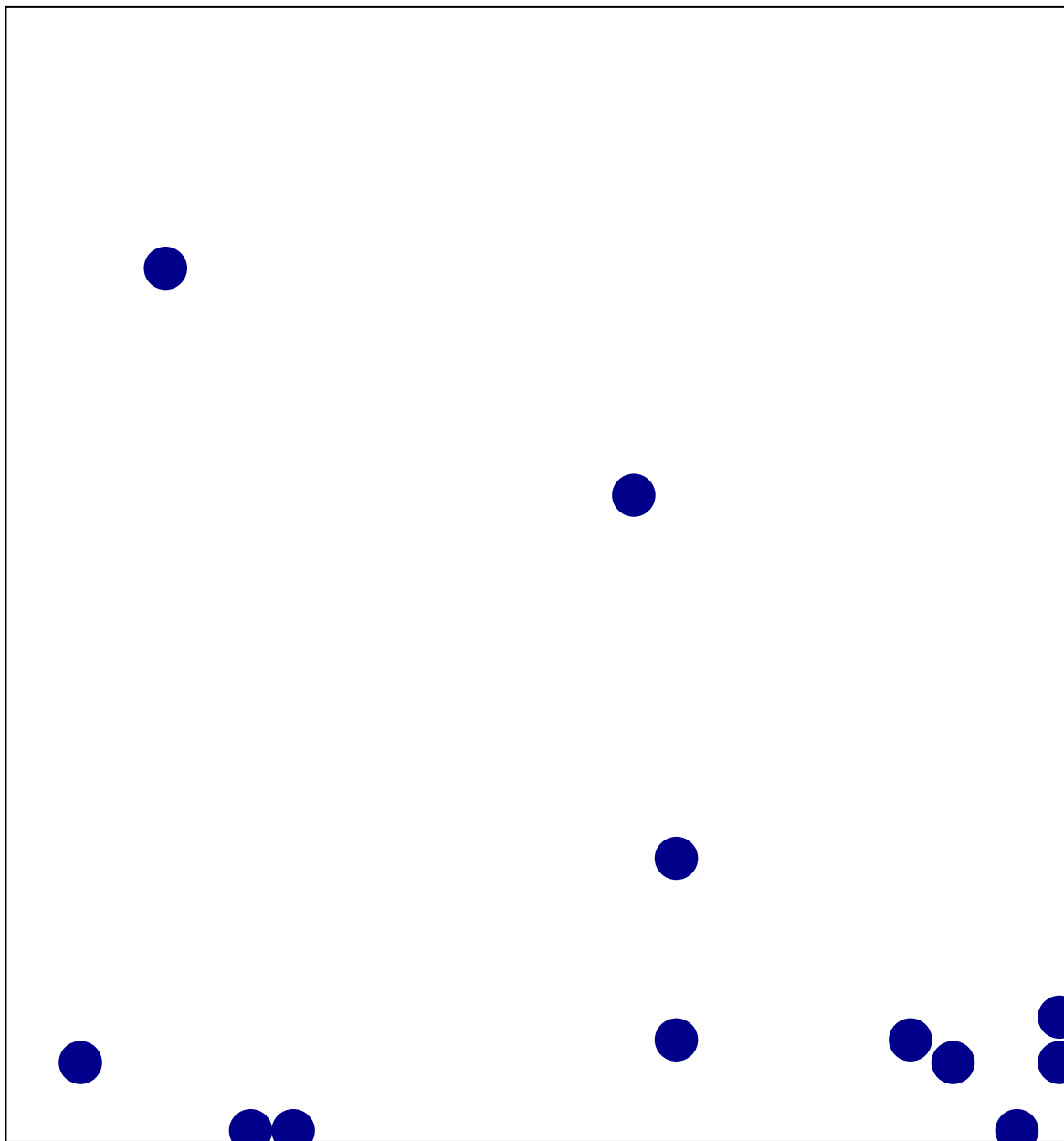


# REACTOME\_CHEMOKINE\_RECEPTORS\_BIND\_CHEMOKINES



# features = 12  
chi-square p = 0.81

# CTOME\_CHEMOKINE\_RECEPTORS\_BIND\_CHEMOK



# features = 12 , max = 1