

**Bioinformatic analysis reveals the importance of epithelial-mesenchymal  
transition in the development of endometriosis**

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## **Supplementary Figure Legends**

### **Supplementary Figure 1. Standardisation of gene expression in GSE7305 dataset.**

Graphs showing the data before (left) and after (right) normalisation.

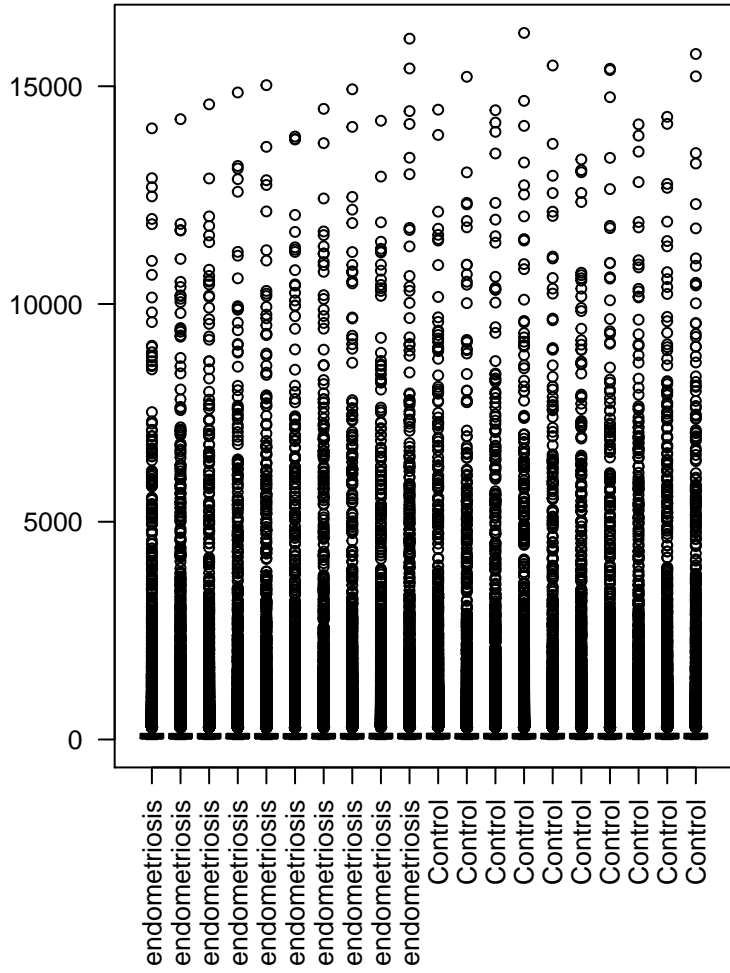
### **Supplementary Figure 2. Standardisation of gene expression in GSE11691**

**dataset.** Graphs showing the data before (left) and after (right) normalisation.

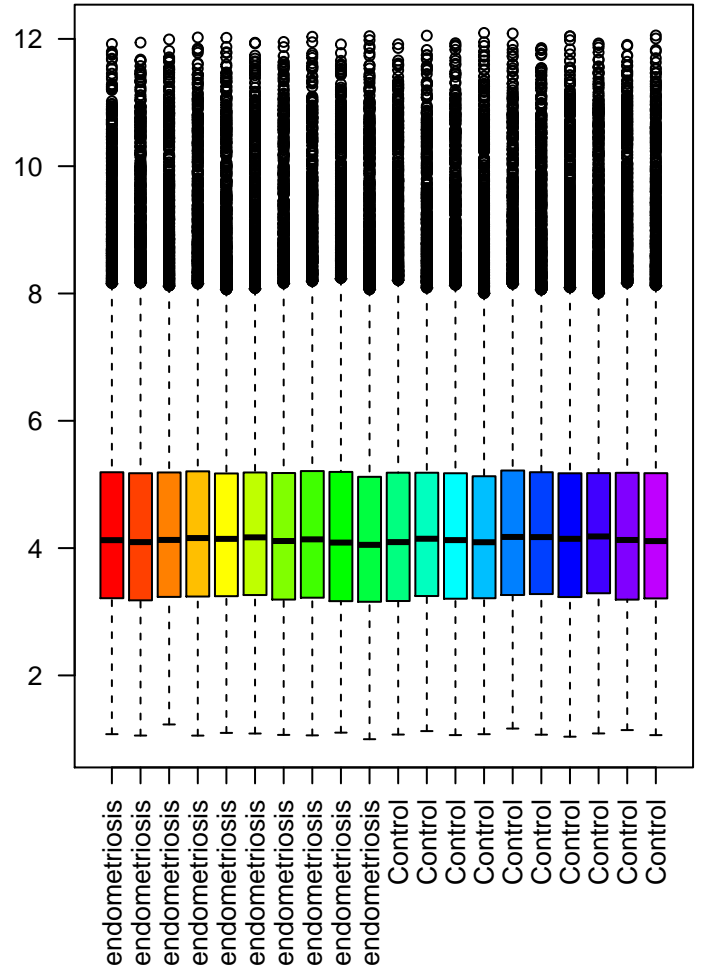
### **Supplementary Figure 3. Standardisation of gene expression in GSE12768**

**dataset.** Graphs showing the data before (left) and after (right) normalisation.

expression value

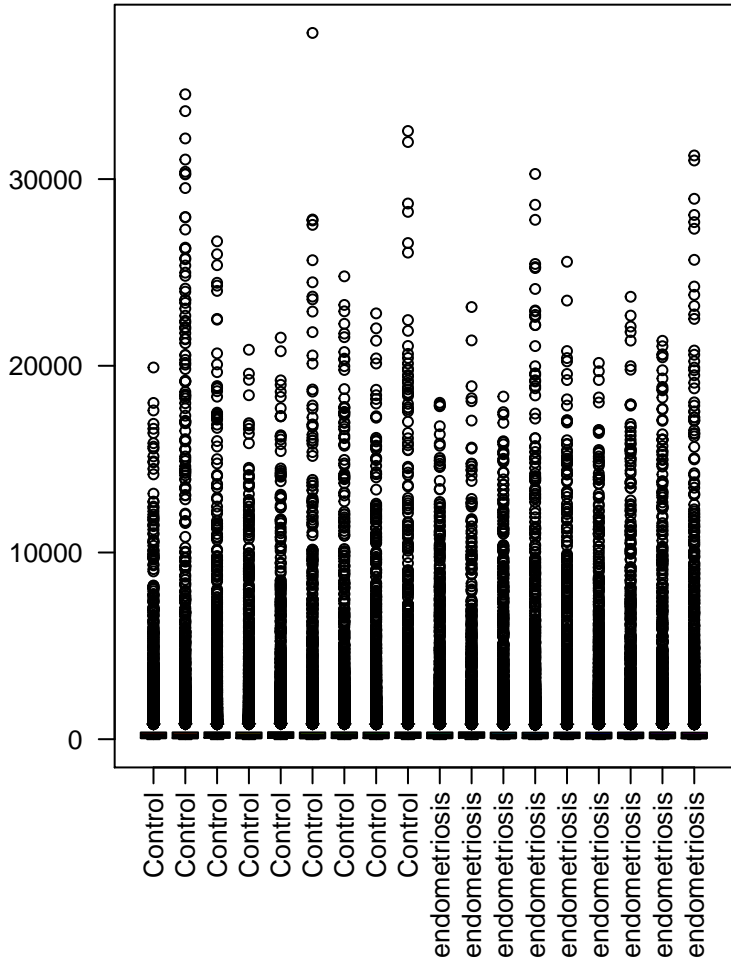


expression value

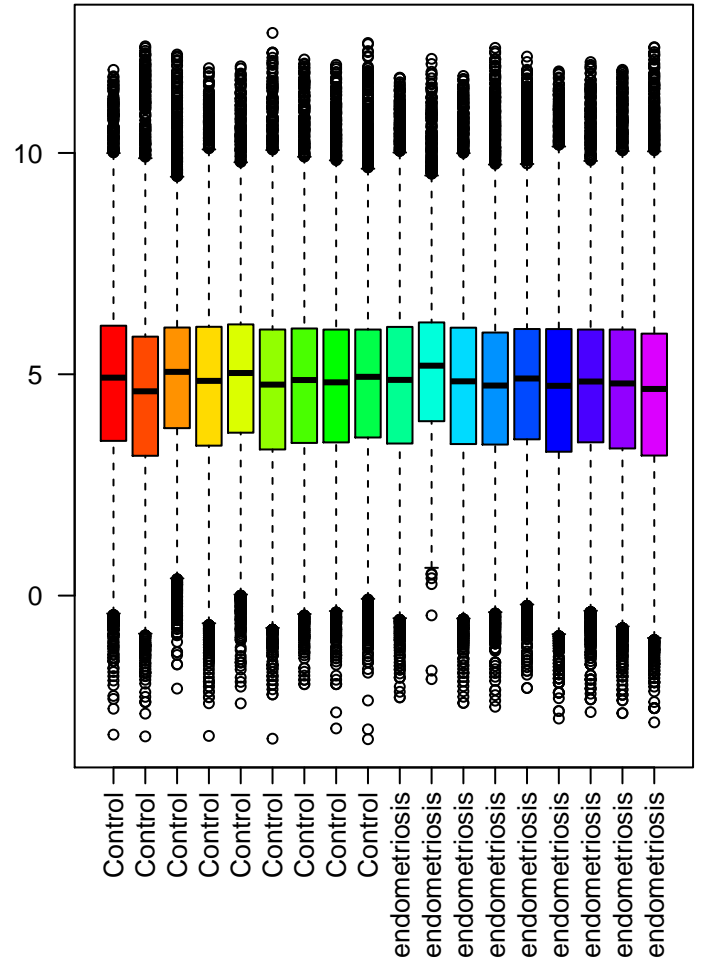


Supplementary Figure 1. Standardisation of gene expression in GSE7305 dataset.

expression value

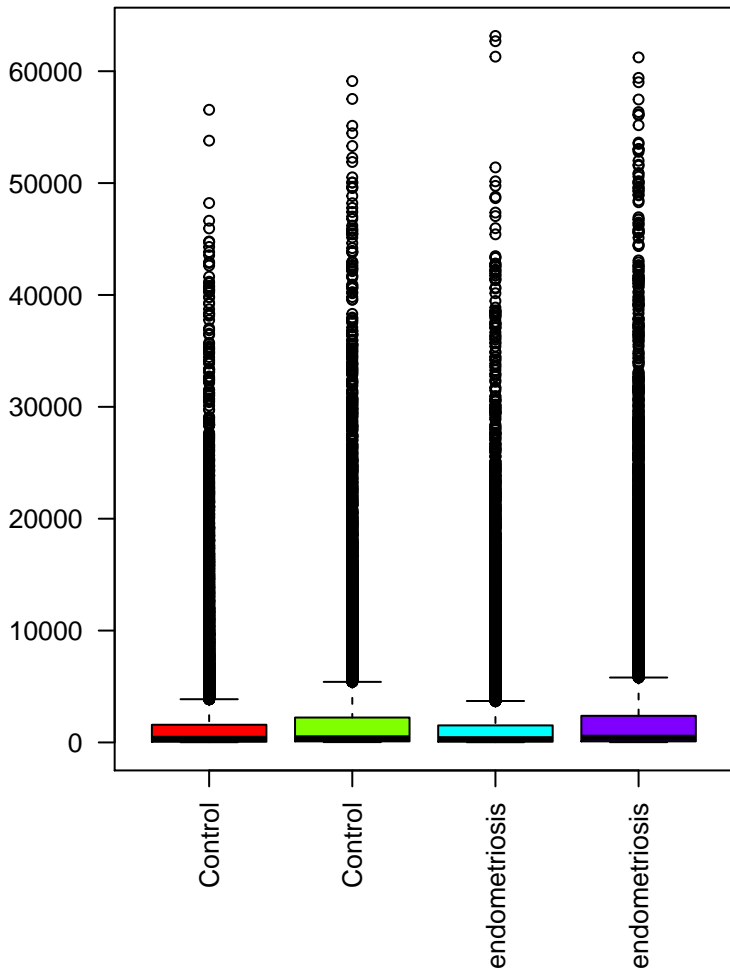


expression value

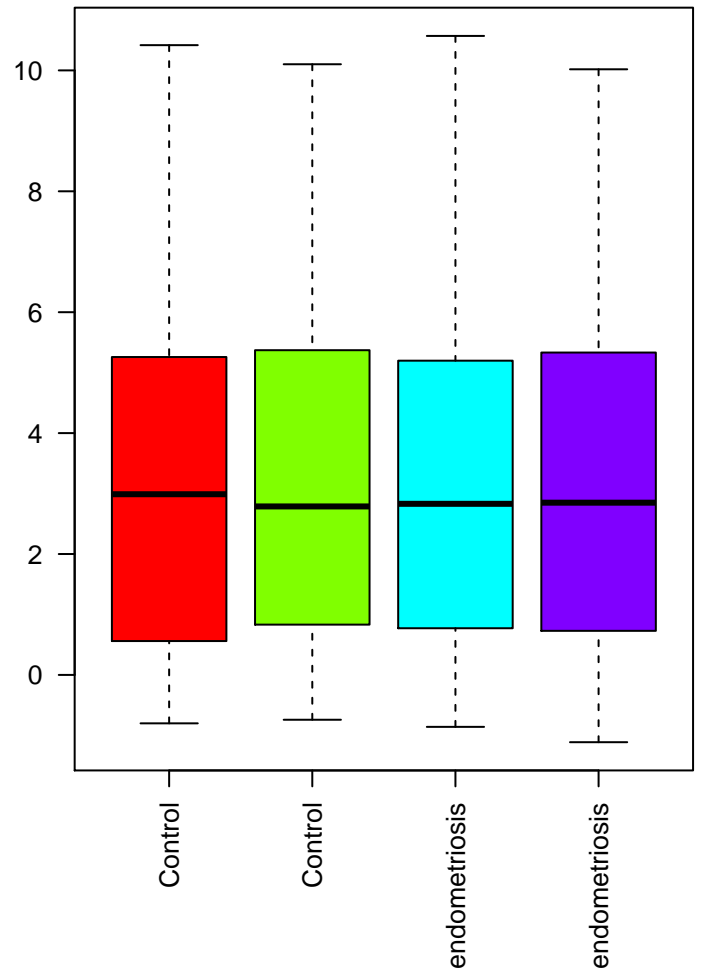


Supplementary Figure 2. Standardisation of gene expression in GSE11691 dataset.

expression value



expression value



Supplementary Figure 3. Standardisation of gene expression in GSE12768 dataset.