

Adam B Olshen

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

2,659
citations

840776

11
h-index

888059

17
g-index

22
all docs

22
docs citations

22
times ranked

7079
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular and phenotypic diversity of CBL-mutated juvenile myelomonocytic leukemia. <i>Haematologica</i> , 2022, 107, 178-186.	3.5	25
2	Discovering dominant tumor immune archetypes in a pan-cancer census. <i>Cell</i> , 2022, 185, 184-203.e19.	28.9	70
3	Ferrous iron-activatable drug conjugate achieves potent MAPK blockade in KRAS-driven tumors. <i>Journal of Experimental Medicine</i> , 2022, 219, .	8.5	15
4	COVIDNearTerm: A simple method to forecast COVID-19 hospitalizations. <i>Journal of Clinical and Translational Science</i> , 2022, 6, .	0.6	5
5	Prediction of evening fatigue severity in outpatients receiving chemotherapy: less may be more. <i>Fatigue: Biomedicine, Health and Behavior</i> , 2021, 9, 14-32.	1.9	8
6	Exploring the genetic and epigenetic origins of juvenile myelomonocytic leukemia using newborn screening samples. <i>Leukemia</i> , 2021, , .	7.2	9
7	International Consensus Definition of DNA Methylation Subgroups in Juvenile Myelomonocytic Leukemia. <i>Clinical Cancer Research</i> , 2021, 27, 158-168.	7.0	35
8	Next-Generation Sequencing of Retinoblastoma Identifies Pathogenic Alterations beyond RB1 Inactivation That Correlate with Aggressive Histopathologic Features. <i>Ophthalmology</i> , 2020, 127, 804-813.	5.2	39
9	Human pediatric B-cell acute lymphoblastic leukemias can be classified as B-1 or B-2-like based on a minimal transcriptional signature. <i>Experimental Hematology</i> , 2020, 90, 65-71.e1.	0.4	7
10	Genetic Alterations Precede DNA Methylation Changes in Juvenile Myelomonocytic Leukemia. <i>Blood</i> , 2020, 136, 19-20.	1.4	0
11	Identifying a potential biomarker for primary focal segmental glomerulosclerosis and its association with recurrence after transplantation. <i>Clinical Transplantation</i> , 2019, 33, e13487.	1.6	7
12	GENE-47. A 3D ATLAS TO EVALUATE THE SPATIAL PATTERNING OF GENETIC ALTERATIONS AND TUMOR CELL STATES IN GLIOMA. <i>Neuro-Oncology</i> , 2019, 21, vi107-vi108.	1.2	0
13	Novel Aggregate Deletion/Substitution/Addition Learning Algorithms for Recursive Partitioning. <i>Journal of Computational and Graphical Statistics</i> , 2018, 27, 146-156.	1.7	0
14	Glucocorticoids Paradoxically Induce Intrinsic Steroid Resistance through a STAT5-Mediated Survival Mechanism in T-Cell Acute Lymphoblastic Leukemia. <i>Blood</i> , 2018, 132, 913-913.	1.4	0
15	Establishment and characterization of an oral tongue squamous cell carcinoma cell line from a never-smoking patient. <i>Oral Oncology</i> , 2017, 69, 1-10.	1.5	8
16	Clonal expansion and epigenetic reprogramming following deletion or amplification of mutant IDH1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 10743-10748.	7.1	109
17	Quantitative proteomic analyses of mammary organoids reveals distinct signatures after exposure to environmental chemicals. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E1343-51.	7.1	45
18	Identifying recurrent mutations in cancer reveals widespread lineage diversity and mutational specificity. <i>Nature Biotechnology</i> , 2016, 34, 155-163.	17.5	634

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19	Assessing gene-level translational control from ribosome profiling. <i>Bioinformatics</i> , 2013, 29, 2995-3002.	4.1	79
20	Subtypes of pancreatic ductal adenocarcinoma and their differing responses to therapy. <i>Nature Medicine</i> , 2011, 17, 500-503.	30.7	1,460
21	Parent-specific copy number in paired tumor-normal studies using circular binary segmentation. <i>Bioinformatics</i> , 2011, 27, 2038-2046.	4.1	100