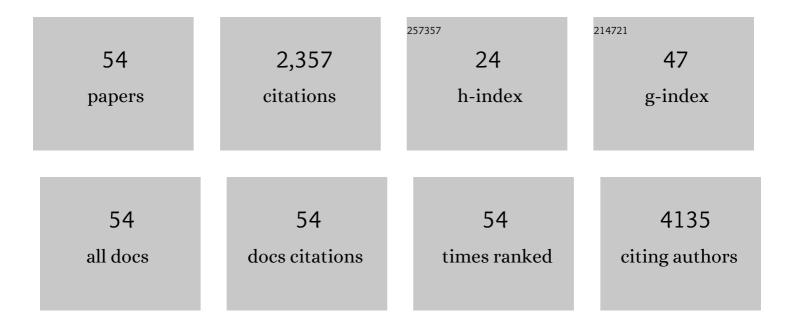
## Jose Luis Costa

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Gastric microbial community profiling reveals a dysbiotic cancer-associated microbiota. Gut, 2018, 67, 226-236.	6.1	496
2	High-resolution aCGH and expression profiling identifies a novel genomic subtype of ER negative breast cancer. Genome Biology, 2007, 8, R215.	13.9	275
3	Alterations in Vitamin D signalling and metabolic pathways in breast cancer progression: a study of VDR, CYP27B1 and CYP24A1 expression in benign and malignant breast lesions Vitamin D pathways unbalanced in breast lesions. BMC Cancer, 2010, 10, 483.	1.1	164
4	Integration of next-generation sequencing in clinical diagnostic molecular pathology laboratories for analysis of solid tumours; an expert opinion on behalf of IQN Path ASBL. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2017, 470, 5-20.	1.4	82
5	Cyanobiont diversity within coralloid roots of selected cycad species. FEMS Microbiology Ecology, 1999, 28, 85-91.	1.3	80
6	Diversity of Cyanobacterial Hydrogenases, a Molecular Approach. Current Microbiology, 2000, 40, 356-361.	1.0	76
7	Genetic Diversity of Nostoc Symbionts Endophytically Associated with Two Bryophyte Species. Applied and Environmental Microbiology, 2001, 67, 4393-4396.	1.4	75
8	Apocrine carcinoma of the breast: a comprehensive review. Histology and Histopathology, 2013, 28, 1393-409.	0.5	67
9	1Alpha,25-dihydroxyvitamin D3 induces de novo E-cadherin expression in triple-negative breast cancer cells by CDH1-promoter demethylation. Anticancer Research, 2012, 32, 249-57.	0.5	63
10	Prevalence of BRCA1/BRCA2 mutations in a Brazilian population sample at-risk for hereditary breast cancer and characterization of its genetic ancestry. Oncotarget, 2016, 7, 80465-80481.	0.8	62
11	Heterozygous germline mutations in A2ML1 are associated with a disorder clinically related to Noonan syndrome. European Journal of Human Genetics, 2015, 23, 317-324.	1.4	61
12	Vitamin D and the mammary gland: a review on its role in normal development and breast cancer. Breast Cancer Research, 2012, 14, 211.	2.2	55
13	The Role of Liquid Biopsy in Early Diagnosis of Lung Cancer. Frontiers in Oncology, 2021, 11, 634316.	1.3	50
14	P-cadherin, vimentin and CK14 for identification of basal-like phenotype in breast carcinomas: an immunohistochemical study. Histology and Histopathology, 2010, 25, 963-74.	0.5	46
15	Array Comparative Genomic Hybridization Copy Number Profiling: A New Tool for Translational Research in Solid Malignancies. Seminars in Radiation Oncology, 2008, 18, 98-104.	1.0	45
16	Liquid Biopsy beyond Circulating Tumor Cells and Cell-Free DNA. Acta Cytologica, 2019, 63, 479-488.	0.7	42
17	Anti-proliferative action of vitamin D in MCF7 is still active after siRNA-VDR knock-down. BMC Genomics, 2009, 10, 499.	1.2	41
18	Identifying relationships between imaging phenotypes and lung cancer-related mutation status: EGFR and KRAS. Scientific Reports, 2020, 10, 3625.	1.6	41

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19	PIKing the right isoform: the emergent role of the p110β subunit in breast cancer. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2010, 456, 235-243.	1.4	37
20	Nonoptical Massive Parallel DNA Sequencing of <i>BRCA1</i> and <i>BRCA2</i> Genes in a Diagnostic Setting. Human Mutation, 2013, 34, 629-635.	1.1	37
21	Sequence based data supports a single Nostoc strain in individual coralloid roots of cycads. FEMS Microbiology Ecology, 2004, 49, 481-487.	1.3	36
22	The Cyanobacterial tRNALeu (UAA) Intron: Evolutionary Patterns in a Genetic Marker. Molecular Biology and Evolution, 2002, 19, 850-857.	3.5	34
23	Very small mobile repeated elements in cyanobacterial genomes. Genome Research, 2008, 18, 1484-1499.	2.4	27
24	Expression Microarray Analysis and Oligo Array Comparative Genomic Hybridization of Acquired Gemcitabine Resistance in Mouse Colon Reveals Selection for Chromosomal Aberrations. Cancer Research, 2005, 65, 10208-10213.	0.4	26
25	<i>EGFR</i> Assessment in Lung Cancer CT Images: Analysis of Local and Holistic Regions of Interest Using Deep Unsupervised Transfer Learning. IEEE Access, 2021, 9, 58667-58676.	2.6	24
26	Targeted Gene Next-Generation Sequencing Panel in Patients with Advanced Lung Adenocarcinoma: Paving the Way for Clinical Implementation. Cancers, 2019, 11, 1229.	1.7	23
27	Machine Learning and Feature Selection Methods for EGFR Mutation Status Prediction in Lung Cancer. Applied Sciences (Switzerland), 2021, 11, 3273.	1.3	21
28	Multicenter Evaluation of the Idylla NRAS-BRAF Mutation Test in Metastatic Colorectal Cancer. Journal of Molecular Diagnostics, 2018, 20, 664-676.	1.2	19
29	Simultaneous detection of lung fusions using a multiplex RT-PCR next generation sequencing-based approach: a multi-institutional research study. BMC Cancer, 2018, 18, 828.	1.1	19
30	Towards Machine Learning-Aided Lung Cancer Clinical Routines: Approaches and Open Challenges. Journal of Personalized Medicine, 2022, 12, 480.	1.1	19
31	New massive parallel sequencing approach improves the genetic characterization of congenital myopathies. Journal of Human Genetics, 2016, 61, 497-505.	1.1	15
32	Benign and malignant apocrine lesions of the breast. Expert Review of Anticancer Therapy, 2012, 12, 215-221.	1.1	14
33	Comprehensive massive parallel DNA sequencing strategy for the genetic diagnosis of the neuro-cardio-facio-cutaneous syndromes. European Journal of Human Genetics, 2015, 23, 347-353.	1.4	14
34	Comprehensive Perspective for Lung Cancer Characterisation Based on Al Solutions Using CT Images. Journal of Clinical Medicine, 2021, 10, 118.	1.0	14
35	Circulating Tumor DNA: A Step into the Future of Cancer Management. Acta Cytologica, 2019, 63, 456-465.	0.7	13
36	BBIT20 inhibits homologous DNA repair with disruption of the BRCA1–BARD1 interaction in breast and ovarian cancer. British Journal of Pharmacology, 2021, 178, 3627-3647.	2.7	13

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37	Liquid Biopsy for Disease Monitoring in Non-Small Cell Lung Cancer: The Link between Biology and the Clinic. Cells, 2021, 10, 1912.	1.8	13
38	Molecular characterization of CD44+/CD24â^'/Ck+/CD45â^' cells in benign and malignant breast lesions. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2017, 470, 311-322.	1.4	12
39	Induction of apoptosis increases sensitivity to detect cancer mutations in plasma. European Journal of Cancer, 2020, 127, 130-138.	1.3	11
40	Targeting p53 for Melanoma Treatment: Counteracting Tumour Proliferation, Dissemination and Therapeutic Resistance. Cancers, 2021, 13, 1648.	1.7	11
41	Genetic Heterogeneity in Colorectal Cancer and its Clinical Implications. Acta Medica Portuguesa, 2015, 28, 370-375.	0.2	10
42	Pre-Training Autoencoder for Lung Nodule Malignancy Assessment Using CT Images. Applied Sciences (Switzerland), 2020, 10, 7837.	1.3	10
43	The value of cell-free circulating tumour DNA profiling in advanced non-small cell lung cancer (NSCLC) management. Cancer Cell International, 2021, 21, 675.	1.8	9
44	Molecular alterations in endometrial archived liquidâ€based cytology. Diagnostic Cytopathology, 2013, 41, 492-496.	0.5	8
45	Clinical Application of Next-Generation Sequencing of Plasma Cell-Free DNA for Genotyping Untreated Advanced Non-Small Cell Lung Cancer. Cancers, 2021, 13, 2707.	1.7	8
46	Sharing Biomedical Data: Strengthening Al Development in Healthcare. Healthcare (Switzerland), 2021, 9, 827.	1.0	8
47	An Interpretable Approach for Lung Cancer Prediction and Subtype Classification using Gene Expression. , 2021, 2021, 1707-1710.		7
48	Liquid Biopsy: A New Tool in Oncology. Acta Cytologica, 2019, 63, 448-448.	0.7	6
49	RAF-1 promotes survival of thyroid cancer cells harboring RET/PTC1 rearrangement independently of ERK activation. Molecular and Cellular Endocrinology, 2015, 415, 64-75.	1.6	5
50	Utility of Circulating Tumor DNA in Different Clinical Scenarios of Breast Cancer. Cancers, 2020, 12, 3797.	1.7	4
51	Multiple instance learning for lung pathophysiological findings detection using CT scans. Medical and Biological Engineering and Computing, 2022, 60, 1569-1584.	1.6	4
52	The Adaptive Immune Landscape of the Colorectal Adenoma–Carcinoma Sequence. International Journal of Molecular Sciences, 2021, 22, 9791.	1.8	3
53	Validation of a Targeted Next-Generation Sequencing Panel for Tumor Mutation Burden Analysis. Journal of Molecular Diagnostics, 2021, 23, 882-893.	1.2	2
54	Differential Gene Expression Analysis ofÂtheÂMost Relevant Genes forÂLung Cancer Prediction andÂSub-type Classification. Lecture Notes in Computer Science, 2022, , 182-191.	1.0	0