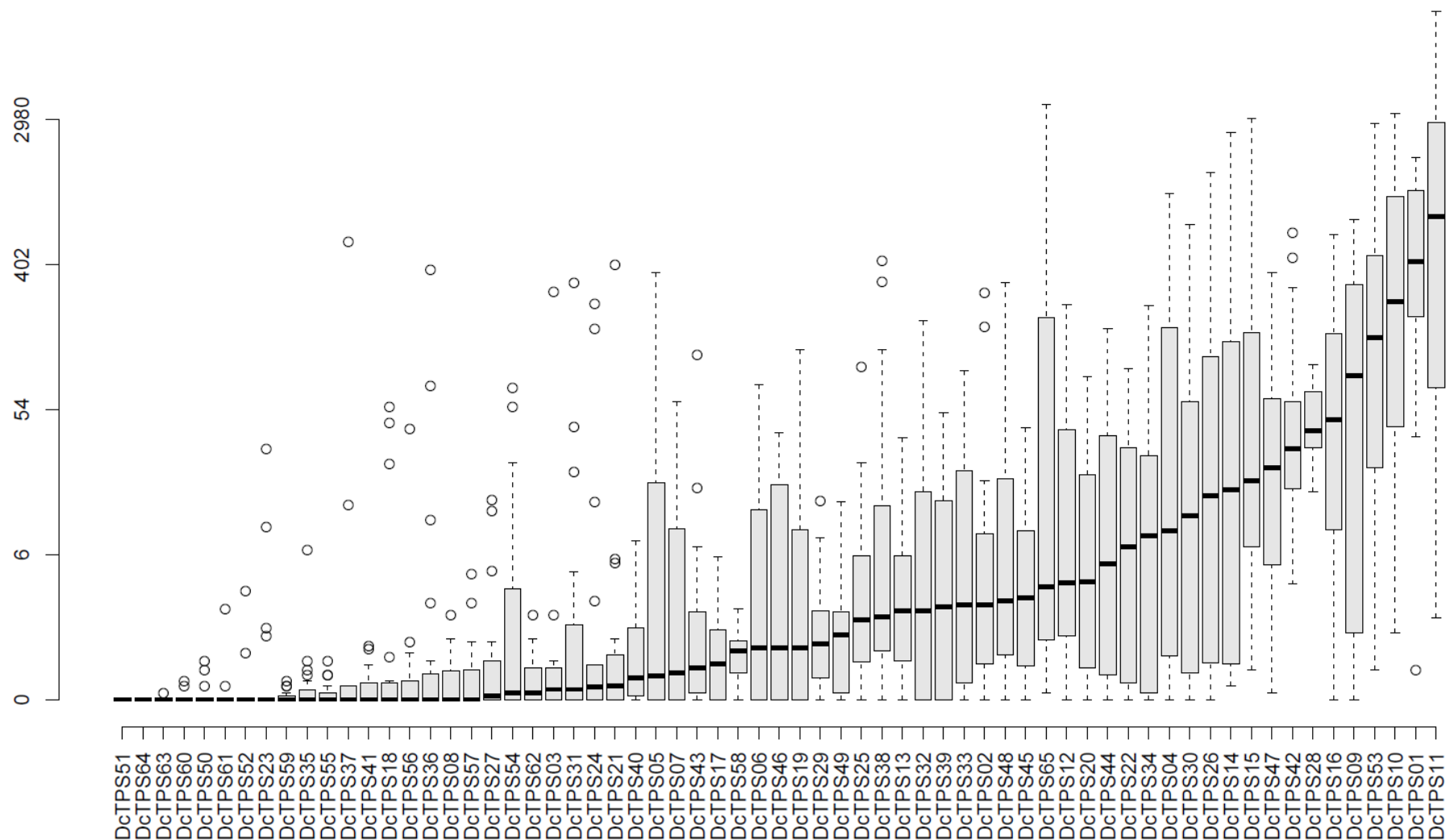


Supplementary Figure 1



Boxplot of fpkm values for the expression of the 65 TPS candidate genes in 20 RNA-seq samples (Iorizzo et al. 2016). The upper side of a box displays the 25% quartile, the bold line displays the median, the lower side of a box displays the 75% quartile, whiskers indicate the variability outside the 25% and 75% quartile, while dots display outliers. The candidate genes have been sorted according to the median fpkm value.