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Erratum: A cross comparison of technologies for the detection of microRNAs in clinical FFPE samples of hepatoblastoma patients

Aniruddha Chatterjee, Anna L. Leichter, Vicky Fan, Peter Tsai, Rachel V. Purcell, Michael J. Sullivan & Michael R Eccles

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In this Article, the Accession codes were omitted from the Additional Information section:

Data availability: Accession codes: All the data generated and described in this article have been submitted to the NCBI Gene Expression Omnibus (GEO; <http://www.ncbi.nlm.nih.gov/geo/>). The NGS data were submitted under accession number GSE62010 (URL: <http://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE62017>), The microarray data were submitted under accession number GSE62011 (URL: <http://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE62011>) and the NanoString data were submitted under accession number GSE62017 (URL: <http://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE62017>). These three accession numbers are linked as super series.



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