

Genome sequencing of normal cells reveals development processes

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Citation Report

#	ARTICLE	IF	CITATIONS
1	From regenerative dentistry to regenerative medicine: progress, challenges, and potential applications of oral stem cells. <i>Stem Cells and Cloning: Advances and Applications</i> , 2014, 7, 89.	2.3	62
2	Postzygotic single-nucleotide mosaicism in whole-genome sequences of clinically unremarkable individuals. <i>Cell Research</i> , 2014, 24, 1311-1327.	5.7	54
3	Chromosomale Mosaik in der klinischen Zytogenetik. <i>Medizinische Genetik</i> , 2014, 26, 302-308.	0.1	3
4	Quantitative Single-Cell Approaches to Stem Cell Research. <i>Cell Stem Cell</i> , 2014, 15, 546-558.	5.2	112
5	The Role of Constitutional Copy Number Variants in Breast Cancer. <i>Microarrays (Basel, Switzerland)</i> , 2015, 4, 407-423.	1.4	9
6	Analyzing Fluctuating Asymmetry with Geometric Morphometrics: Concepts, Methods, and Applications. <i>Symmetry</i> , 2015, 7, 843-934.	1.1	295
7	Genetic Mosaics and the Germ Line Lineage. <i>Genes</i> , 2015, 6, 216-237.	1.0	28
8	A new conceptual framework for investigating complex genetic disease. <i>Frontiers in Genetics</i> , 2015, 6, 327.	1.1	6
9	High burden and pervasive positive selection of somatic mutations in normal human skin. <i>Science</i> , 2015, 348, 880-886.	6.0	1,431
10	The Technology and Biology of Single-Cell RNA Sequencing. <i>Molecular Cell</i> , 2015, 58, 610-620.	4.5	1,014
11	From Mutational Mechanisms in Single Cells to Mutational Patterns in Cancer Genomes. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2015, 80, 117-137.	2.0	11
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17	Using tumour phylogenetics to identify the roots of metastasis in humans. <i>Nature Reviews Clinical Oncology</i> , 2015, 12, 258-272.	12.5	122
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20	Somatic mosaicism: implications for disease and transmission genetics. <i>Trends in Genetics</i> , 2015, 31, 382-392.	2.9	234
21	PhyloWGS: Reconstructing subclonal composition and evolution from whole-genome sequencing of tumors. <i>Genome Biology</i> , 2015, 16, 35.	3.8	351
22	On the identification of low allele frequency mosaic mutations in the brains of Alzheimer's disease patients. <i>Alzheimer's and Dementia</i> , 2015, 11, 1265-1276.	0.4	57
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