

Thejaswini Venkatesh

List of PR Articles by Year in descending order

Source: [//exaly.com/author-pdf/5187282/publications.pdf](https://exaly.com/author-pdf/5187282/publications.pdf)

Version: 2025-02-01

33

PR articles

513

PR citations

712385

12

PR h-index

649231

21

g-index

35

documents

549

doc citations

772842

12

h-index

1117

citing authors

#	ARTICLE	IF	PR CITATIONS
1	Study on the interaction of the bromodomain inhibitor JQ1 with human serum albumin by spectroscopic and molecular docking studies. <i>Journal of Molecular Structure</i> , 2023, 1273, 134374.	4.2	9
2	YBX1/lncRNA SBF2-AS1 interaction regulates proliferation and tamoxifen sensitivity via PI3K/AKT/MTOR signaling in breast cancer cells. <i>Molecular Biology Reports</i> , 2023, 50, 3413-3428.	2.6	20
3	Deciphering the role of MitomiRs in cancer: A comprehensive review. <i>Mitochondrion</i> , 2023, 70, 118-130.	4.1	8
4	LncRNAâ€“miRNAâ€“mRNA regulatory axes in endometrial cancer: a comprehensive overview. <i>Archives of Gynecology and Obstetrics</i> , 2022, 306, 1431-1447.	2.2	49
5	Review of: "Deep sequencing of yeast and mouse tRNAs and tRNA fragments using OTTR". <i>Qeios</i> , 2022, , .	3.6	0
6	Current Update on Nanotechnology-Based Approaches in Ovarian Cancer Therapy. <i>Reproductive Sciences</i> , 2022, 30, 335-349.	2.2	16
7	In silico identification of rice and human miRNAs sequences within tRFs of <i>Oryza sativa</i> and expression analysis of rice tRFs with their corresponding miRNAs. <i>South African Journal of Botany</i> , 2022, 149, 487-495.	2.7	0
8	Landscape of Clinically Relevant Exosomal tRNA-Derived Non-coding RNAs. <i>Molecular Biotechnology</i> , 2022, 65, 300-310.	2.1	12
9	Regulation of processing bodies: From viruses to cancer epigenetic machinery. <i>Cell Biology International</i> , 2021, 45, 708-719.	3.0	10
10	Long Non-coding RNA NEAT1 as an Emerging Biomarker in Breast and Gynecologic Cancers: a Systematic Overview. <i>Reproductive Sciences</i> , 2021, 28, 2436-2447.	2.2	14
11	Breast cancer fibroblasts and cross-talk. <i>Clinica Chimica Acta</i> , 2021, 521, 158-169.	1.6	10
12	Comparative expression analysis of tRF-3001a and tRF-1003 with corresponding miRNAs (miR-1260a and) Tj ETQq0 0 0 rgBT /Overlock 1 48, 7313-7324.	2.6	9
13	Deciphering piRNA biogenesis through cytoplasmic granules, mitochondria and exosomes. <i>Archives of Biochemistry and Biophysics</i> , 2020, 695, 108597.	2.9	26
14	Liquid biopsy in ovarian cancer. <i>Clinica Chimica Acta</i> , 2020, 510, 28-34.	1.6	21
15	In silico analysis of COSMIC retrieved P body gene mutations in breast cancer. <i>Gene Reports</i> , 2020, 19, 100617.	0.7	1
16	In silico prediction of UCLH1 disease-causing SNPs and its effects on protein stability. <i>Gene Reports</i> , 2020, 19, 100677.	0.7	0
17	Regulated expression of Gemin5, Xrn1, Cpeb and Stau1 in the uterus and ovaries after superovulation and the effect of exogenous estradiol and leptin in rodents. <i>Molecular Biology Reports</i> , 2019, 46, 2533-2540.	2.6	1
18	PTPH1 immunohistochemical expression and promoter methylation in breast cancer patients from India: A retrospective study. <i>Journal of Cellular Physiology</i> , 2019, 234, 1071-1079.	4.2	5

#	ARTICLE	IF	PR CITATIONS
19	Increased virulence in <i>Pseudomonas aeruginosa</i> at pathological glucose levels. Infectious Diseases, 2019, 51, 153-156.	1.8	5
20	Gene expression changes and promoter methylation with the combined effects of estradiol and leptin in uterine tissue of the ovariectomized mice model of menopause. Molecular Biology Reports, 2019, 47, 151-168.	2.6	6
21	YBX1 at the crossroads of non-coding transcriptome, exosomal, and cytoplasmic granular signaling. European Journal of Cell Biology, 2018, 97, 163-167.	4.0	85
22	Next-generation sequencing for endocrine cancers: Recent advances and challenges. Tumor Biology, 2017, 39, 101042831769837.	1.1	8
23	Exploration of acute genotoxic effects and antigenotoxic potential of gambogic acid using <i>Allium cepa</i> assay. Plant Physiology and Biochemistry, 2017, 118, 643-652.	5.5	9
24	tRFs: miRNAs in disguise. Gene, 2016, 579, 133-138.	2.4	90
25	Mining of single nucleotide polymorphisms in the 3' untranslated region of liver cancer-implicated miR-122 target genes. Annals of Translational Medicine, 2016, 4, 102-102.	1.8	2
26	Non-coding RNAs: Functions and applications in endocrine-related cancer. Molecular and Cellular Endocrinology, 2015, 416, 88-96.	3.5	35
27	HER2 and uPAR cooperativity contribute to metastatic phenotype of HER2-positive breast cancer. Oncoscience, 2015, 2, 207-224.	1.3	26
28	New insights into the genetic basis of infertility. The Application of Clinical Genetics, 2014, , 235.	2.7	30
29	Emerging roles of MCPH1: Expedition from primary microcephaly to cancer. European Journal of Cell Biology, 2014, 93, 98-105.	4.0	28
30	Exploration of deleterious single nucleotide polymorphisms in the components of human P bodies: An in silico approach. Gene, 2013, 528, 360-363.	2.4	4
31	Computational interrogation of cis-regulatory elements of genes that are common targets of luteotropin and luteolysin in the primate corpus luteum. Gene, 2013, 515, 403-409.	2.4	1
32	Primary Microcephaly Gene MCPH1 Shows Signatures of Tumor Suppressors and Is Regulated by miR-27a in Oral Squamous Cell Carcinoma. PLoS ONE, 2013, 8, e54643.	2.4	44
33	Single nucleotide polymorphisms in genes that are common targets of luteotropin and luteolysin in primate corpus luteum: Computational exploration. Gene, 2012, 511, 353-357.	2.4	5