Yoshiaki Tanaka

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9640270/publications.pdf

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31	2,417	17 h-index	27
papers	citations		g-index
35	35	35	3733
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Engineering of human brain organoids with a functional vascular-like system. Nature Methods, 2019, 16, 1169-1175.	19.0	551
2	Fusion of Regionally Specified hPSC-Derived Organoids Models Human Brain Development and Interneuron Migration. Cell Stem Cell, 2017, 21, 383-398.e7.	11.1	508
3	hESC-Derived Thalamic Organoids Form Reciprocal Projections When Fused with Cortical Organoids. Cell Stem Cell, 2019, 24, 487-497.e7.	11.1	305
4	Synthetic Analyses of Single-Cell Transcriptomes from Multiple Brain Organoids and Fetal Brain. Cell Reports, 2020, 30, 1682-1689.e3.	6.4	150
5	Histone Deacetylases Positively Regulate Transcription through the Elongation Machinery. Cell Reports, 2015, 13, 1444-1455.	6.4	138
6	Two methods for full-length RNA sequencing for low quantities of cells and single cells. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 594-599.	7.1	103
7	Direct Reprogramming of Human Dermal Fibroblasts Into Endothelial Cells Using ER71/ETV2. Circulation Research, 2017, 120, 848-861.	4.5	90
8	Human induced pluripotent stem cells and neurodegenerative disease. Current Opinion in Neurology, 2012, 25, 125-130.	3.6	64
9	Dysregulation of BRD4 Function Underlies the Functional Abnormalities of MeCP2 Mutant Neurons. Molecular Cell, 2020, 79, 84-98.e9.	9.7	53
10	Expression of the transcription factor PU.1 induces the generation of microglia-like cells in human cortical organoids. Nature Communications, 2022, 13, 430.	12.8	49
11	Effects of Alu elements on global nucleosome positioning in the human genome. BMC Genomics, 2010, 11, 309.	2.8	46
12	Single cell transcriptomics reveals unanticipated features of early hematopoietic precursors. Nucleic Acids Research, 2017, 45, gkw1214.	14.5	40
13	Uhrf1 regulates active transcriptional marks at bivalent domains in pluripotent stem cells through Setd1a. Nature Communications, 2018, 9, 2583.	12.8	35
14	The RNA exosome nuclease complex regulates human embryonic stem cell differentiation. Journal of Cell Biology, 2019, 218, 2564-2582.	5.2	35
15	X Chromosome of Female Cells Shows Dynamic Changes in Status during Human Somatic Cell Reprogramming. Stem Cell Reports, 2014, 2, 896-909.	4.8	33
16	Transcriptional regulation in pluripotent stem cells by methyl CpG-binding protein 2 (MeCP2). Human Molecular Genetics, 2014, 23, 1045-1055.	2.9	32
17	Regulation of the DNA Methylation Landscape in Human Somatic Cell Reprogramming by the miR-29 Family. Stem Cell Reports, 2016, 7, 43-54.	4.8	31
18	Exploration of alcohol use disorder-associated brain miRNA–mRNA regulatory networks. Translational Psychiatry, 2021, 11, 504.	4.8	23

#	Article	IF	Citations
19	Transcriptome Signature and Regulation in Human Somatic Cell Reprogramming. Stem Cell Reports, 2015, 4, 1125-1139.	4.8	19
20	Salivary microRNAs identified by small RNA sequencing and machine learning as potential biomarkers of alcohol dependence. Epigenomics, 2019, 11, 739-749.	2.1	19
21	Identification and validation of a five-IncRNA signature for predicting survival with targeted drug candidates in ovarian cancer. Bioengineered, 2021, 12, 3263-3274.	3.2	19
22	Mural Cell-Specific Deletion of Cerebral Cavernous Malformation 3 in the Brain Induces Cerebral Cavernous Malformations. Arteriosclerosis, Thrombosis, and Vascular Biology, 2020, 40, 2171-2186.	2.4	18
23	Regional specification and complementation with non-neuroectodermal cells in human brain organoids. Journal of Molecular Medicine, 2021, 99, 489-500.	3.9	14
24	Positional variations among heterogeneous nucleosome maps give dynamical information on chromatin. Chromosoma, 2010, 119, 391-404.	2.2	8
25	An assessment of prediction algorithms for nucleosome positioning. Genome Informatics, 2009, 23, 169-78.	0.4	7
26	AN ASSESSMENT OF PREDICTION ALGORITHMS FOR NUCLEOSOME POSITIONING. , 2009, , .		5
27	Impact of Retrotransposons in Pluripotent Stem Cells. Molecules and Cells, 2012, 34, 509-516.	2.6	5
28	Challenges for Computational Stem Cell Biology: A Discussion for the Field. Stem Cell Reports, 2021, 16, 3-9.	4.8	4
29	New Advances in Human X Chromosome Status from a Developmental and Stem Cell Biology. Tissue Engineering and Regenerative Medicine, 2017, 14, 643-652.	3.7	0
30	EFFECT OF SALIVA-PAROTIN UPON THE PATHOLOGICAL CHANGEOF THE PARADENTIUM IN RATS DURING THE COURSEOF THE EXPERIMENTAL LATHYRISM. Koku Eisei Gakkai Zasshi, 1960, 10, 184-195.	0.0	0
31	3 Genetic and Epigenetic Considerations in iPSC Technology. , 2017, , 51-86.		O