

Table S10 Details of raw CEL (microarray chip) expression data (obtained from GEO database) have been used in present study. This data are obtained from studies on breast, colorectal, endometrial, gastric, liver, lung, ovarian, pancreatic, prostate, testicular, bladder, intestine neuroendocrine, cervical and renal cancers as well as glioblastoma

	Accession numbers	Cancer name	Study
^a Transcript microarray data	GSE17025	Endometrial cancer	Day <i>et al.</i> , 2011 ¹
	GSE26910	Breast and prostate cancer	Planche <i>et al.</i> , 2011 ²
	GSE18520	Ovarian cancer	Mok <i>et al.</i> , 2009 ³
	GSE15220	Testicular cancer	Cheung <i>et al.</i> , 2010 ⁴
	GSE29721	Liver cancer	Stefanska <i>et al.</i> , 2011 ⁵
	GSE19826	Gastric cancer	Wang <i>et al.</i> , 2012 ⁶
	GSE22780	Pancreatic adenocarcinoma	Balasenthil <i>et al.</i> , 2010 ⁷
	GSE23878	Colorectal cancer	Uddin <i>et al.</i> , 2011 ⁸
	GSE15824	Glioblastoma	Grzmil <i>et al.</i> , 2011 ⁹
	GSE18842	Lung adenocarcinoma	Landi <i>et al.</i> , 2008 ¹⁰
^a MicroRNA microarray data	GSE23383	Ovarian cancer	Shahab <i>et al.</i> , 2011 ¹¹
	GSE23022	Prostate carcinoma	Wach <i>et al.</i> , 2012 ¹²
	GSE39015	Cervical cancer	Hall <i>et al.</i> , 2012 ¹³
	GSE41282	Renal cancer	Wach <i>et al.</i> , 2012
	GSE39093	Bladder tumor	Xu <i>et al.</i> , 2013 ¹⁴
	GSE33568	Intestinal neuroendocrine tumor	Li <i>et al.</i> , 2013 ¹⁵

^aWe analyzed total of cancerous cel files vs. total of normal s in each study

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