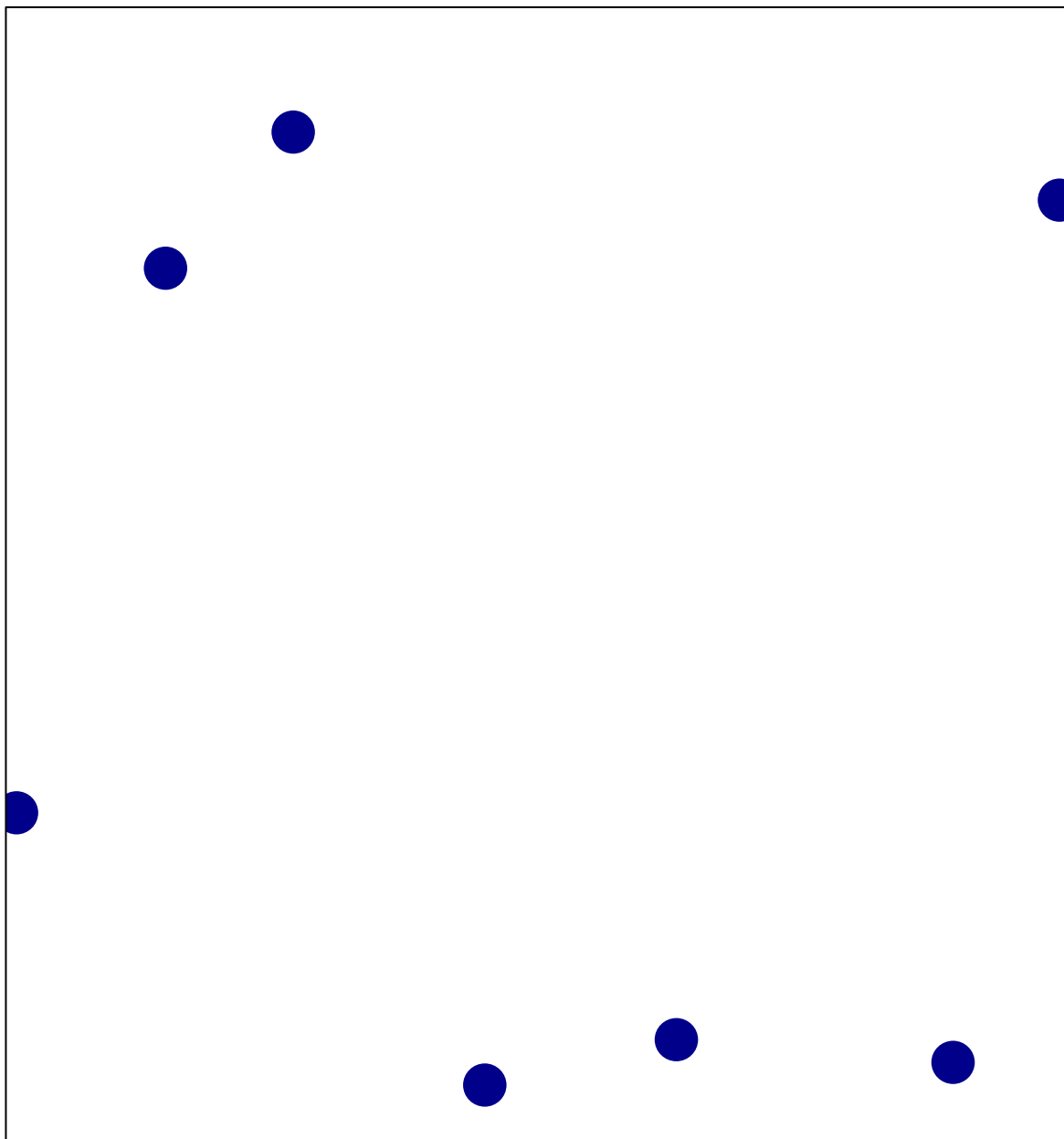
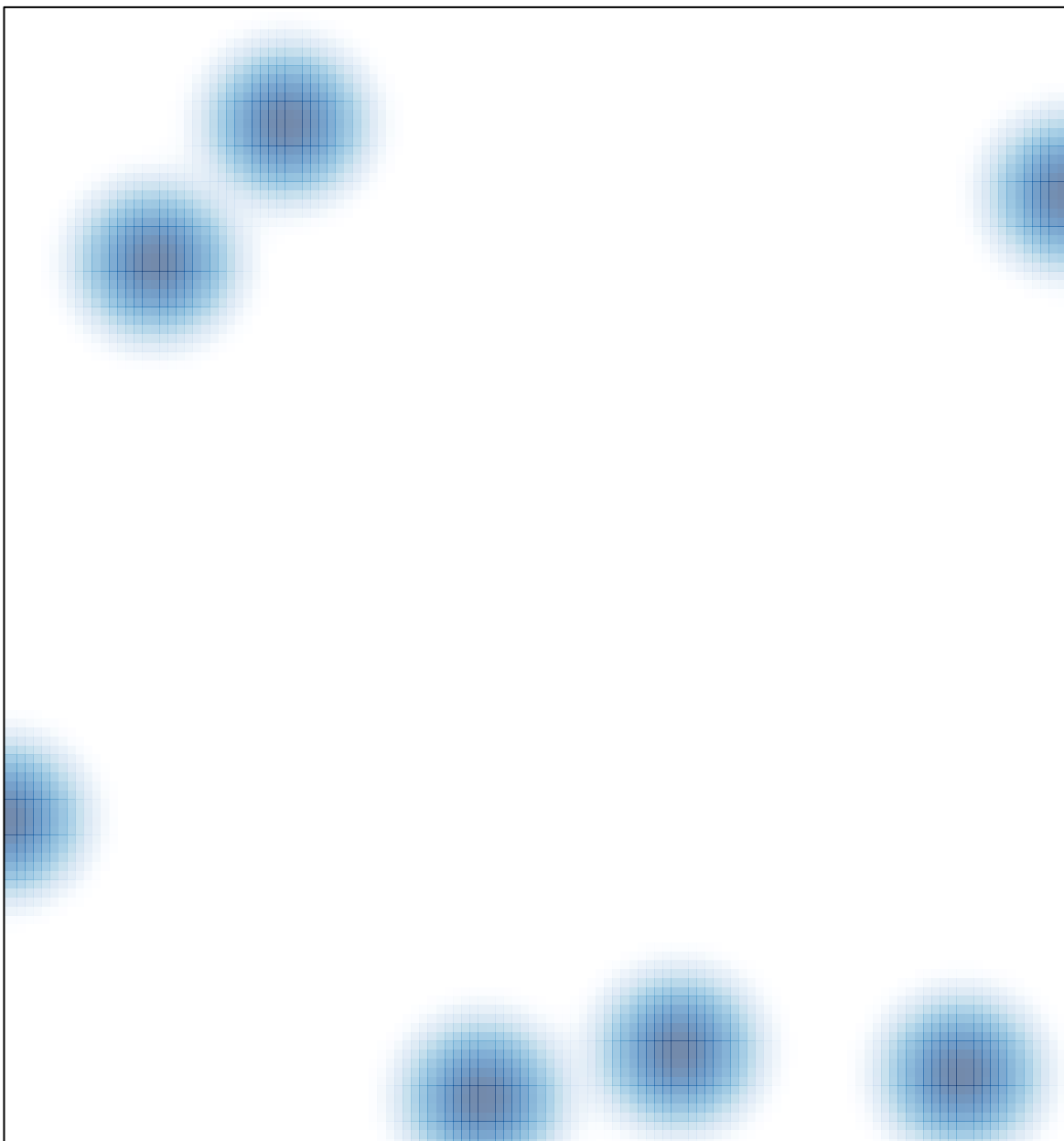


# REACTOME\_PEPTIDE\_LIGAND\_BINDING\_RECEPTORS



# features = 7  
chi-square p = 0.85

# REACTOME\_PEPTIDE\_LIGAND\_BINDING\_RECEPTOR



# features = 7 , max = 1