Supplemental Figure 1. Validation of TLDA findings using individual RT-qPCR arrays.
Expression profiles of the four significantly altered miRNAs identified in plasma from women without (Con.) and with (Endo.) endometriosis (n = 8 in each group) across the three phases of the menstrual cycle. The panels on the left show all samples combined irrespective of menstrual cycle phase whereas panels on the right show individual menstrual cycle phase comparisons, where T1 = early proliferative phase, T2 = late proliferative phase and T3 = mid luteal phase. Data are presented as relative expression values normalised to spiked in cel-miR-54. The median (middle line) and interquartile ranges (top and bottom lines) are shown while statistical significance was inferred when $P < 0.05$. * = $P < 0.05$, ** = $P \leq 0.01$, *** = $P \leq 0.001$ and NS = non-significant.