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