

Figure S1. Comparison of eight-gene signature model with the other existing models. ROC and K-M curves generated using the model by (A) Xie *et al* (45), (B) LV *et al* (46) and (C) Quist *et al* (47). (D) RMS curve comparing the three models with the eight-gene signature; (E) DCA curve comparing the three models with eight-gene signature. ROC, receiver operating characteristic; AUC, area under the curve; K-M, Kaplan-Meier; CI, confidence interval; RMS, root mean square; DCA, decision curve analysis.

