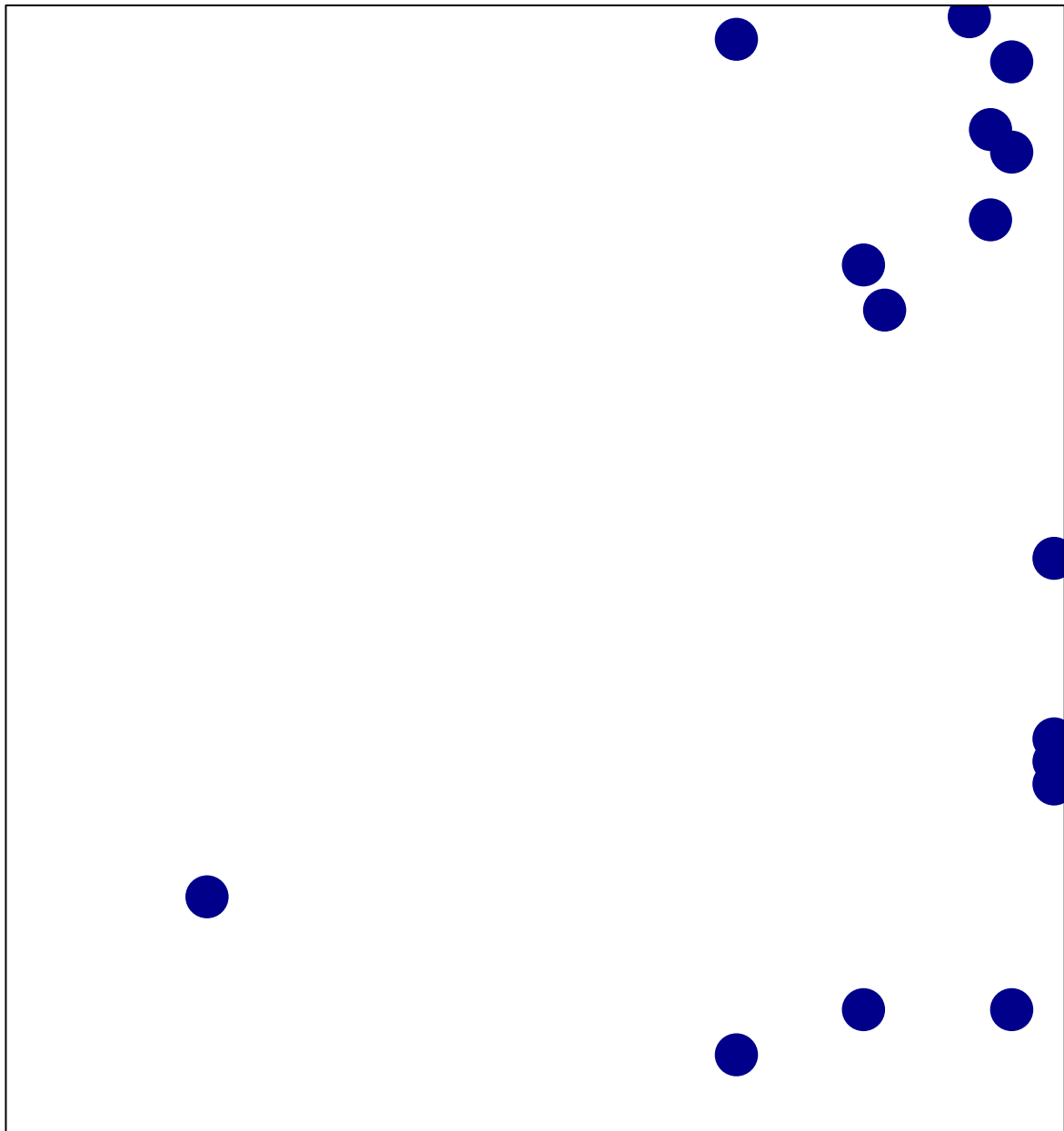
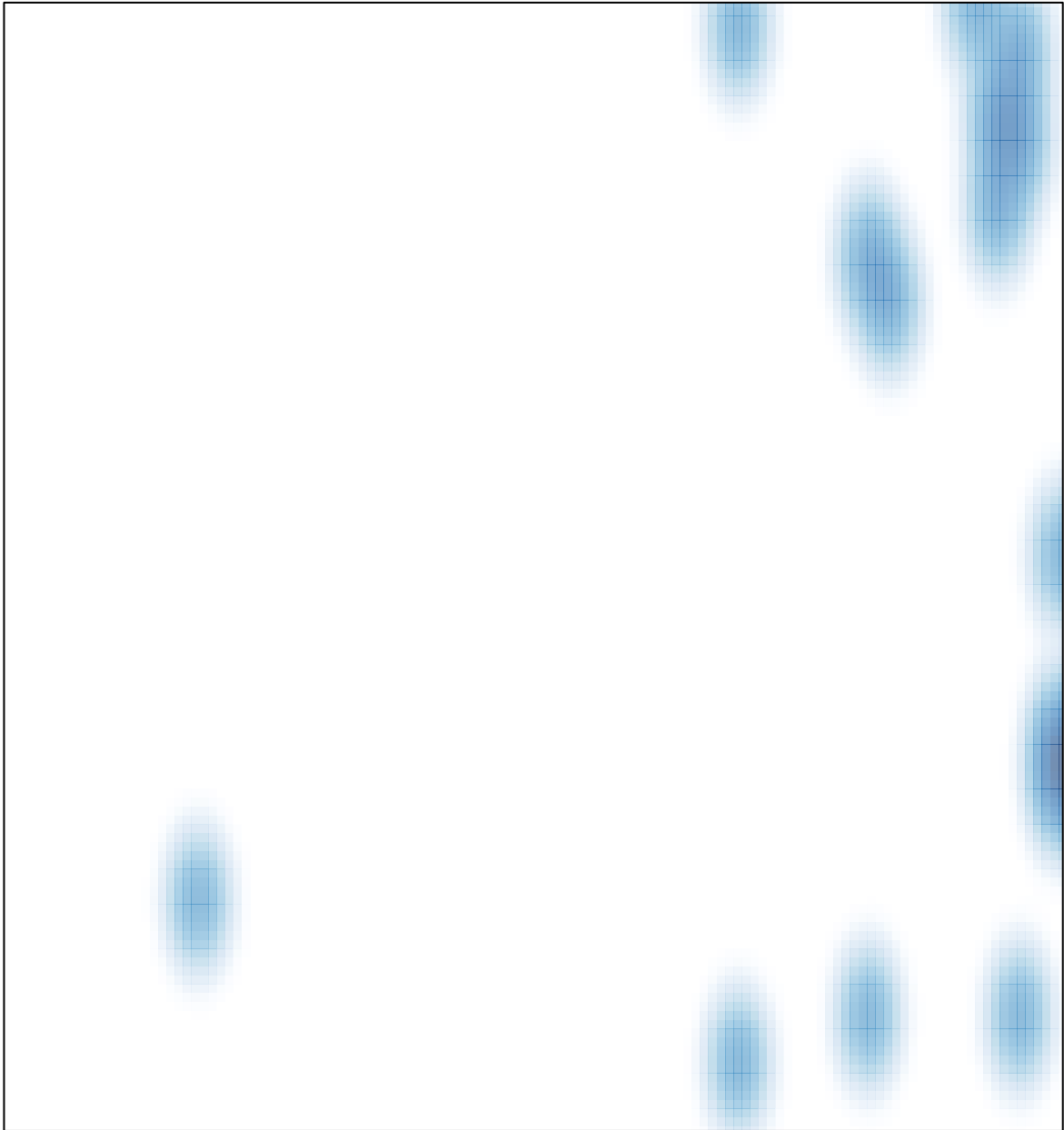


# RICKMAN\_TUMOR\_DIFFERENTIATED\_WELL\_VS\_POORLY\_UP



# features = 16  
chi-square p = 0.81

# KMAN\_TUMOR\_DIFFERENTIATED\_WELL\_VS\_POOR



# features = 16 , max = 1