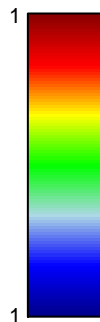
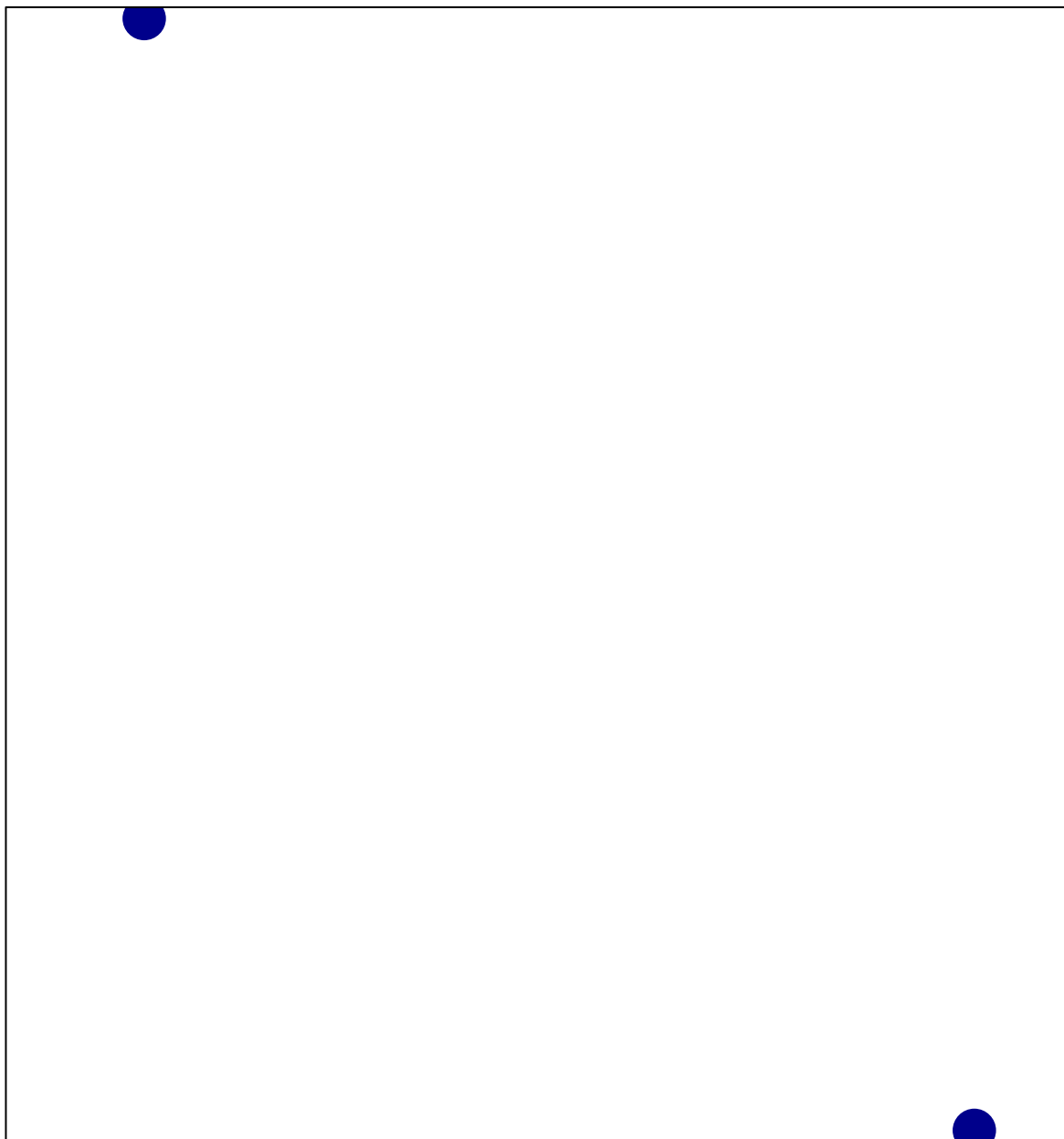
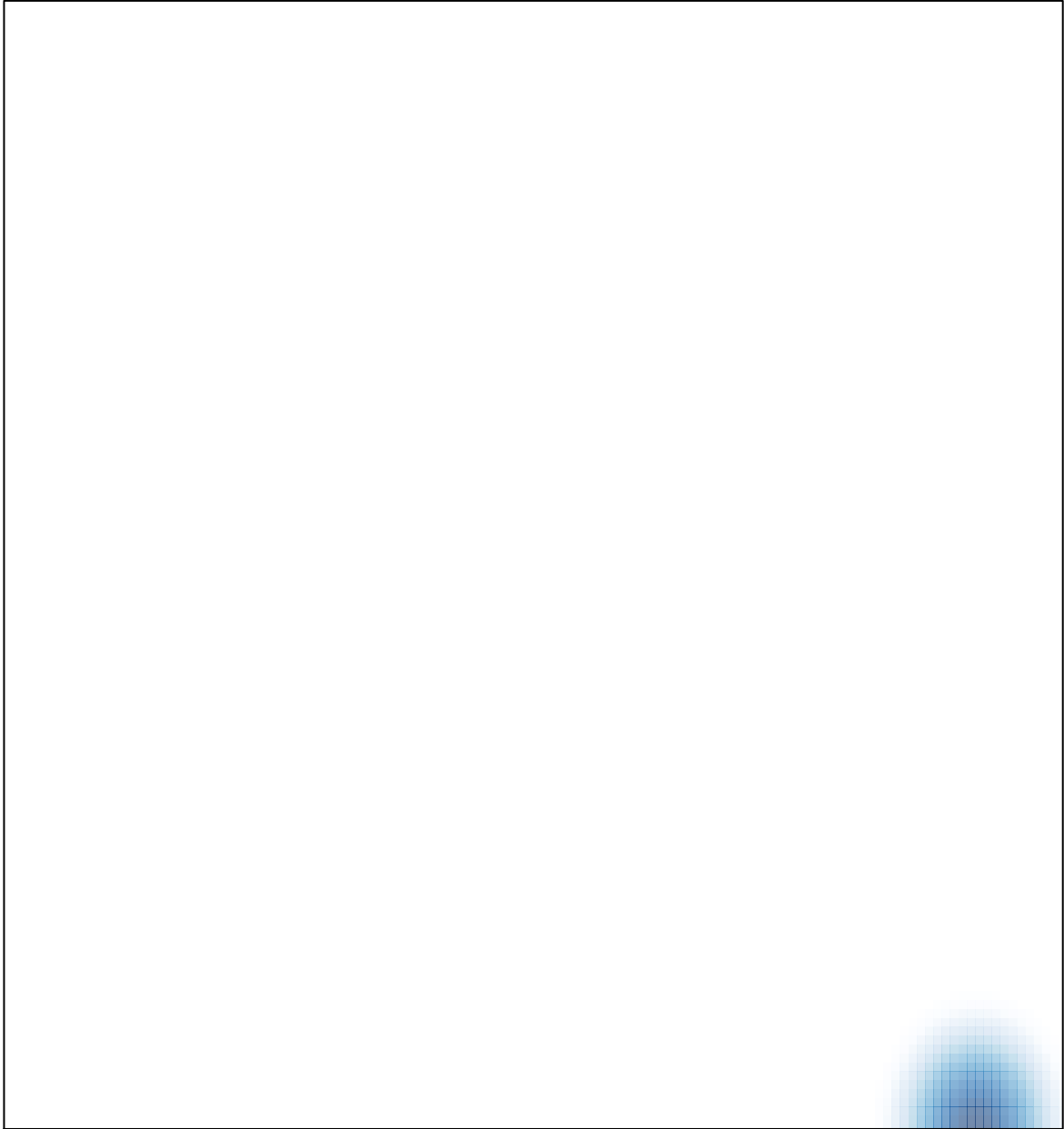


# Hewish\_dMMR-secondary-mutations\_Cell-motility



# features = 2  
chi-square p = 0.84

# Hewish\_dMMR-secondary-mutations\_Cell-motility



# features = 2 , max = 1