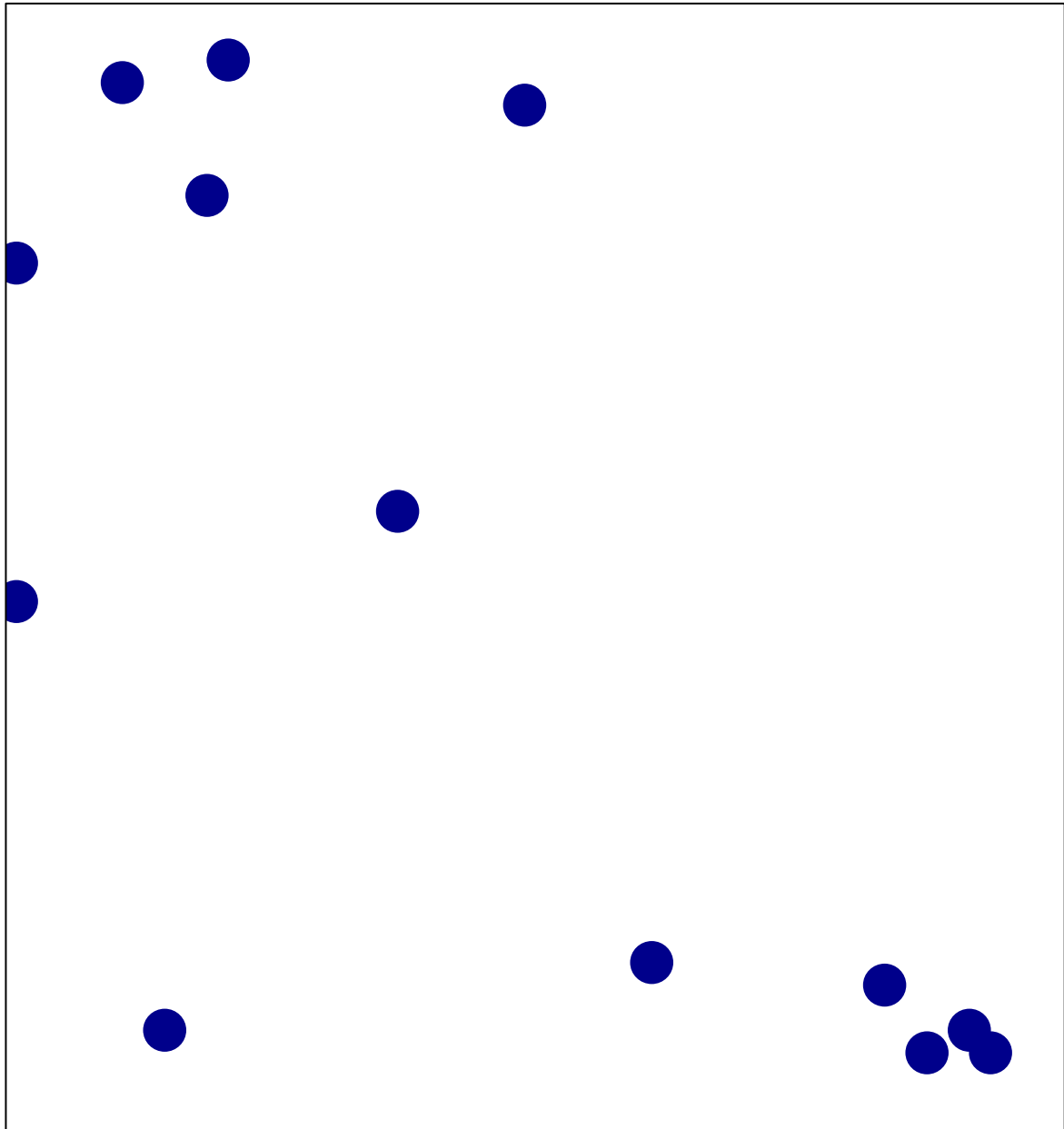


# phosphotyrosine binding



# features = 13  
chi-square p = 0.84

# phosphotyrosine binding



# features = 13 , max = 1