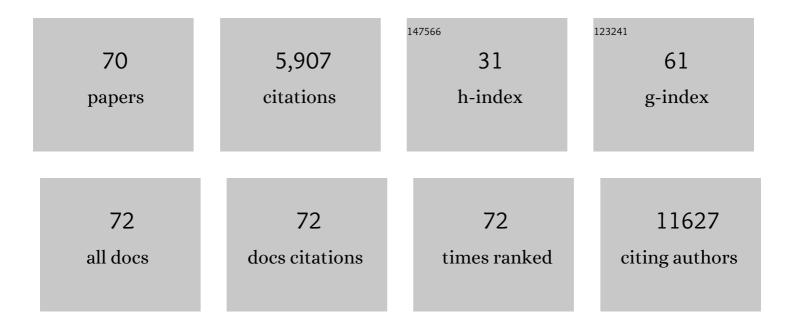
Joseph F Costello

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	PI3K/AKT/mTOR signaling pathway activity in IDH-mutant diffuse glioma and clinical implications. Neuro-Oncology, 2022, 24, 1471-1481.	0.6	14
2	<i>TERT</i> promoter C228T mutation in neural progenitors confers growth advantage following telomere shortening <i>in vivo</i> . Neuro-Oncology, 2022, 24, 2063-2075.	0.6	9
3	Imaging biomarkers of TERT or GABPB1 silencing in TERT-positive glioblastoma. Neuro-Oncology, 2022, , ·	0.6	3
4	Meningioma DNA methylation groups identify biological drivers and therapeutic vulnerabilities. Nature Genetics, 2022, 54, 649-659.	9.4	93
5	Deuterium Metabolic Imaging Reports on TERT Expression and Early Response to Therapy in Cancer. Clinical Cancer Research, 2022, 28, 3526-3536.	3.2	15
6	Functional analysis of low-grade glioma genetic variants predicts key target genes and transcription factors. Neuro-Oncology, 2021, 23, 638-649.	0.6	9
7	Early Noninvasive Metabolic Biomarkers of Mutant IDH Inhibition in Glioma. Metabolites, 2021, 11, 109.	1.3	15
8	Cancer-specific loss of <i>TERT</i> activation sensitizes glioblastoma to DNA damage. Proceedings of the United States of America, 2021, 118, .	3.3	28
9	miRNA-independent function of long noncoding pri-miRNA loci. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	18
10	Temozolomide-induced hypermutation is associated with distant recurrence and reduced survival after high-grade transformation of low-grade <i>IDH</i> mutant gliomas. Neuro-Oncology, 2021, 23, 1872-1884.	0.6	48
11	Metabolic imaging detects elevated glucose flux through the pentose phosphate pathway associated with TERT expression in low-grade gliomas. Neuro-Oncology, 2021, 23, 1509-1522.	0.6	15
12	Spatial concordance of DNA methylation classification in diffuse glioma. Neuro-Oncology, 2021, 23, 2054-2065.	0.6	19
13	Non-invasive assessment of telomere maintenance mechanisms in brain tumors. Nature Communications, 2021, 12, 92.	5.8	21
14	Unique challenges for glioblastoma immunotherapy—discussions across neuro-oncology and non-neuro-oncology experts in cancer immunology. Meeting Report from the 2019 SNO Immuno-Oncology Think Tank. Neuro-Oncology, 2021, 23, 356-375.	0.6	59
15	Epigenomic contributions to tumor cell heterogeneity and plasticity. Nature Genetics, 2021, 53, 1403-1404.	9.4	3
16	The DNA methylation landscape of advanced prostate cancer. Nature Genetics, 2020, 52, 778-789.	9.4	198
17	Epigenomic programming in early fetal brain development. Epigenomics, 2020, 12, 1053-1070.	1.0	9
18	Prostate Cancer Progression and the Epigenome. New England Journal of Medicine, 2020, 383, 2287-2290.	13.9	5

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19	Glutamate Is a Noninvasive Metabolic Biomarker of IDH1-Mutant Glioma Response to Temozolomide Treatment. Cancer Research, 2020, 80, 5098-5108.	0.4	18
20	Multiplatform genomic profiling and magnetic resonance imaging identify mechanisms underlying intratumor heterogeneity in meningioma. Nature Communications, 2020, 11, 4803.	5.8	56
21	Patient-derived cells from recurrent tumors that model the evolution of IDH-mutant glioma. Neuro-Oncology Advances, 2020, 2, vdaa088.	0.4	16
22	Gliomas arising in the setting of Li-Fraumeni syndrome stratify into two molecular subgroups with divergent clinicopathologic features. Acta Neuropathologica, 2020, 139, 953-957.	3.9	18
23	MGMT promoter methylation level in newly diagnosed low-grade glioma is a predictor of hypermutation at recurrence. Neuro-Oncology, 2020, 22, 1580-1590.	0.6	55
24	Multiplatform Molecular Profiling Reveals Epigenomic Intratumor Heterogeneity in Ependymoma. Cell Reports, 2020, 30, 1300-1309.e5.	2.9	11
25	PATH-12. TEMOZOLOMIDE-INDUCED HYPERMUTATION IS ASSOCIATED WITH HIGH-GRADE TRANSFORMATION, DISTANT RECURRENCE AND REDUCED SURVIVAL IN INITIALLY LOW GRADE IDH-MUTANT GLIOMAS. Neuro-Oncology, 2020, 22, ii166-ii166.	0.6	2
26	EPCO-36. GENOMIC INSTABILITY AND TRANSCRIPTOMIC SIGNATURES UNDERLYING EPIGENETIC MENINGIOMA SUBGROUPS REVEALS MECHANISMS OF IMMUNE INFILTRATION AND THERAPEUTIC VULNERABILITIES. Neuro-Oncology, 2020, 22, ii77-ii77.	0.6	0
27	BIOM-19. METABOLIC ALTERATION INDUCED BY SELECTIVE KNOCK DOWN OF GABPB1L IN U251 CELLS. Neuro-Oncology, 2020, 22, ii5-ii6.	0.6	0
28	TAMI-08. A TALE OF TWO TELOMERE MAINTENANCE MECHANISMS: TERT EXPRESSION AND THE ALT PATHWAY INDUCE UNIQUE MRS-DETECTABLE METABOLIC REPROGRAMMING IN LOW-GRADE GLIOMAS. Neuro-Oncology, 2020, 22, ii214-ii214.	0.6	0
29	BIOM-38. PI3K/AKT/mTOR SIGNALING PATHWAY ACTIVITY IN IDH-MUTANT DIFFUSE GLIOMA. Neuro-Oncology, 2020, 22, ii9-ii10.	0.6	0
30	EPCO-31. EPIGENOMIC INTRATUMORAL HETEROGENEITY OF GLIOBLASTOMA IN THREE-DIMENSIONAL SPACE. Neuro-Oncology, 2020, 22, ii76-ii76.	0.6	2
31	Recurrent KBTBD4 small in-frame insertions and absence of DROSHA deletion or DICER1 mutation differentiate pineal parenchymal tumor of intermediate differentiation (PPTID) from pineoblastoma. Acta Neuropathologica, 2019, 137, 851-854.	3.9	45
32	TMOD-27. A NEURAL CREST CELL SUBPOPULATION UNDERLIES INTRATUMOR HETEROGENEITY IN MENINGIOMA. Neuro-Oncology, 2019, 21, vi268-vi268.	0.6	1
33	CSIG-24. GABP LINKS AMPK SIGNALING TO TERT REGULATION IN A TERT PROMOTER MUTATION DEPENDENT MANNER. Neuro-Oncology, 2019, 21, vi49-vi49.	0.6	1
34	IMMU-11. SPATIOTEMPORAL IMMUNOGENOMIC ANALYSIS OF THE T-CELL REPERTOIRE IN IDH-MUTANT LOWER GRADE GLIOMAS. Neuro-Oncology, 2019, 21, vi121-vi121.	0.6	0
35	GENE-47. A 3D ATLAS TO EVALUATE THE SPATIAL PATTERNING OF GENETIC ALTERATIONS AND TUMOR CELL STATES IN GLIOMA. Neuro-Oncology, 2019, 21, vi107-vi108.	0.6	0
36	GENE-43. TARGETING GABPb1L INHIBITS IN VIVO GROWTH OF TERT PROMOTER MUTANT GLIOBLASTOMA. Neuro-Oncology, 2019, 21, vi106-vi107.	0.6	0

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#	Article	IF	CITATIONS
37	Genomic Profiling of BDE-47 Effects on Human Placental Cytotrophoblasts. Toxicological Sciences, 2019, 167, 211-226.	1.4	32
38	Temozolomide-associated hypermutation in gliomas. Neuro-Oncology, 2018, 20, 1300-1309.	0.6	130
39	Glioma through the looking GLASS: molecular evolution of diffuse gliomas and the Glioma Longitudinal Analysis Consortium. Neuro-Oncology, 2018, 20, 873-884.	0.6	119
40	Antigenic expression and spontaneous immune responses support the use of a selected peptide set from the IMA950 glioblastoma vaccine for immunotherapy of grade II and III glioma. Oncolmmunology, 2018, 7, e1391972.	2.1	42
41	Genomic analysis of the origins and evolution of multicentric diffuse lower-grade gliomas. Neuro-Oncology, 2018, 20, 632-641.	0.6	33
42	The long non-coding RNA <i>HOTAIR</i> is transcriptionally activated by HOXA9 and is an independent prognostic marker in patients with malignant glioma. Oncotarget, 2018, 9, 15740-15756.	0.8	28
43	Disruption of the β1L Isoform of GABP Reverses Glioblastoma Replicative Immortality in a TERT Promoter Mutation-Dependent Manner. Cancer Cell, 2018, 34, 513-528.e8.	7.7	103
44	Chemotherapy for adult low-grade gliomas: clinical outcomes by molecular subtype in a phase II study of adjuvant temozolomide. Neuro-Oncology, 2017, 19, now176.	0.6	70
45	DNA methylation: an epigenetic mark of cellular memory. Experimental and Molecular Medicine, 2017, 49, e322-e322.	3.2	286
46	Treatment-Induced Mutagenesis and Selective Pressures Sculpt Cancer Evolution. Cold Spring Harbor Perspectives in Medicine, 2017, 7, a026617.	2.9	59
47	Comprehensive Analysis of Hypermutation in Human Cancer. Cell, 2017, 171, 1042-1056.e10.	13.5	596
48	Clonal expansion and epigenetic reprogramming following deletion or amplification of mutant <i>IDH1</i> . Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 10743-10748.	3.3	109
49	Probing the phosphatidylinositol 3â€kinase/mammalian target of rapamycin pathway in gliomas: A phase 2 study of everolimus for recurrent adult lowâ€grade gliomas. Cancer, 2017, 123, 4631-4639.	2.0	43
50	Somatic and Germline <i>TP53</i> Alterations in Second Malignant Neoplasms from Pediatric Cancer Survivors. Clinical Cancer Research, 2017, 23, 1852-1861.	3.2	29
51	Genome-wide DNA methylation is predictive of outcome in juvenile myelomonocytic leukemia. Nature Communications, 2017, 8, 2127.	5.8	75
52	Isocitrate dehydrogenase mutations suppress STAT1 and CD8+ T cell accumulation in gliomas. Journal of Clinical Investigation, 2017, 127, 1425-1437.	3.9	334
53	CBIO-02. MUTANT IDH EXPRESSION DRIVES TERT PROMOTER REACTIVATION AS PART OF THE CELLULAR TRANSFORMATION PROCESS. Neuro-Oncology, 2016, 18, vi35-vi35.	0.6	0
54	Intratumoral Heterogeneity of the Epigenome. Cancer Cell, 2016, 29, 440-451.	7.7	172

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#	Article	IF	CITATIONS
55	Mutant IDH1 Expression Drives <i>TERT</i> Promoter Reactivation as Part of the Cellular Transformation Process. Cancer Research, 2016, 76, 6680-6689.	0.4	55
56	Understanding TERT Promoter Mutations: A Common Path to Immortality. Molecular Cancer Research, 2016, 14, 315-323.	1.5	222
57	Oncogene brought into the loop. Nature, 2016, 529, 34-35.	13.7	6
58	Mutant IDH1 expression is associated with down-regulation of monocarboxylate transporters. Oncotarget, 2016, 7, 34942-34955.	0.8	32
59	MPTH-29CONNECTING MUTANT GENOTYPES TO ABERRANT TRANSCRIPTIONAL SIGNATURES ACROSS SERIAL SECTIONS OF A HUMAN TUMOR. Neuro-Oncology, 2015, 17, v144.4-v145.	0.6	0
60	HIGD1A Regulates Oxygen Consumption, ROS Production, and AMPK Activity during Glucose Deprivation to Modulate Cell Survival and Tumor Growth. Cell Reports, 2015, 10, 891-899.	2.9	79
61	Epigenetic and transcriptional determinants of the human breast. Nature Communications, 2015, 6, 6351.	5.8	56
62	Evolution of DNA repair defects during malignant progression of low-grade gliomas after temozolomide treatment. Acta Neuropathologica, 2015, 129, 597-607.	3.9	143
63	The transcription factor GABP selectively binds and activates the mutant TERT promoter in cancer. Science, 2015, 348, 1036-1039.	6.0	451
64	The genomic landscape of juvenile myelomonocytic leukemia. Nature Genetics, 2015, 47, 1326-1333.	9.4	233
65	DNA Methylation and Somatic Mutations Converge on the Cell Cycle and Define Similar Evolutionary Histories in Brain Tumors. Cancer Cell, 2015, 28, 307-317.	7.7	221
66	Regulatory network decoded from epigenomes of surface ectoderm-derived cell types. Nature Communications, 2014, 5, 5442.	5.8	25
67	Transcription Restores DNA Repair to Heterochromatin, Determining Regional Mutation Rates in Cancer Genomes. Cell Reports, 2014, 9, 1228-1234.	2.9	104
68	Mutational Analysis Reveals the Origin and Therapy-Driven Evolution of Recurrent Glioma. Science, 2014, 343, 189-193.	6.0	1,147
69	Restriction Landmark Genomic Scanning: Analysis of CpG Islands in Genomes by 2D Gel Electrophoresis. Methods in Molecular Biology, 2009, 507, 131-148.	0.4	17
70	DNA methylation in brain development and gliomagenesis. Frontiers in Bioscience - Landmark, 2003, 8, s175-184.	3.0	44