

Dmitriy A Bolotin

List of Publications by Citations

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27
papers

3,148
citations

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h-index

28
g-index

28
ext. papers

4,580
ext. citations

13.9
avg, IF

4.69
L-index

#	Paper	IF	Citations
27	MiXCR: software for comprehensive adaptive immunity profiling. <i>Nature Methods</i> , 2015 , 12, 380-1	21.6	696
26	VDJtools: Unifying Post-analysis of T Cell Receptor Repertoires. <i>PLoS Computational Biology</i> , 2015 , 11, e1004503	5	282
25	Towards error-free profiling of immune repertoires. <i>Nature Methods</i> , 2014 , 11, 653-5	21.6	267
24	Age-related decrease in TCR repertoire diversity measured with deep and normalized sequence profiling. <i>Journal of Immunology</i> , 2014 , 192, 2689-98	5.3	249
23	Local fitness landscape of the green fluorescent protein. <i>Nature</i> , 2016 , 533, 397-401	50.4	232
22	tcR: an R package for T cell receptor repertoire advanced data analysis. <i>BMC Bioinformatics</i> , 2015 , 16, 175	3.6	156
21	MiTCR: software for T-cell receptor sequencing data analysis. <i>Nature Methods</i> , 2013 , 10, 813-4	21.6	138
20	High-throughput identification of antigen-specific TCRs by TCR gene capture. <i>Nature Medicine</i> , 2013 , 19, 1534-41	50.5	127
19	Antigen receptor repertoire profiling from RNA-seq data. <i>Nature Biotechnology</i> , 2017 , 35, 908-911	44.5	125
18	Next generation sequencing for TCR repertoire profiling: platform-specific features and correction algorithms. <i>European Journal of Immunology</i> , 2012 , 42, 3073-83	6.1	121
17	High-quality full-length immunoglobulin profiling with unique molecular barcoding. <i>Nature Protocols</i> , 2016 , 11, 1599-616	18.8	109
16	Preparing unbiased T-cell receptor and antibody cDNA libraries for the deep next generation sequencing profiling. <i>Frontiers in Immunology</i> , 2013 , 4, 456	8.4	104
15	Pairing of T-cell receptor chains via emulsion PCR. <i>European Journal of Immunology</i> , 2013 , 43, 2507-15	6.1	95
14	Dynamics of Individual T Cell Repertoires: From Cord Blood to Centenarians. <i>Journal of Immunology</i> , 2016 , 196, 5005-13	5.3	94
13	Distinctive properties of identical twins' TCR repertoires revealed by high-throughput sequencing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 5980-5	11.5	86
12	Quantitative profiling of immune repertoires for minor lymphocyte counts using unique molecular identifiers. <i>Journal of Immunology</i> , 2015 , 194, 6155-63	5.3	58
11	Quantitative tracking of T cell clones after haematopoietic stem cell transplantation. <i>EMBO Molecular Medicine</i> , 2011 , 3, 201-7	12	53

10	Mother and child T cell receptor repertoires: deep profiling study. <i>Frontiers in Immunology</i> , 2013 , 4, 463	8.4	36
9	Huge Overlap of Individual TCR Beta Repertoires. <i>Frontiers in Immunology</i> , 2013 , 4, 466	8.4	32
8	Comparative analysis of murine T-cell receptor repertoires. <i>Immunology</i> , 2018 , 153, 133-144	7.8	29
7	First autologous hematopoietic SCT for ankylosing spondylitis: a case report and clues to understanding the therapy. <i>Bone Marrow Transplantation</i> , 2012 , 47, 1479-81	4.4	20
6	Can we see PIP(3) and hydrogen peroxide with a single probe?. <i>Antioxidants and Redox Signaling</i> , 2012 , 17, 505-12	8.4	18
5	Redberry: a computer algebra system designed for tensor manipulation. <i>Journal of Physics: Conference Series</i> , 2015 , 608, 012060	0.3	7
4	Rhodobase, a meta-analytical tool for reconstructing gene regulatory networks in a model photosynthetic bacterium. <i>BioSystems</i> , 2011 , 103, 125-31	1.9	5
3	Reply to "Evaluation of immune repertoire inference methods from RNA-seq data". <i>Nature Biotechnology</i> , 2018 , 36, 1035-1036	44.5	4
2	Application of nonsense-mediated primer exclusion (NOPE) for preparation of unique molecular barcoded libraries. <i>BMC Genomics</i> , 2017 , 18, 440	4.5	2
1	Distinct organization of adaptive immunity in the long-lived rodent <i>Spalax galili</i> . <i>Nature Aging</i> , 2021 , 1, 179-189		1