## Manjie Xing

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10545861/publications.pdf

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17	880	11	18
papers	citations	h-index	g-index
19	19	19	1910 citing authors
all docs	docs citations	times ranked	

#	Article	IF	CITATIONS
1	Genomic and Epigenomic Profiling of High-Risk Intestinal Metaplasia Reveals Molecular Determinants of Progression to Gastric Cancer. Cancer Cell, 2018, 33, 137-150.e5.	16.8	175
2	Epigenomic profiling of primary gastric adenocarcinoma reveals super-enhancer heterogeneity. Nature Communications, 2016, 7, 12983.	12.8	123
3	<i>VHL</i> Deficiency Drives Enhancer Activation of Oncogenes in Clear Cell Renal Cell Carcinoma. Cancer Discovery, 2017, 7, 1284-1305.	9.4	111
4	Mutation hotspots at CTCF binding sites coupled to chromosomal instability in gastrointestinal cancers. Nature Communications, 2018, 9, 1520.	12.8	109
5	Nanoscale chromatin profiling of gastric adenocarcinoma reveals cancer-associated cryptic promoters and somatically acquired regulatory elements. Nature Communications, 2014, 5, 4361.	12.8	72
6	Cross-species chromatin interactions drive transcriptional rewiring in Epstein–Barr virus–positive gastric adenocarcinoma. Nature Genetics, 2020, 52, 919-930.	21.4	65
7	Epigenomic Promoter Alterations Amplify Gene Isoform and Immunogenic Diversity in Gastric Adenocarcinoma. Cancer Discovery, 2017, 7, 630-651.	9.4	48
8	HoxC5 and miR-615-3p target newly evolved genomic regions to repress hTERT and inhibit tumorigenesis. Nature Communications, 2018, 9, 100.	12.8	38
9	Integrated paired-end enhancer profiling and whole-genome sequencing reveals recurrent <i>CCNE1</i> and <i>IGF2</i> enhancer hijacking in primary gastric adenocarcinoma. Gut, 2020, 69, 1039-1052.	12.1	36
10	$\langle i \rangle$ HNF4α $\langle i \rangle$ pathway mapping identifies wild-type $\langle i \rangle$ IDH1 $\langle i \rangle$ as a targetable metabolic node in gastric cancer. Gut, 2020, 69, 231-242.	12.1	27
11	Epigenetic promoter alterations in GI tumour immune-editing and resistance to immune checkpoint inhibition. Gut, 2022, 71, 1277-1288.	12.1	23
12	Genomic and epigenomic EBF1 alterations modulate TERT expression in gastric cancer. Journal of Clinical Investigation, 2020, 130, 3005-3020.	8.2	12
13	Highly recurrent CBS epimutations in gastric cancer CpG island methylator phenotypes and inflammation. Genome Biology, 2021, 22, 167.	8.8	10
14	Epigenomic Consequences of Coding and Noncoding Driver Mutations. Trends in Cancer, 2016, 2, 585-605.	7.4	8
15	Integrative epigenomic and high-throughput functional enhancer profiling reveals determinants of enhancer heterogeneity in gastric cancer. Genome Medicine, 2021, 13, 158.	8.2	7
16	Chromatin Rewiring by Mismatch Repair Protein MSH2 Alters Cell Adhesion Pathways and Sensitivity to BET Inhibition in Gastric Cancer. Cancer Research, 2022, 82, 2538-2551.	0.9	7
17	Regulatory enhancer profiling of mesenchymal-type gastric cancer reveals subtype-specific epigenomic landscapes and targetable vulnerabilities. Gut, 2023, 72, 226-241.	12.1	6