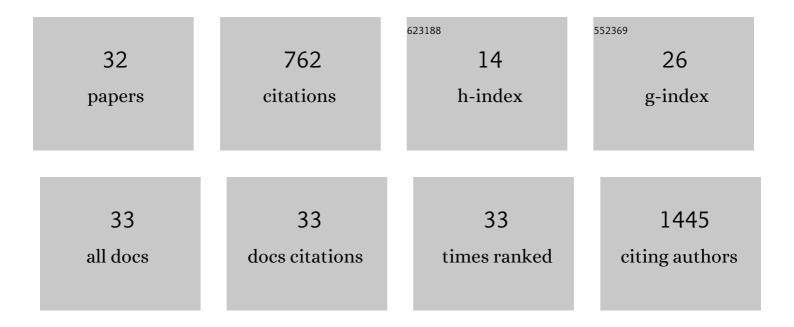
## Arvind Y M Sundaram

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3835893/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Optimization of enzymatic fragmentation is crucial to maximize genome coverage: a comparison of library preparation methods for Illumina sequencing. BMC Genomics, 2022, 23, 92.	1.2	4
2	Single-Cell RNA Sequencing of <i>In Vitro</i> Expanded Chondrocytes: MSC-Like Cells With No Evidence of Distinct Subsets. Cartilage, 2021, 13, 774S-784S.	1.4	4
3	Small-Scale Comparative Genomic Analysis of Listeria monocytogenes Isolated from Environments of Salmon Processing Plants and Human Cases in Norway. Hygiene, 2021, 1, 43-55.	0.5	8
4	Rapid SARS-CoV-2 variant monitoring using PCR confirmed by whole genome sequencing in a high-volume diagnostic laboratory. Journal of Clinical Virology, 2021, 141, 104906.	1.6	21
5	Gene Expression in Embryos From Norwegian Red Bulls With High or Low Non Return Rate: An RNA-Seq Study of in vivo-Produced Single Embryos. Frontiers in Genetics, 2021, 12, 780113.	1.1	6
6	Systematic assessment of commercially available low-input miRNA library preparation kits. RNA Biology, 2020, 17, 75-86.	1.5	28
7	An Illumina approach to MHC typing of Atlantic salmon. Immunogenetics, 2020, 72, 89-100.	1.2	7
8	Selective Stimulation of Duplicated Atlantic Salmon MHC Pathway Genes by Interferon-Gamma. Frontiers in Immunology, 2020, 11, 571650.	2.2	3
9	Candidate genes for monitoring hydrogen peroxide resistance in the salmon louse, Lepeophtheirus salmonis. Parasites and Vectors, 2020, 13, 344.	1.0	3
10	Gene expression and gastrointestinal function is altered in piglet small intestine by weaning and inclusion of Cyberlindnera jadinii yeast as a protein source. Journal of Functional Foods, 2020, 73, 104118.	1.6	12
11	Distinct Pattern of Endoplasmic Reticulum Protein Processing and Extracellular Matrix Proteins in Functioning and Silent Corticotroph Pituitary Adenomas. Cancers, 2020, 12, 2980.	1.7	9
12	Transcriptome profiling of human thymic CD4+ and CD8+ T cells compared to primary peripheral T cells. BMC Genomics, 2020, 21, 350.	1.2	11
13	An extension to: Systematic assessment of commercially available low-input miRNA library preparation kits. RNA Biology, 2020, 17, 1284-1292.	1.5	4
14	Effects of long-term feeding of rapeseed meal on skeletal muscle transcriptome, production efficiency and meat quality traits in Norwegian Landrace growing-finishing pigs. PLoS ONE, 2019, 14, e0220441.	1.1	21
15	Transcriptomes of antigen presenting cells in human thymus. PLoS ONE, 2019, 14, e0218858.	1.1	16
16	Exposure to Broad-Spectrum Visible Light Causes Major Transcriptomic Changes in Listeria monocytogenes EGDe. Applied and Environmental Microbiology, 2019, 85, .	1.4	1
17	Postovulatory maternal transcriptome in Atlantic salmon and its relation to developmental potential of embryos. BMC Genomics, 2019, 20, 315.	1.2	10
18	Towards population genomics in non-model species with large genomes: a case study of the marine zooplankton <i>Calanus finmarchicus</i> . Royal Society Open Science, 2019, 6, 180608.	1.1	34

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#	Article	IF	CITATIONS
19	Gene expression profiling of fast- and slow-growing non-functioning gonadotroph pituitary adenomas. European Journal of Endocrinology, 2018, 178, 295-307.	1.9	29
20	Monocytes accumulate in the airways of children with fatal asthma. Clinical and Experimental Allergy, 2018, 48, 1631-1639.	1.4	30
21	Deltamethrin resistance in the salmon louse, Lepeophtheirus salmonis (KrÃ,yer): Maternal inheritance and reduced apoptosis. Scientific Reports, 2018, 8, 8450.	1.6	13
22	Issues with RNA-seq analysis in non-model organisms: A salmonid example. Developmental and Comparative Immunology, 2017, 75, 38-47.	1.0	7
23	A c-Myb mutant causes deregulated differentiation due to impaired histone binding and abrogated pioneer factor function. Nucleic Acids Research, 2017, 45, 7681-7696.	6.5	28
24	Full genome survey and dynamics of gene expression in the greater amberjack Seriola dumerili. GigaScience, 2017, 6, 1-13.	3.3	16
25	A novel ultra high-throughput 16S rRNA gene amplicon sequencing library preparation method for the Illumina HiSeq platform. Microbiome, 2017, 5, 68.	4.9	93
26	A comparative study of ChIP-seq sequencing library preparation methods. BMC Genomics, 2016, 17, 816.	1.2	25
27	Contrasting transcriptome response to thermal stress in two key zooplankton species, Calanus finmarchicus and C. glacialis. Marine Ecology - Progress Series, 2015, 534, 79-93.	0.9	30
28	Thermal plasticity of the miRNA transcriptome during Senegalese sole development. BMC Genomics, 2014, 15, 525.	1.2	58
29	Diversification of the expanded teleost-specific toll-like receptor family in Atlantic cod, Gadus morhua. BMC Evolutionary Biology, 2012, 12, 256.	3.2	65
30	Positive selection pressure within teleost toll-like receptors tlr21 and tlr22 subfamilies and their response to temperature stress and microbial components in zebrafish. Molecular Biology Reports, 2012, 39, 8965-8975.	1.0	54
31	LSH and G9a/GLP complex are required for developmentally programmed DNA methylation. Genome Research, 2011, 21, 83-94.	2.4	108
32	Tetraploid Ancestry Provided Atlantic Salmon With Two Paralogue Functional T Cell Receptor Beta Regions Whereof One Is Completely Novel. Frontiers in Immunology, 0, 13, .	2.2	4