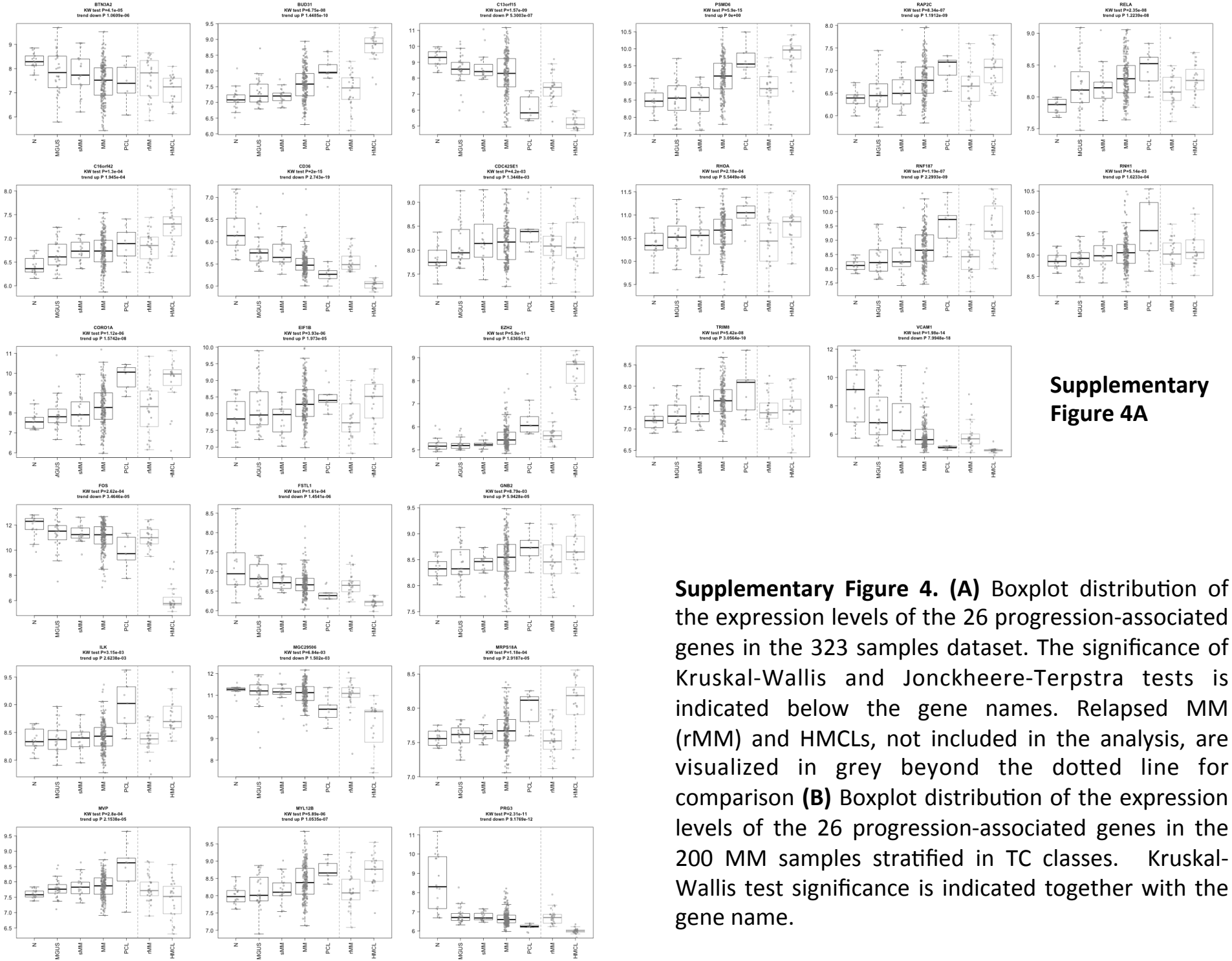
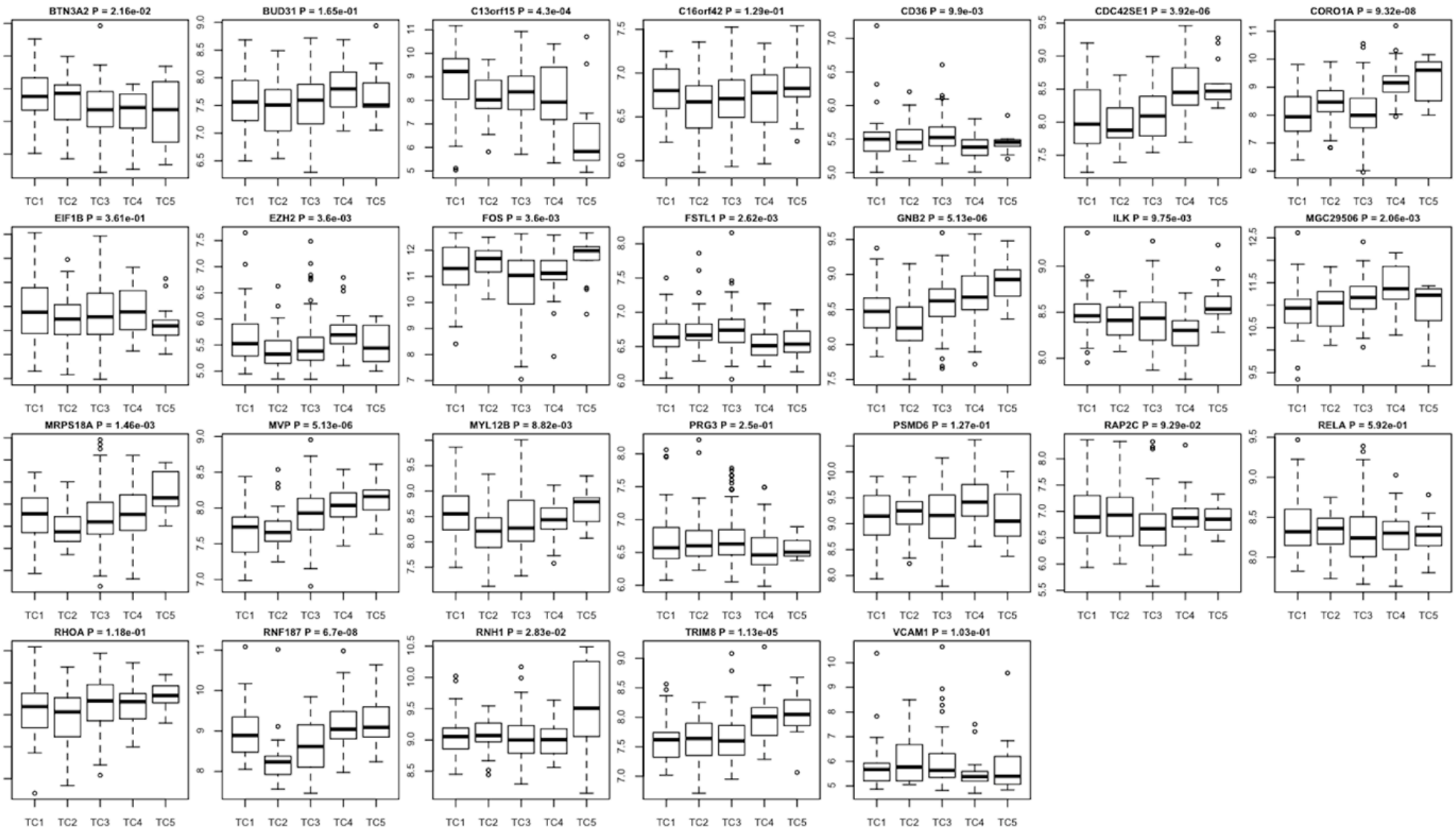


Supplementary Figure 4A

Supplementary Figure 4. (A) Boxplot distribution of the expression levels of the 26 progression-associated genes in the 323 samples dataset. The significance of Kruskal-Wallis and Jonckheere-Terpstra tests is indicated below the gene names. Relapsed MM (rMM) and HMCLs, not included in the analysis, are visualized in grey beyond the dotted line for comparison **(B)** Boxplot distribution of the expression levels of the 26 progression-associated genes in the 200 MM samples stratified in TC classes. Kruskal-Wallis test significance is indicated together with the gene name.



Supplementary Figure 4B