

# Giovanni Crisafulli

## List of Publications by Year in descending order

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Version: 2024-02-01

39  
papers

3,482  
citations

361413

20  
h-index

526287

27  
g-index

41  
all docs

41  
docs citations

41  
times ranked

6336  
citing authors

#	ARTICLE	IF	CITATIONS
1	Clonal evolution and resistance to EGFR blockade in the blood of colorectal cancer patients. <i>Nature Medicine</i> , 2015, 21, 795-801.	30.7	809
2	Inactivation of DNA repair triggers neoantigen generation and impairs tumour growth. <i>Nature</i> , 2017, 552, 116-120.	27.8	480
3	Tumor Heterogeneity and Lesion-Specific Response to Targeted Therapy in Colorectal Cancer. <i>Cancer Discovery</i> , 2016, 6, 147-153.	9.4	338
4	Adaptive mutability of colorectal cancers in response to targeted therapies. <i>Science</i> , 2019, 366, 1473-1480.	12.6	290
5	Acquired Resistance to the TRK Inhibitor Entrectinib in Colorectal Cancer. <i>Cancer Discovery</i> , 2016, 6, 36-44.	9.4	258
6	Emergence of Multiple <i>EGFR</i> Extracellular Mutations during Cetuximab Treatment in Colorectal Cancer. <i>Clinical Cancer Research</i> , 2015, 21, 2157-2166.	7.0	227
7	Acquired RAS or EGFR mutations and duration of response to EGFR blockade in colorectal cancer. <i>Nature Communications</i> , 2016, 7, 13665.	12.8	170
8	Radiologic and Genomic Evolution of Individual Metastases during HER2 Blockade in Colorectal Cancer. <i>Cancer Cell</i> , 2018, 34, 148-162.e7.	16.8	129
9	Molecular Landscape of Acquired Resistance to Targeted Therapy Combinations in <i>BRAF</i> -Mutant Colorectal Cancer. <i>Cancer Research</i> , 2016, 76, 4504-4515.	0.9	91
10	Genotyping tumour DNA in cerebrospinal fluid and plasma of a HER2-positive breast cancer patient with brain metastases. <i>ESMO Open</i> , 2017, 2, e000253.	4.5	56
11	Loss of AXIN1 drives acquired resistance to <i>WNT</i> pathway blockade in colorectal cancer cells carrying <i>RSPO</i> 3 fusions. <i>EMBO Molecular Medicine</i> , 2017, 9, 293-303.	6.9	54
12	Werner Helicase Is a Synthetic-Lethal Vulnerability in Mismatch Repair-Deficient Colorectal Cancer Refractory to Targeted Therapies, Chemotherapy, and Immunotherapy. <i>Cancer Discovery</i> , 2021, 11, 1923-1937.	9.4	48
13	Temozolomide Treatment Alters Mismatch Repair and Boosts Mutational Burden in Tumor and Blood of Colorectal Cancer Patients. <i>Cancer Discovery</i> , 2022, 12, 1656-1675.	9.4	48
14	Genetic Evolution of Glioblastoma Stem-Like Cells From Primary to Recurrent Tumor. <i>Stem Cells</i> , 2017, 35, 2218-2228.	3.2	47
15	Liquid biopsies to monitor and direct cancer treatment in colorectal cancer. <i>British Journal of Cancer</i> , 2022, 127, 394-407.	6.4	41
16	Tracking <i>CAD-ALK</i> gene rearrangement in urine and blood of a colorectal cancer patient treated with an ALK inhibitor. <i>Annals of Oncology</i> , 2017, 28, 1302-1308.	1.2	32
17	Emergence of MET hyper-amplification at progression to MET and BRAF inhibition in colorectal cancer. <i>British Journal of Cancer</i> , 2017, 117, 347-352.	6.4	31
18	A Genomic Analysis Workflow for Colorectal Cancer Precision Oncology. <i>Clinical Colorectal Cancer</i> , 2019, 18, 91-101.e3.	2.3	29

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19	Whole exome sequencing analysis of urine trans-renal tumour DNA in metastatic colorectal cancer patients. <i>ESMO Open</i> , 2019, 4, e000572.	4.5	27
20	A modified fluctuation-test framework characterizes the population dynamics and mutation rate of colorectal cancer persister cells. <i>Nature Genetics</i> , 2022, 54, 976-984.	21.4	23
21	Pembrolizumab in MMR-proficient metastatic colorectal cancer pharmacologically primed to trigger dynamic hypermutation status: The ARETHUSA trial.. <i>Journal of Clinical Oncology</i> , 2019, 37, TPS2659-TPS2659.	1.6	10
22	An extended multi-locus molecular typing schema for <i>Streptococcus pneumoniae</i> demonstrates that a limited number of capsular switch events is responsible for serotype heterogeneity of closely related strains from different countries. <i>Infection, Genetics and Evolution</i> , 2013, 13, 151-161.	2.3	9
23	Sequence Analysis of 96 Genomic Regions Identifies Distinct Evolutionary Lineages within CC156, the Largest <i>Streptococcus pneumoniae</i> Clonal Complex in the MLST Database. <i>PLoS ONE</i> , 2013, 8, e61003.	2.5	8
24	Tracking colorectal cancer evolution in time and space. <i>Annals of Oncology</i> , 2017, 28, 1163-1165.	1.2	5
25	Abstract 5723: Inactivation of DNA repair triggers neoantigen generation and impairs tumor growth. <i>Cancer Research</i> , 2018, 78, 5723-5723.	0.9	5
26	Assessment of HER2 ( <i>ERBB2</i> ) amplification (HER2amp) using blood-based circulating tumor DNA (ctDNA) next generation sequencing (NGS) and correlation with tissue-based testing in metastatic colorectal cancer (mCRC).. <i>Journal of Clinical Oncology</i> , 2021, 39, 3589-3589.	1.6	2
27	Abstract PR01: Acquisition of resistance to anti-EGFR therapy drives genomic heterogeneity and lesion-specific responses in colorectal cancer. , 2015, , .		1
28	Abstract 878: Tumor heterogeneity and lesion-specific response to targeted therapy in colorectal cancer. , 2016, , .		1
29	Abstract 2848: Radiographic and genomic evolution of individual metastases during HER2 blockade in colorectal cancer. , 2018, , .		1
30	Raise and decline of KRAS mutant clones in colorectal cancers (CRCs) treated with multiple rounds of anti-EGFR antibodies.. <i>Journal of Clinical Oncology</i> , 2015, 33, 11073-11073.	1.6	0
31	Abstract 616: Blood-based molecular landscapes of resistance to EGFR blockade in colorectal cancer patients. , 2015, , .		0
32	Abstract 3588: Emergence of multiple EGFR extracellular mutations during cetuximab treatment in colorectal cancer. , 2015, , .		0
33	Abstract 3834: Tracking CAD-ALK gene translocation in urine and plasma of a colorectal cancer patient treated with ALK blockade. , 2017, , .		0
34	Abstract 2913: Emergence of RAS or EGFR mutant clones affects duration of response to EGFR blockade in colorectal cancers. , 2017, , .		0
35	Abstract 2743: Accumulation of predicted neoantigens by MMR deficiency triggered by temozolomide treatment of human colorectal cancer. , 2018, , .		0
36	Abstract B069: Temozolomide drives mismatch repair deficiency and fosters neoantigen generation in tumor cells. , 2019, , .		0

#	ARTICLE	IF	CITATIONS
37	Abstract A120: Adaptive mutability of colorectal cancers in response to targeted therapies. , 2019, , .		0
38	Abstract B11: Whole-exome sequencing analysis of urine transrenal tumor DNA in metastatic colorectal cancer patients. , 2020, , .		0
39	PARP1 Inhibitor and Trabectedin Combination Does Not Increase Tumor Mutational Burden in Advanced Sarcomasâ€™A Preclinical and Translational Study. Cancers, 2021, 13, 6295.	3.7	0