

Gianmarco Contino

List of Publications by Year in descending order

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Version: 2024-02-01

26
papers

4,042
citations

430874

18
h-index

526287

27
g-index

54
all docs

54
docs citations

54
times ranked

8457
citing authors

#	ARTICLE	IF	CITATIONS
1	Rearrangement processes and structural variations show evidence of selection in oesophageal adenocarcinomas. <i>Communications Biology</i> , 2022, 5, 335.	4.4	8
2	Identifying Cancer Drivers Using DRIVE: A Feature-Based Machine Learning Model for a Pan-Cancer Assessment of Somatic Missense Mutations. <i>Cancers</i> , 2021, 13, 2779.	3.7	4
3	Comparison of outcomes between neoadjuvant chemoradiotherapy and neoadjuvant chemotherapy in patients with locally advanced esophageal cancer: A network meta-analysis. <i>EClinicalMedicine</i> , 2021, 42, 101183.	7.1	17
4	Identification of Subtypes of Barrett's Esophagus and Esophageal Adenocarcinoma Based on DNA Methylation Profiles and Integration of Transcriptome and Genome Data. <i>Gastroenterology</i> , 2020, 158, 1682-1697.e1.	1.3	58
5	Pan-cancer analysis of whole genomes identifies driver rearrangements promoted by LINE-1 retrotransposition. <i>Nature Genetics</i> , 2020, 52, 306-319.	21.4	275
6	Transcriptomic profiling reveals three molecular phenotypes of adenocarcinoma at the gastroesophageal junction. <i>International Journal of Cancer</i> , 2019, 145, 3389-3401.	5.1	17
7	The landscape of selection in 551 esophageal adenocarcinomas defines genomic biomarkers for the clinic. <i>Nature Genetics</i> , 2019, 51, 506-516.	21.4	166
8	The Evolving Genomic Landscape of Barrett's Esophagus and Esophageal Adenocarcinoma. <i>Gastroenterology</i> , 2017, 153, 657-673.e1.	1.3	69
9	Mutational signatures in esophageal adenocarcinoma define etiologically distinct subgroups with therapeutic relevance. <i>Nature Genetics</i> , 2016, 48, 1131-1141.	21.4	332
10	Whole-genome sequencing of nine esophageal adenocarcinoma cell lines. <i>F1000Research</i> , 2016, 5, 1336.	1.6	23
11	Combined MEK and PI3K Inhibition in a Mouse Model of Pancreatic Cancer. <i>Clinical Cancer Research</i> , 2015, 21, 396-404.	7.0	121
12	Two-year follow-up of wound complications associated with laparoendoscopic single-site adjustable gastric banding. <i>Surgery for Obesity and Related Diseases</i> , 2013, 9, 696-700.	1.2	5
13	KDM2B promotes pancreatic cancer via Polycomb-dependent and -independent transcriptional programs. <i>Journal of Clinical Investigation</i> , 2013, 123, 727-39.	8.2	144
14	TGF- β 2 and α 26 Integrin Act in a Common Pathway to Suppress Pancreatic Cancer Progression. <i>Cancer Research</i> , 2012, 72, 4840-4845.	0.9	82
15	Recurrence and prognostic factors in patients with aggressive fibromatosis. The role of radical surgery and its limitations. <i>World Journal of Surgical Oncology</i> , 2012, 10, 184.	1.9	32
16	RNA sequencing of pancreatic circulating tumour cells implicates WNT signalling in metastasis. <i>Nature</i> , 2012, 487, 510-513.	27.8	439
17	Pancreatic cancers require autophagy for tumor growth. <i>Genes and Development</i> , 2011, 25, 717-729.	5.9	1,224
18	Aberrant Overexpression of Satellite Repeats in Pancreatic and Other Epithelial Cancers. <i>Science</i> , 2011, 331, 593-596.	12.6	452

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19	<i>STAT3</i> Plays a Critical Role in <i>KRAS</i> -Induced Pancreatic Tumorigenesis. <i>Cancer Research</i> , 2011, 71, 5020-5029.	0.9	358
20	Lysine-specific Demethylase 2B (KDM2B)-let-7-Enhancer of Zester Homolog 2 (EZH2) Pathway Regulates Cell Cycle Progression and Senescence in Primary Cells. <i>Journal of Biological Chemistry</i> , 2011, 286, 33061-33069.	3.4	106
21	Aggressive Treatment Approach for Cloacogenic Carcinoma of the Anorectum: Report from a Single Cancer Center. <i>Digestive Surgery</i> , 2010, 27, 297-301.	1.2	8
22	Molecular imaging agents: impact on diagnosis and therapeutics in oncology. <i>Expert Reviews in Molecular Medicine</i> , 2010, 12, e20.	3.9	28
23	Metachronous Colon Metastases from Gastric Adenocarcinoma: A Case Report. <i>Case Reports in Oncology</i> , 2009, 2, 92-96.	0.7	19
24	Hereditary spastic paraplegia: clinical genomics and pharmacogenetic perspectives. <i>Expert Opinion on Pharmacotherapy</i> , 2006, 7, 1849-1856.	1.8	10
25	Expression analysis of the gene encoding for the U-box-type ubiquitin ligase UBE4A in human tissues. <i>Gene</i> , 2004, 328, 69-74.	2.2	22
26	Role of genetics in prevention of coronary atherosclerosis. <i>Current Opinion in Cardiology</i> , 2003, 18, 368-371.	1.8	18