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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Scalable whole-exome sequencing of cell-free DNA reveals high concordance with metastatic tumors. Nature Communications, 2017, 8, 1324.	12.8	584
2	Structural Alterations Driving Castration-Resistant Prostate Cancer Revealed by Linked-Read Genome Sequencing. Cell, 2018, 174, 433-447.e19.	28.9	258
3	Association of Cell-Free DNA Tumor Fraction and Somatic Copy Number Alterations With Survival in Metastatic Triple-Negative Breast Cancer. Journal of Clinical Oncology, 2018, 36, 543-553.	1.6	162
4	Sensitive Detection of Minimal Residual Disease in Patients Treated for Early-Stage Breast Cancer. Clinical Cancer Research, 2020, 26, 2556-2564.	7.0	109
5	Tumor fraction in cell-free DNA as a biomarker in prostate cancer. JCI Insight, 2018, 3, .	5.0	94
6	Massively parallel enrichment of low-frequency alleles enables duplex sequencing at low depth. Nature Biomedical Engineering, 2022, 6, 257-266.	22.5	32
7	Liquid biopsy detection of genomic alterations in pediatric brain tumors from cell-free DNA in peripheral blood, CSF, and urine. Neuro-Oncology, 2022, 24, 1352-1363.	1.2	29
8	Modeling clonal structure over narrow time frames via circulating tumor DNA in metastatic breast cancer. Genome Medicine, 2021, 13, 89.	8.2	10
9	Duplex-Repair enables highly accurate sequencing, despite DNA damage. Nucleic Acids Research, 2022, 50, e1-e1.	14.5	10
10	Circulating Cell-Free DNA as Biomarker of Taxane Resistance in Metastatic Castration-Resistant Prostate Cancer. Cancers, 2021, 13, 4055.	3.7	1