

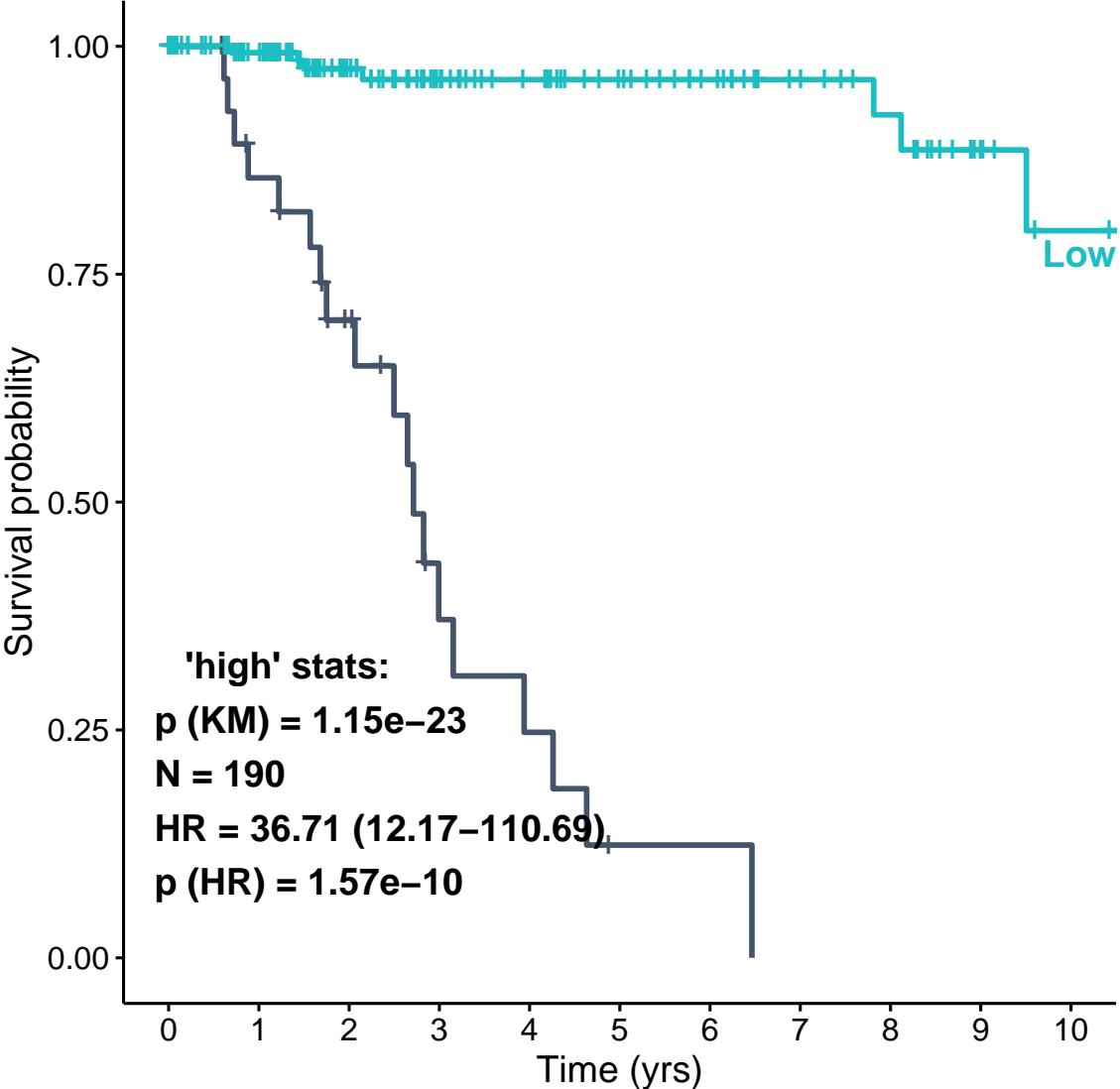


planA_BOTHfcUP_progWORSE (343 genes)

Strata  Row_mode=high  Row_mode=low



Number at risk

