

Baris Tursun

List of Publications by Year in descending order

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Version: 2024-02-01

36
papers

1,472
citations

394421

19
h-index

414414

32
g-index

41
all docs

41
docs citations

41
times ranked

2050
citing authors

#	ARTICLE	IF	CITATIONS
1	Robust co-immunoprecipitation with mass spectrometry for <i>Caenorhabditis elegans</i> using solid-phase enhanced sample preparation. <i>BioTechniques</i> , 2022, 72, 175-184.	1.8	2
2	SUMOylation of the chromodomain factor MRG-1 in <i>C. elegans</i> affects chromatin-regulatory dynamics. <i>BioTechniques</i> , 2022, 73, 5-17.	1.8	6
3	The CONJUDOR pipeline for multiplexed knockdown of gene pairs identifies RBBP-5 as a germ cell reprogramming barrier in <i>C. elegans</i> . <i>Nucleic Acids Research</i> , 2021, 49, e22-e22.	14.5	2
4	The pseudogene is required to safeguard germ cells against reprogramming. <i>MicroPublication Biology</i> , 2021, 2021, .	0.1	0
5	Induced Neurons From Germ Cells in <i>Caenorhabditis elegans</i> . <i>Frontiers in Neuroscience</i> , 2021, 15, 771687.	2.8	6
6	Conversion of Germ Cells to Somatic Cell Types in <i>C. elegans</i> . <i>Journal of Developmental Biology</i> , 2020, 8, 24.	1.7	3
7	Epigenetic chaperoning of aging. <i>Aging</i> , 2020, 12, 1044-1046.	3.1	1
8	The conserved histone chaperone LIN-53 is required for normal lifespan and maintenance of muscle integrity in <i>Caenorhabditis elegans</i> . <i>Aging Cell</i> , 2019, 18, e13012.	6.7	13
9	Strategies for in vivo reprogramming. <i>Current Opinion in Cell Biology</i> , 2019, 61, 9-15.	5.4	19
10	CRISPR-activation-based screen reveals neuronal fate promotion by polycomb repressive complex 2 during direct reprogramming. <i>Stem Cell Investigation</i> , 2019, 6, 32-32.	3.0	0
11	HOT or not: examining the basis of high-occupancy target regions. <i>Nucleic Acids Research</i> , 2019, 47, 5735-5745.	14.5	41
12	<i>Caenorhabditis elegans</i> and systems biology – it is systems™ all the way down!. <i>Current Opinion in Systems Biology</i> , 2019, 13, iv-vi.	2.6	0
13	MRG-1/MRG15 Is a Barrier for Germ Cell to Neuron Reprogramming in <i>Caenorhabditis elegans</i> . <i>Genetics</i> , 2019, 211, 121-139.	2.9	38
14	Application of RNAi and Heat-shock-induced Transcription Factor Expression to Reprogram Germ Cells to Neurons in <i>C. elegans</i> . <i>Journal of Visualized Experiments</i> , 2018, , .	0.3	5
15	Determinants of promoter and enhancer transcription directionality in metazoans. <i>Nature Communications</i> , 2018, 9, 4472.	12.8	22
16	FACT Sets a Barrier for Cell Fate Reprogramming in <i>Caenorhabditis elegans</i> and Human Cells. <i>Developmental Cell</i> , 2018, 46, 611-626.e12.	7.0	89
17	Transdifferentiation: do transition states lie on the path of development?. <i>Current Opinion in Systems Biology</i> , 2018, 11, 18-23.	2.6	17
18	Neuronal inhibition of the autophagy nucleation complex extends life span in post-reproductive <i>C. elegans</i> . <i>Genes and Development</i> , 2017, 31, 1561-1572.	5.9	67

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19	A tissue-specific protein purification approach in <i>Caenorhabditis elegans</i> identifies novel interaction partners of DLG-1/Discs large. <i>BMC Biology</i> , 2016, 14, 66.	3.8	40
20	Increasing Notch signaling antagonizes PRC2-mediated silencing to promote reprogramming of germ cells into neurons. <i>ELife</i> , 2016, 5, .	6.0	45
21	Two distinct types of neuronal asymmetries are controlled by the <i>Caenorhabditis elegans</i> zinc finger transcription factor <i>die-1</i> . <i>Genes and Development</i> , 2014, 28, 34-43.	5.9	29
22	Cellular reprogramming processes in <i>Drosophila</i> and <i>C. elegans</i> . <i>Current Opinion in Genetics and Development</i> , 2012, 22, 475-484.	3.3	6
23	Removal of Polycomb Repressive Complex 2 Makes <i>C. elegans</i> Germ Cells Susceptible to Direct Conversion into Specific Somatic Cell Types. <i>Cell Reports</i> , 2012, 2, 1178-1186.	6.4	119
24	Direct Conversion of <i>C. elegans</i> Germ Cells into Specific Neuron Types. <i>Science</i> , 2011, 331, 304-308.	12.6	219
25	The Groucho ortholog UNC-37 interacts with the short Groucho-like protein LSY-22 to control developmental decisions in <i>C. elegans</i> . <i>Development (Cambridge)</i> , 2010, 137, 1799-1805.	2.5	31
26	Neuron-type specific regulation of a 3'UTR through redundant and combinatorially acting <i>cis</i> -regulatory elements. <i>Rna</i> , 2010, 16, 349-363.	3.5	16
27	A Toolkit and Robust Pipeline for the Generation of Fosmid-Based Reporter Genes in <i>C. elegans</i> . <i>PLoS ONE</i> , 2009, 4, e4625.	2.5	160
28	The <i>C. elegans</i> Tailless/TLX transcription factor <i>nhr-67</i> controls neuronal identity and left/right asymmetric fate diversification. <i>Development (Cambridge)</i> , 2009, 136, 2933-2944.	2.5	42
29	Regulation of Estrogen-Dependent Transcription by the LIM Cofactors CLIM and RLIM in Breast Cancer. <i>Cancer Research</i> , 2009, 69, 128-136.	0.9	57
30	Proteasomal selection of multiprotein complexes recruited by LIM homeodomain transcription factors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 15000-15005.	7.1	46
31	Dynamic expression of LIM cofactors in the developing mouse neural tube. <i>Developmental Dynamics</i> , 2006, 235, 786-791.	1.8	29
32	The ubiquitin ligase Rnf6 regulates local LIM kinase 1 levels in axonal growth cones. <i>Genes and Development</i> , 2005, 19, 2307-2319.	5.9	98
33	Mitochondrial signal peptidases of yeast: The rhomboid peptidase Pcp1 and its substrate cytochrome c peroxidase. <i>Gene</i> , 2005, 354, 58-63.	2.2	24
34	Comparing protein stabilities during zebrafish embryogenesis. <i>Cytotechnology</i> , 2003, 25, 85-89.	0.7	5
35	A Novel Two-step Mechanism for Removal of a Mitochondrial Signal Sequence Involves the mAAA Complex and the Putative Rhomboid Protease Pcp1. <i>Journal of Molecular Biology</i> , 2002, 323, 835-843.	4.2	169
36	The UNC-83/UNC-84 LINC members are required for body wall muscle nuclei positioning in <i>C. elegans</i> . <i>Matters Select</i> , 0, , .	3.0	0