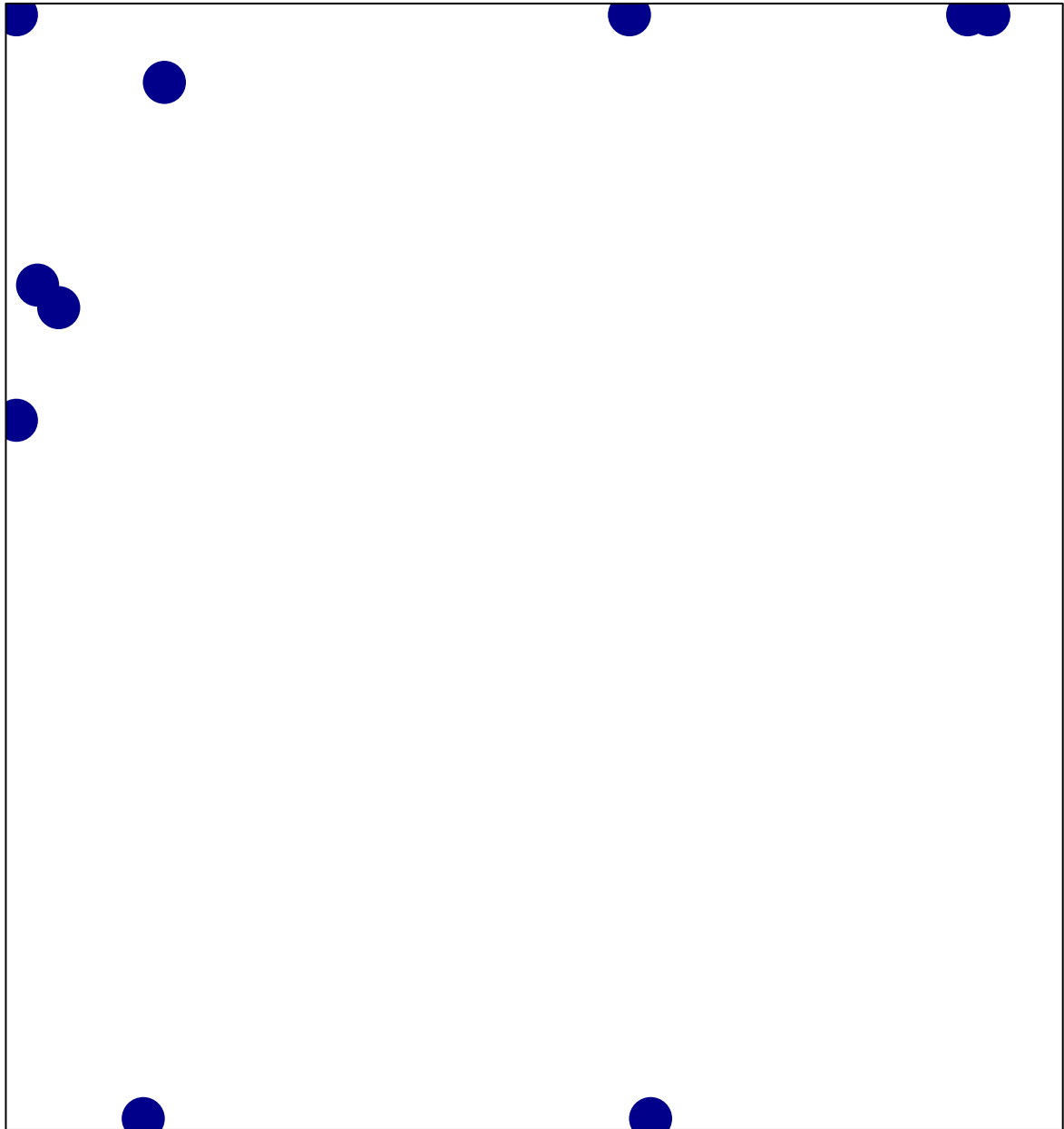


# nuclear exosome (RNase complex)



# features = 10  
chi-square p = 0.8

# nuclear exosome (RNase complex)



# features = 10 , max = 1