Giuseppe Rospo

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Clonal evolution and resistance to EGFR blockade in the blood of colorectal cancer patients. Nature Medicine, 2015, 21, 795-801.	30.7	809
2	Inactivation of DNA repair triggers neoantigen generation and impairs tumour growth. Nature, 2017, 552, 116-120.	27.8	480
3	Acquired Resistance to the TRK Inhibitor Entrectinib in Colorectal Cancer. Cancer Discovery, 2016, 6, 36-44.	9.4	258
4	Radiologic and Genomic Evolution of Individual Metastases during HER2 Blockade in Colorectal Cancer. Cancer Cell, 2018, 34, 148-162.e7.	16.8	129
5	Efficacy of Sym004 in Patients With Metastatic Colorectal Cancer With Acquired Resistance to Anti-EGFR Therapy and Molecularly Selected by Circulating Tumor DNA Analyses. JAMA Oncology, 2018, 4, e175245.	7.1	98
6	The Clinical Impact of the Genomic Landscape of Mismatch Repair–Deficient Cancers. Cancer Discovery, 2018, 8, 1518-1528.	9.4	77
7	A Subset of Colorectal Cancers with Cross-Sensitivity to Olaparib and Oxaliplatin. Clinical Cancer Research, 2020, 26, 1372-1384.	7.0	66
8	Werner Helicase Is a Synthetic-Lethal Vulnerability in Mismatch Repair–Deficient Colorectal Cancer Refractory to Targeted Therapies, Chemotherapy, and Immunotherapy. Cancer Discovery, 2021, 11, 1923-1937.	9.4	48
9	Evolving neoantigen profiles in colorectal cancers with DNA repair defects. Genome Medicine, 2019, 11, 42.	8.2	42
10	CD4 T Cell–Dependent Rejection of Beta-2 Microglobulin Null Mismatch Repair–Deficient Tumors. Cancer Discovery, 2021, 11, 1844-1859.	9.4	37
11	Emergence of MET hyper-amplification at progression to MET and BRAF inhibition in colorectal cancer. British Journal of Cancer, 2017, 117, 347-352.	6.4	31
12	A Genomic Analysis Workflow for Colorectal Cancer Precision Oncology. Clinical Colorectal Cancer, 2019, 18, 91-101.e3.	2.3	29
13	Two main mutational processes operate in the absence of DNA mismatch repair. DNA Repair, 2020, 89, 102827.	2.8	19
14	Reliance upon ancestral mutations is maintained in colorectal cancers that heterogeneously evolve during targeted therapies. Nature Communications, 2018, 9, 2287.	12.8	18
15	DeCoClu: Density consensus clustering approach for public transport data. Information Sciences, 2016, 328, 378-388.	6.9	16
16	TRKA expression and <i>NTRK1</i> gene copy number across solid tumours. Journal of Clinical Pathology, 2018, 71, 926-931.	2.0	12
17	Tracking colorectal cancer evolution in time and space. Annals of Oncology, 2017, 28, 1163-1165.	1.2	5