

Predicting effective microRNA target sites in mammalia

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Citation Report

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1	The evolution of Homo sapiens denisova and Homo sapiens neanderthalensis miRNA targeting genes in the prenatal and postnatal brain. BMC Genomics, 2015, 16, S4.	1.2	3
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20	Identification of More Feasible MicroRNA-mRNA Interactions within Multiple Cancers Using Principal Component Analysis Based Unsupervised Feature Extraction. <i>International Journal of Molecular Sciences</i> , 2016, 17, 696.	1.8	35
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1810	Upregulation of miR-9 and miR-193b over human Th17 cell differentiation. <i>Molecular Genetics & Genomic Medicine</i> , 2020, 8, e1538.	0.6	5
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1812	Contribution of miRNAs, tRNAs and tRFs to Aberrant Signaling and Translation Deregulation in Lung Cancer. <i>Cancers</i> , 2020, 12, 3056.	1.7	7
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1814	High CENPM mRNA expression and its prognostic significance in hepatocellular carcinoma: a study based on data mining. <i>Cancer Cell International</i> , 2020, 20, 406.	1.8	7
1815	The impact of miR-9 in osteosarcoma. <i>Medicine (United States)</i> , 2020, 99, e21902.	0.4	5
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1818	Identification of circular RNA hsa_circ_0044556 and its effect on the progression of colorectal cancer. <i>Cancer Cell International</i> , 2020, 20, 427.	1.8	13
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1836	STAT3-Induced lncRNA SNHG17 Exerts Oncogenic Effects on Ovarian Cancer through Regulating CDK6. <i>Molecular Therapy - Nucleic Acids</i> , 2020, 22, 38-49.	2.3	24
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2647	Profound downregulation of neural transcription factor Npas4 and Nr4a family in fetal mice neurons infected with Zika virus. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009425.	1.3	5
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2671	TSCCA: A tensor sparse CCA method for detecting microRNA-gene patterns from multiple cancers. <i>PLoS Computational Biology</i> , 2021, 17, e1009044.	1.5	10
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2690	Tumor-Derived Extracellular Vesicles Promote Activation of Carcinoma-Associated Fibroblasts and Facilitate Invasion and Metastasis of Ovarian Cancer by Carrying miR-630. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 652322.	1.8	33
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2698	A microRNA expression and regulatory element activity atlas of the mouse immune system. <i>Nature Immunology</i> , 2021, 22, 914-927.	7.0	19
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2700	Regulatory Mechanisms and Functional Roles of Hypoxia-Induced Long Non-Coding RNA MTORT1 in Breast Cancer Cells. <i>Frontiers in Oncology</i> , 2021, 11, 663114.	1.3	10
2701	Immune System and Neuroinflammation in Idiopathic Parkinson's Disease: Association Analysis of Genetic Variants and miRNAs Interactions. <i>Frontiers in Genetics</i> , 2021, 12, 651971.	1.1	8
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2708	miR-130b and miR-128a are essential lineage-specific codrivers of t(4;11) MLL-AF4 acute leukemia. <i>Blood</i> , 2021, 138, 2066-2092.	0.6	19
2709	An Integrative Transcriptomic and Methylation Approach for Identifying Differentially Expressed Circular RNAs Associated with DNA Methylation Change. <i>Biomedicines</i> , 2021, 9, 657.	1.4	6
2711	Identifying miRNA-mRNA Integration Set Associated With Survival Time. <i>Frontiers in Genetics</i> , 2021, 12, 634922.	1.1	1
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2714	Identification of Potential ceRNA Network and Patterns of Immune Cell Infiltration in Systemic Sclerosis-Associated Interstitial Lung Disease. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 622021.	1.8	8
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