

Jonathan J Havel

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

21
papers

7,956
citations

15
h-index

22
g-index

22
ext. papers

10,410
ext. citations

18.5
avg. IF

6.08
L-index

#	Paper	IF	Citations
21	Cancer immunology. Mutational landscape determines sensitivity to PD-1 blockade in non-small cell lung cancer. <i>Science</i> , 2015 , 348, 124-8	33.3	5003
20	The evolving landscape of biomarkers for checkpoint inhibitor immunotherapy. <i>Nature Reviews Cancer</i> , 2019 , 19, 133-150	31.3	996
19	Tumor and Microenvironment Evolution during Immunotherapy with Nivolumab. <i>Cell</i> , 2017 , 171, 934-943	30.16	831
18	The head and neck cancer immune landscape and its immunotherapeutic implications. <i>JCI Insight</i> , 2016 , 1, e89829	9.9	356
17	Genetic diversity of tumors with mismatch repair deficiency influences anti-PD-1 immunotherapy response. <i>Science</i> , 2019 , 364, 485-491	33.3	228
16	Immunogenic neoantigens derived from gene fusions stimulate T cell responses. <i>Nature Medicine</i> , 2019 , 25, 767-775	50.5	149
15	The role of neoantigens in response to immune checkpoint blockade. <i>International Immunology</i> , 2016 , 28, 411-9	4.9	88
14	Recurrent SERPINB3 and SERPINB4 mutations in patients who respond to anti-CTLA4 immunotherapy. <i>Nature Genetics</i> , 2016 , 48, 1327-1329	36.3	84
13	Multi-dimensional genomic analysis of myoepithelial carcinoma identifies prevalent oncogenic gene fusions. <i>Nature Communications</i> , 2017 , 8, 1197	17.4	46
12	Beta2-microglobulin signaling blockade inhibited androgen receptor axis and caused apoptosis in human prostate cancer cells. <i>Clinical Cancer Research</i> , 2008 , 14, 5341-7	12.9	37
11	Nuclear PRAS40 couples the Akt/mTORC1 signaling axis to the RPL11-HDM2-p53 nucleolar stress response pathway. <i>Oncogene</i> , 2015 , 34, 1487-98	9.2	35
10	Commensal bacteria stimulate antitumor responses via T cell cross-reactivity. <i>JCI Insight</i> , 2020 , 5,	9.9	30
9	ImmunoMap: A Bioinformatics Tool for T-cell Repertoire Analysis. <i>Cancer Immunology Research</i> , 2018 , 6, 151-162	12.5	25
8	Enabling systematic interrogation of protein-protein interactions in live cells with a versatile ultra-high-throughput biosensor platform. <i>Journal of Molecular Cell Biology</i> , 2016 , 8, 271-81	6.3	18
7	AKT1, LKB1, and YAP1 Revealed as MYC Interactors with NanoLuc-Based Protein-Fragment Complementation Assay. <i>Molecular Pharmacology</i> , 2017 , 91, 339-347	4.3	17
6	High-resolution genomic analysis: the tumor-immune interface comes into focus. <i>Genome Biology</i> , 2015 , 16, 65	18.3	3
5	ProteinProtein Interactions. <i>Springer Protocols</i> , 2008 , 463-494	0.3	3

4	Qa-1 Modulates Resistance to Anti-PD-1 Immune Checkpoint Blockade in Tumors with Defects in Antigen Processing. <i>Molecular Cancer Research</i> , 2021 , 19, 1076-1084	6.6	3
3	Time-Resolved Fluorescence Resonance Energy Transfer Technologies in HTS198-214		2
2	MEK Inhibitors in Lung Cancer-You Can Teach an Old Drug New Tricks. <i>Cancer Research</i> , 2019 , 79, 5699-5701	10.1	2
1	Immunogenomics 2019 , 99-110		