

Supplementary Files:

468 **Figure S1.** Potential microRNA biomarkers used for differentiating MPM from other
PE 469 causing diseases were identified in PE supernatants by OpenArray profiling. The ex-
pression 470 profiles for (a) miR-186, (b) miR-29a, (c) miR-31 and (d) miR-342-3p did not vali-
date when 471 the microRNAs were measured by qRT-PCR. MicroRNA expression was
plotted on a 472 logarythmic scale and U6snRNA was used as an endogenous control. The
line within the 473 boxes represents the median values and the top and bottom of the
boxes indicates the 474 interquartile ranges. The whiskers demonstrate the upper and lower ad-
jacent values for each 475 sample group. ** $p < 0.01$. * $p < 0.05$.

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Figure s1

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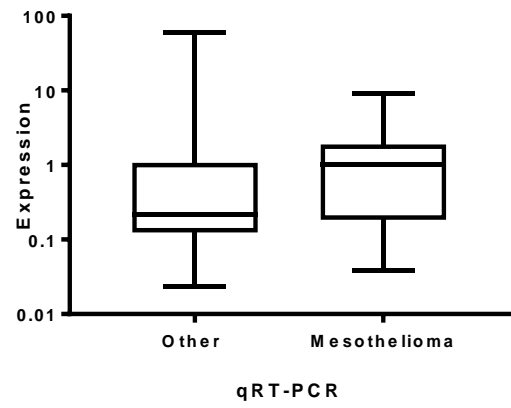
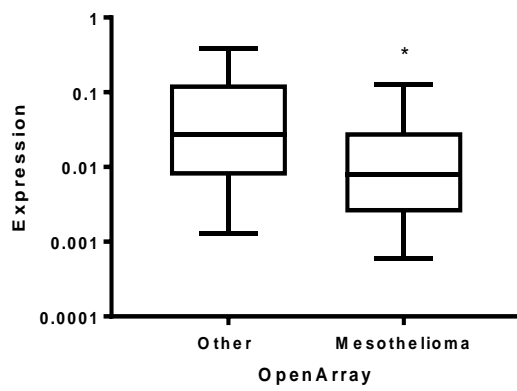
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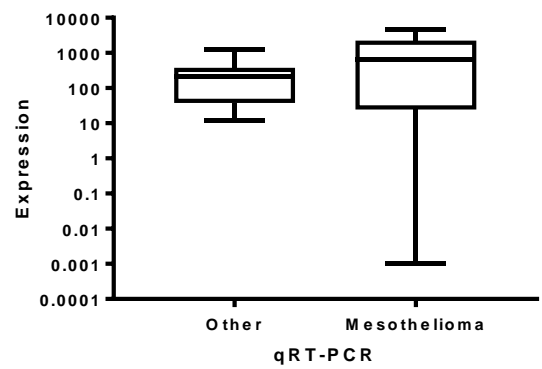
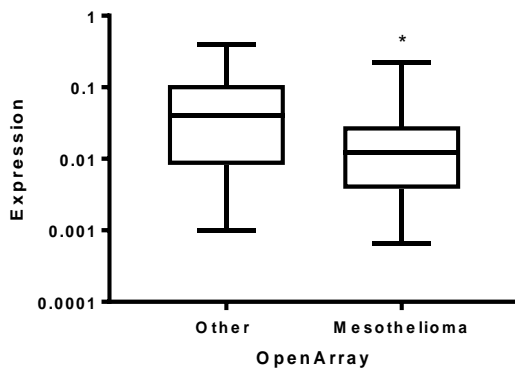
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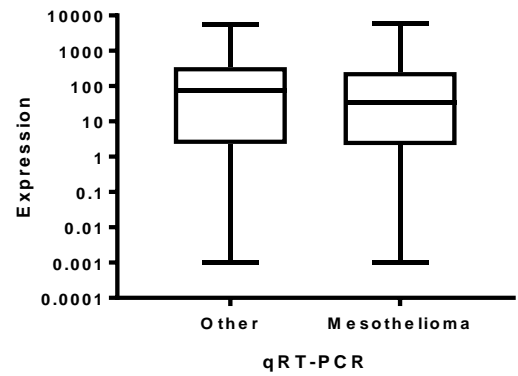
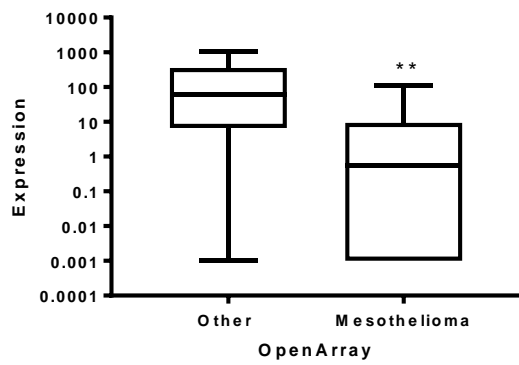
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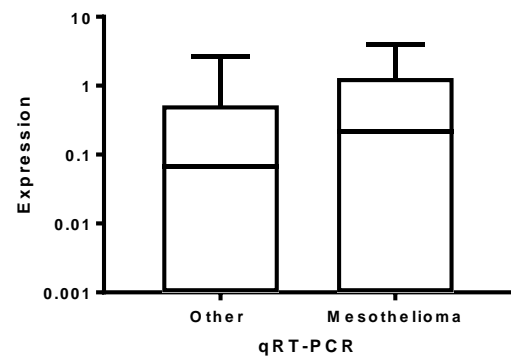
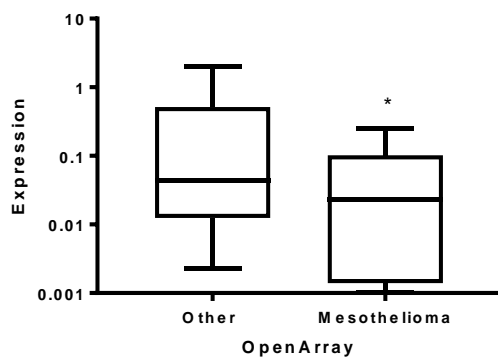
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