

Andy Greenfield

List of Publications by Year in descending order

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72
papers

4,596
citations

109137

35
h-index

102304

66
g-index

79
all docs

79
docs citations

79
times ranked

6040
citing authors

#	ARTICLE	IF	CITATIONS
1	The Sry-related gene Sox9 is expressed during chondrogenesis in mouse embryos. <i>Nature Genetics</i> , 1995, 9, 15-20.	9.4	627
2	The UTX gene escapes X inactivation in mice and humans. <i>Human Molecular Genetics</i> , 1998, 7, 737-742.	1.4	218
3	An H α -YDb epitope is encoded by a novel mouse Y chromosome gene. <i>Nature Genetics</i> , 1996, 14, 474-478.	9.4	176
4	Expression of a linear Sry transcript in the mouse genital ridge. <i>Nature Genetics</i> , 1995, 10, 480-482.	9.4	165
5	Pkd11l1 establishes left-right asymmetry and physically interacts with Pkd2. <i>Development (Cambridge)</i> , 2011, 138, 1131-1142.	1.2	156
6	Mutations in MAP3K1 Cause 46,XY Disorders of Sex Development and Implicate a Common Signal Transduction Pathway in Human Testis Determination. <i>American Journal of Human Genetics</i> , 2010, 87, 898-904.	2.6	155
7	SCRIB expression is deregulated in human prostate cancer, and its deficiency in mice promotes prostate neoplasia. <i>Journal of Clinical Investigation</i> , 2011, 121, 4257-4267.	3.9	153
8	A gene-driven ENU-based approach to generating an allelic series in any gene. <i>Mammalian Genome</i> , 2004, 15, 585-591.	1.0	148
9	The PCP genes Celsr1 and Vangl2 are required for normal lung branching morphogenesis. <i>Human Molecular Genetics</i> , 2010, 19, 2251-2267.	1.4	146
10	Secreted frizzled-related protein 5 suppresses adipocyte mitochondrial metabolism through WNT inhibition. <i>Journal of Clinical Investigation</i> , 2012, 122, 2405-2416.	3.9	141
11	Dissecting Cell Lineage Specification and Sex Fate Determination in Gonadal Somatic Cells Using Single-Cell Transcriptomics. <i>Cell Reports</i> , 2019, 26, 3272-3283.e3.	2.9	137
12	ISSCR Guidelines for Stem Cell Research and Clinical Translation: The 2021 update. <i>Stem Cell Reports</i> , 2021, 16, 1398-1408.	2.3	134
13	Loss of Mitogen-Activated Protein Kinase Kinase Kinase 4 (MAP3K4) Reveals a Requirement for MAPK Signalling in Mouse Sex Determination. <i>PLoS Biology</i> , 2009, 7, e1000196.	2.6	130
14	Regulation of hepatic metabolic pathways by the orphan nuclear receptor SHP. <i>EMBO Journal</i> , 2005, 24, 2624-2633.	3.5	129
15	Gadd45 ³ and Map3k4 Interactions Regulate Mouse Testis Determination via p38 MAPK-Mediated Control of Sry Expression. <i>Developmental Cell</i> , 2012, 23, 1020-1031.	3.1	122
16	Zic2 is required for neural crest formation and hindbrain patterning during mouse development. <i>Developmental Biology</i> , 2003, 264, 391-406.	0.9	107
17	The Mouse Y Chromosome Interval Necessary for Spermatogonial Proliferation is Gene Dense with Syntenic Homology to the Human AZFa Region. <i>Human Molecular Genetics</i> , 1998, 7, 1713-1724.	1.4	96
18	Mouse hitchhiker mutants have spina bifida, dorso-ventral patterning defects and polydactyly: identification of Tulp3 as a novel negative regulator of the Sonic hedgehog pathway. <i>Human Molecular Genetics</i> , 2009, 18, 1719-1739.	1.4	88

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19	SW-ARRAY: a dynamic programming solution for the identification of copy-number changes in genomic DNA using array comparative genome hybridization data. <i>Nucleic Acids Research</i> , 2005, 33, 3455-3464.	6.5	87
20	Assisted reproductive technologies to prevent human mitochondrial disease transmission. <i>Nature Biotechnology</i> , 2017, 35, 1059-1068.	9.4	87
21	Sfrp1 and Sfrp2 are required for normal male sexual development in mice. <i>Developmental Biology</i> , 2009, 326, 273-284.	0.9	84
22	Genetic analyses reveal a requirement for Dicer1 in the mouse urogenital tract. <i>Mammalian Genome</i> , 2009, 20, 140-151.	1.0	82
23	Cloning, Mapping, and Expression Analysis of a Gene Encoding a Novel Mammalian EGF-Related Protein (SCUBE1). <i>Genomics</i> , 2000, 70, 74-81.	1.3	72
24	Scribble is required for normal epithelial cell-cell contacts and lumen morphogenesis in the mammalian lung. <i>Developmental Biology</i> , 2013, 373, 267-280.	0.9	71
25	Haploinsufficiency of the murine Col3a1 locus causes aortic dissection: a novel model of the vascular type of Ehlers-Danlos syndrome. <i>Cardiovascular Research</i> , 2011, 90, 182-190.	1.8	68
26	Characterizing the bipotential mammalian gonad. <i>Current Topics in Developmental Biology</i> , 2019, 134, 167-194.	1.0	63
27	ZNRF3 functions in mammalian sex determination by inhibiting canonical WNT signaling. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 5474-5479.	3.3	62
28	Single primer amplification (SPA) of cDNA for microarray expression analysis. <i>Nucleic Acids Research</i> , 2003, 31, 9e-9.	6.5	55
29	Minor Abnormalities of Testis Development in Mice Lacking the Gene Encoding the MAPK Signalling Component, MAP3K1. <i>PLoS ONE</i> , 2011, 6, e19572.	1.1	55
30	Groucho/transducin-like Enhancer of split (TLE) family members interact with the yeast transcriptional co-repressor SSN6 and mammalian SSN6-related proteins: implications for evolutionary conservation of transcription repression mechanisms. <i>Biochemical Journal</i> , 1999, 337, 13-17.	1.7	51
31	The molecular and cellular basis of gonadal sex reversal in mice and humans. <i>Wiley Interdisciplinary Reviews: Developmental Biology</i> , 2012, 1, 559-577.	5.9	51
32	DNA microarrays and development. <i>Human Molecular Genetics</i> , 2003, 12, 1R-8.	1.4	45
33	Expression of a novel mammalian epidermal growth factor-related gene during mouse neural development. <i>Mechanisms of Development</i> , 2001, 102, 209-211.	1.7	44
34	Sexually dimorphic expression of Gata-2 during mouse gonad development. <i>Mechanisms of Development</i> , 2002, 111, 159-162.	1.7	42
35	Loss of p300 and CBP disrupts histone acetylation at the mouse Sry promoter and causes XY gonadal sex reversal. <i>Human Molecular Genetics</i> , 2018, 27, 190-198.	1.4	39
36	Novel gene expression patterns along the proximo-distal axis of the mouse embryo before gastrulation. <i>BMC Developmental Biology</i> , 2007, 7, 8.	2.1	34

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37	Pathogenic variants in the DEAH-box RNA helicase DHX37 are a frequent cause of 46,XY gonadal dysgenesis and 46,XY testicular regression syndrome. <i>Genetics in Medicine</i> , 2020, 22, 150-159.	1.1	34
38	Origin, specification and differentiation of a rare supporting-like lineage in the developing mouse gonad. <i>Science Advances</i> , 2022, 8, .	4.7	32
39	Addressing gaps in care of people with conditions affecting sex development and maturation. <i>Nature Reviews Endocrinology</i> , 2019, 15, 615-622.	4.3	30
40	Dissecting the genetic complexity of human 6p deletion syndromes by using a region-specific, phenotype-driven mouse screen. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 12477-12482.	3.3	28
41	A Novel Mouse <i>Fgfr2</i> Mutant, Hobbyhorse (<i>hob</i>), Exhibits Complete XY Gonadal Sex Reversal. <i>PLoS ONE</i> , 2014, 9, e100447.	1.1	26
42	Candidate testis-determining gene, Maestro (<i>Mro</i>), encodes a novel HEAT repeat protein. <i>Developmental Dynamics</i> , 2003, 227, 600-607.	0.8	25
43	Transgenic expression of <i>Map3k4</i> rescues T-associated sex reversal (<i>Tas</i>) in mice. <i>Human Molecular Genetics</i> , 2014, 23, 3035-3044.	1.4	24
44	Genetic Disruption of 21-Hydroxylase in Zebrafish Causes Interrenal Hyperplasia. <i>Endocrinology</i> , 2017, 158, 4165-4173.	1.4	24
45	Characterising Novel Pathways in Testis Determination Using Mouse Genetics. <i>Sexual Development</i> , 2014, 8, 199-207.	1.1	23
46	Sexually dimorphic expression of secreted frizzled-related (<i>SFRP</i>) genes in the developing mouse Müllerian duct. <i>Molecular Reproduction and Development</i> , 2006, 73, 1008-1016.	1.0	21
47	Groucho/transducin-like Enhancer of split (<i>TLE</i>) family members interact with the yeast transcriptional co-repressor <i>SSN6</i> and mammalian <i>SSN6</i> -related proteins: implications for evolutionary conservation of transcription repression mechanisms. <i>Biochemical Journal</i> , 1999, 337, 13.	1.7	20
48	Genetic Analyses Reveal Functions for <i>MAP2K3</i> and <i>MAP2K6</i> in Mouse Testis Determination1. <i>Biology of Reproduction</i> , 2016, 94, 103.	1.2	18
49	1 Sry and Mammalian Sex Determination. <i>Current Topics in Developmental Biology</i> , 1996, 34, 1-23.	1.0	17
50	Arrest of WNT/ β -catenin signaling enables the transition from pluripotent to differentiated germ cells in mouse ovaries. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	17
51	Understanding sex determination in the mouse: genetics, epigenetics and the story of mutual antagonisms. <i>Journal of Genetics</i> , 2015, 94, 585-590.	0.4	16
52	Applications of DNA microarrays to the transcriptional analysis of mammalian genomes. <i>Mammalian Genome</i> , 2000, 11, 609-613.	1.0	14
53	The Gonadal Supporting Cell Lineage and Mammalian Sex Determination: The Differentiation of Sertoli and Granulosa Cells. <i>Results and Problems in Cell Differentiation</i> , 2016, 58, 47-66.	0.2	14
54	ENU mutagenesis as a tool for understanding lung development and disease. <i>Biochemical Society Transactions</i> , 2009, 37, 838-842.	1.6	12

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55	Carry on editing. British Medical Bulletin, 2018, 127, 23-31.	2.7	10
56	Editing mammalian genomes: ethical considerations. Mammalian Genome, 2017, 28, 388-393.	1.0	8
57	Male mice lacking ADAMTS-16 are fertile but exhibit testes of reduced weight. Scientific Reports, 2019, 9, 17195.	1.6	8
58	The Maestro (Mro) Gene Is Dispensable for Normal Sexual Development and Fertility in Mice. PLoS ONE, 2008, 3, e4091.	1.1	7
59	Protection Against XY Gonadal Sex Reversal by a Variant Region on Mouse Chromosome 13. Genetics, 2020, 214, 467-477.	1.2	6
60	Genes, cells and organs: recent developments in the molecular genetics of mammalian sex determination. Mammalian Genome, 1998, 9, 683-687.	1.0	5
61	Using DNA Microarrays. Methods in Molecular Biology, 2008, 461, 605-629.	0.4	5
62	Gadd45g is required for timely Sry expression independently of RSPO1 activity. Reproduction, 2022, 163, 333-340.	1.1	5
63	LIMaS: the JAVA-based application and database for microarray experiment tracking. Mammalian Genome, 2004, 15, 740-747.	1.0	4
64	CRISPR-Cas9-Mediated Mutagenesis: Mind the Gap?. CRISPR Journal, 2018, 1, 263-264.	1.4	3
65	Making sense of heritable human genome editing: Scientific and ethical considerations. Progress in Molecular Biology and Translational Science, 2021, 182, 1-28.	0.9	3
66	The molecular genetic basis of fetal granulosa cell development. Current Opinion in Endocrine and Metabolic Research, 2021, 18, 1-7.	0.6	3
67	Characterisation and use of a functional Gadd45g bacterial artificial chromosome. Scientific Reports, 2018, 8, 17318.	1.6	2
68	Cloning, mitochondrial replacement and genome editing: 25 years of ethical debate since Dolly. Reproduction, 2021, 162, F69-F78.	1.1	2
69	Sexual development. , 2013, , 8-17.		1
70	Broad-spectrum XX and XY gonadal dysgenesis in patients with a homozygous L193S variant in PPP2R3C. European Journal of Endocrinology, 2021, 186, 65-72.	1.9	1
71	The Reproductive System. , 2016, , 121-132.		0
72	The Molecular Control of Testis Determination. , 2018, , 93-99.		0