Supplementary Figure S2 Transcriptional changes associated with *in vitro* ovulation. Analysis of follicles before and after hCG treatment showing qRT–PCR results of transcripts associated with *in vitro* ovulation: progesterone receptor (*Pgr*), tumor necrosis factor, alpha-induced protein 6 (*Tnfaip6*), prostaglandin-endoperoxidase protein synthase 2 (*Ptgs2*), plasminogen activator, urokinase (*Plau*) and a disintegrin and metalloproteinase with thrombospondin type 1 motif 1 (*Adamts-1*). Transcript abundance is expressed as fold change relative to pre-hCG follicles (FC = 1). Data are expressed as mean ± SEM, *n* = 3 cultures; *P < 0.05, **P < 0.01, ***P < 0.001 according to one-way ANOVA followed by the Bonferroni's Multiple Comparison Test.