James E Hadfield

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	The genomic and transcriptomic architecture of 2,000 breast tumours reveals novel subgroups. Nature, 2012, 486, 346-352.	27.8	4,708
2	Non-invasive analysis of acquired resistance to cancer therapy by sequencing of plasma DNA. Nature, 2013, 497, 108-112.	27.8	1,443
3	RNA sequencing: the teenage years. Nature Reviews Genetics, 2019, 20, 631-656.	16.3	1,192
4	Noninvasive Identification and Monitoring of Cancer Mutations by Targeted Deep Sequencing of Plasma DNA. Science Translational Medicine, 2012, 4, 136ra68.	12.4	1,086
5	Systematic comparison of microarray profiling, real-time PCR, and next-generation sequencing technologies for measuring differential microRNA expression. Rna, 2010, 16, 991-1006.	3.5	588
6	The androgen receptor fuels prostate cancer by regulating central metabolism and biosynthesis. EMBO Journal, 2011, 30, 2719-2733.	7.8	530
7	Progesterone receptor modulates ERα action in breast cancer. Nature, 2015, 523, 313-317.	27.8	504
8	ChIP-seq: Using high-throughput sequencing to discover protein–DNA interactions. Methods, 2009, 48, 240-248.	3.8	455
9	Differentiation dynamics of mammary epithelial cells revealed by single-cell RNAÂsequencing. Nature Communications, 2017, 8, 2128.	12.8	234
10	Germline pathogenic variants in PALB2 and other cancer-predisposing genes in families with hereditary diffuse gastric cancer without CDH1 mutation: a whole-exome sequencing study. The Lancet Gastroenterology and Hepatology, 2018, 3, 489-498.	8.1	87
11	The pitfalls of platform comparison: DNA copy number array technologies assessed. BMC Genomics, 2009, 10, 588.	2.8	80
12	Transducin-like enhancer protein 1 mediates estrogen receptor binding and transcriptional activity in breast cancer cells. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 2748-2753.	7.1	47
13	Introduction to miRNA Profiling Technologies and Cross-Platform Comparison. Methods in Molecular Biology, 2012, 822, 19-31.	0.9	34
14	The role of high-throughput technologies in clinical cancer genomics. Expert Review of Molecular Diagnostics, 2013, 13, 167-181.	3.1	26
15	Independence of HIF1a and androgen signaling pathways in prostate cancer. BMC Cancer, 2020, 20, 469.	2.6	25
16	A profusion of confusion in NGS methods naming. Nature Methods, 2018, 15, 7-8.	19.0	9
17	Large-scale mutagenesis directed at specific chromosomes in wheat. Genome, 2001, 44, 45-49.	2.0	8
18	A differential PCR assay for the detection of c-erbB 2 amplification used in a prospective study of breast cancer Journal of Clinical Pathology, 1997, 50, 254-256.	1.9	7

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19	The cost of reducing starting RNA quantity for Illumina BeadArrays: A bead-level dilution experiment. BMC Genomics, 2010, 11, 540.	2.8	4
20	Characterization of DNA-Protein Interactions: Design and Analysis of ChIP-Seq Experiments. , 2016, , 223-260		3

223-260.