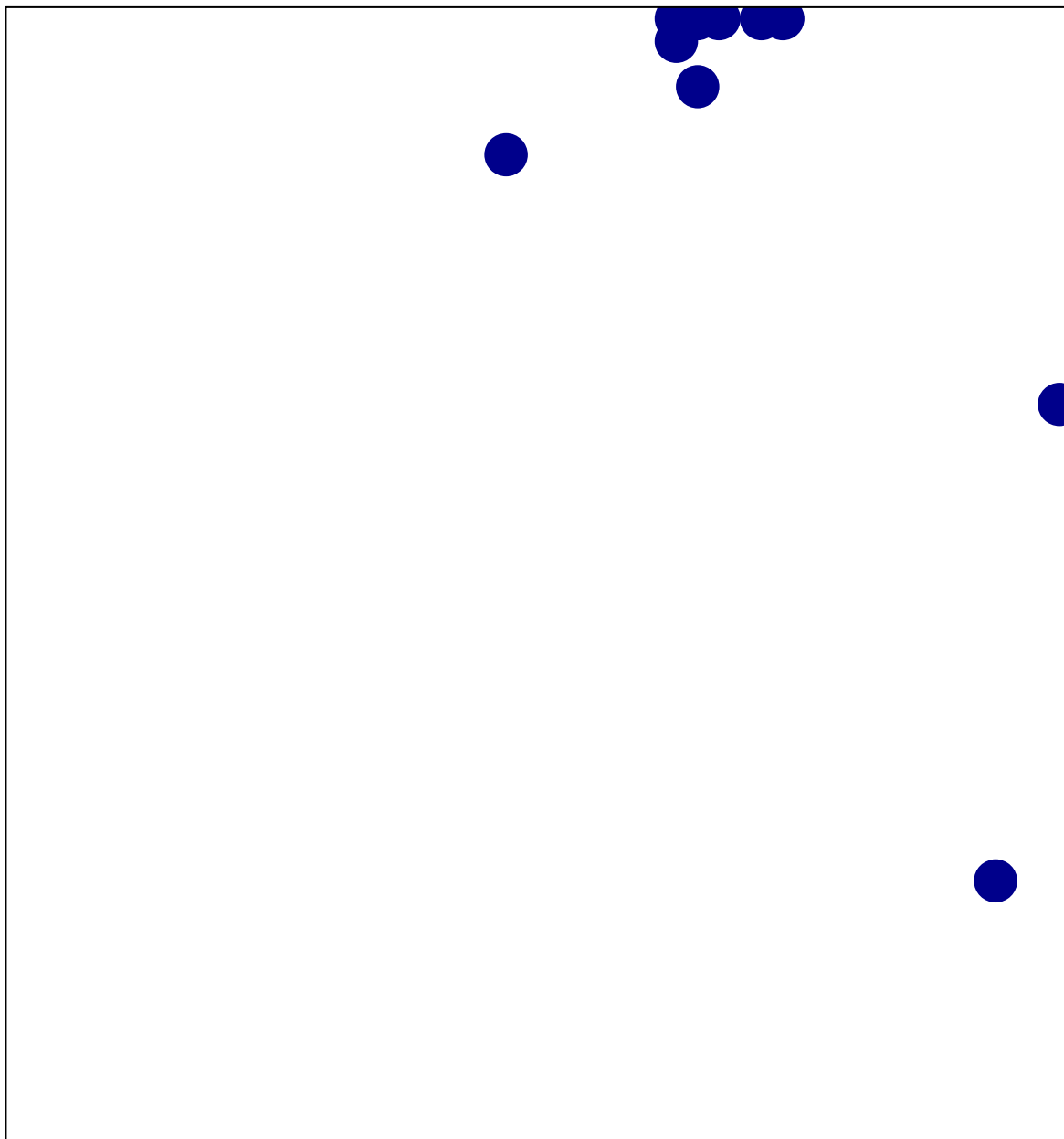


# large ribosomal subunit



# features = 10  
chi-square p = 0.78

# large ribosomal subunit



# features = 10 , max = 1