

Srikant Ambatipudi

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

19 papers	1,065 citations	13 h-index	19 g-index
19 ext. papers	1,381 ext. citations	5.9 avg, IF	3.06 L-index

#	Paper	IF	Citations
19	Epigenome-wide association meta-analysis of DNA methylation with coffee and tea consumption. <i>Nature Communications</i> , 2021 , 12, 2830	17.4	9
18	Blood DNA methylation and breast cancer risk: a meta-analysis of four prospective cohort studies. <i>Breast Cancer Research</i> , 2019 , 21, 62	8.3	20
17	Epigenome-wide association study for lifetime estrogen exposure identifies an epigenetic signature associated with breast cancer risk. <i>Clinical Epigenetics</i> , 2019 , 11, 66	7.7	12
16	Smoking-associated DNA methylation changes: no smoke without fire. <i>Epigenomics</i> , 2019 , 11, 1117-1119	4.4	3
15	Identifying and correcting epigenetics measurements for systematic sources of variation. <i>Clinical Epigenetics</i> , 2018 , 10, 38	7.7	13
14	Roadmap for investigating epigenome deregulation and environmental origins of cancer. <i>International Journal of Cancer</i> , 2018 , 142, 874-882	7.5	46
13	DNA methylation derived systemic inflammation indices are associated with head and neck cancer development and survival. <i>Oral Oncology</i> , 2018 , 85, 87-94	4.4	9
12	Assessing the Role of DNA Methylation-Derived Neutrophil-to-Lymphocyte Ratio in Rheumatoid Arthritis. <i>Journal of Immunology Research</i> , 2018 , 2018, 2624981	4.5	8
11	DNA methylome analysis identifies accelerated epigenetic ageing associated with postmenopausal breast cancer susceptibility. <i>European Journal of Cancer</i> , 2017 , 75, 299-307	7.5	104
10	Chromosomal Alterations and Gene Expression Changes Associated with the Progression of Leukoplakia to Advanced Gingivobuccal Cancer. <i>Translational Oncology</i> , 2017 , 10, 396-409	4.9	36
9	Integrated data analysis reveals potential drivers and pathways disrupted by DNA methylation in papillary thyroid carcinomas. <i>Clinical Epigenetics</i> , 2017 , 9, 45	7.7	40
8	Prognostic Classifier Based on Genome-Wide DNA Methylation Profiling in Well-Differentiated Thyroid Tumors. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2017 , 102, 4089-4099	5.6	33
7	Epigenetic Signatures of Cigarette Smoking. <i>Circulation: Cardiovascular Genetics</i> , 2016 , 9, 436-447		442
6	Tobacco smoking-associated genome-wide DNA methylation changes in the EPIC study. <i>Epigenomics</i> , 2016 , 8, 599-618	4.4	145
5	The LPA1/ZEB1/miR-21-activation pathway regulates metastasis in basal breast cancer. <i>Oncotarget</i> , 2015 , 6, 20604-20	3.3	46
4	14-3-3 σ -Mediated transport of plakoglobin to the cell border is required for the initiation of desmosome assembly in vitro and in vivo. <i>Journal of Cell Science</i> , 2014 , 127, 2174-88	5.3	13
3	Downregulation of keratin 76 expression during oral carcinogenesis of human, hamster and mouse. <i>PLoS ONE</i> , 2013 , 8, e70688	3.7	16

2	Genome-wide expression and copy number analysis identifies driver genes in gingivobuccal cancers. <i>Genes Chromosomes and Cancer</i> , 2012 , 51, 161-73	5	28
1	Genomic profiling of advanced-stage oral cancers reveals chromosome 11q alterations as markers of poor clinical outcome. <i>PLoS ONE</i> , 2011 , 6, e17250	3-7	42