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Comprehensive landscape of subtype-specific coding and non-coding RNA transcripts in breast cancer

DOI: 10.18632/oncotarget.11998
Oncotarget, 2016, 7, 68851-68863.

Source: <https://exaly.com/paper-pdf/63308477/citation-report.pdf>

Version: 2024-04-28

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#	Paper	IF	Citations
18	DNA methylation alterations induced by transient exposure of MCF-7 cells to maghemite nanoparticles. <i>Nanomedicine</i> , 2017 , 12, 2637-2649	5.6	4
17	A Ten-Long Non-Coding RNA Model Improves Prognosis Prediction of Hepatocellular Carcinoma Patients. 2018 ,		0
16	lncRNA Gene Signatures for Prediction of Breast Cancer Intrinsic Subtypes and Prognosis. <i>Genes</i> , 2018 , 9,	4.2	26
15	Breast cancer genome analysis in time and space: biomarker development strategy. <i>Biomarkers in Medicine</i> , 2018 , 12, 547-550	2.3	1
14	A long non-coding RNA signature to improve prognostic prediction in clear cell renal cell carcinoma. <i>Biomedicine and Pharmacotherapy</i> , 2019 , 118, 109079	7.5	14
13	is a critical oncogenic driver and contributes to epithelial-mesenchymal transition in papillary thyroid carcinoma. <i>Cancer Management and Research</i> , 2019 , 11, 2059-2072	3.6	19
12	Identification of lncRNAs associated with early-stage breast cancer and their prognostic implications. <i>Molecular Oncology</i> , 2019 , 13, 1342-1355	7.9	28
11	Upregulated lncRNA-CCAT1 promotes hepatocellular carcinoma progression by functioning as miR-30c-2-3p sponge. <i>Cell Biochemistry and Function</i> , 2019 , 37, 84-92	4.2	20
10	Dynamic genome and transcriptional network-based biomarkers and drugs: precision in breast cancer therapy. <i>Medicinal Research Reviews</i> , 2019 , 39, 1205-1227	14.4	15
9	Are long non-coding RNAs involved in the interaction circuit between estrogen receptor and vitamin D receptor?. <i>Meta Gene</i> , 2019 , 19, 1-9	0.7	3
8	DSCAM-AS1-Driven Proliferation of Breast Cancer Cells Involves Regulation of Alternative Exon Splicing and 3eEnd Usage. <i>Cancers</i> , 2020 , 12,	6.6	12
7	The transcriptome-wide landscape of molecular subtype-specific mRNA expression profiles in acute myeloid leukemia. <i>American Journal of Hematology</i> , 2021 , 96, 580-588	7.1	2
6	A five-long non-coding RNA signature to improve prognosis prediction of clear cell renal cell carcinoma. <i>Oncotarget</i> , 2017 , 8, 58699-58708	3.3	26
5	Identification of lncRNAs associated with early stage breast cancer and their prognostic implications.		
4	A comprehensive study of mRNA and long noncoding RNAs in Indian Breast cancer patients using transcriptomics approach.		
3	Discovery of druggable cancer-specific pathways with application in acute myeloid leukemia. 2022 , 11,		0
2	Transcriptomic profiling of Indian breast cancer patients revealed subtype-specific mRNA and lncRNA signatures. 13,		0

- 1 A Comprehensive Landscape of Imaging Feature-Associated RNA Expression Profiles in Human Breast Tissue. **2023**, 23, 1432

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