

# DIANA-microT web server v5.0: service integration into workflows

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Regulation of the unfolded protein response by microRNAs. Cellular and Molecular Biology Letters, 2013, 18, 555-78.	7.0	49
2	Genome-Wide Analysis of miRNA Signature in the APP <sup>swe</sup> /PS1 <sup>E9</sup> Mouse Model of Alzheimer's Disease. PLoS ONE, 2014, 9, e101725.	2.5	23
3	Pathway Analysis of MicroRNA Expression Profile during Murine Osteoclastogenesis. PLoS ONE, 2014, 9, e107262.	2.5	35
4	Frequent Co-Expression of miRNA-5p and -3p Species and Cross-Targeting in Induced Pluripotent Stem Cells. International Journal of Medical Sciences, 2014, 11, 824-833.	2.5	36
5	MR-microT. , 2014, , .		9
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9	MicroRNA Dysregulation, Gene Networks, and Risk for Schizophrenia in 22q11.2 Deletion Syndrome. Frontiers in Neurology, 2014, 5, 238.	2.4	42
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17	Resources for Small Regulatory RNAs. Current Protocols in Molecular Biology, 2014, 107, 19.8.1-14.	2.9	2
18	New Approaches to Comparative and Animal Stress Biology Research in the Post-genomic Era: A Contextual Overview. Computational and Structural Biotechnology Journal, 2014, 11, 138-146.	4.1	8

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20	An integrated analysis of the SOX2 microRNA response program in human pluripotent and nullipotent stem cell lines. <i>BMC Genomics</i> , 2014, 15, 711.	2.8	19
21	MicroRNA-375 inhibits colorectal cancer growth by targeting PIK3CA. <i>Biochemical and Biophysical Research Communications</i> , 2014, 444, 199-204.	2.1	103
22	Computational Methods for MicroRNA Target Prediction. <i>Genes</i> , 2014, 5, 671-683.	2.4	92
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