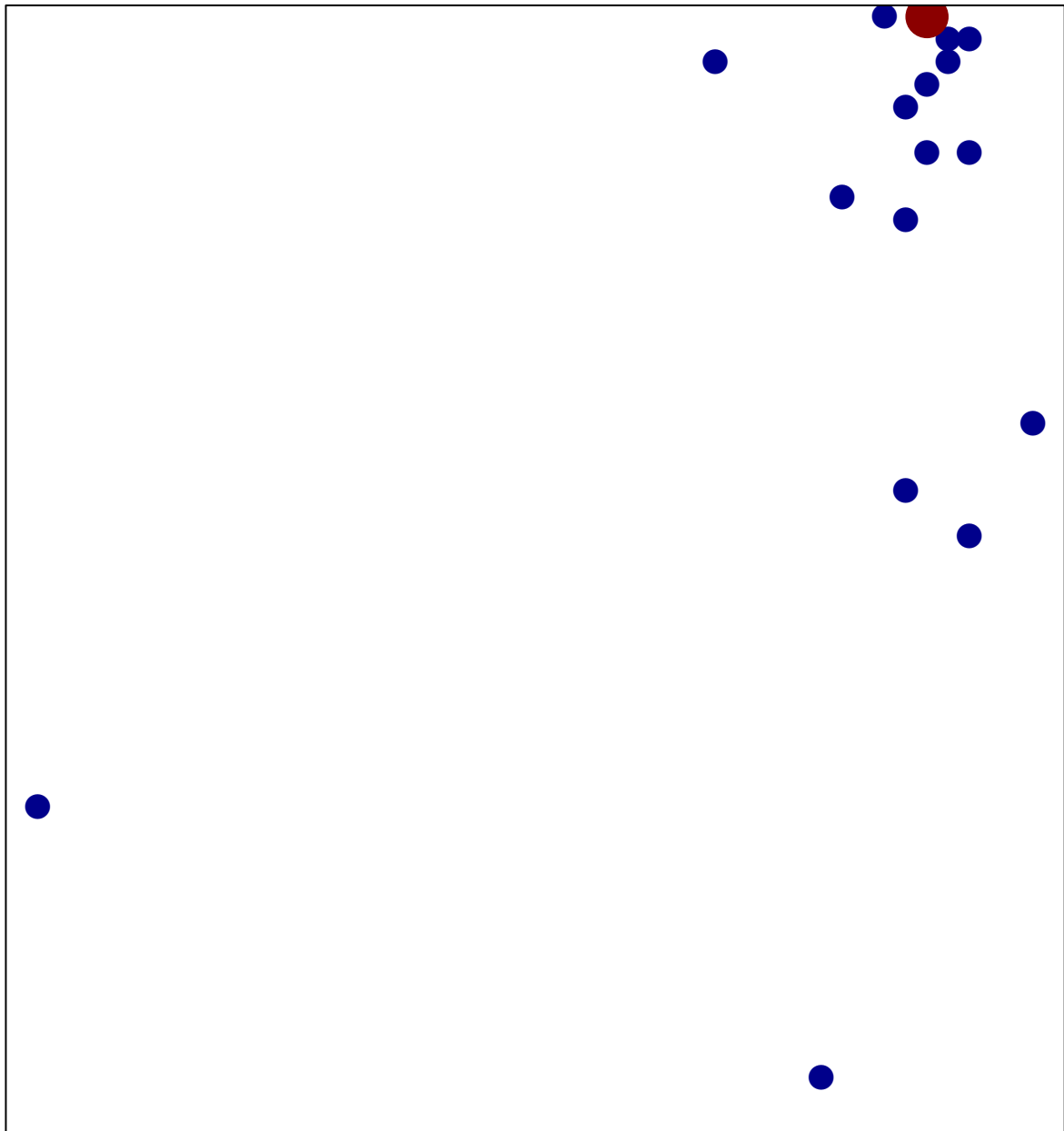


# mitotic sister chromatid segregation



# features = 18  
chi-square p = 0.79

# mitotic sister chromatid segregation



# features = 18 , max = 2