

The Cancer Genome Atlas Pan-Cancer analysis project

Nature Genetics

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Citation Report

#	ARTICLE	IF	CITATIONS
1	La formalisation de la GRH dans une PME comme enjeu d'une certification RSE. Revue De Gestion Des Ressources Humaines, 2012, NÂ° 83, 20-30.	0.1	8
2	Comparisons across cancers. Nature, 2013, 502, 306-307.	13.7	10
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5	Comprehensive identification of mutational cancer driver genes across 12 tumor types. Scientific Reports, 2013, 3, 2650.	1.6	437
6	Exploring TCGA Pan-Cancer Data at the UCSC Cancer Genomics Browser. Scientific Reports, 2013, 3, 2652.	1.6	235
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8	Localized structural frustration for evaluating the impact of sequence variants. Nucleic Acids Research, 2013, 44, 10062-10073.	6.5	13
9	Molecular Pathology of Lung Cancer: Current Status and Future Directions. Tuberculosis and Respiratory Diseases, 2014, 77, 49.	0.7	12
10	A Pan-Cancer Analysis of Transcriptome Changes Associated with Somatic Mutations in U2AF1 Reveals Commonly Altered Splicing Events. PLoS ONE, 2014, 9, e87361.	1.1	168
11	A Robust and Accurate Method for Feature Selection and Prioritization from Multi-Class OMICs Data. PLoS ONE, 2014, 9, e107801.	1.1	32
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20	SummonChimera infers integrated viral genomes with nucleotide precision from NGS data. <i>BMC Bioinformatics</i> , 2014, 15, 348.	1.2	16
21	ExpressionData - A public resource of high quality curated datasets representing gene expression across anatomy, development and experimental conditions. <i>BioData Mining</i> , 2014, 7, 18.	2.2	22
22	Global optimization of somatic variant identification in cancer genomes with a global community challenge. <i>Nature Genetics</i> , 2014, 46, 318-319.	9.4	42
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24	Identifying Cancer Subtypes based on Somatic Mutation Profile. , 2014, , .		7
25	Genotoxic Anti-Cancer Agents and Their Relationship to DNA Damage, Mitosis, and Checkpoint Adaptation in Proliferating Cancer Cells. <i>International Journal of Molecular Sciences</i> , 2014, 15, 3403-3431.	1.8	155
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