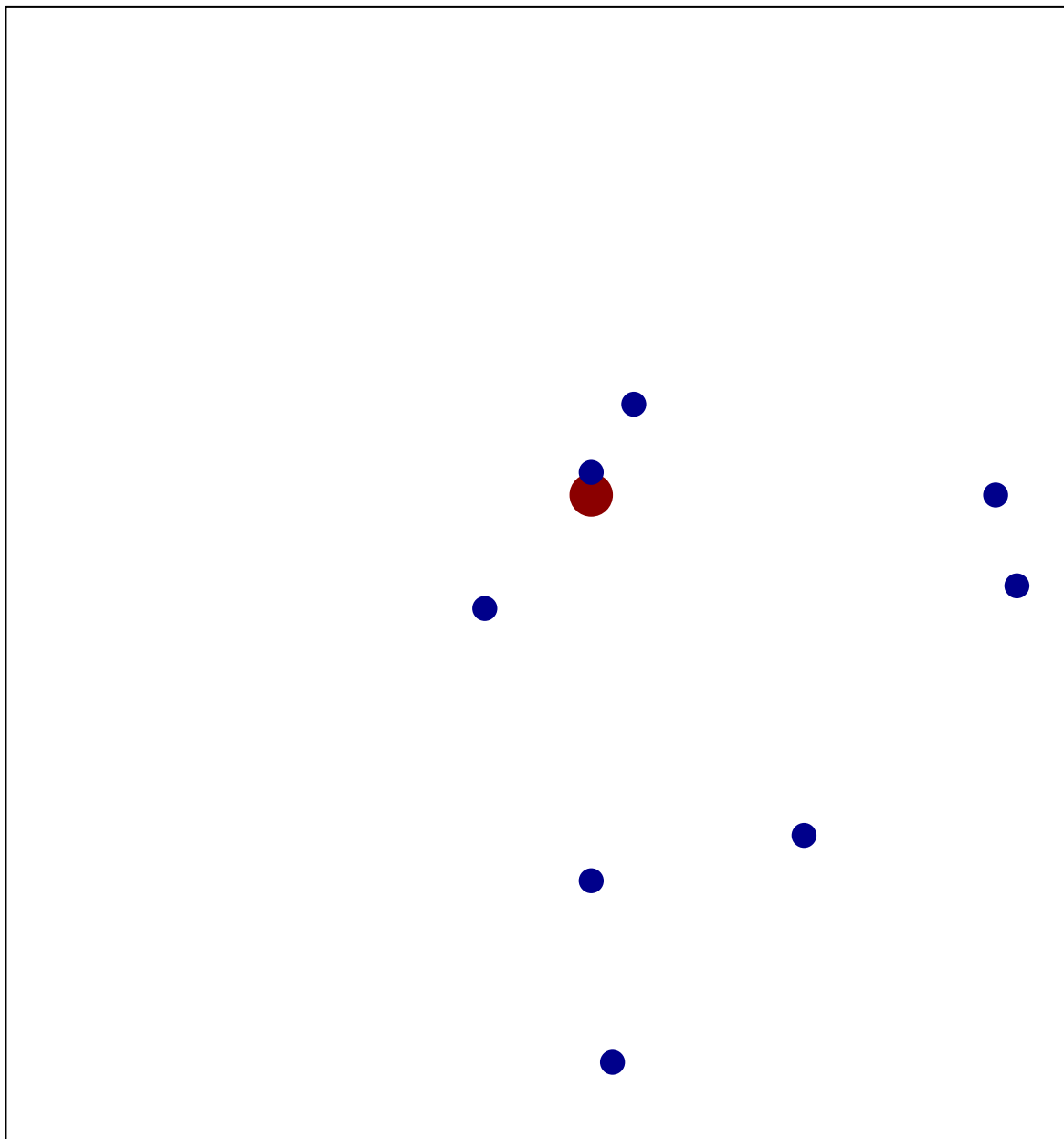


# REACTOME\_EICOSANOID\_LIGAND\_BINDING\_RECEPTORS



# features = 10  
chi-square p = 0.79

# ACTOME\_EICOSANOID\_LIGAND\_BINDING\_RECEPT



# features = 10 , max = 2