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

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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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