

Supplementary Information for Integrated Bioinformatic Analysis of Differentially Expressed Genes Associated with Wound Healing

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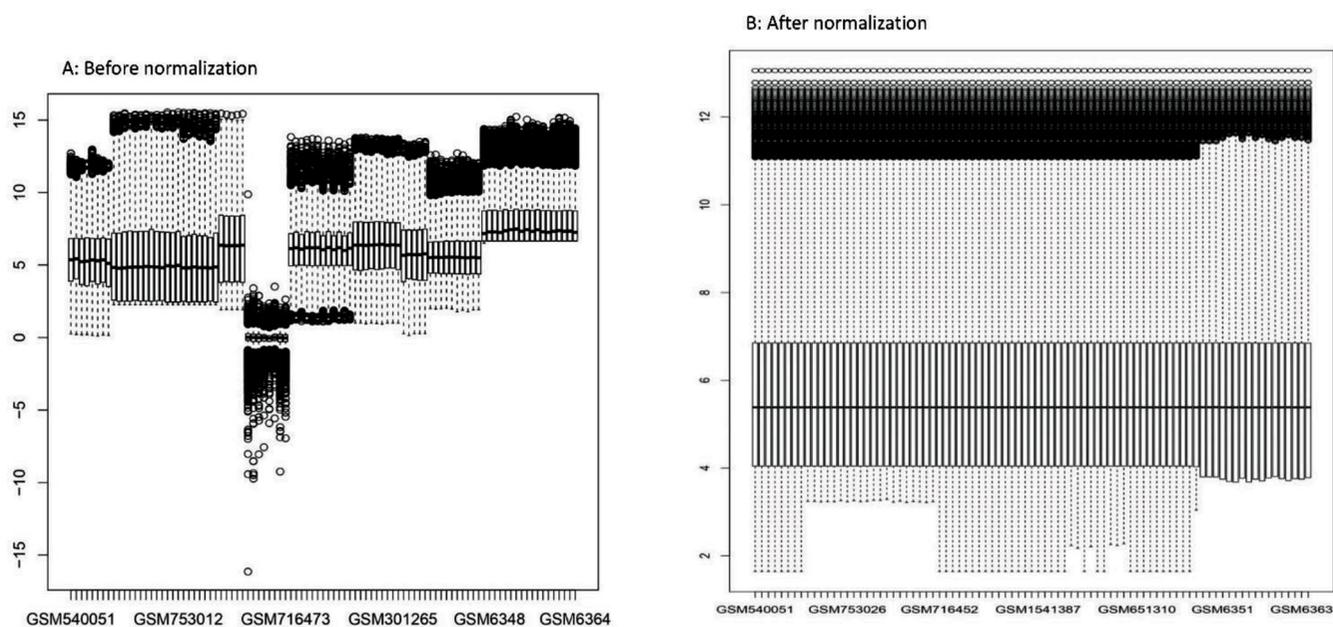


Fig.S1: Box plots of selected gene expression data before and after normalization. Similar medians of expression levels confirm that normalization was carried out properly.

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