David M Kingsley

List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/5459438/david-m-kingsley-publications-by-year.pdf

Version: 2024-04-25

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

59	9,420	38	66
papers	citations	h-index	g-index
66	10,988 ext. citations	16.8	5.63
ext. papers		avg, IF	L-index

#	Paper	IF	Citations
59	Genetic studies of human-chimpanzee divergence using stem cell fusions <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	3
58	Characterization of mouse Bmp5 regulatory injury element in zebrafish wound models. <i>Bone</i> , 2021 , 155, 116263	4.7	0
57	Predicting future from past: The genomic basis of recurrent and rapid stickleback evolution. <i>Science Advances</i> , 2021 , 7,	14.3	14
56	Fitness maps to a large-effect locus in introduced stickleback populations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	11
55	Assembly of the threespine stickleback Y chromosome reveals convergent signatures of sex chromosome evolution. <i>Genome Biology</i> , 2020 , 21, 177	18.3	30
54	DNA fragility in the parallel evolution of pelvic reduction in stickleback fish. <i>Science</i> , 2019 , 363, 81-84	33.3	90
53	Efficient CRISPR-Cas9 editing of major evolutionary loci in sticklebacks <i>Evolutionary Ecology Research</i> , 2019 , 20, 107-132		1
52	Characterization of a Human-Specific Tandem Repeat Associated with Bipolar Disorder and Schizophrenia. <i>American Journal of Human Genetics</i> , 2018 , 103, 421-430	11	48
51	A novel enhancer near the gene influences development and evolution of pelvic appendages in vertebrates. <i>ELife</i> , 2018 , 7,	8.9	18
50	Detecting differential copy number variation between groups of samples. <i>Genome Research</i> , 2018 , 28, 256-265	9.7	4
49	Experimental evidence for rapid genomic adaptation to a new niche in an adaptive radiation. <i>Nature Ecology and Evolution</i> , 2018 , 2, 1128-1138	12.3	43
48	Genetic Coupling of Female Mate Choice with Polygenic Ecological Divergence Facilitates Stickleback Speciation. <i>Current Biology</i> , 2017 , 27, 3344-3349.e4	6.3	28
47	Convergent evolution of SWS2 opsin facilitates adaptive radiation of threespine stickleback into different light environments. <i>PLoS Biology</i> , 2017 , 15, e2001627	9.7	38
46	Dorsal spine evolution in threespine sticklebacks via a splicing change in MSX2A. <i>BMC Biology</i> , 2017 , 15, 115	7.3	13
45	An Unexpectedly Complex Architecture for Skin Pigmentation in Africans. <i>Cell</i> , 2017 , 171, 1340-1353.e1	1 4 56.2	85
44	Ancient selection for derived alleles at a GDF5 enhancer influencing human growth and osteoarthritis risk. <i>Nature Genetics</i> , 2017 , 49, 1202-1210	36.3	53
43	Genomic dissection of conserved transcriptional regulation in intestinal epithelial cells. <i>PLoS Biology</i> , 2017 , 15, e2002054	9.7	47

(2012-2017)

42	Beautiful Piles of Bones: An Interview with 2017 Genetics Society of America Medal Recipient David M. Kingsley. <i>Genetics</i> , 2017 , 207, 1221-1222	4	
41	Evolving New Skeletal Traits by cis-Regulatory Changes in Bone Morphogenetic Proteins. <i>Cell</i> , 2016 , 164, 45-56	56.2	97
40	Heads, Shoulders, Elbows, Knees, and Toes: Modular Gdf5 Enhancers Control Different Joints in the Vertebrate Skeleton. <i>PLoS Genetics</i> , 2016 , 12, e1006454	6	39
39	A distinct regulatory region of the Bmp5 locus activates gene expression following adult bone fracture or soft tissue injury. <i>Bone</i> , 2015 , 77, 31-41	4.7	20
38	Extent of QTL Reuse During Repeated Phenotypic Divergence of Sympatric Threespine Stickleback. <i>Genetics</i> , 2015 , 201, 1189-200	4	40
37	A recurrent regulatory change underlying altered expression and Wnt response of the stickleback armor plates gene EDA. <i>ELife</i> , 2015 , 4, e05290	8.9	71
36	Mesenchymal cells. Defining a mesenchymal progenitor niche at single-cell resolution. <i>Science</i> , 2014 , 346, 1258810	33.3	99
35	A molecular basis for classic blond hair color in Europeans. <i>Nature Genetics</i> , 2014 , 46, 748-52	36.3	122
34	Genetics of ecological divergence during speciation. <i>Nature</i> , 2014 , 511, 307-11	50.4	192
33	The phosphate exporter xpr1b is required for differentiation of tissue-resident macrophages. <i>Cell Reports</i> , 2014 , 8, 1659-1667	10.6	36
32	Evolved tooth gain in sticklebacks is associated with a cis-regulatory allele of Bmp6. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 13912-7	11.5	56
31	Modular skeletal evolution in sticklebacks is controlled by additive and clustered quantitative trait Loci. <i>Genetics</i> , 2014 , 197, 405-20	4	86
30	Phylogeography and adaptation genetics of stickleback from the Haida Gwaii archipelago revealed using genome-wide single nucleotide polymorphism genotyping. <i>Molecular Ecology</i> , 2013 , 22, 1917-32	5.7	44
29	A penile spine/vibrissa enhancer sequence is missing in modern and extinct humans but is retained in multiple primates with penile spines and sensory vibrissae. <i>PLoS ONE</i> , 2013 , 8, e84258	3.7	15
28	Genetic signature of adaptive peak shift in threespine stickleback. <i>Evolution; International Journal of Organic Evolution</i> , 2012 , 66, 2439-50	3.8	59
27	The genomic basis of adaptive evolution in threespine sticklebacks. <i>Nature</i> , 2012 , 484, 55-61	50.4	1187
26	Population genomics of parallel phenotypic evolution in stickleback across stream-lake ecological transitions. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2012 , 279, 1277-86	4.4	101
25	A genome-wide SNP genotyping array reveals patterns of global and repeated species-pair divergence in sticklebacks. <i>Current Biology</i> , 2012 , 22, 83-90	6.3	172

24	When evolution hurts: height, arthritis risk, and the regulatory architecture of GDF5 function. <i>FASEB Journal</i> , 2012 , 26, 457.1	0.9	
23	Three periods of regulatory innovation during vertebrate evolution. <i>Science</i> , 2011 , 333, 1019-24	33.3	92
22	Adaptive evolution of pelvic reduction in sticklebacks by recurrent deletion of a Pitx1 enhancer. <i>Science</i> , 2010 , 327, 302-5	33.3	709
21	From atoms to traits. <i>Scientific American</i> , 2009 , 300, 52-9	0.5	8
20	Shaping skeletal growth by modular regulatory elements in the Bmp5 gene. <i>PLoS Genetics</i> , 2008 , 4, e10	006308	45
19	Dual hindlimb control elements in the Tbx4 gene and region-specific control of bone size in vertebrate limbs. <i>Development (Cambridge)</i> , 2008 , 135, 2543-53	6.6	71
18	The genetics of adaptive shape shift in stickleback: pleiotropy and effect size. <i>Evolution; International Journal of Organic Evolution</i> , 2008 , 62, 76-85	3.8	203
17	Parallel genetic origins of pelvic reduction in vertebrates. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 13753-8	11.5	170
16	Widespread parallel evolution in sticklebacks by repeated fixation of Ectodysplasin alleles. <i>Science</i> , 2005 , 307, 1928-33	33.3	1083
15	A Simple and Efficient Microinjection Protocol for Making Transgenic Sticklebacks. <i>Behaviour</i> , 2004 , 141, 1345-1355	1.4	20
14	Genetic and developmental basis of evolutionary pelvic reduction in threespine sticklebacks. <i>Nature</i> , 2004 , 428, 717-23	50.4	645
13	The genetic architecture of parallel armor plate reduction in threespine sticklebacks. <i>PLoS Biology</i> , 2004 , 2, E109	9.7	281
12	A general approach for identifying distant regulatory elements applied to the Gdf6 gene. <i>Genome Research</i> , 2003 , 13, 2069-81	9.7	73
11	Genetic control of bone and joint formation. <i>Novartis Foundation Symposium</i> , 2001 , 232, 213-22; discussion 222-34, 272-82		19
10	The genetic architecture of divergence between threespine stickleback species. <i>Nature</i> , 2001 , 414, 901	-5 0.4	413
9	Reciprocal mouse and human limb phenotypes caused by gain- and loss-of-function mutations affecting Lmbr1. <i>Genetics</i> , 2001 , 159, 715-26	4	27
8	Role of the mouse ank gene in control of tissue calcification and arthritis. <i>Science</i> , 2000 , 289, 265-70	33.3	565
7	An extensive 3th egulatory region controls expression of Bmp5 in specific anatomical structures of the mouse embryo. <i>Genetics</i> , 1998 , 148, 401-8	4	70

LIST OF PUBLICATIONS

6	Spectrum of Bmp5 mutations from germline mutagenesis experiments in mice. <i>Genetics</i> , 1997 , 145, 43	5- <u>4</u> 3	47
5	The mouse Snell's waltzer deafness gene encodes an unconventional myosin required for structural integrity of inner ear hair cells. <i>Nature Genetics</i> , 1995 , 11, 369-75	36.3	423
4	What do BMPs do in mammals? Clues from the mouse short-ear mutation. <i>Trends in Genetics</i> , 1994 , 10, 16-21	8.5	280
3	Limb alterations in brachypodism mice due to mutations in a new member of the TGF beta-superfamily. <i>Nature</i> , 1994 , 368, 639-43	50.4	773
2	The mouse short ear skeletal morphogenesis locus is associated with defects in a bone morphogenetic member of the TGF beta superfamily. <i>Cell</i> , 1992 , 71, 399-410	56.2	432
1	Mouse chromosome 9. <i>Mammalian Genome</i> , 1992 , 3 Spec No, S136-52	3.2	6