## Marco Gerlinger

List of Publications by Year in descending order

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201385 223531 12,255 53 27 46 citations h-index g-index papers 61 61 61 20186 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Intratumor Heterogeneity and Branched Evolution Revealed by Multiregion Sequencing. New England Journal of Medicine, 2012, 366, 883-892.	13.9	6,769
2	Genomic architecture and evolution of clear cell renal cell carcinomas defined by multiregion sequencing. Nature Genetics, 2014, 46, 225-233.	9.4	1,103
3	Spatial and temporal diversity in genomic instability processes defines lung cancer evolution. Science, 2014, 346, 251-256.	6.0	962
4	Intratumor Heterogeneity: Seeing the Wood for the Trees. Science Translational Medicine, 2012, 4, 127ps10.	5.8	443
5	How Darwinian models inform therapeutic failure initiated by clonal heterogeneity in cancer medicine. British Journal of Cancer, 2010, 103, 1139-1143.	2.9	381
6	Classifying the evolutionary and ecological features of neoplasms. Nature Reviews Cancer, 2017, 17, 605-619.	12.8	303
7	Cancer Evolution and the Limits of Predictability in Precision Cancer Medicine. Trends in Cancer, 2016, 2, 49-63.	3.8	222
8	Cancer: Evolution Within a Lifetime. Annual Review of Genetics, 2014, 48, 215-236.	3.2	196
9	Genomic and Transcriptomic Determinants of Therapy Resistance and Immune Landscape Evolution during Anti-EGFR Treatment in Colorectal Cancer. Cancer Cell, 2019, 36, 35-50.e9.	7.7	179
10	Systematic Evaluation of the Prognostic Impact and Intratumour Heterogeneity of Clear Cell Renal Cell Carcinoma Biomarkers. European Urology, 2014, 66, 936-948.	0.9	141
11	Assessment of an RNA interference screen-derived mitotic and ceramide pathway metagene as a predictor of response to neoadjuvant paclitaxel for primary triple-negative breast cancer: a retrospective analysis of five clinical trials. Lancet Oncology, The, 2010, 11, 358-365.	5.1	116
12	Immunopeptidomics of colorectal cancer organoids reveals a sparse HLA class I neoantigen landscape and no increase in neoantigens with interferon or MEK-inhibitor treatment., 2019, 7, 309.		112
13	Intratumour Heterogeneity in Urologic Cancers: From Molecular Evidence to Clinical Implications. European Urology, 2015, 67, 729-737.	0.9	100
14	Ultraâ€deep T cell receptor sequencing reveals the complexity and intratumour heterogeneity of T cell clones in renal cell carcinomas. Journal of Pathology, 2013, 231, 424-432.	2.1	93
15	Genomeâ€wide RNA interference analysis of renal carcinoma survival regulators identifies MCT4 as a Warburg effect metabolic target. Journal of Pathology, 2012, 227, 146-156.	2.1	92
16	Egr-1 Induces the Expression of Its Corepressor Nab2 by Activation of the Nab2 Promoter Thereby Establishing a Negative Feedback Loop. Journal of Biological Chemistry, 2005, 280, 42785-42793.	1.6	83
17	Parallel evolution of tumour subclones mimics diversity between tumours. Journal of Pathology, 2013, 230, 356-364.	2.1	79
18	Anti-cancer drug resistance: Understanding the mechanisms through the use of integrative genomics and functional RNA interference. European Journal of Cancer, 2010, 46, 2166-2177.	1.3	71

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19	Development of synchronous VHL syndrome tumors reveals contingencies and constraints to tumor evolution. Genome Biology, 2014, 15, 433.	3.8	69
20	CEA expression heterogeneity and plasticity confer resistance to the CEA-targeting bispecific immunotherapy antibody cibisatamab (CEA-TCB) in patient-derived colorectal cancer organoids., 2019, 7, 101.		65
21	The Effect of VEGF-Targeted Therapy on Biomarker Expression in Sequential Tissue from Patients with Metastatic Clear Cell Renal Cancer. Clinical Cancer Research, 2013, 19, 6924-6934.	3.2	62
22	Identifying key questions in the ecology and evolution of cancer. Evolutionary Applications, 2021, 14, 877-892.	1.5	58
23	Dissecting cancer evolution at the macro-heterogeneity and micro-heterogeneity scale. Current Opinion in Genetics and Development, $2015$ , $30$ , $1-6$ .	1.5	57
24	The promise of circulating tumor cell analysis in cancer management. Genome Biology, 2014, 15, 448.	3.8	47
25	Ultra-Sensitive Mutation Detection and Genome-Wide DNA Copy Number Reconstruction by Error-Corrected Circulating Tumor DNA Sequencing. Clinical Chemistry, 2018, 64, 1626-1635.	1.5	46
26	Extreme intratumour heterogeneity and driver evolution in mismatch repair deficient gastro-oesophageal cancer. Nature Communications, 2020, 11, 139.	5.8	44
27	Predictive biomarker discovery through the parallel integration of clinical trial and functional genomics datasets. Genome Medicine, 2010, 2, 53.	3.6	43
28	Circulating tumour DNA, a promising biomarker for the management of colorectal cancer. Critical Reviews in Oncology/Hematology, 2018, 122, 72-82.	2.0	40
29	Defining the true impact of coronavirus disease 2019 in the at-risk population of patients with cancer. European Journal of Cancer, 2020, 136, 99-106.	1.3	31
30	Heterogeneous response and progression patterns reveal phenotypic heterogeneity of tyrosine kinase inhibitor response in metastatic renal cell carcinoma. BMC Medicine, 2016, 14, 185.	2.3	29
31	Spontaneous CD8 T Cell Responses against the Melanocyte Differentiation Antigen RAB38/NY-MEL-1 in Melanoma Patients. Journal of Immunology, 2006, 177, 8212-8218.	0.4	24
32	Computational optimisation of targeted DNA sequencing for cancer detection. Scientific Reports, 2013, 3, 3309.	1.6	20
33	Immunotherapy Sensitivity of Mismatch Repair-Deficient Cancer: Mutation Load Is Not Enough. Cancer Cell, 2021, 39, 16-18.	7.7	18
34	Elevated LDH predicts poor outcome of recurrent germ cell tumours treated with dose dense chemotherapy. European Journal of Cancer, 2010, 46, 2913-2918.	1.3	17
35	Mutational signatures impact the evolution of anti-EGFR antibody resistance in colorectal cancer. Nature Ecology and Evolution, 2021, 5, 1024-1032.	3.4	16
36	Detecting and Tracking Circulating Tumour DNA Copy Number Profiles during First Line Chemotherapy in Oesophagogastric Adenocarcinoma. Cancers, 2019, 11, 736.	1.7	15

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37	Computational Image Analysis of T-Cell Infiltrates in Resectable Gastric Cancer: Association with Survival and Molecular Subtypes. Journal of the National Cancer Institute, 2021, 113, 88-98.	3.0	15
38	Multiplexed single cell protein expression analysis in solid tumours using a miniaturised microfluidic assay. Convergent Science Physical Oncology, 2017, 3, 024003.	2.6	13
39	Targeted drugs ramp up cancer mutability. Science, 2019, 366, 1452-1453.	6.0	13
40	Surveillance investigations after high-dose therapy with stem cell rescue for recurrent follicular lymphoma have no impact on management. Haematologica, 2010, 95, 1130-1135.	1.7	10
41	Efficacy and Cardiotoxic Safety Profile of Raltitrexed in Fluoropyrimidines-Pretreated or High-Risk Cardiac Patients With GI Malignancies: Large Single-Center Experience. Clinical Colorectal Cancer, 2019, 18, 64-71.e1.	1.0	10
42	Metastasis Seeding Cells: Lone Invaders or Mass Migrators?. Clinical Cancer Research, 2018, 24, 2032-2034.	3.2	7
43	Questions to guide cancer evolution as a framework for furthering progress in cancer research and sustainable patient outcomes. , 2022, 39, .		7
44	Prognostic and Predictive Markers in Metastatic Renal Cell Carcinoma. Journal of Clinical Oncology, 2013, 31, 971-972.	0.8	6
45	Cancer (r)evolution. Nature Ecology and Evolution, 2017, 1, 1051-1052.	3.4	6
46	Circulating Tumour DNA Sequencing Identifies a Genetic Resistance-Gap in Colorectal Cancers with Acquired Resistance to EGFR-Antibodies and Chemotherapy. Cancers, 2020, 12, 3736.	1.7	6
47	Effect of perioperative FLOT <i>versus</i> ECF/ECX on short-term outcomes after surgery for resectable oesophagogastric adenocarcinoma: propensity score-matched study. BJS Open, 2022, 6, .	0.7	3
48	Genetic Intratumor Heterogeneity., 2015,, 571-593.		2
49	Diagnostic Accuracy and Safety of Coaxial System in Oncology Patients Treated in a Specialist Cancer Center With Prospective Validation Within Clinical Trial Data. Frontiers in Oncology, 2020, 10, 1634.	1.3	2
50	Translating Seminoma Genomic Landscapes into Clinical Practice. European Urology, 2015, 68, 84-85.	0.9	0
51	Active Surveillance Strategies Have Neither Clinical or Survival Benefit Following High-Dose Chemotherapy and Progenitor Cell Rescue in Follicular Lymphoma Blood, 2008, 112, 3251-3251.	0.6	0
52	Abstract PR012: Genetic and immune landscape evolution defines subtypes of MMR deficient colorectal cancer. Cancer Research, 2022, 82, PR012-PR012.	0.4	0
53	Abstract A002: Genetic and immune landscape evolution defines subtypes of MMR deficient colorectal cancer. Cancer Research, 2022, 82, A002-A002.	0.4	0