## Boris P Hejblum

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7490253/publications.pdf

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623734 526287 1,273 29 14 27 citations g-index h-index papers 40 40 40 3094 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Systems analysis of sex differences reveals an immunosuppressive role for testosterone in the response to influenza vaccination. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 869-874.	7.1	542
2	Systems Vaccinology Identifies an Early Innate Immune Signature as a Correlate of Antibody Responses to the Ebola Vaccine rVSV-ZEBOV. Cell Reports, 2017, 20, 2251-2261.	6.4	107
3	CD177, a specific marker of neutrophil activation, is associated with coronavirus disease 2019 severity and death. IScience, 2021, 24, 102711.	4.1	79
4	Time-Course Gene Set Analysis for Longitudinal Gene Expression Data. PLoS Computational Biology, 2015, 11, e1004310.	3.2	60
5	Pattern of polyphenol intake and the long-term risk of dementia in older persons. Neurology, 2018, 90, e1979-e1988.	1.1	55
6	Group and sparse group partial least square approaches applied in genomics context. Bioinformatics, 2016, 32, 35-42.	4.1	50
7	Immune Alterations in a Patient with SARS-CoV-2-Related Acute Respiratory Distress Syndrome. Journal of Clinical Immunology, 2020, 40, 1082-1092.	3.8	48
8	Dietâ€Related Metabolites Associated with Cognitive Decline Revealed by Untargeted Metabolomics in a Prospective Cohort. Molecular Nutrition and Food Research, 2019, 63, e1900177.	3.3	40
9	Predicting Progression to Advanced Age-Related Macular Degeneration from Clinical, Genetic, and Lifestyle Factors UsingÂMachine Learning. Ophthalmology, 2021, 128, 587-597.	<b>5.</b> 2	34
10	Long-lasting severe immune dysfunction in Ebola virus disease survivors. Nature Communications, 2020, 11, 3730.	12.8	33
11	Phenomeâ€Wide Association Study of Autoantibodies to Citrullinated and Noncitrullinated Epitopes in Rheumatoid Arthritis. Arthritis and Rheumatology, 2017, 69, 742-749.	5 <b>.</b> 6	26
12	cytometree: A binary tree algorithm for automatic gating in cytometry analysis. Cytometry Part A: the Journal of the International Society for Analytical Cytology, 2018, 93, 1132-1140.	1.5	21
13	Probabilistic record linkage of de-identified research datasets with discrepancies using diagnosis codes. Scientific Data, 2019, 6, 180298.	5.3	21
14	PheProb: probabilistic phenotyping using diagnosis codes to improve power for genetic association studies. Journal of the American Medical Informatics Association: JAMIA, 2018, 25, 1359-1365.	4.4	18
15	dearseq: a variance component score test for RNA-seq differential analysis that effectively controls the false discovery rate. NAR Genomics and Bioinformatics, 2020, 2, Iqaa093.	3.2	18
16	Benefits of dimension reduction in penalized regression methods for high-dimensional grouped data: a case study in low sample size. Bioinformatics, 2019, 35, 3628-3634.	4.1	15
17	Sequential Dirichlet process mixtures of multivariate skew \$t\$-distributions for model-based clustering of flow cytometry data. Annals of Applied Statistics, 2019, 13, .	1.1	14
18	Predicting the retinal content in omegaâ€3 fatty acids for ageâ€related macularâ€degeneration. Clinical and Translational Medicine, 2021, 11, e404.	4.0	12

#	Article	IF	CITATIONS
19	Gene Expression Signatures Associated With Immune and Virological Responses to Therapeutic Vaccination With Dendritic Cells in HIV-Infected Individuals. Frontiers in Immunology, 2019, 10, 874.	4.8	11
20	Variance component score test for time-course gene set analysis of longitudinal RNA-seq data. Biostatistics, 2017, 18, 589-604.	1.5	10
21	Association Between Anti–Citrullinated Fibrinogen Antibodies and Coronary Artery Disease in Rheumatoid Arthritis. Arthritis Care and Research, 2018, 70, 1113-1117.	3.4	9
22	T Cell Immunogenicity, Gene Expression Profile, and Safety of Four Heterologous Prime-Boost Combinations of HIV Vaccine Candidates in Healthy Volunteers: Results of the Randomized Multi-Arm Phase I/II ANRS VRI01 Trial. Journal of Immunology, 2022, 208, 2663-2674.	0.8	8
23	Evidence synthesis through a degradation model applied to myocardial infarction. Lifetime Data Analysis, 2013, 19, 1-18.	0.9	5
24	Kernel machine score test for pathway analysis in the presence of semi-competing risks. Statistical Methods in Medical Research, 2018, 27, 1099-1114.	1.5	2
25	Analyzing cellular immunogenicity in vaccine clinical trials: a new statistical method including non-specific responses for accurate estimation of vaccine effect. Journal of Immunological Methods, 2020, 477, 112711.	1.4	2
26	Bayesian mixture models for cytometry data analysis. Wiley Interdisciplinary Reviews: Computational Statistics, 2021, 13, e1535.	3.9	1
27	Semi-supervised estimation of covariance with application to phenome-wide association studies with electronic medical records data. Statistical Methods in Medical Research, 2020, 29, 455-465.	1.5	0
28	ATLAS: an automated association test using probabilistically linked health records with application to genetic studies. Journal of the American Medical Informatics Association: JAMIA, 2021, 28, 2582-2592.	4.4	0
29	Doubly robust evaluation of high-dimensional surrogate markers. Biostatistics, 2023, 24, 985-999.	1.5	O