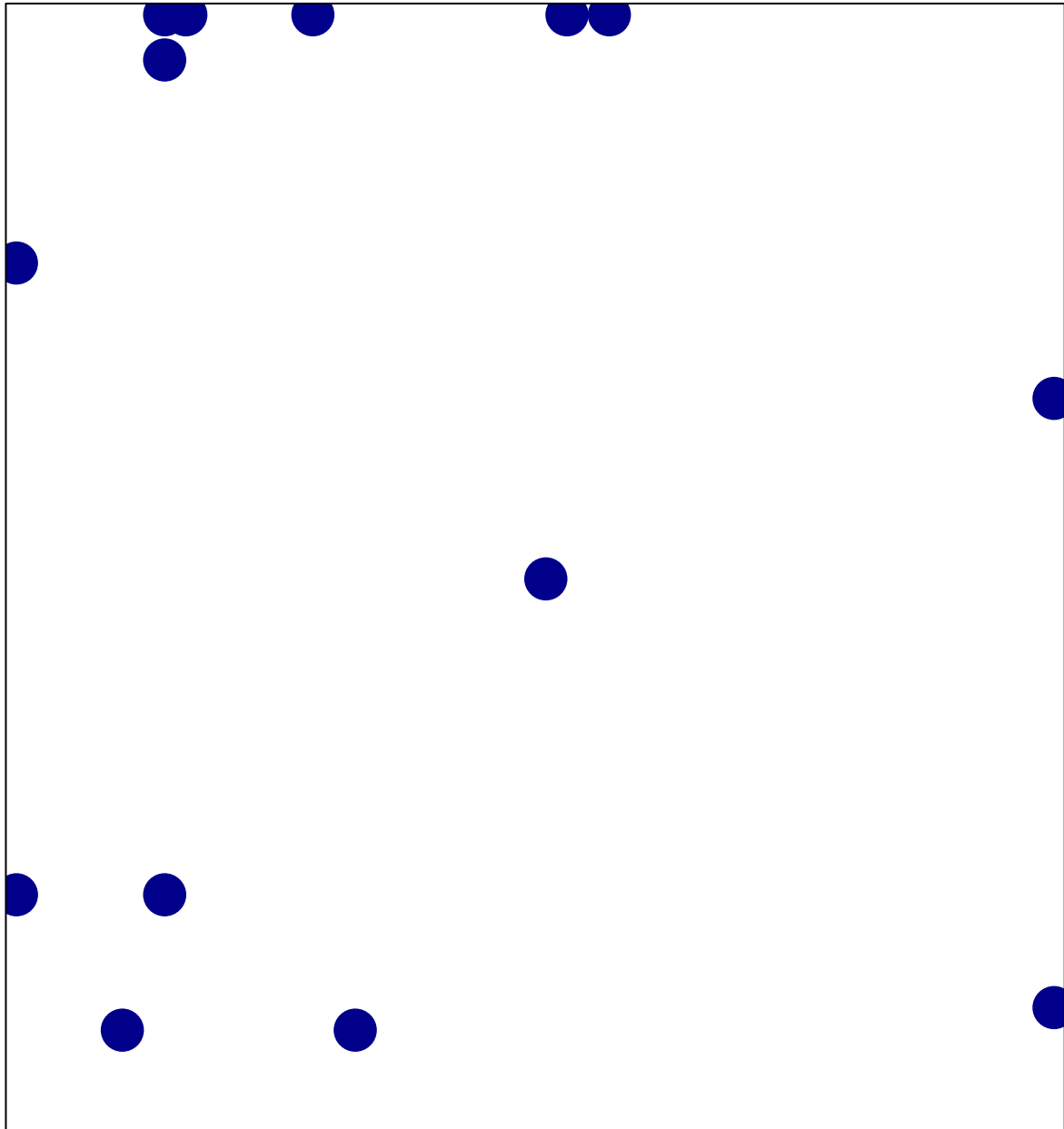


# centrosome localization



# features = 14  
chi-square p = 0.78

# centrosome localization



# features = 14 , max = 1