

Supplemental figure for:

**Differential Effects of the Hormonal and Copper Intrauterine Device
on the Endometrial Transcriptome**

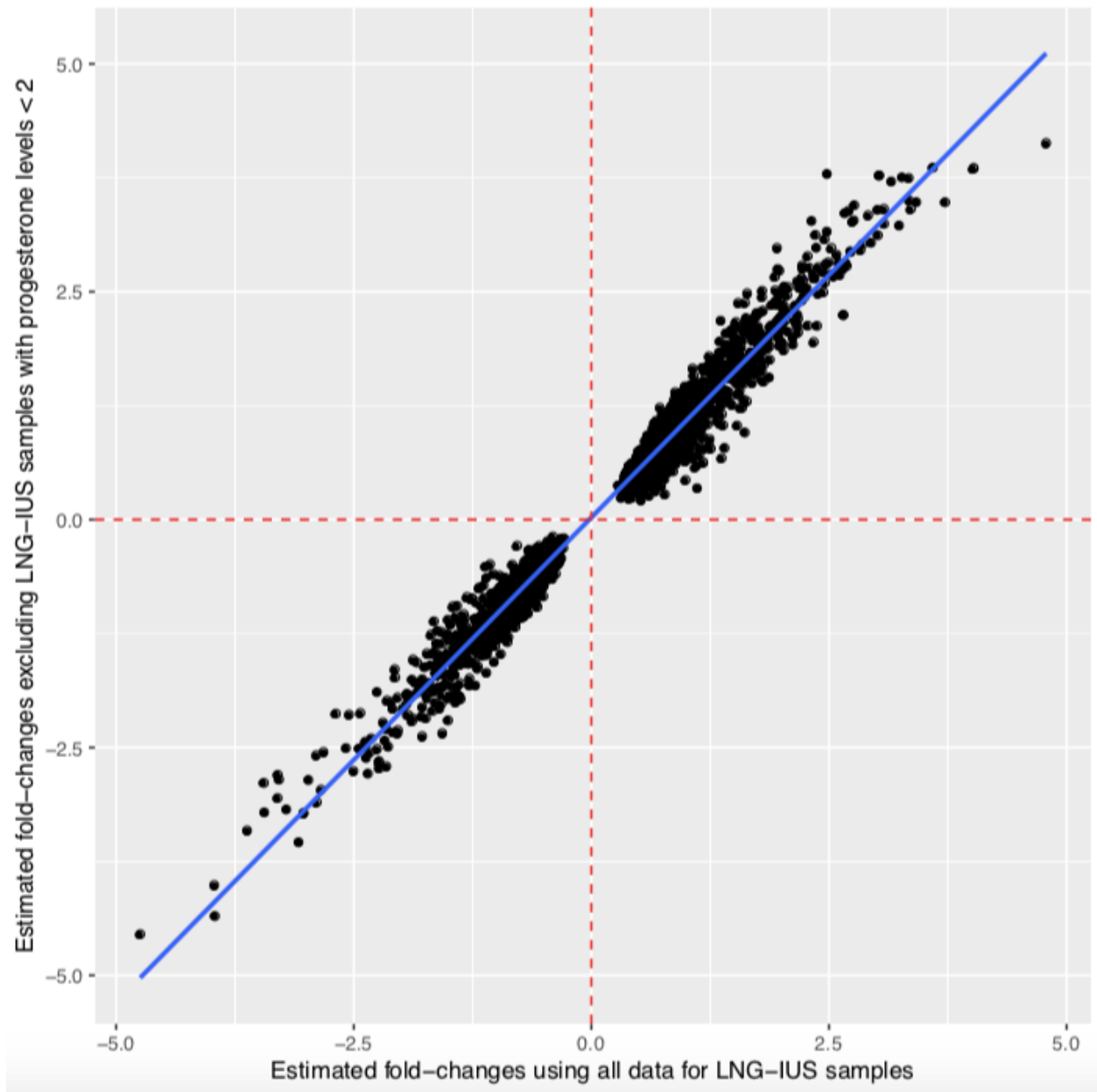
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Supplemental Figure Legend: Concordance of results of gene expression in LNG-IUS samples overall with results from ovulatory samples.

Scatter plot with best fit line demonstrates the estimated fold-changes of the expression of 2509 genes identified using all the data (x-axis) plotted against the estimated fold-changes of expression of these genes using data for only those LNG-IUS samples who had ovulated (progesterone >2) (y-axis).