Standalone markers, BCR-free survival, mRNA

model inference
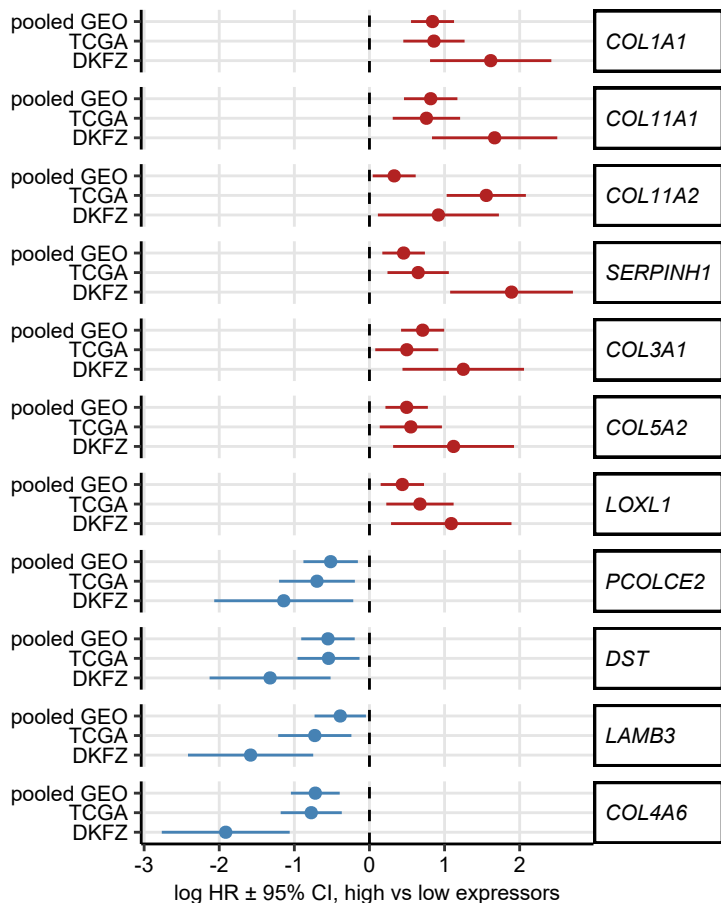
performance statistics

Genes (top to bottom): COL1A1, COL11A1, COL11A2, SERPINH1, COL3A1, COL5A2, LOXL1, PCOLCE2, DST, LAMB3, COL4A6

Each panel rows: pooled GEO, TCGA, DKFZ

x-axis (left): log HR ± 95% CI, high vs low expressors

x-axis (right): C-index
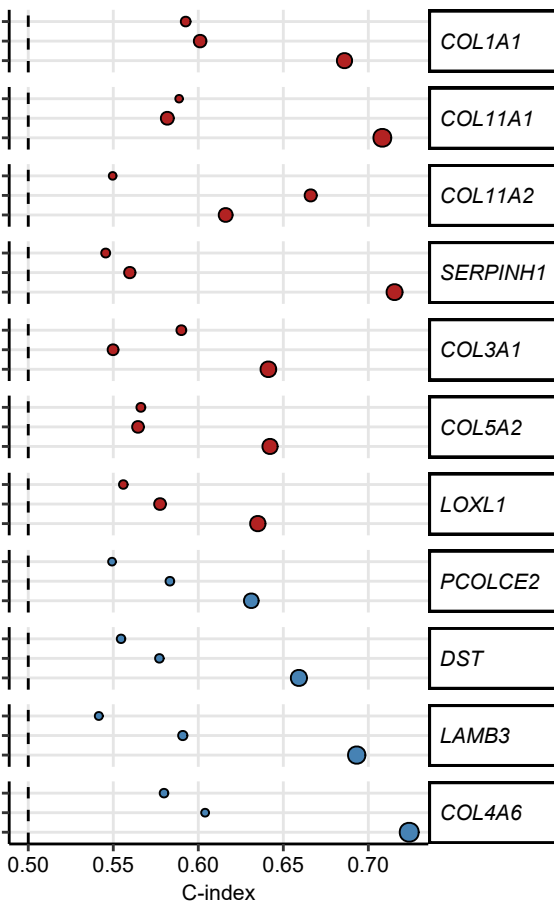
1 - IBS   ○ 0.80   ○ 0.82   ○ 0.84   ○ 0.86   ○ 0.88

association with risk   ● unfavorable   ● favorable