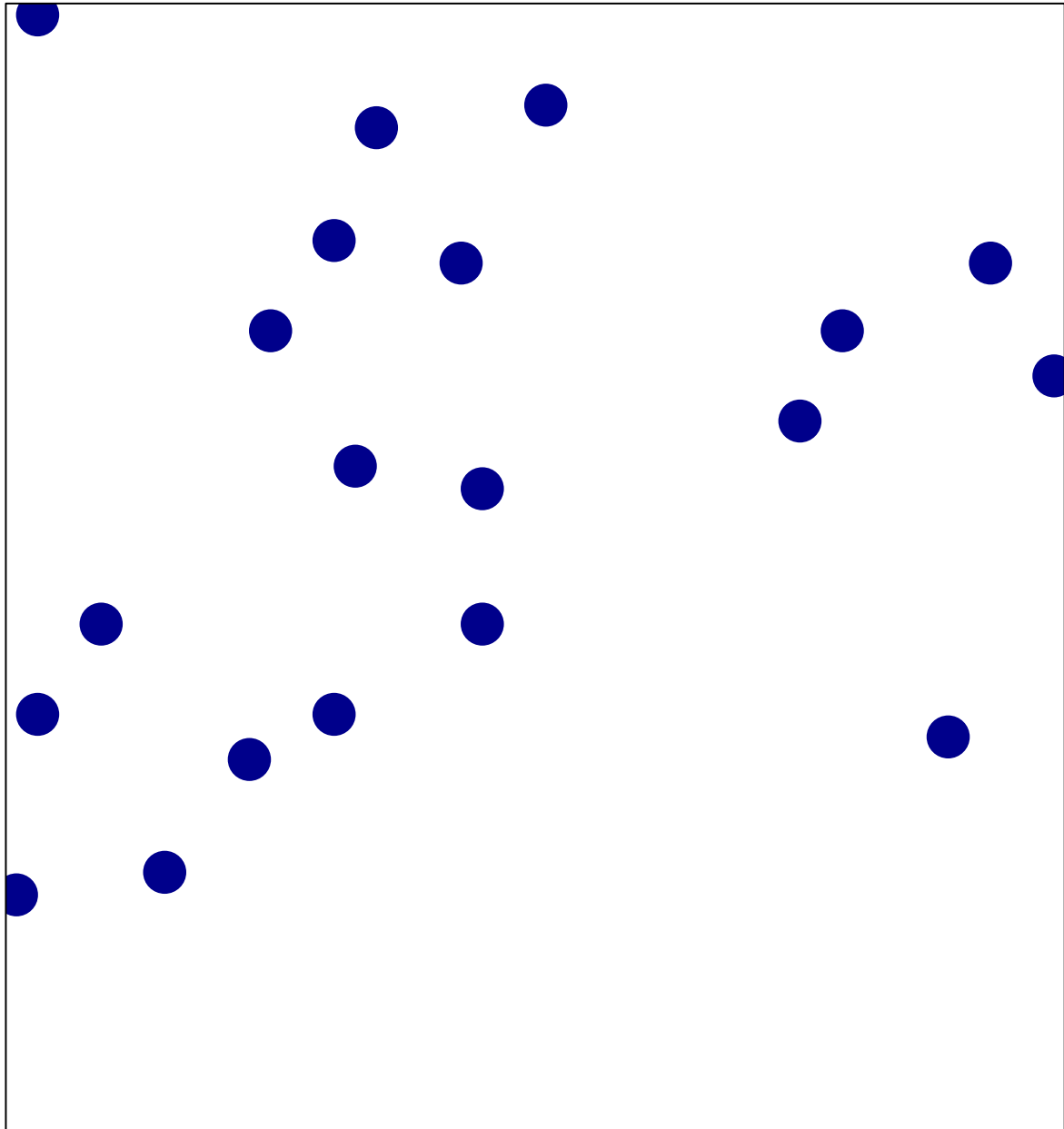


# histone deacetylation



# features = 20  
chi-square p = 0.85

# histone deacetylation



# features = 20 , max = 1