

Figure 7

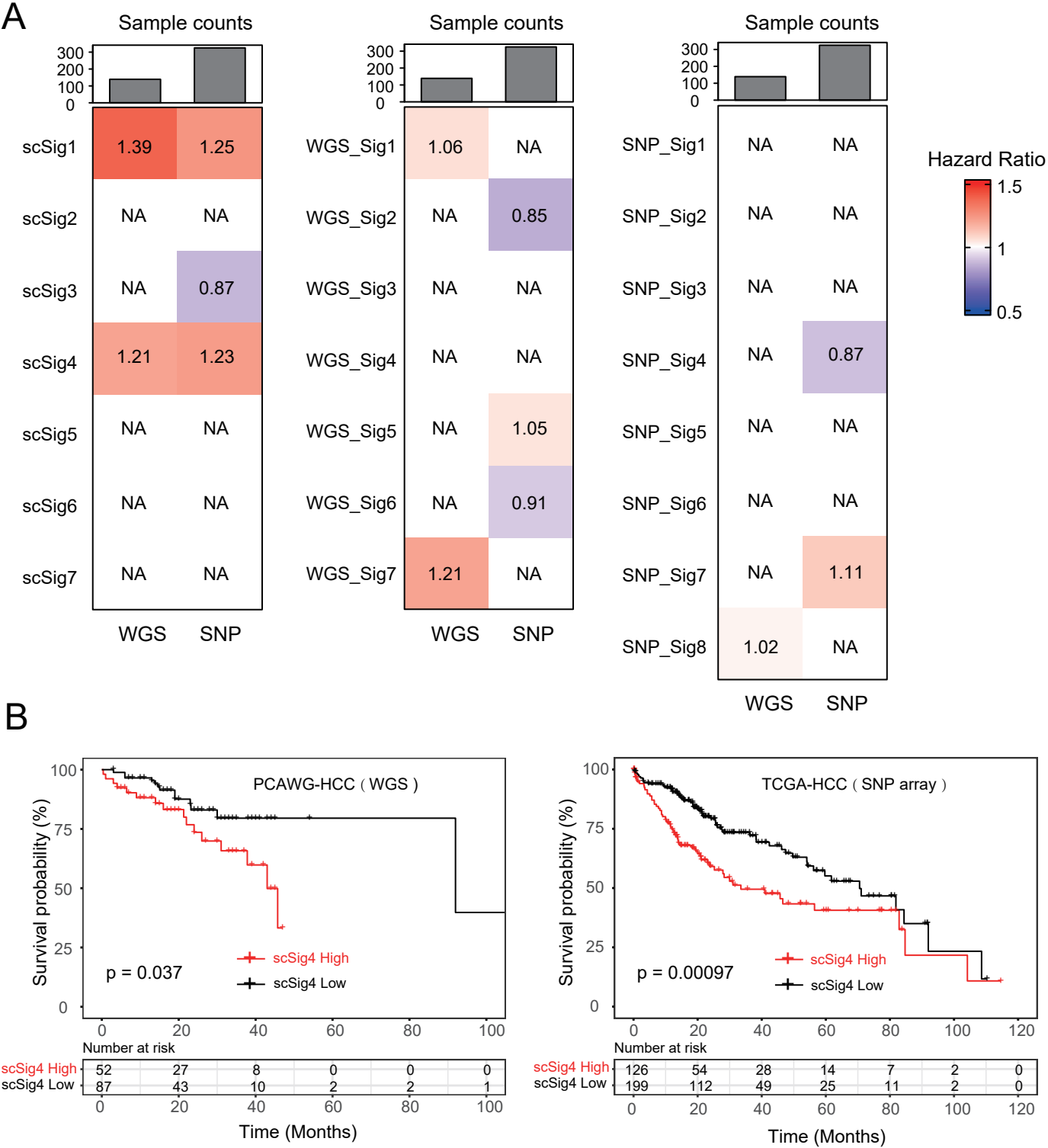


Figure 7. Single-cell CNA signatures show robust performance in prognosis prediction compared with CNA signatures derived from bulk tissues.

A, The prognosis performance of scSig, WGS_Sig, and SNP_Sig in WGS and SNP array data was analyzed. Shown are the HR values of Signature that are significantly related to prognosis (FDR < 0.05), where HR > 1 indicates an association with poor prognosis, and HR < 1 indicates an association with better prognosis.

B, The prognosis performance of the scSig4 in both WGS and SNP datasets. Consistent prognostic results in two different data sets.