Drosophila</i> homologs of baculovirus inhibitor of apoptosis proteins function to block cell death. <i>Cell</i> , 1995, 83, 1253-1262.	27.8	736
155	KSR, a novel protein kinase required for RAS signal transduction. <i>Cell</i> , 1995, 83, 879-888.	27.8	383
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