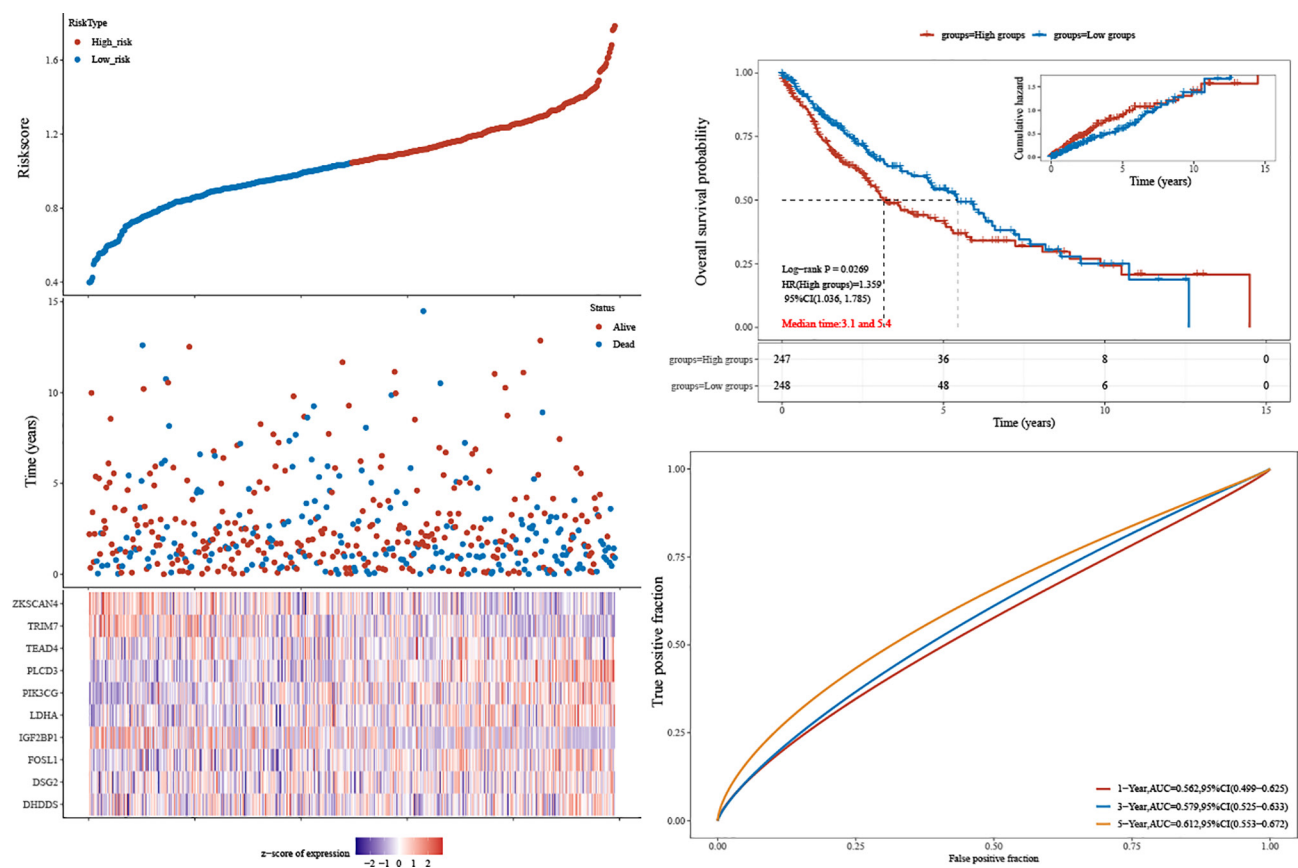**Figure S1** Construction of a telomere gene-related prognostic model in SCC database. HR, hazard ratio; CI, confidence interval; AUC, area under the curve; SCC, squamous cell carcinoma.