Table S2

Normality assessment of log-transformed expression data of candidate reference genes by Shapiro-Wilk test

Gen Symbol	W	p-value
Candidate reference genes for microRNA normalization		
5S	0.9494	0.1088
miR-16	0.9683	0.3985
miR-103	0.9839	0.8769
miR-191	0.9771	0.6642
miR-Let7a	0.9576	0.193
RNU48	0.9634	0.2877
Candidate reference genes for mRNA normalization		
18S	0.9817	0.814
B2M	0.9898	0.9825
HPRT1	0.967	0.3672
SDHA	0.9817	0.8118

W: the value of the Shapiro-Wilk statistic, p.value: an approximate p-value for the test.