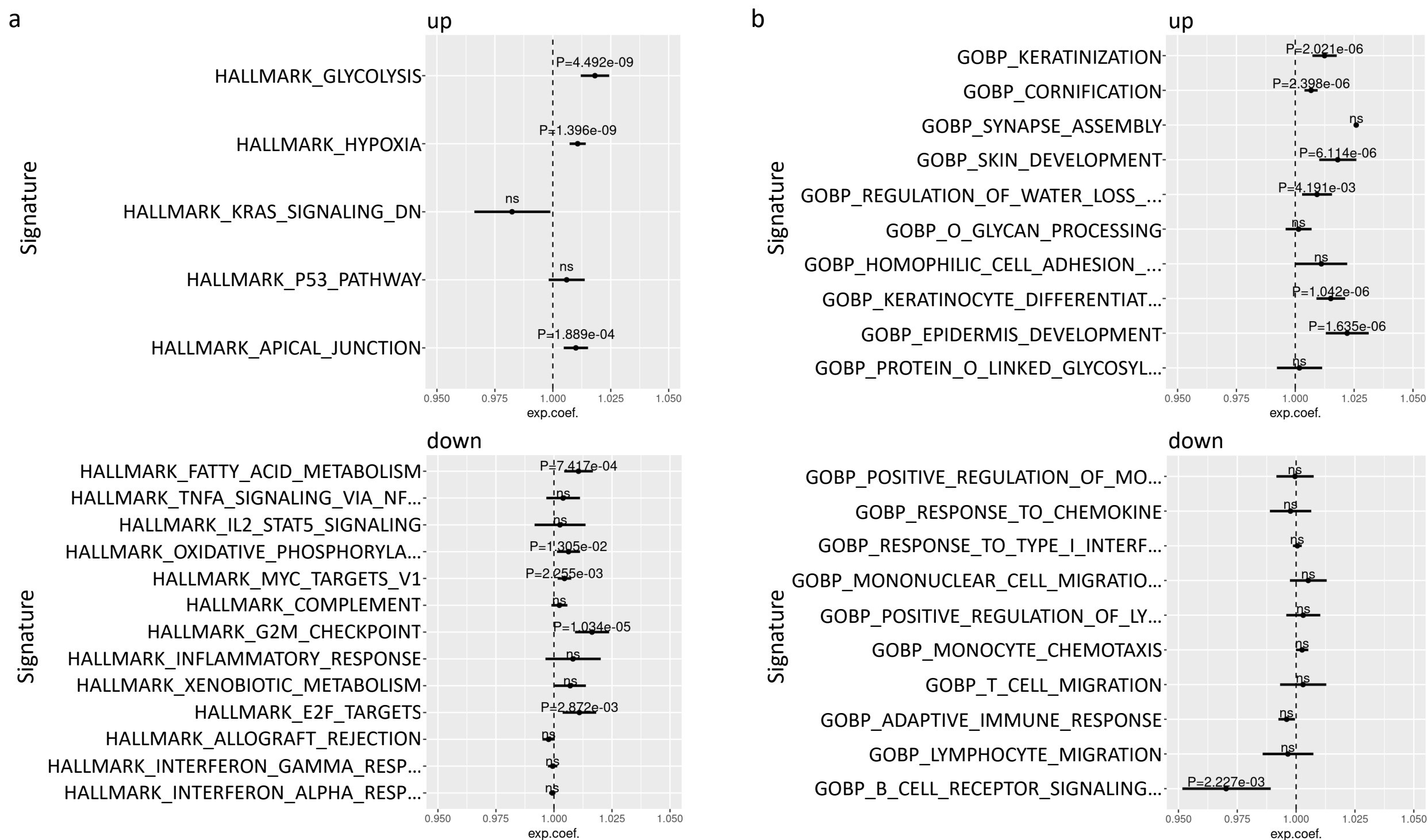


Fig. S15



**Figure S15.** Hazard ratios with deviation for specific signatures detected in the myeloid subtype of LUAD. The relationship between increased and decreased expression of signatures and event-free survival using transcriptome data from TCGA NSCLC was plotted. (a) hallmark gene set. (b) GO set.