## Hagen U Tilgner

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5773696/publications.pdf

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687220 940416 2,357 15 13 16 citations h-index g-index papers 22 22 22 3953 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	A single-molecule long-read survey of the human transcriptome. Nature Biotechnology, 2013, 31, 1009-1014.	9.4	600
2	SynGO: An Evidence-Based, Expert-Curated Knowledge Base for the Synapse. Neuron, 2019, 103, 217-234.e4.	3.8	518
3	Defining a personal, allele-specific, and single-molecule long-read transcriptome. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 9869-9874.	3.3	259
4	Single-cell isoform RNA sequencing characterizes isoforms in thousands of cerebellar cells. Nature Biotechnology, 2018, 36, 1197-1202.	9.4	253
5	Comprehensive transcriptome analysis using synthetic long-read sequencing reveals molecular co-association of distant splicing events. Nature Biotechnology, 2015, 33, 736-742.	9.4	205
6	A spatially resolved brain region- and cell type-specific isoform atlas of the postnatal mouse brain. Nature Communications, 2021, 12, 463.	5.8	109
7	Microfluidic isoform sequencing shows widespread splicing coordination in the human transcriptome. Genome Research, 2018, 28, 231-242.	2.4	64
8	Accurate Identification and Analysis of Human mRNA Isoforms Using Deep Long Read Sequencing. G3: Genes, Genomes, Genetics, 2013, 3, 387-397.	0.8	59
9	Schizophrenia is defined by cell-specific neuropathology and multiple neurodevelopmental mechanisms in patient-derived cerebral organoids. Molecular Psychiatry, 2022, 27, 1416-1434.	4.1	57
10	Single-nuclei isoform RNA sequencing unlocks barcoded exon connectivity in frozen brain tissue. Nature Biotechnology, 2022, 40, 1082-1092.	9.4	52
11	Mapping genetic variations to three-dimensional protein structures to enhance variant interpretation: a proposed framework. Genome Medicine, 2017, 9, 113.	3.6	47
12	Getting the Entire Message: Progress in Isoform Sequencing. Frontiers in Genetics, 2019, 10, 709.	1.1	39
13	Targeted, High-Resolution RNA Sequencing of Non-coding Genomic Regions Associated With Neuropsychiatric Functions. Frontiers in Genetics, 2019, 10, 309.	1.1	28
14	Sequencing of individual barcoded cDNAs using Pacific Biosciences and Oxford Nanopore Technologies reveals platform-specific error patterns. Genome Research, 2022, 32, 726-737.	2.4	13
15	ScisorWiz: visualizing differential isoform expression in single-cell long-read data. Bioinformatics, 2022, 38, 3474-3476.	1.8	11