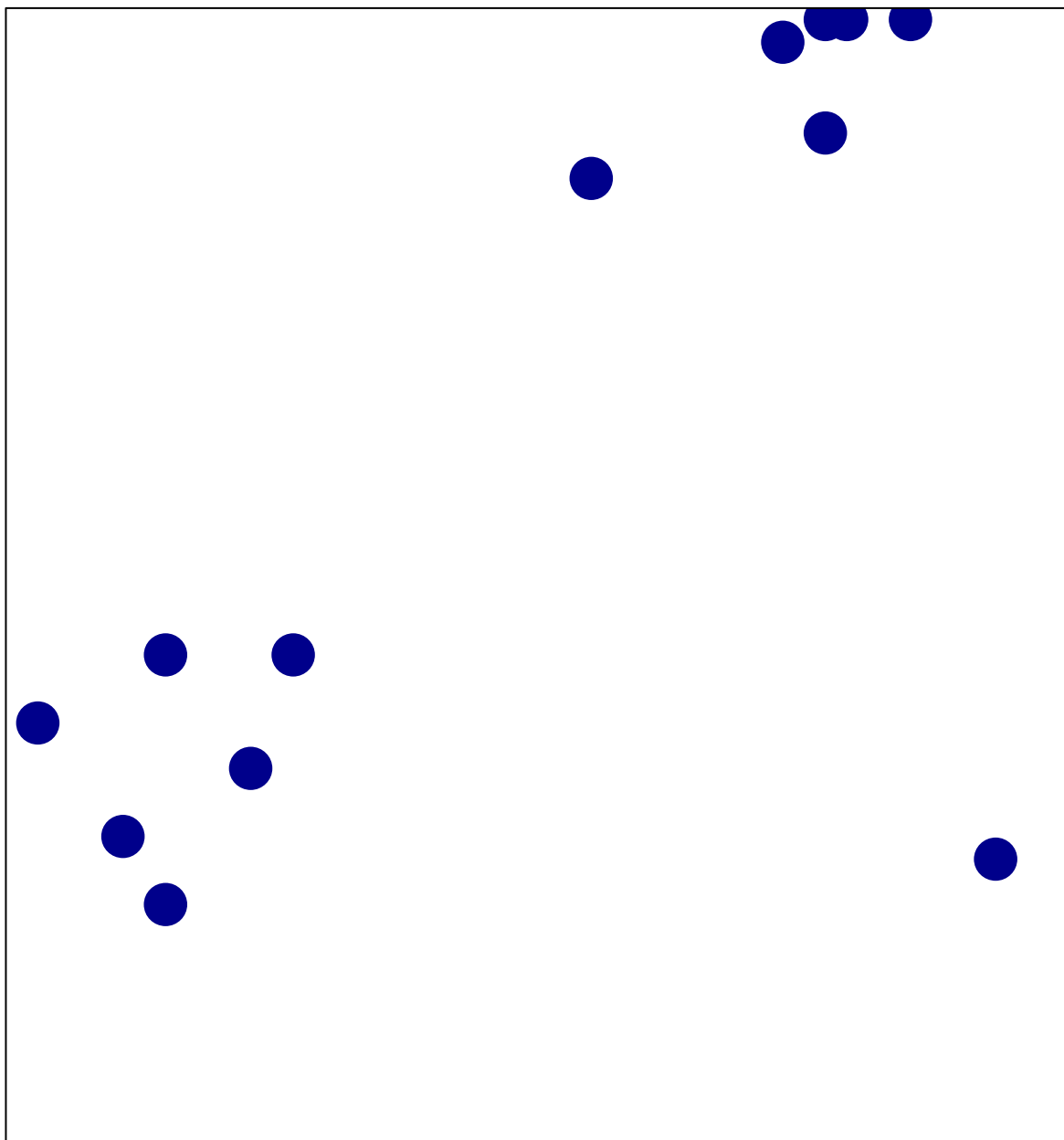


# tRNA aminoacylation



# features = 13  
chi-square p = 0.83

# tRNA aminoacylation



# features = 13 , max = 1