A Pan-Cancer Compendium of Genes Deregulated by So More Than 1,400 Cases

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Citation Report

#	Article	IF	CITATIONS
1	The clinical applications of The Cancer Genome Atlas project for bladder cancer. Expert Review of Anticancer Therapy, 2018, 18, 973-980.	1.1	12
2	Technologies to study spatial genome organization: beyond 3C. Briefings in Functional Genomics, 2019, 18, 395-401.	1.3	4
3	Global impact of somatic structural variation on the DNA methylome of human cancers. Genome Biology, 2019, 20, 209.	3.8	40
4	Regional perturbation of gene transcription is associated with intrachromosomal rearrangements and gene fusion transcripts in high grade ovarian cancer. Scientific Reports, 2019, 9, 3590.	1.6	8
5	CDK12 phosphorylates 4E-BP1 to enable mTORC1-dependent translation and mitotic genome stability. Genes and Development, 2019, 33, 418-435.	2.7	50
6	Large-scale compound screens and pharmacogenomic interactions in cancer. Current Opinion in Genetics and Development, 2019, 54, 12-16.	1.5	6
7	Cancer mutational burden is shaped by G4 DNA, replication stress and mitochondrial dysfunction. Progress in Biophysics and Molecular Biology, 2019, 147, 47-61.	1.4	35
8	<scp>CDK</scp> 12 drives breast tumor initiation and trastuzumab resistance via <scp>WNT</scp> and <scp>IRS</scp> 1â€ErbB― <scp>PI</scp> 3K signaling. EMBO Reports, 2019, 20, e48058.	2.0	76
9	Telomere Maintenance in Pediatric Cancer. International Journal of Molecular Sciences, 2019, 20, 5836.	1.8	10
10	Therapeutic targeting of transcriptional cyclin-dependent kinases. Transcription, 2019, 10, 118-136.	1.7	78
11	Somatic gene mutation signatures predict cancer type and prognosis in multiple cancers with pan-cancer 1000 gene panel. Cancer Letters, 2020, 470, 181-190.	3.2	29
12	Epigenomic technologies for precision oncology. Seminars in Cancer Biology, 2022, 84, 60-68.	4.3	6
13	SV-HotSpot: detection and visualization of hotspots targeted by structural variants associated with gene expression. Scientific Reports, 2020, 10, 15890.	1.6	3
14	DNA mismatch repair promotes APOBEC3-mediated diffuse hypermutation in human cancers. Nature Genetics, 2020, 52, 958-968.	9.4	76
15	Revealing the Impact of Structural Variants in Multiple Myeloma. Blood Cancer Discovery, 2020, 1, 258-273.	2.6	81
16	A Practical Guide for Structural Variation Detection in the Human Genome. Current Protocols in Human Genetics, 2020, 107, e103.	3.5	12
17	CDK12 and PAK2 as novel therapeutic targets for human gastric cancer. Theranostics, 2020, 10, 6201-6215.	4.6	44
18	Cancer-associated mutations in endometriosis: shedding light on the pathogenesis and pathophysiology. Human Reproduction Update, 2020, 26, 423-449.	5.2	57

#	ARTICLE	IF	CITATIONS
19	Genome rearrangements associated with aberrant telomere maintenance. Current Opinion in Genetics and Development, 2020, 60, 31-40.	1.5	17
20	Dysregulation of cancer genes by recurrent intergenic fusions. Genome Biology, 2020, 21, 166.	3.8	22
21	Three-dimensional chromatin in disease: What holds us together and what drives us apart?. Current Opinion in Cell Biology, 2020, 64, 1-9.	2.6	51
22	Larotrectinib in patients with TRK fusion-positive solid tumours: a pooled analysis of three phase $1/2$ clinical trials. Lancet Oncology, The, 2020, 21, 531-540.	5.1	608
23	PI4KIIIβ is a therapeutic target in chromosome $1q\hat{a}\in$ amplified lung adenocarcinoma. Science Translational Medicine, 2020, 12, .	5.8	41
24	High-coverage whole-genome analysis of 1220 cancers reveals hundreds of genes deregulated by rearrangement-mediated cis-regulatory alterations. Nature Communications, 2020, 11, 736.	5.8	50
25	The mutational landscape of normal human endometrial epithelium. Nature, 2020, 580, 640-646.	13.7	338
26	Immuno-Oncology Biomarkers for Personalized Immunotherapy in Breast Cancer. Frontiers in Cell and Developmental Biology, 2020, 8, 162.	1.8	21
27	Detection of somatic structural variants from short-read next-generation sequencing data. Briefings in Bioinformatics, 2021, 22, .	3.2	34
28	Integrating proteomics into precision oncology. International Journal of Cancer, 2021, 148, 1438-1451.	2.3	15
29	Target Validationâ€"Prosecuting the Target. , 2021, , .		0
30	A pediatric brain tumor atlas of genes deregulated by somatic genomic rearrangement. Nature Communications, 2021, 12, 937.	5.8	23
31	Systematic identification of non-coding somatic single nucleotide variants associated with altered transcription and DNA methylation in adult and pediatric cancers. NAR Cancer, 2021, 3, zcab001.	1.6	7
32	Cancer regulatory variation. Current Opinion in Genetics and Development, 2021, 66, 41-49.	1.5	6
33	Loss of $TGF\hat{l}^2$ signaling increases alternative end-joining DNA repair that sensitizes to genotoxic therapies across cancer types. Science Translational Medicine, 2021, 13, .	5.8	33
34	SVExpress: identifying gene features altered recurrently in expression with nearby structural variant breakpoints. BMC Bioinformatics, 2021, 22, 135.	1.2	3
35	Gene Fusions Create Partner and Collateral Dependencies Essential to Cancer Cell Survival. Cancer Research, 2021, 81, 3971-3984.	0.4	11
36	Breast Cancer and Anaesthesia: Genetic Influence. International Journal of Molecular Sciences, 2021, 22, 7653.	1.8	8

#	ARTICLE	IF	Citations
37	Pan-cancer analysis of pathway-based gene expression pattern at the individual level reveals biomarkers of clinical prognosis. Cell Reports Methods, 2021, 1, 100050.	1.4	10
38	Impact of Disease Evolution on Efficacy Outcomes From Larotrectinib in Patients With Locally Advanced or Metastatic Tropomyosin Receptor Kinase Fusion–Positive Solid Tumors. JCO Precision Oncology, 2021, 5, 1458-1465.	1.5	4
40	svMIL: predicting the pathogenic effect of TAD boundary-disrupting somatic structural variants throughÂmultiple instance learning. Bioinformatics, 2020, 36, i692-i699.	1.8	4
44	Breaking point: the genesis and impact of structural variation in tumours. F1000Research, 2018, 7, 1814.	0.8	7
45	Characterization of Structural Variations in the Context of 3D Chromatin Structure. Molecules and Cells, 2019, 42, 512-522.	1.0	12
46	Tumor suppressor genes and allele-specific expression: mechanisms and significance. Oncotarget, 2020, 11, 462-479.	0.8	5
48	Pan-Cancer Analysis of Pathway-Based Gene Expression Pattern at the Individual Level Reveals Novel Biomarkers of Clinical Prognosis. SSRN Electronic Journal, 0, , .	0.4	0
49	QNBC Is Associated with High Genomic Instability Characterized by Copy Number Alterations and miRNA Deregulation. International Journal of Molecular Sciences, 2021, 22, 11548.	1.8	10
51	Anaesthesia and cancer: can anaesthetic drugs modify gene expression?. Ecancermedicalscience, 2020, 14, 1080.	0.6	2
52	Rearrangement-mediated cis-regulatory alterations in advanced patient tumors reveal interactions with therapy. Cell Reports, 2021, 37, 110023.	2.9	8
53	Pan-Cancer DNA Methylation Analysis and Tumor Origin Identification of Carcinoma of Unknown Primary Site Based on Multi-Omics. Frontiers in Genetics, 2021, 12, 798748.	1.1	2
54	Towards accurate and reliable resolution of structural variants for clinical diagnosis. Genome Biology, 2022, 23, 68.	3.8	34
55	A nine-month-old boy with regression of milestones and severe constipation: an unusual case of a large spinal NTRK1 fusion pilocytic astrocytoma. Child's Nervous System, 0, , .	0.6	0
56	Modeling tissue-specific breakpoint proximity of structural variations from whole-genomes to identify cancer drivers. Nature Communications, 2022, 13, .	5.8	1
57	Can abnormal chromatin folding cause high-penetrance cancer predisposition?. Physiological Genomics, 2022, 54, 380-388.	1.0	0
58	Clinical proteomics towards multiomics in cancer. Mass Spectrometry Reviews, 0, , .	2.8	6
59	Survival Outcomes of Patients With Tropomyosin Receptor Kinase Fusion-Positive Cancer Receiving Larotrectinib Versus Standard of Care: A Matching-Adjusted Indirect Comparison Using Real-World Data. JCO Precision Oncology, 2023, , .	1.5	1
60	Long-read sequencing identifies novel structural variations in colorectal cancer. PLoS Genetics, 2023, 19, e1010514.	1.5	7

ARTICLE IF CITATIONS

Function and Evolution of the Loop Extrusion Machinery in Animals. International Journal of Molecular Sciences, 2023, 24, 5017.

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