

Specific miRNA signatures are associated with metastas renal cell carcinoma

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Citation Report

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1	microRNA-30c negatively regulates endometrial cancer cells by targeting metastasis-associated gene-1. <i>Oncology Reports</i> , 2012, 27, 807-12.	1.2	53
2	Reference genes for the relative quantification of microRNAs in renal cell carcinomas and their metastases. <i>Analytical Biochemistry</i> , 2011, 417, 233-241.	1.1	78
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4	A Lasso regression model for the construction of microRNA-target regulatory networks. <i>Bioinformatics</i> , 2011, 27, 2406-2413.	1.8	110
5	miRNA profiling in metastatic renal cell carcinoma reveals a tumour-suppressor effect for miR-215. <i>British Journal of Cancer</i> , 2011, 105, 1741-1749.	2.9	152
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7	Need for Rigor in Design, Reporting, and Interpretation of Transcriptomic Biomarker Studies. <i>Journal of Clinical Microbiology</i> , 2012, 50, 4190-4191.	1.8	0
8	Identification of Metastamirs as Metastasis-associated MicroRNAs in Clear Cell Renal Cell Carcinomas. <i>International Journal of Biological Sciences</i> , 2012, 8, 1363-1374.	2.6	92
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20	MicroRNA-30a inhibits epithelial-to-mesenchymal transition by targeting Snai1 and is downregulated in non-small cell lung cancer. <i>International Journal of Cancer</i> , 2012, 130, 2044-2053.	2.3	266
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