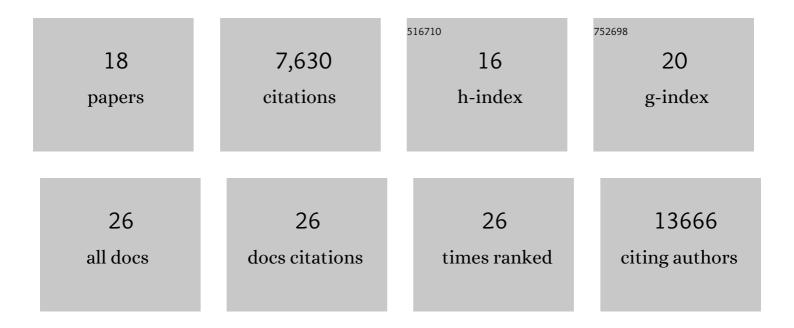
Eduard Porta-Pardo

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1389384/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	The structural coverage of the human proteome before and after AlphaFold. PLoS Computational Biology, 2022, 18, e1009818.	3.2	72
2	Detection of oncogenic and clinically actionable mutations in cancer genomes critically depends on variant calling tools. Bioinformatics, 2022, 38, 3181-3191.	4.1	9
3	Germline genetic contribution to the immune landscape of cancer. Immunity, 2021, 54, 367-386.e8.	14.3	95
4	Neoantigen prediction and computational perspectives towards clinical benefit: recommendations from the ESMO Precision Medicine Working Group. Annals of Oncology, 2020, 31, 978-990.	1.2	87
5	Understanding oncogenicity of cancer driver genes and mutations in the cancer genomics era. FEBS Letters, 2020, 594, 4233-4246.	2.8	20
6	Cancer3D 2.0: interactive analysis of 3D patterns of cancer mutations in cancer subsets. Nucleic Acids Research, 2019, 47, D895-D899.	14.5	12
7	Comprehensive Characterization of Cancer Driver Genes and Mutations. Cell, 2018, 173, 371-385.e18.	28.9	1,670
8	Perspective on Oncogenic Processes at the End of the Beginning of Cancer Genomics. Cell, 2018, 173, 305-320.e10.	28.9	272
9	The Immune Landscape of Cancer. Immunity, 2018, 48, 812-830.e14.	14.3	3,706
10	The Functional Impact of Alternative Splicing in Cancer. Cell Reports, 2017, 20, 2215-2226.	6.4	517
11	Comparison of algorithms for the detection of cancer drivers at subgene resolution. Nature Methods, 2017, 14, 782-788.	19.0	72
12	AIRE genetic variants and predisposition to polygenic autoimmune disease: The case of Graves' disease and a systematic literature review. Human Immunology, 2016, 77, 643-651.	2.4	20
13	Mutation Drivers of Immunological Responses to Cancer. Cancer Immunology Research, 2016, 4, 789-798.	3.4	32
14	A Pan-Cancer Catalogue of Cancer Driver Protein Interaction Interfaces. PLoS Computational Biology, 2015, 11, e1004518.	3.2	122
15	Analysis of Individual Protein Regions Provides Novel Insights on Cancer Pharmacogenomics. PLoS Computational Biology, 2015, 11, e1004024.	3.2	10
16	Cancer3D: understanding cancer mutations through protein structures. Nucleic Acids Research, 2015, 43, D968-D973.	14.5	46
17	Autoimmune Predisposition in Down Syndrome May Result from a Partial Central Tolerance Failure due to Insufficient Intrathymic Expression of <i>AIRE</i> and Peripheral Antigens. Journal of Immunology, 2014, 193, 3872-3879.	0.8	88
18	e-Driver: a novel method to identify protein regions driving cancer. Bioinformatics, 2014, 30, 3109-3114.	4.1	116