

DIANA miRPath v.2.0: investigating the combinatorial e

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Tuning the engine. <i>RNA Biology</i> , 2012, 9, 1224-1232.	1.5	12
2	MicroRNA-mediated regulation of target genes in several brain regions is correlated to both microRNA-targeting-specific promoter methylation and differential microRNA expression. <i>BioData Mining</i> , 2013, 6, 11.	2.2	15
3	Signalink 2 – a signaling pathway resource with multi-layered regulatory networks. <i>BMC Systems Biology</i> , 2013, 7, 7.	3.0	169
4	A proteomic approach of pediatric astrocytomas: MiRNAs and network insight. <i>Journal of Proteomics</i> , 2013, 94, 162-175.	1.2	20
5	MicroRNA signatures in hereditary breast cancer. <i>Breast Cancer Research and Treatment</i> , 2013, 142, 19-30.	1.1	40
6	Extremely Complex Populations of Small RNAs in the Mouse Retina and RPE/Choroid. , 2013, 54, 8140.		22
7	Using context-specific effect of miRNAs to identify functional associations between miRNAs and gene signatures. <i>BMC Bioinformatics</i> , 2013, 14, S1.	1.2	50
8	An emerging role for microRNAs in sexually dimorphic neurobiological systems. <i>Pflugers Archiv European Journal of Physiology</i> , 2013, 465, 655-667.	1.3	22
9	Interactions of miR-323/miR-326/miR-329 and miR-130a/miR-155/miR-210 as prognostic indicators for clinical outcome of glioblastoma patients. <i>Journal of Translational Medicine</i> , 2013, 11, 10.	1.8	162
10	Over-expression of the miRNA cluster at chromosome 14q32 in the alcoholic brain correlates with suppression of predicted target mRNA required for oligodendrocyte proliferation. <i>Gene</i> , 2013, 526, 356-363.	1.0	31
11	miR-143 decreases COX-2 mRNA stability and expression in pancreatic cancer cells. <i>Biochemical and Biophysical Research Communications</i> , 2013, 439, 6-11.	1.0	64
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15	Signature MicroRNA expression patterns identified in humans with 22q11.2 deletion/DiGeorge syndrome. <i>Clinical Immunology</i> , 2013, 147, 11-22.	1.4	58
16	Opposing actions of environmental enrichment and Alzheimer's disease on the expression of hippocampal microRNAs in mouse models. <i>Translational Psychiatry</i> , 2013, 3, e304-e304.	2.4	73
17	Identification of active transcription factor and miRNA regulatory pathways in Alzheimer's disease. <i>Bioinformatics</i> , 2013, 29, 2596-2602.	1.8	66
18	MicroRNA-322 (miR-322) and Its Target Protein Tob2 Modulate Osterix (Osx) mRNA Stability. <i>Journal of Biological Chemistry</i> , 2013, 288, 14264-14275.	1.6	77

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20	Hypoxia Induces Mesenchymal Gene Expression in Renal Tubular Epithelial Cells: An in vitro Model of Kidney Transplant Fibrosis. <i>Nephron Extra</i> , 2013, 3, 50-58.	1.1	34
21	Differential Expression of miR-145 in Children with Kawasaki Disease. <i>PLoS ONE</i> , 2013, 8, e58159.	1.1	60
22	In-Depth Characterization of microRNA Transcriptome in Melanoma. <i>PLoS ONE</i> , 2013, 8, e72699.	1.1	109
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