

Alex E Lash

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

25,153
citations

304602

22
h-index

642610

23
g-index

26
all docs

26
docs citations

26
times ranked

47483
citing authors

#	ARTICLE	IF	CITATIONS
1	Damaging de novo mutations diminish motor skills in children on the autism spectrum. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E1859-E1866.	3.3	49
2	Phase II Study of a Non-Platinum-Containing Doublet of Paclitaxel and Pemetrexed with Bevacizumab as Initial Therapy for Patients with Advanced Lung Adenocarcinomas. Journal of Thoracic Oncology, 2016, 11, 890-899.	0.5	4
3	Genome-wide prediction and functional characterization of the genetic basis of autism spectrum disorder. Nature Neuroscience, 2016, 19, 1454-1462.	7.1	359
4	Robust classification of protein variation using structural modelling and large-scale data integration. Nucleic Acids Research, 2016, 44, 2501-2513.	6.5	52
5	The Somatic Genomic Landscape of Glioblastoma. Cell, 2013, 155, 462-477.	13.5	3,979
6	Harnessing Technology to Improve Clinical Trials: Study of Real-Time Informatics to Collect Data, Toxicities, Image Response Assessments, and Patient-Reported Outcomes in a Phase II Clinical Trial. Journal of Clinical Oncology, 2013, 31, 2004-2009.	0.8	38
7	Germline mutations in BAP1 predispose to melanocytic tumors. Nature Genetics, 2011, 43, 1018-1021.	9.4	662
8	Integrative Genomic Profiling of Human Prostate Cancer. Cancer Cell, 2010, 18, 11-22.	7.7	3,151
9	Subtype-specific genomic alterations define new targets for soft-tissue sarcoma therapy. Nature Genetics, 2010, 42, 715-721.	9.4	642
10	Genomic and Biological Characterization of Exon 4 KRAS Mutations in Human Cancer. Cancer Research, 2010, 70, 5901-5911.	0.4	245
11	Comprehensive Genomic Analysis Reveals Clinically Relevant Molecular Distinctions between Thymic Carcinomas and Thymomas. Clinical Cancer Research, 2009, 15, 6790-6799.	3.2	176
12	Comparisons of tyrosine phosphorylated proteins in cells expressing lung cancer-specific alleles of EGFR and KRAS. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 14112-14117.	3.3	113
13	CancerGenes: a gene selection resource for cancer genome projects. Nucleic Acids Research, 2007, 35, D721-D726.	6.5	158
14	Mutational Analysis of EGFR and Related Signaling Pathway Genes in Lung Adenocarcinomas Identifies a Novel Somatic Kinase Domain Mutation in FGFR4. PLoS ONE, 2007, 2, e426.	1.1	77
15	Characterizing the cancer genome in lung adenocarcinoma. Nature, 2007, 450, 893-898.	13.7	1,020
16	A FoxO-Smad synexpression group in human keratinocytes. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 12747-12752.	3.3	221
17	Frequent Mutation of the PIK3CA Gene in Ovarian and Breast Cancers. Clinical Cancer Research, 2005, 11, 2875-2878.	3.2	397
18	NCBI GEO: mining millions of expression profiles--database and tools. Nucleic Acids Research, 2004, 33, D562-D566.	6.5	972

#	ARTICLE	IF	CITATIONS
19	Database resources of the National Center for Biotechnology. <i>Nucleic Acids Research</i> , 2003, 31, 28-33.	6.5	879
20	Gene Expression Omnibus: NCBI gene expression and hybridization array data repository. <i>Nucleic Acids Research</i> , 2002, 30, 207-210.	6.5	10,953
21	A systematic, high-resolution linkage of the cytogenetic and physical maps of the human genome. <i>Nature Genetics</i> , 2000, 24, 339-340.	9.4	52
22	Analysis of human transcriptomes. <i>Nature Genetics</i> , 1999, 23, 387-388.	9.4	719
23	Monoclonal Origin of Multicentric Kaposi's Sarcoma Lesions. <i>New England Journal of Medicine</i> , 1997, 336, 988-993.	13.9	226