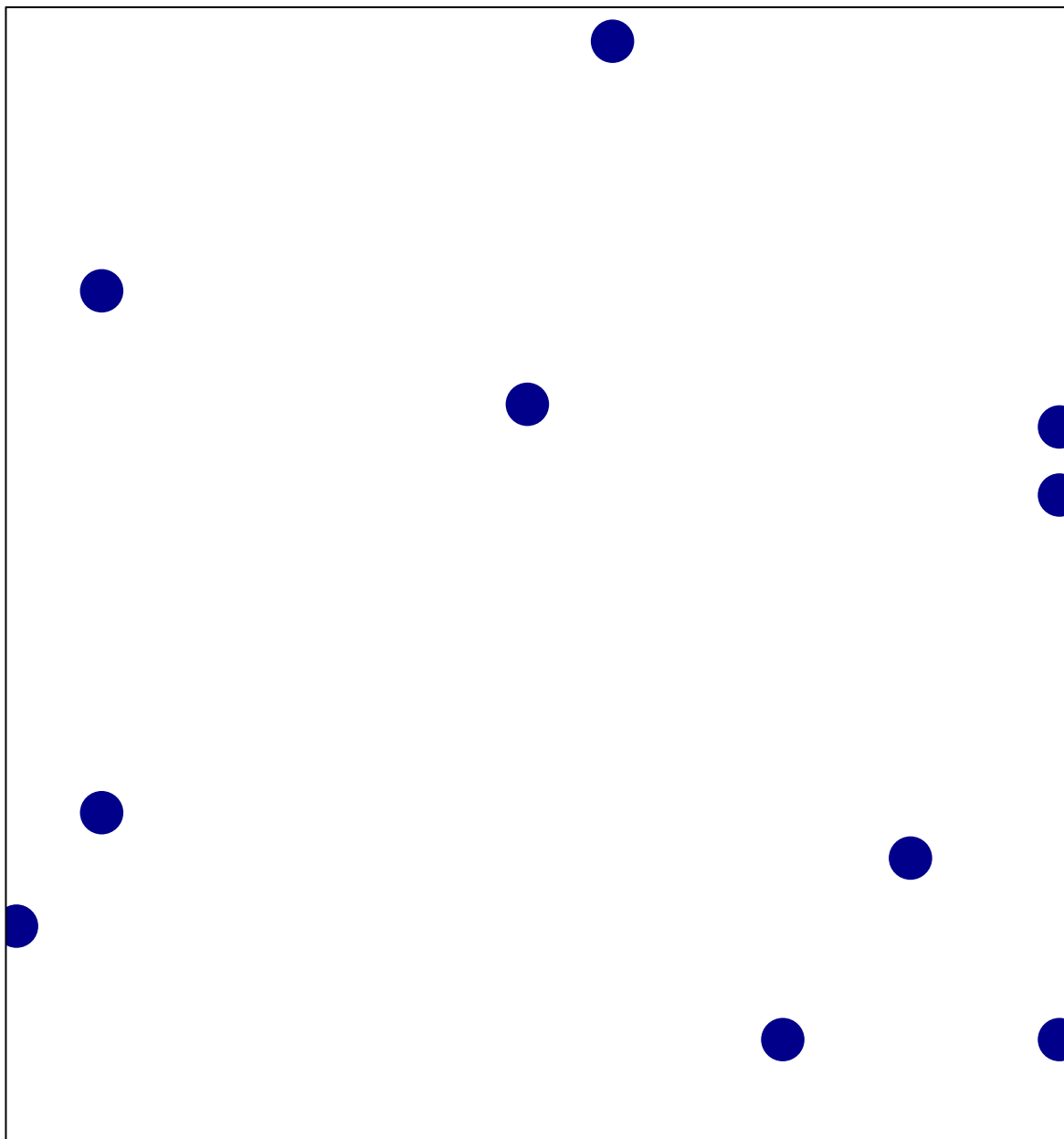
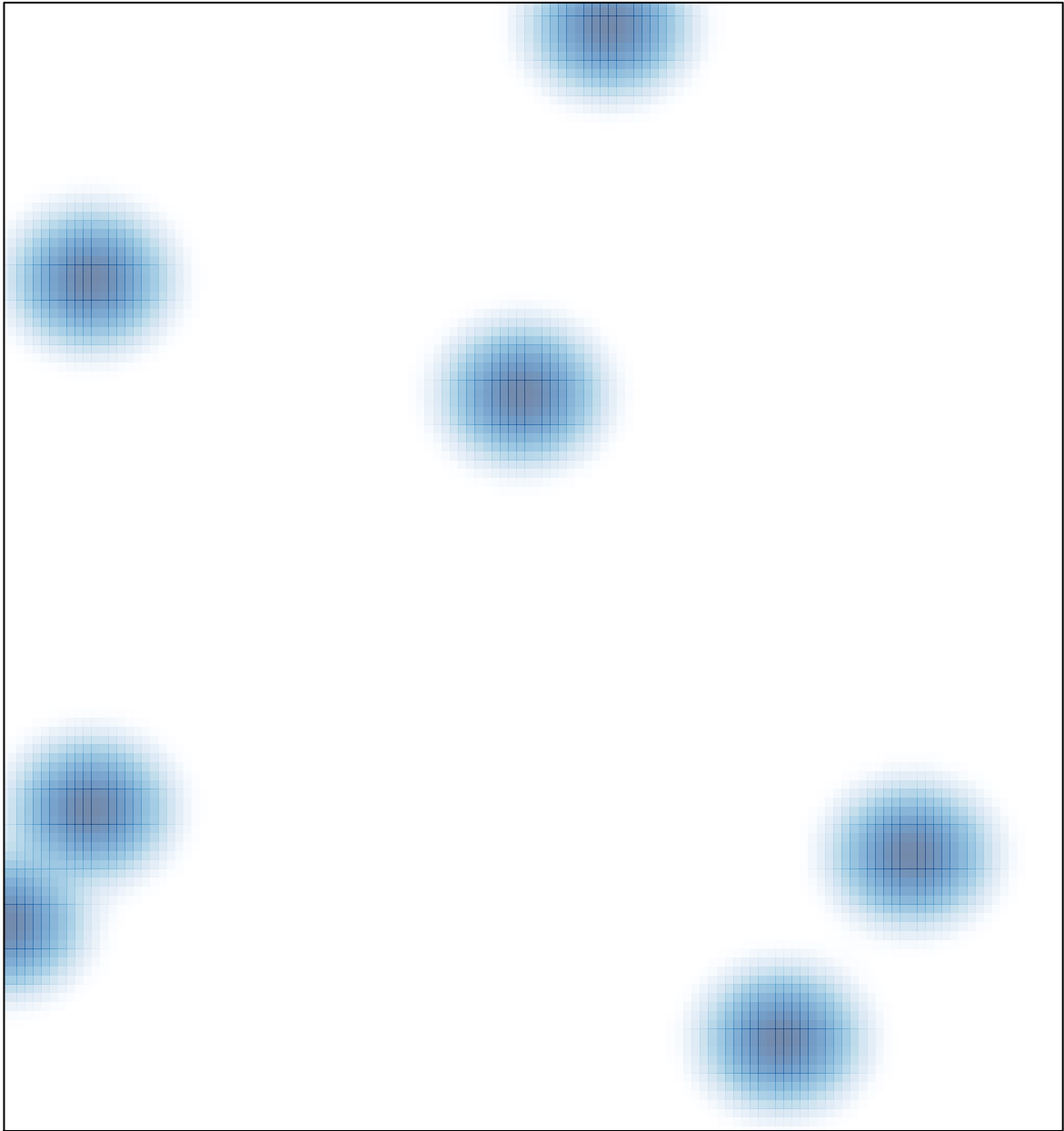


# REACTOME\_VEGF\_LIGAND\_RECEPTOR\_INTERACTIONS



# features = 10  
chi-square p = 0.82

# EACTOME\_VEGF\_LIGAND\_RECEPTOR\_INTERACTIO



# features = 10 , max = 1