Yuki Kato

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3126214/publications.pdf

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14 papers	323 citations	1163117 8 h-index	1058476 14 g-index
F F 0 2 0	5		8
15 all docs	15 docs citations	15 times ranked	229 citing authors

#	Article	IF	CITATIONS
1	Mutations in the adenosine deaminase ADAR1 that prevent endogenous Z-RNA binding induce Aicardi-Goutià res-syndrome-like encephalopathy. Immunity, 2021, 54, 1976-1988.e7.	14.3	56
2	<scp>ADAR</scp> 1â€mediated <scp>RNA</scp> editing is required for thymic selfâ€tolerance and inhibition of autoimmunity. EMBO Reports, 2018, 19, .	4.5	47
3	RNA editing at a limited number of sites is sufficient to prevent MDA5 activation in the mouse brain. PLoS Genetics, 2021, 17, e1009516.	3. 5	42
4	DAFS: simultaneous aligning and folding of RNA sequences via dual decomposition. Bioinformatics, 2012, 28, 3218-3224.	4.1	40
5	A comparative analysis of ADAR mutant mice reveals site-specific regulation of RNA editing. Rna, 2020, 26, 454-469.	3. 5	38
6	Prediction of RNA secondary structure including pseudoknots for long sequences. Briefings in Bioinformatics, 2022, 23, .	6.5	36
7	Age-dependent decline in remyelination capacity is mediated by apelin–APJ signaling. Nature Aging, 2021, 1, 284-294.	11.6	18
8	ADAR1 Regulates Early T Cell Development via MDA5-Dependent and -Independent Pathways. Journal of Immunology, 2020, 204, 2156-2168.	0.8	17
9	An Aicardi-GoutiÃ"res Syndromeâ€"Causative Point Mutation in <i>Adar1</i> Gene Invokes Multiorgan Inflammation and Late-Onset Encephalopathy in Mice. Journal of Immunology, 2021, 207, 3016-3027.	0.8	11
10	An accessibility-incorporated method for accurate prediction of RNA–RNA interactions from sequence data. Bioinformatics, 2017, 33, 202-209.	4.1	7
11	Dimethylarginine dimethylaminohydrolase 1 as a novel regulator of oligodendrocyte differentiation in the central nervous system remyelination. Glia, 2021, 69, 2591-2604.	4.9	4
12	Myotube-derived factor promotes oligodendrocyte precursor cell proliferation. Biochemical and Biophysical Research Communications, 2018, 500, 609-613.	2.1	3
13	Bivartect: accurate and memory-saving breakpoint detection by direct read comparison. Bioinformatics, 2020, 36, 2725-2730.	4.1	3
14	Alignment-free comparative genomic screen for structured RNAs using coarse-grained secondary structure dot plots. BMC Genomics, 2017, 18, 935.	2.8	1