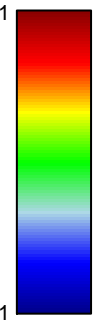
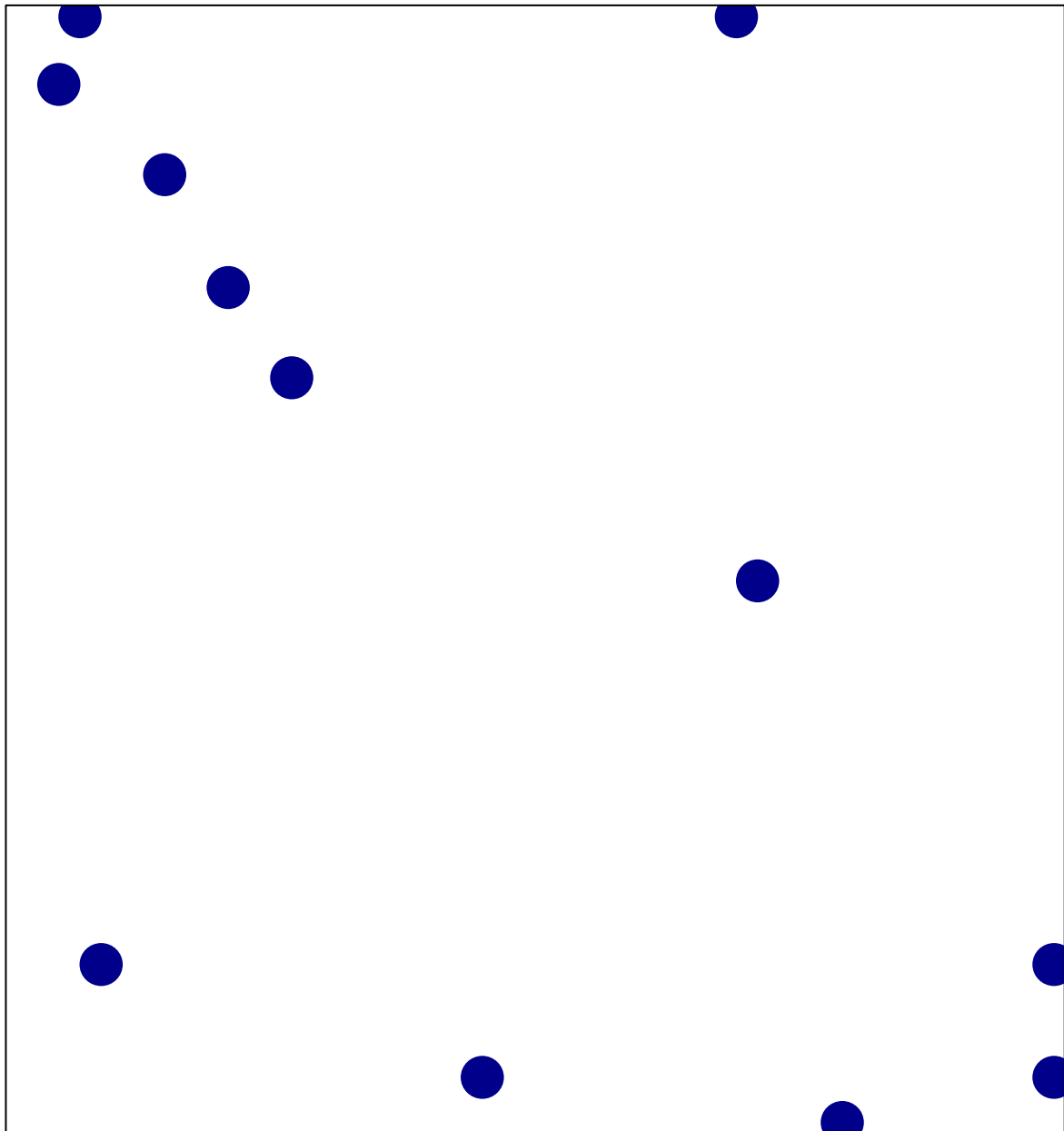
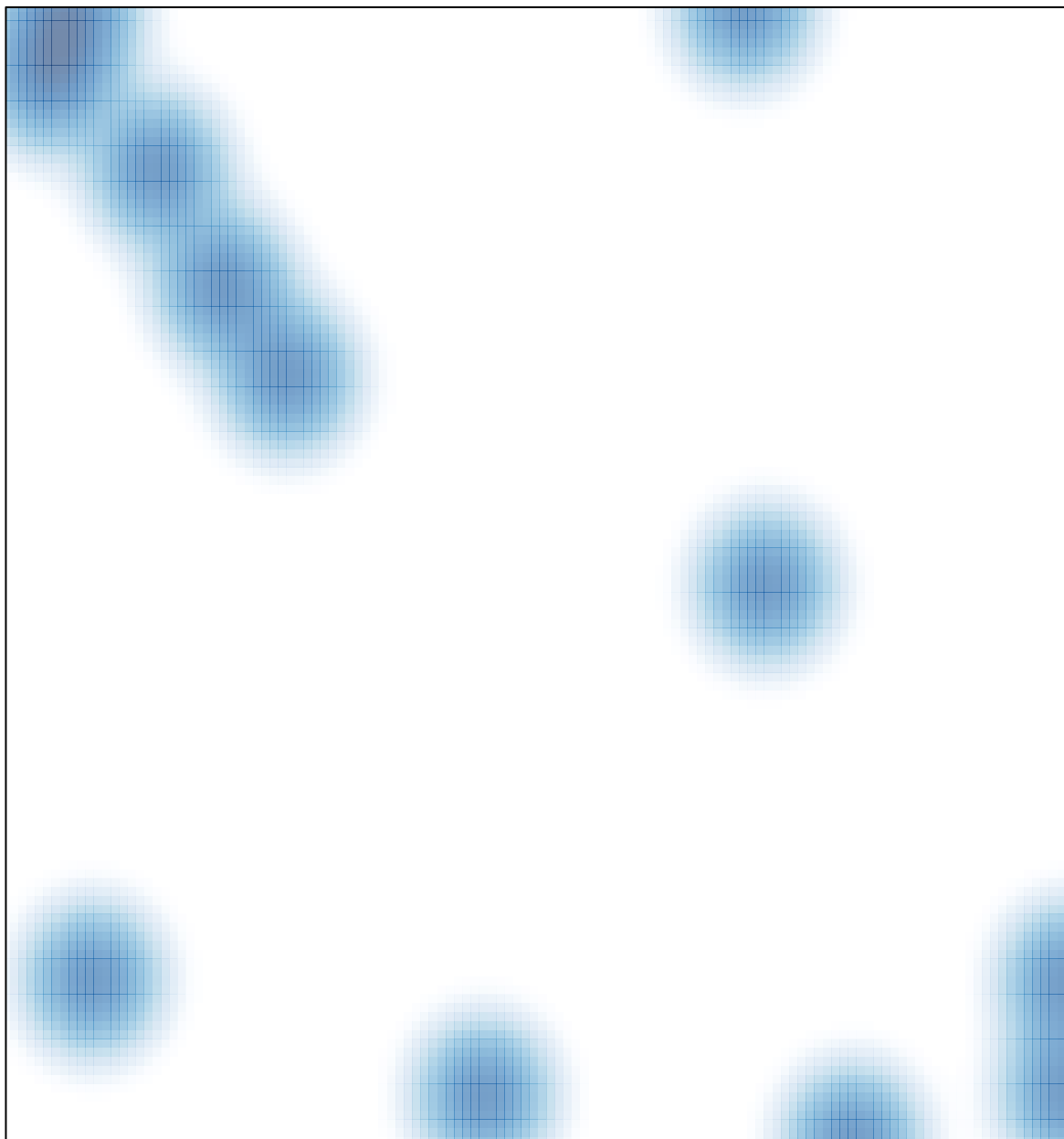


phosphatidylinositol acyl-chain remodeling



features = 12
chi-square p = 0.83

phosphatidylinositol acyl-chain remodeling



features = 12 , max = 1