## David R Jones

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2589434/publications.pdf

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840776 1281871 5,224 11 11 11 citations h-index g-index papers 12 12 12 11058 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Landscape of somatic mutations in 560 breast cancer whole-genome sequences. Nature, 2016, 534, 47-54.	27.8	1,760
2	Mutational Processes Molding the Genomes of 21 Breast Cancers. Cell, 2012, 149, 979-993.	28.9	1,673
3	Analysis of the genetic phylogeny of multifocal prostate cancer identifies multiple independent clonal expansions in neoplastic and morphologically normal prostate tissue. Nature Genetics, 2015, 47, 367-372.	21.4	380
4	Characterizing Mutational Signatures in Human Cancer Cell Lines Reveals Episodic APOBEC Mutagenesis. Cell, 2019, 176, 1282-1294.e20.	28.9	298
5	A comprehensive assessment of somatic mutation detection in cancer using whole-genome sequencing. Nature Communications, 2015, 6, 10001.	12.8	266
6	Somatic mutation landscapes at single-molecule resolution. Nature, 2021, 593, 405-410.	27.8	254
7	Somatic mutation rates scale with lifespan across mammals. Nature, 2022, 604, 517-524.	27.8	211
8	cgpCaVEManWrapper: Simple Execution of CaVEMan in Order to Detect Somatic Single Nucleotide Variants in NGS Data. Current Protocols in Bioinformatics, 2016, 56, 15.10.1-15.10.18.	25.8	155
9	ascatNgs: Identifying Somatically Acquired Copyâ€Number Alterations from Wholeâ€Genome Sequencing Data. Current Protocols in Bioinformatics, 2016, 56, 15.9.1-15.9.17.	25.8	111
10	Mutational signatures in esophageal squamous cell carcinoma from eight countries with varying incidence. Nature Genetics, 2021, 53, 1553-1563.	21.4	71
11	Appraising the relevance of DNA copy number loss and gain in prostate cancer using whole genome DNA sequence data. PLoS Genetics, 2017, 13, e1007001.	3.5	34