

Sebastian Carrasco Pro

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8802807/publications.pdf>

Version: 2024-02-01

11
papers

468
citations

1040056

9
h-index

1281871

11
g-index

14
all docs

14
docs citations

14
times ranked

934
citing authors

#	ARTICLE	IF	CITATIONS
1	Development and validation of a broad scheme for prediction of HLA class II restricted T cell epitopes. <i>Journal of Immunological Methods</i> , 2015, 422, 28-34.	1.4	171
2	Automatic Generation of Validated Specific Epitope Sets. <i>Journal of Immunology Research</i> , 2015, 2015, 1-11.	2.2	90
3	Global landscape of mouse and human cytokine transcriptional regulation. <i>Nucleic Acids Research</i> , 2018, 46, 9321-9337.	14.5	56
4	Differential Recognition of <i>Mycobacterium tuberculosis</i> Specific Epitopes as a Function of Tuberculosis Disease History. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2017, 196, 772-781.	5.6	39
5	Microbiota epitope similarity either dampens or enhances the immunogenicity of disease-associated antigenic epitopes. <i>PLoS ONE</i> , 2018, 13, e0196551.	2.5	31
6	Comprehensive mapping of the human cytokine gene regulatory network. <i>Nucleic Acids Research</i> , 2020, 48, 12055-12073.	14.5	20
7	Identification of Single Nucleotide Non-coding Driver Mutations in Cancer. <i>Frontiers in Genetics</i> , 2018, 9, 16.	2.3	19
8	Prediction of genome-wide effects of single nucleotide variants on transcription factor binding. <i>Scientific Reports</i> , 2020, 10, 17632.	3.3	11
9	TsAg5, a <i>Taenia solium</i> cysticercus protein with a marginal trypsin-like activity in the diagnosis of human neurocysticercosis. <i>Molecular and Biochemical Parasitology</i> , 2011, 180, 115-119.	1.1	10
10	Improved pan-specific MHC class I peptide-binding predictions using a novel representation of the MHC-binding cleft environment. <i>Tissue Antigens</i> , 2014, 83, 94-100.	1.0	10
11	Discovering human transcription factor physical interactions with genetic variants, novel DNA motifs, and repetitive elements using enhanced yeast one-hybrid assays. <i>Genome Research</i> , 2019, 29, 1533-1544.	5.5	10