

A

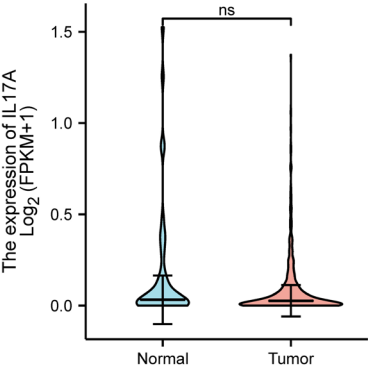


Figure 1. The expression of IL17A between normal and HNSC tumor tissues

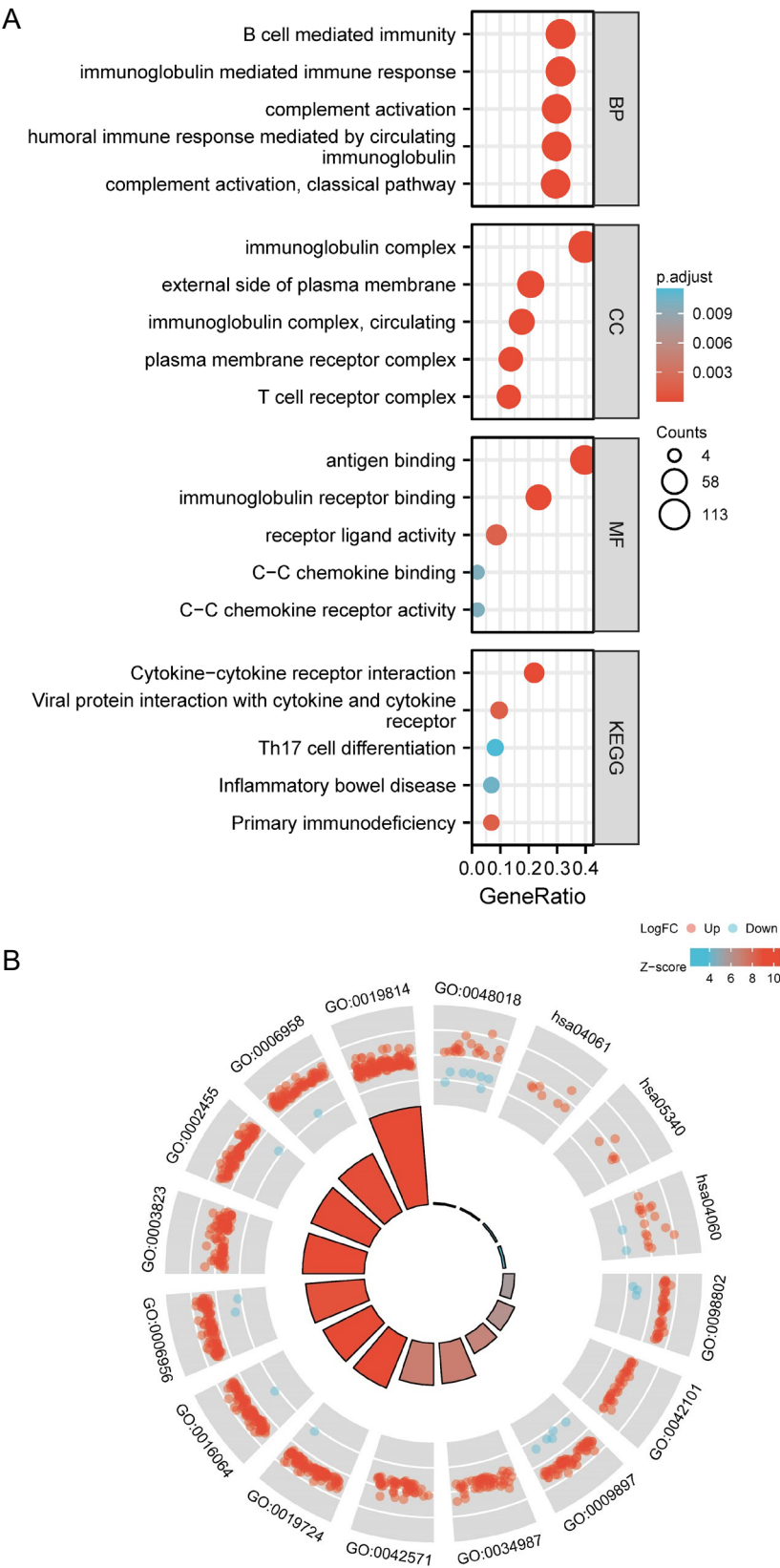


Figure 2. Pathway enrichment results for differentially expressed genes between high and low expression of IL17A in HNSC patients from KEGG and GO analysis (A, B).

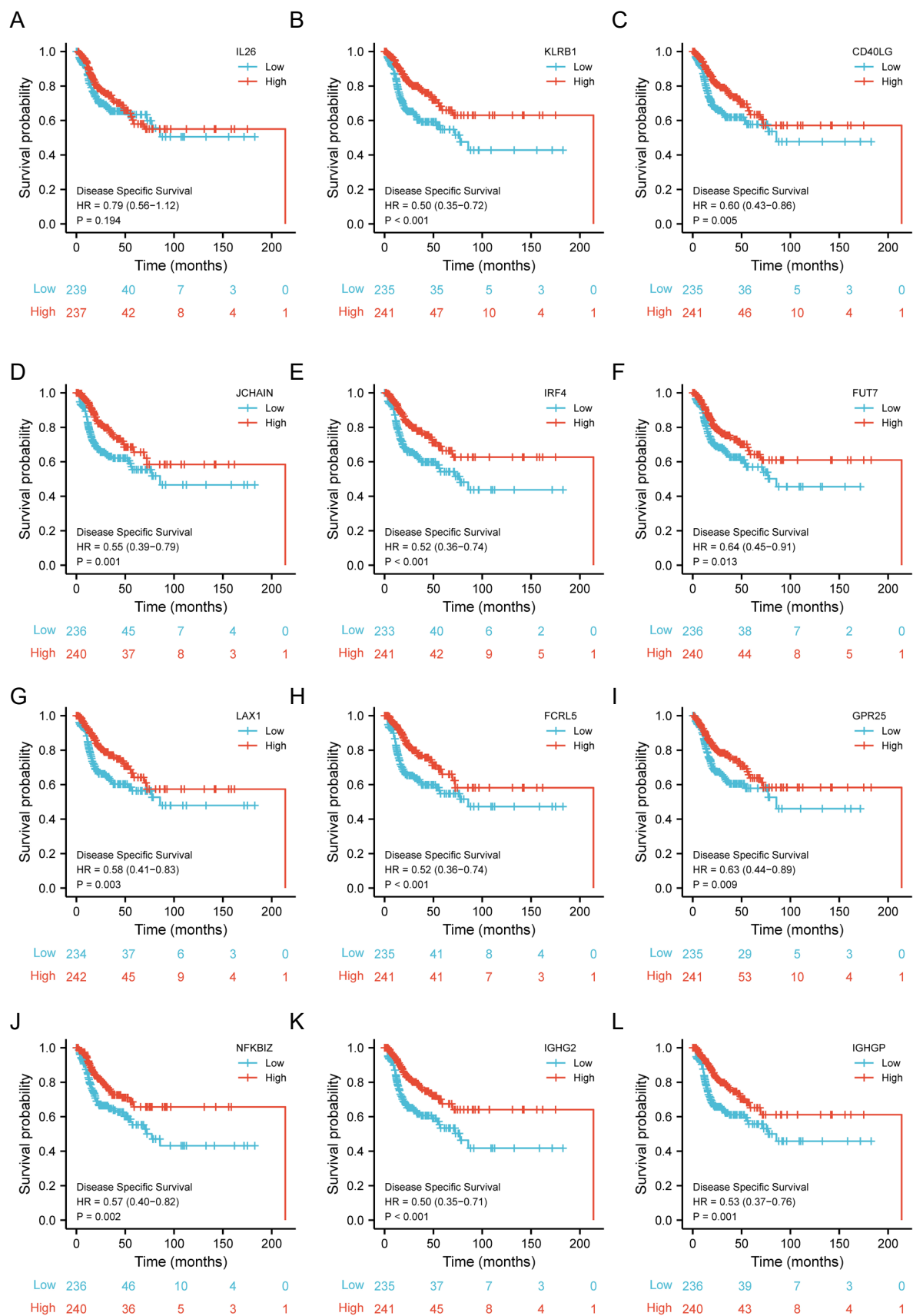


Figure 3. Kaplan–Meier survival curves (disease specific survival) comparing the high and low expression of the IL17A correlated 12 genes in HNSC patients (A-L).