

Table S1. Ranking of the most stable miRNAs according to the Normfinder algorithm.

miRNA	C_t (CONTROL) (mean \pm SEM)	C_t (DFD) (mean \pm SEM)	p-value
mmu-let-7d	28.55 \pm 0.7	28.96 \pm 0.8	0.313
hsa-miR-125b	24.79 \pm 0.7	25.03 \pm 0.9	0.580
bta-miR-425-5p	29.93 \pm 0.8	30.25 \pm 0.9	0.486
bta-miR-660	27.87 \pm 0.8	28.22 \pm 0.8	0.430
hsa-miR-148b	28.93 \pm 0.6	29.02 \pm 0.8	0.814
hsa-miR-151-3p	28.55 \pm 0.7	28.96 \pm 0.8	0.640
hsa-miR-10b	24.67 \pm 0.7	25.04 \pm 0.9	0.417
bta-miR-342	26.41 \pm 0.6	26.76 \pm 0.8	0.348

DFD: dark, firm and dry.