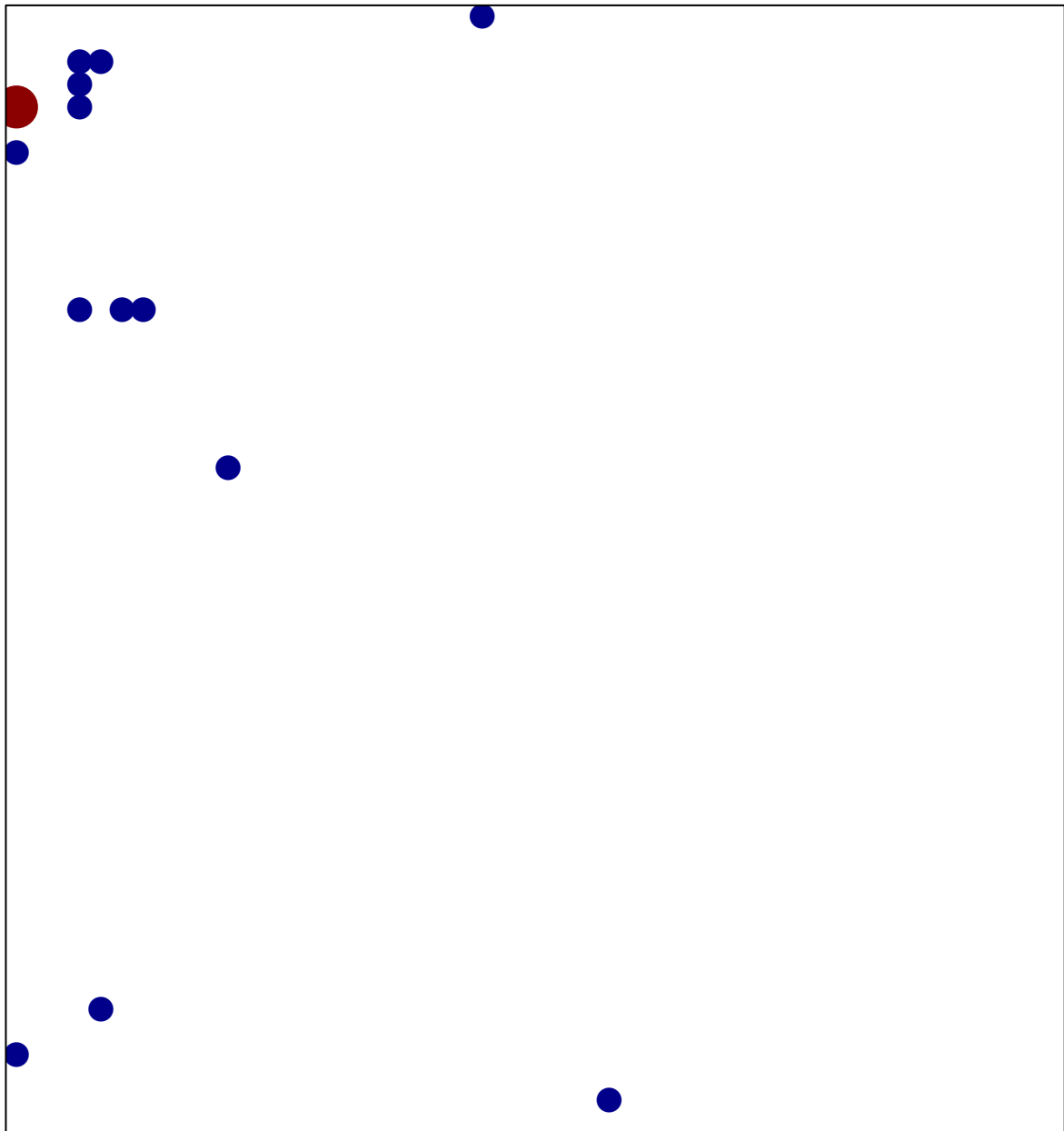# RICKMAN_TUMOR_DIFFERENTIATED_WELL_VS_POORLY_DN

features = 15
chi−square p = 0.78

XMAN_TUMOR_DIFFERENTIATED_WELL_VS_POORL

features = 15 , max = 2