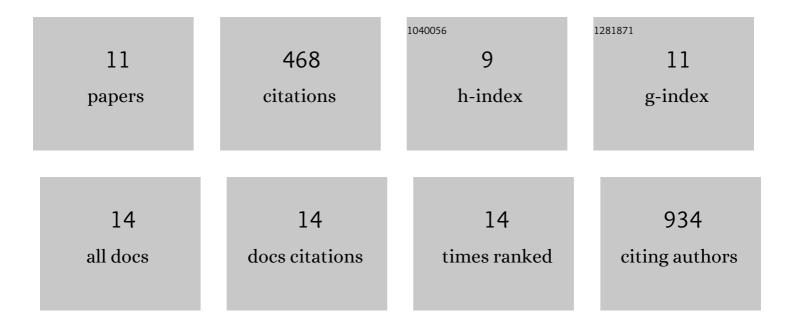
Sebastian Carrasco Pro

List of Publications by Year in descending order

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

Version: 2024-02-01



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