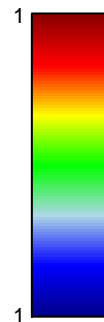
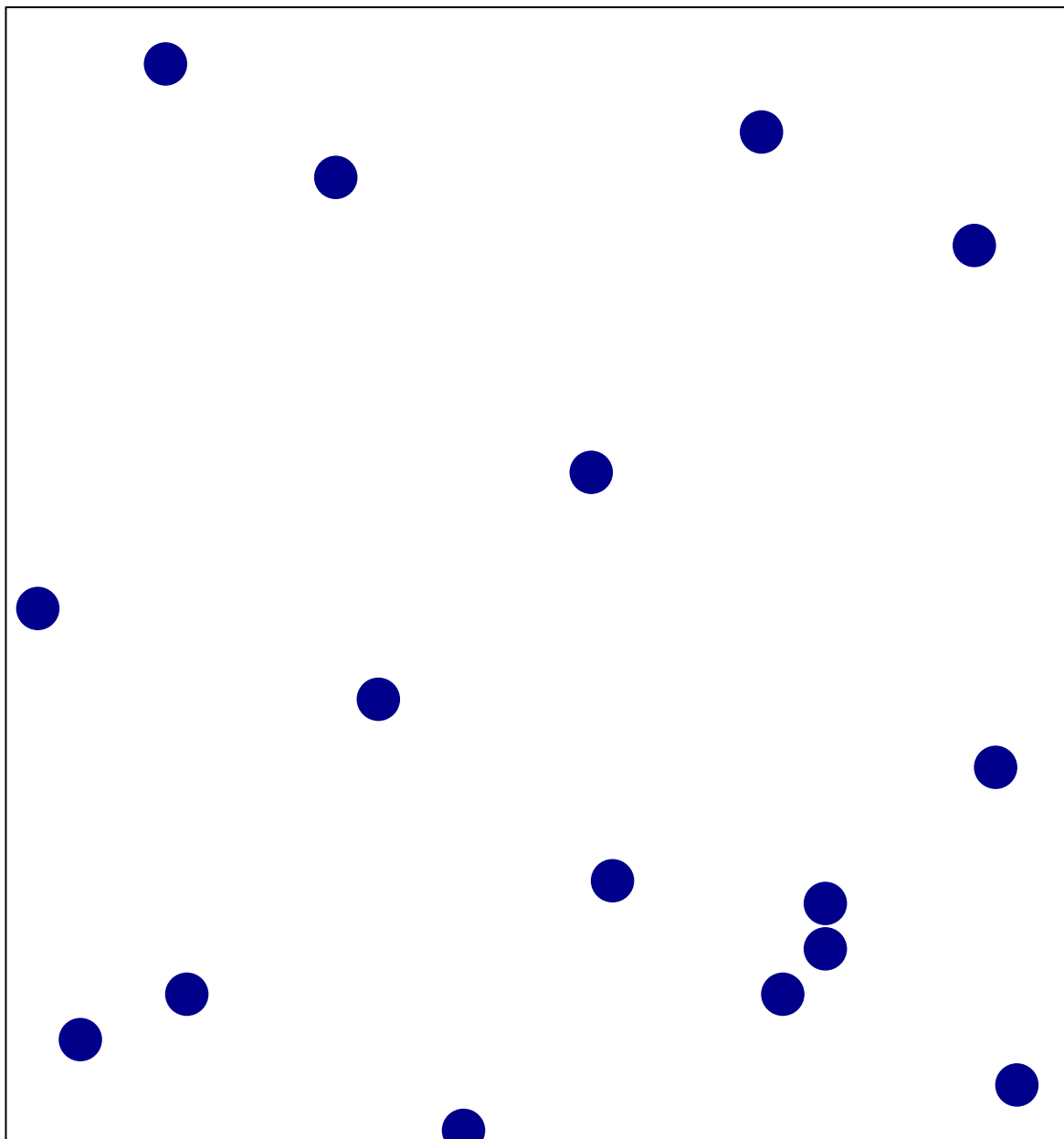
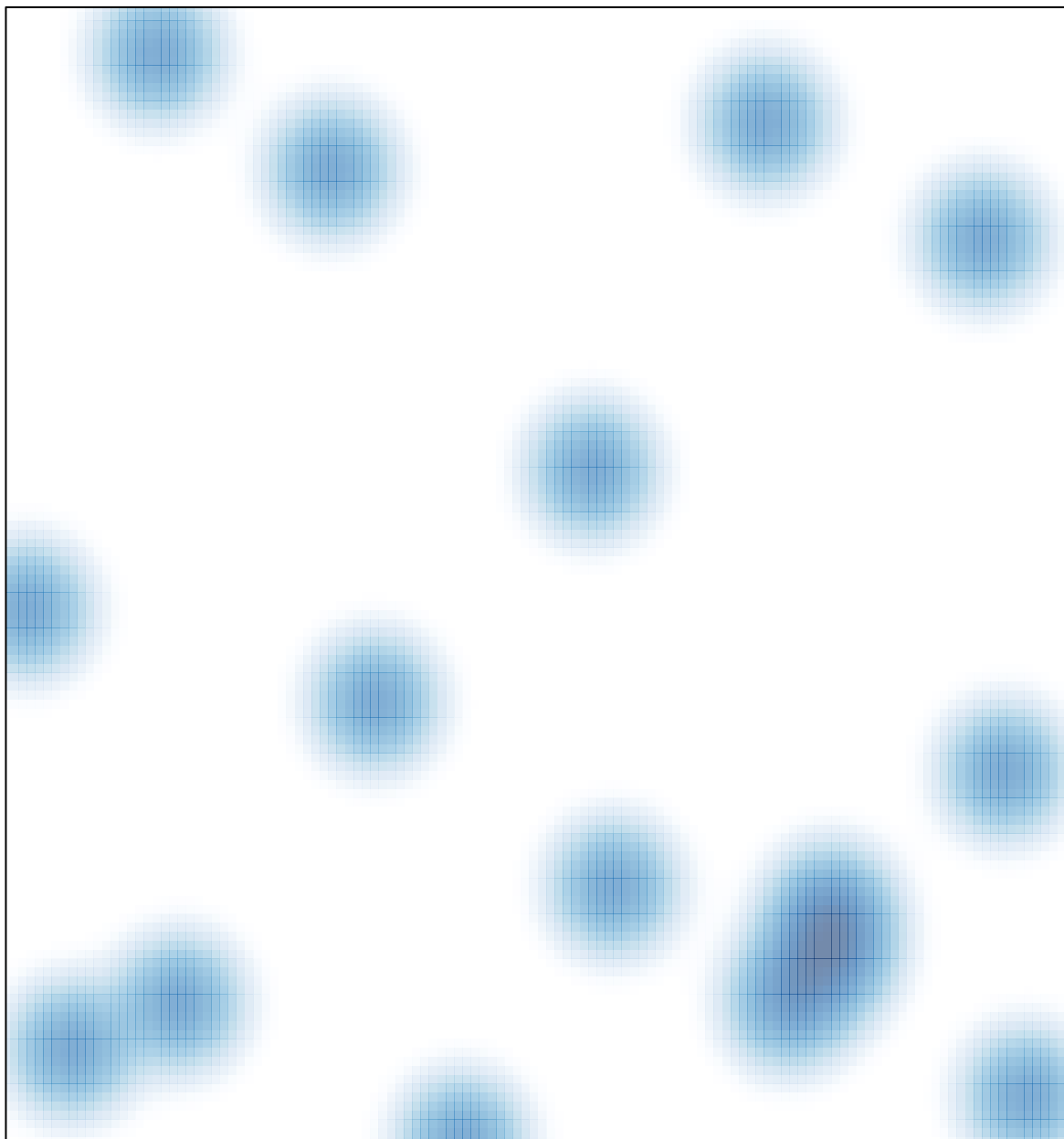


# phosphatidylinositol dephosphorylation



# features = 16  
chi-square p = 0.86

# phosphatidylinositol dephosphorylation



# features = 16 , max = 1