

# Domenec Farre

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1702540/publications.pdf>

Version: 2024-02-01

18  
papers

2,908  
citations

687335

13  
h-index

888047

17  
g-index

18  
all docs

18  
docs citations

18  
times ranked

5667  
citing authors

#	ARTICLE	IF	CITATIONS
1	Divergent Traits and Ligand-Binding Properties of the Cytomegalovirus CD48 Gene Family. <i>Viruses</i> , 2020, 12, 813.	3.3	2
2	Divergent Traits and Ligand-Binding Features of the Cytomegalovirus CD48 Gene Family. <i>Proceedings (mdpi)</i> , 2020, 50, .	0.2	0
3	Subversion of natural killer cell responses by a cytomegalovirus-encoded soluble CD48 decoy receptor. <i>PLoS Pathogens</i> , 2019, 15, e1007658.	4.7	16
4	A Prominent Role of the Human Cytomegalovirus UL8 Glycoprotein in Restraining Proinflammatory Cytokine Production by Myeloid Cells at Late Times during Infection. <i>Journal of Virology</i> , 2018, 92, .	3.4	25
5	Immunoglobulin superfamily members encoded by viruses and their multiple roles in immune evasion. <i>European Journal of Immunology</i> , 2017, 47, 780-796.	2.9	30
6	Elusive Role of the CD94/NKG2C NK Cell Receptor in the Response to Cytomegalovirus: Novel Experimental Observations in a Reporter Cell System. <i>Frontiers in Immunology</i> , 2017, 8, 1317.	4.8	21
7	Novel Role of 3â€™UTR-Embedded Alu Elements as Facilitators of Processed Pseudogene Genesis and Host Gene Capture by Viral Genomes. <i>PLoS ONE</i> , 2016, 11, e0169196.	2.5	13
8	Signaling Lymphocytic Activation Molecule Family Receptor Homologs in New World Monkey Cytomegaloviruses. <i>Journal of Virology</i> , 2015, 89, 11323-11336.	3.4	17
9	Cytomegalovirus m154 Hinders CD48 Cell-Surface Expression and Promotes Viral Escape from Host Natural Killer Cell Control. <i>PLoS Pathogens</i> , 2014, 10, e1004000.	4.7	34
10	The Vertebrate RCAN Gene Family: Novel Insights into Evolution, Structure and Regulation. <i>PLoS ONE</i> , 2014, 9, e85539.	2.5	19
11	Comparative and demographic analysis of orang-utan genomes. <i>Nature</i> , 2011, 469, 529-533.	27.8	541
12	Heterogeneous Patterns of Gene-Expression Diversification in Mammalian Gene Duplicates. <i>Molecular Biology and Evolution</i> , 2010, 27, 325-335.	8.9	40
13	PEAKS: identification of regulatory motifs by their position in DNA sequences. <i>Bioinformatics</i> , 2007, 23, 243-244.	4.1	15
14	Housekeeping genes tend to show reduced upstream sequence conservation. <i>Genome Biology</i> , 2007, 8, R140.	9.6	64
15	Positional bias of general and tissue-specific regulatory motifs in mouse gene promoters. <i>BMC Genomics</i> , 2007, 8, 459.	2.8	19
16	ABS: a database of Annotated regulatory Binding Sites from orthologous promoters. <i>Nucleic Acids Research</i> , 2006, 34, D63-D67.	14.5	56
17	Identification of patterns in biological sequences at the ALGGEN server: PROMO and MALGEN. <i>Nucleic Acids Research</i> , 2003, 31, 3651-3653.	14.5	876
18	PROMO: detection of known transcription regulatory elements using species-tailored searches. <i>Bioinformatics</i> , 2002, 18, 333-334.	4.1	1,120