

# Targeted exploration and analysis of large cross-platform compendia

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Citation Report

#	ARTICLE	IF	CITATIONS
1	PADPIN: protein-protein interaction networks of angiogenesis, arteriogenesis, and inflammation in peripheral arterial disease. <i>Physiological Genomics</i> , 2015, 47, 331-343.	2.3	12
2	Implications of pleiotropy: challenges and opportunities for mining Big Data in biomedicine. <i>Frontiers in Genetics</i> , 2015, 6, 229.	2.3	41
3	Human transient receptor potential (TRP) channel expression profiling in carcinogenesis. <i>International Journal of Developmental Biology</i> , 2015, 59, 399-406.	0.6	26
4	Big Data in <i>Caenorhabditis elegans</i> : <i>quo vadis</i> ?. <i>Molecular Biology of the Cell</i> , 2015, 26, 3909-3914.	2.1	8
5	Cancer: smoother journeys for molecular data. <i>Nature Methods</i> , 2015, 12, 299-302.	19.0	3
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18	Use of Structural Properties of Underlying Graphs in Pathway Enrichment Analysis of Genomic Data. , 2017, , .		3

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19	Selfing in Haploid Plants and Efficacy of Selection: Codon Usage Bias in the Model Moss <i>Physcomitrella patens</i> . <i>Genome Biology and Evolution</i> , 2017, 9, 1528-1546.	2.5	21
20	IFN $\beta$ -Dependent Tissue-Immune Homeostasis Is Co-opted in the Tumor Microenvironment. <i>Cell</i> , 2017, 170, 127-141.e15.	28.9	140
21	Bioinformatic Analysis of Circadian Expression of Oncogenes and Tumor Suppressor Genes. <i>Bioinformatics and Biology Insights</i> , 2017, 11, 1177932217746991.	2.0	3
22	MicroRNA Networks in Breast Cancer Cells. <i>Methods in Molecular Biology</i> , 2018, 1711, 55-81.	0.9	15
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