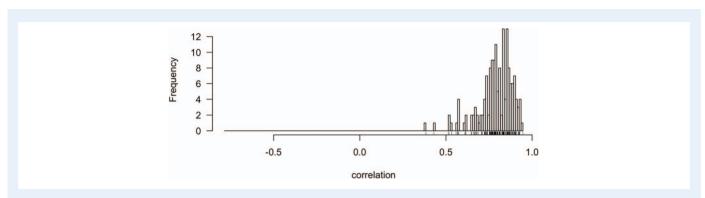
doi:10.1093/humrep/dez279

human SUPPLEMENTARY DATA



**Supplementary Figure S3 Correlation of expression within samples.** The correlation of endometrial gene expression within samples in RNA-Seq and microarray. The correlation was performed in 161 samples using 2162 genes that showed high correlation in both platforms (R > 0.7). The height of the bars represents the frequency of samples with that correlation value.