## Kaia Achim

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8963639/publications.pdf

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1039406 1199166 12 905 9 12 citations h-index g-index papers 15 15 15 1685 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	High-throughput spatial mapping of single-cell RNA-seq data to tissue of origin. Nature Biotechnology, 2015, 33, 503-509.	9.4	380
2	Profiling cellular diversity in sponges informs animal cell type and nervous system evolution. Science, 2021, 374, 717-723.	6.0	111
3	Evolution of neuronal types and families. Current Opinion in Neurobiology, 2019, 56, 144-152.	2.0	94
4	Whole-organism cellular gene-expression atlas reveals conserved cell types in the ventral nerve cord of <i>Platynereis dumerilii</i> . Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 5878-5885.	<b>3.</b> 3	66
5	The role of <i>Tal2</i> and <i>Tal1</i> in the differentiation of midbrain GABAergic neuron precursors. Biology Open, 2013, 2, 990-997.	0.6	57
6	Transcriptional regulatory mechanisms underlying the GABAergic neuron fate in different diencephalic prosomeres. Development (Cambridge), 2012, 139, 3795-3805.	1.2	49
7	Whole-Body Single-Cell Sequencing Reveals Transcriptional Domains in the Annelid Larval Body. Molecular Biology and Evolution, 2018, 35, 1047-1062.	3.5	48
8	Structural evolution of cell types by step-wise assembly of cellular modules. Current Opinion in Genetics and Development, 2014, 27, 102-108.	1.5	41
9	Identifying Cell Types from Spatially Referenced Single-Cell Expression Datasets. PLoS Computational Biology, 2014, 10, e1003824.	1.5	26
10	From spiral cleavage to bilateral symmetry: the developmental cell lineage of the annelid brain. BMC Biology, 2019, 17, 81.	1.7	14
11	Molecular Fingerprint and Developmental Regulation of the Tegmental GABAergic and Glutamatergic Neurons Derived from the Anterior Hindbrain. Cell Reports, 2020, 33, 108268.	2.9	11
12	Spatial Transcriptomics: Constructing a Single-Cell Resolution Transcriptome-Wide Expression Atlas. Methods in Molecular Biology, 2018, 1649, 111-125.	0.4	5