

Christoph Englert

List of Publications by Year in descending order

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Version: 2024-02-01

34
papers

2,639
citations

304602

22
h-index

395590

33
g-index

40
all docs

40
docs citations

40
times ranked

3302
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of adult nephron progenitors capable of kidney regeneration in zebrafish. <i>Nature</i> , 2011, 470, 95-100.	13.7	258
2	The Wilms tumor genes <i>wt1a</i> and <i>wt1b</i> control different steps during formation of the zebrafish pronephros. <i>Developmental Biology</i> , 2007, 309, 87-96.	0.9	254
3	Insights into Sex Chromosome Evolution and Aging from the Genome of a Short-Lived Fish. <i>Cell</i> , 2015, 163, 1527-1538.	13.5	251
4	Focal segmental glomerulosclerosis is induced by microRNA-193a and its downregulation of WT1. <i>Nature Medicine</i> , 2013, 19, 481-487.	15.2	199
5	The Wilms tumor suppressor gene <i>wt1</i> is required for development of the spleen. <i>Current Biology</i> , 1999, 9, 837-S1.	1.8	193
6	Longitudinal RNA-Seq Analysis of Vertebrate Aging Identifies Mitochondrial Complex I as a Small-Molecule-Sensitive Modifier of Lifespan. <i>Cell Systems</i> , 2016, 2, 122-132.	2.9	155
7	The Wilms' tumor gene <i>Wt1</i> is required for normal development of the retina. <i>EMBO Journal</i> , 2002, 21, 1398-1405.	3.5	135
8	Telomeres shorten while Tert expression increases during ageing of the short-lived fish <i>Nothobranchius furzeri</i> . <i>Mechanisms of Ageing and Development</i> , 2009, 130, 290-296.	2.2	115
9	Mitochondrial DNA copy number and function decrease with age in the short-lived fish <i>Nothobranchius furzeri</i> . <i>Aging Cell</i> , 2011, 10, 824-831.	3.0	114
10	Transcriptomic alterations during ageing reflect the shift from cancer to degenerative diseases in the elderly. <i>Nature Communications</i> , 2018, 9, 327.	5.8	94
11	High tandem repeat content in the genome of the short-lived annual fish <i>Nothobranchius furzeri</i> : a new vertebrate model for aging research. <i>Genome Biology</i> , 2009, 10, R16.	13.9	87
12	A highly conserved retinoic acid responsive element controls <i>wt1a</i> expression in the zebrafish pronephros. <i>Development (Cambridge)</i> , 2009, 136, 2883-2892.	1.2	86
13	Identification and comparative expression analysis of a second <i>wt1</i> gene in zebrafish. <i>Developmental Dynamics</i> , 2006, 235, 554-561.	0.8	84
14	Mapping of quantitative trait loci controlling lifespan in the short-lived fish <i>Nothobranchius furzeri</i> – a new vertebrate model for age research. <i>Aging Cell</i> , 2012, 11, 252-261.	3.0	72
15	<i>Nothobranchius furzeri</i> : A Model for Aging Research and More. <i>Trends in Genetics</i> , 2016, 32, 543-552.	2.9	72
16	Mapping Loci Associated With Tail Color and Sex Determination in the Short-Lived Fish <i>Nothobranchius furzeri</i> . <i>Genetics</i> , 2009, 183, 1385-1395.	1.2	67
17	Integration of Cistromic and Transcriptomic Analyses Identifies <i>Nphs2</i> , <i>Mafb</i> , and <i>Magi2</i> as Wilms' Tumor 1 Target Genes in Podocyte Differentiation and Maintenance. <i>Journal of the American Society of Nephrology: JASN</i> , 2015, 26, 2118-2128.	3.0	67
18	Age-dependent decline in fin regenerative capacity in the short-lived fish <i>Nothobranchius furzeri</i> . <i>Aging Cell</i> , 2015, 14, 857-866.	3.0	66

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19	A microinjection protocol for the generation of transgenic killifish (Species: <i>Nothobranchius</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10	0.8	62
20	Wilms Tumor 1b Expression Defines a Pro-regenerative Macrophage Subtype and Is Required for Organ Regeneration in the Zebrafish. <i>Cell Reports</i> , 2019, 28, 1296-1306.e6.	2.9	61
21	Absence of replicative senescence in cultured cells from the short-lived killifish <i>Nothobranchius furzeri</i> . <i>Experimental Gerontology</i> , 2013, 48, 17-28.	1.2	30
22	Neuron-specific inactivation of <i>Wt1</i> alters locomotion in mice and changes interneuron composition in the spinal cord. <i>Life Science Alliance</i> , 2018, 1, e201800106.	1.3	28
23	Wilms Tumor 1b defines a wound-specific sheath cell subpopulation associated with notochord repair. <i>ELife</i> , 2018, 7, .	2.8	21
24	Alternative splicing of Wilms tumor suppressor 1 (<i>Wt1</i>) exon 4 results in protein isoforms with different functions. <i>Developmental Biology</i> , 2014, 393, 24-32.	0.9	12
25	Dispersion/reaggregation in early development of annual killifishes: Phylogenetic distribution and evolutionary significance of a unique feature. <i>Developmental Biology</i> , 2018, 442, 69-79.	0.9	10
26	<i>Wt1</i> Positive dB4 Neurons in the Hindbrain Are Crucial for Respiration. <i>Frontiers in Neuroscience</i> , 2020, 14, 529487.	1.4	8
27	Systems Analysis Reveals Ageing-Related Perturbations in Retinoids and Sex Hormones in Alzheimer's and Parkinson's Diseases. <i>Biomedicines</i> , 2021, 9, 1310.	1.4	8
28	The African turquoise killifish <i>Nothobranchius furzeri</i> as a model for aging research. <i>Drug Discovery Today: Disease Models</i> , 2018, 27, 15-22.	1.2	7
29	Aging Activates the Immune System and Alters the Regenerative Capacity in the Zebrafish Heart. <i>Cells</i> , 2022, 11, 345.	1.8	7
30	Zebrafish <i>Wtx</i> is a negative regulator of Wnt signaling but is dispensable for embryonic development and organ homeostasis. <i>Developmental Dynamics</i> , 2019, 248, 866-881.	0.8	5
31	<i>Wt1</i> transcription factor impairs cardiomyocyte specification and drives a phenotypic switch from myocardium to epicardium. <i>Development (Cambridge)</i> , 2022, 149, .	1.2	5
32	Analysis of Zebrafish Kidney Development with Time-lapse Imaging Using a Dissecting Microscope Equipped for Optical Sectioning. <i>Journal of Visualized Experiments</i> , 2016, , e53921.	0.2	2
33	Temperature throws a developmental switch. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 12553-12555.	3.3	2
34	The Wilms Tumor Gene <i>wt1a</i> Contributes to Blood-Cerebrospinal Fluid Barrier Function in Zebrafish. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 809962.	1.8	0