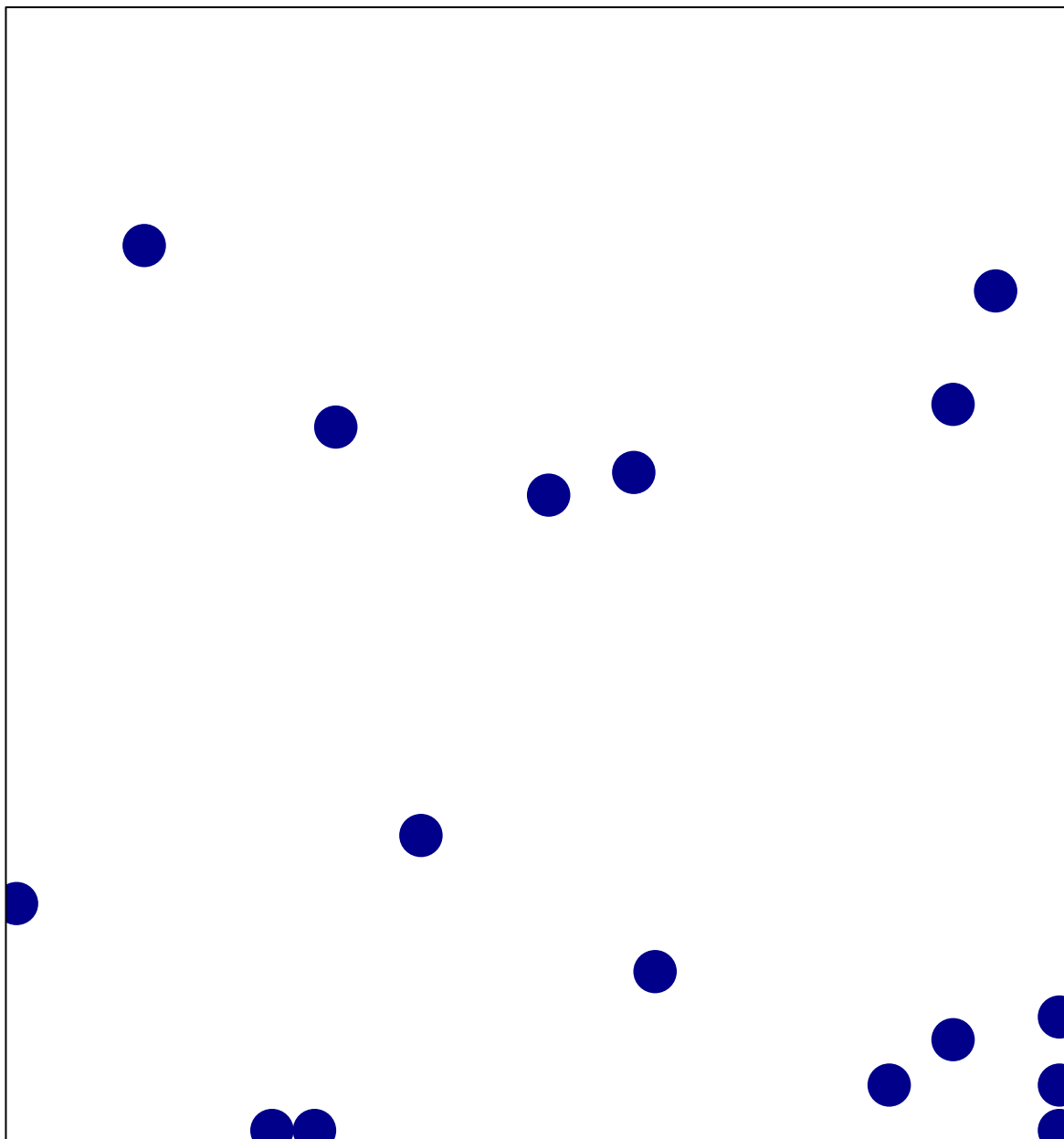
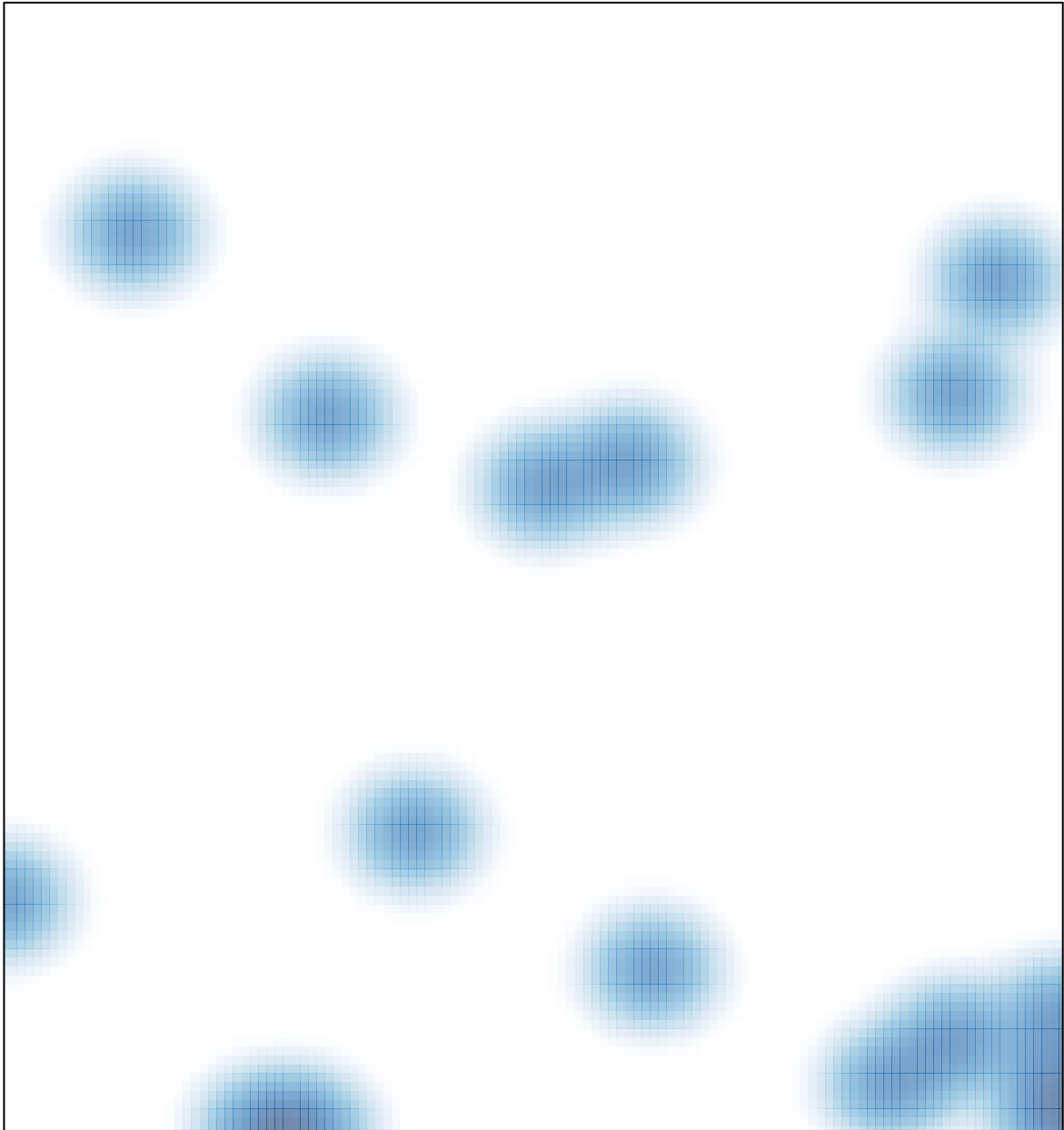


# REACTOME\_INTEGRIN\_ALPHAIIIBETA3\_SIGNALING



# features = 16  
chi-square p = 0.83

# REACTOME\_INTEGRIN\_ALPHAIIIBETA3\_SIGNALIN



# features = 16 , max = 1