

Figure S1 Gene Set Enrichment Analysis.

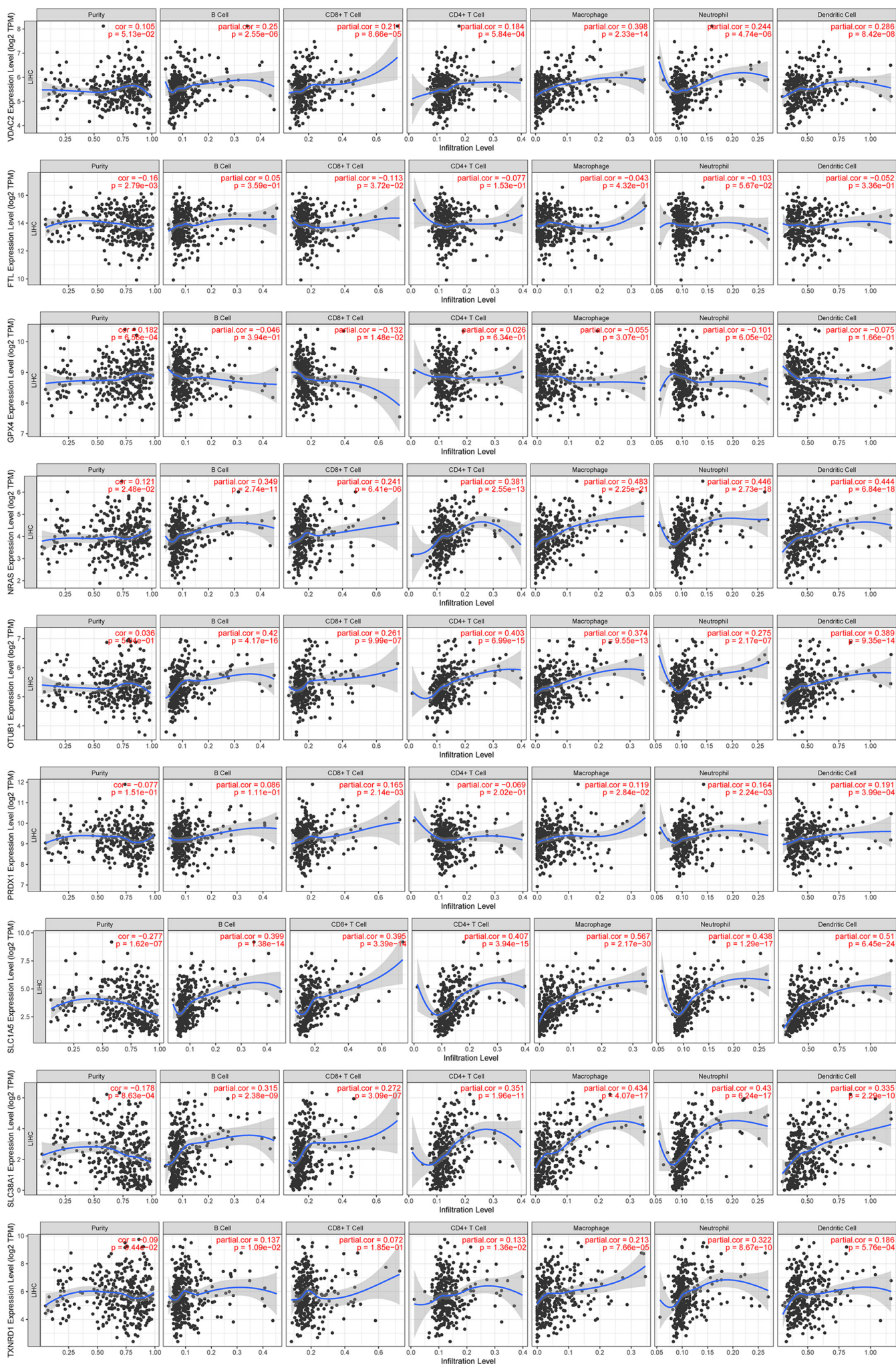


Figure S2 Analyses of immune infiltration based on the TIMER database. TPM, transcripts per million; LIHC, liver hepatocellular carcinoma.

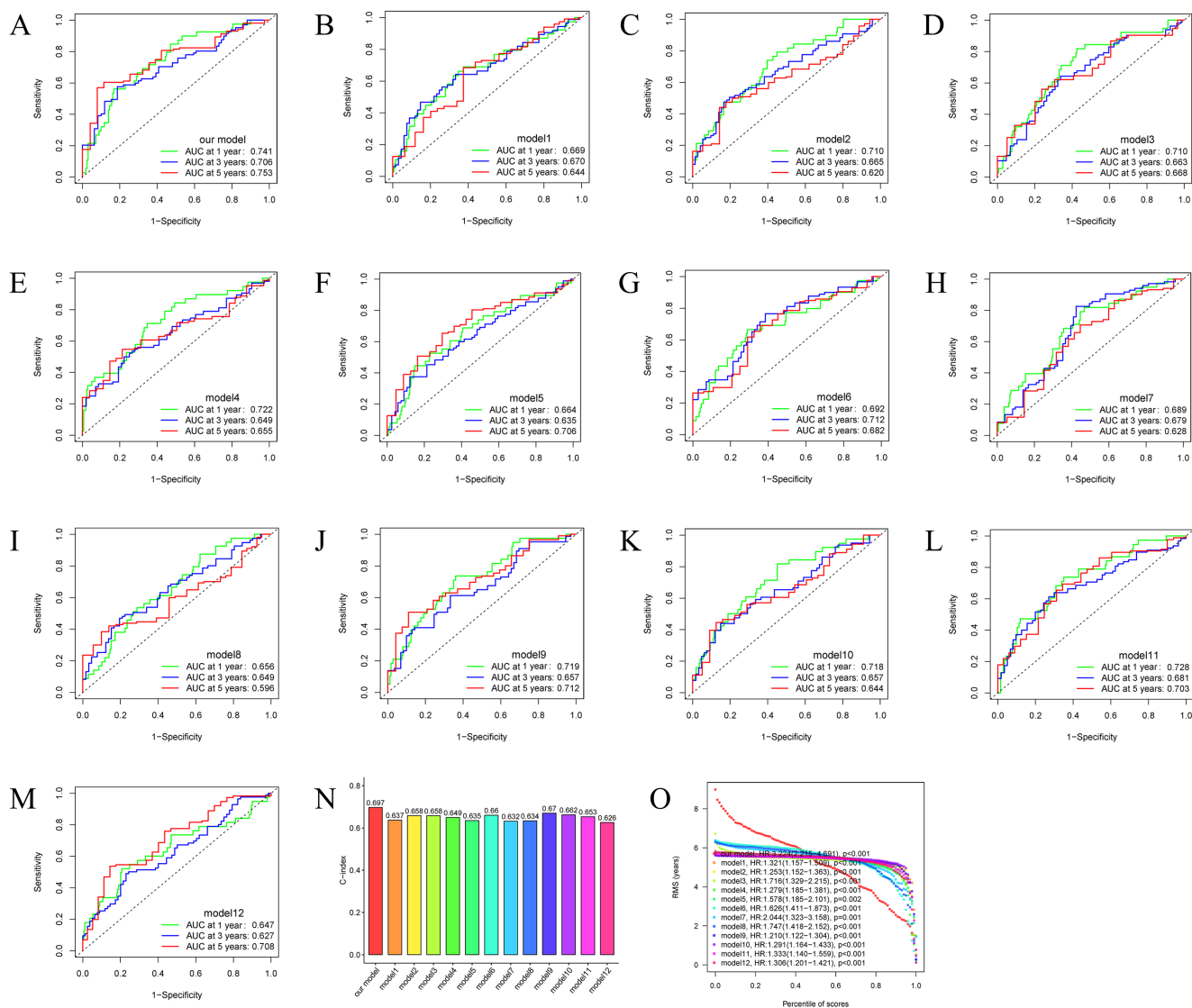


Figure S3 Comparison with other models. (A-M) The ROC curves of 13 models; (N) C-index of 13 models; (O) RMS of 13 models. AUC, area under the curve; HR, hazard ratio; ROC, receiver operating characteristic; C-index, concordance index; RMS, restricted mean survival.