histone acetyl–lysine binding
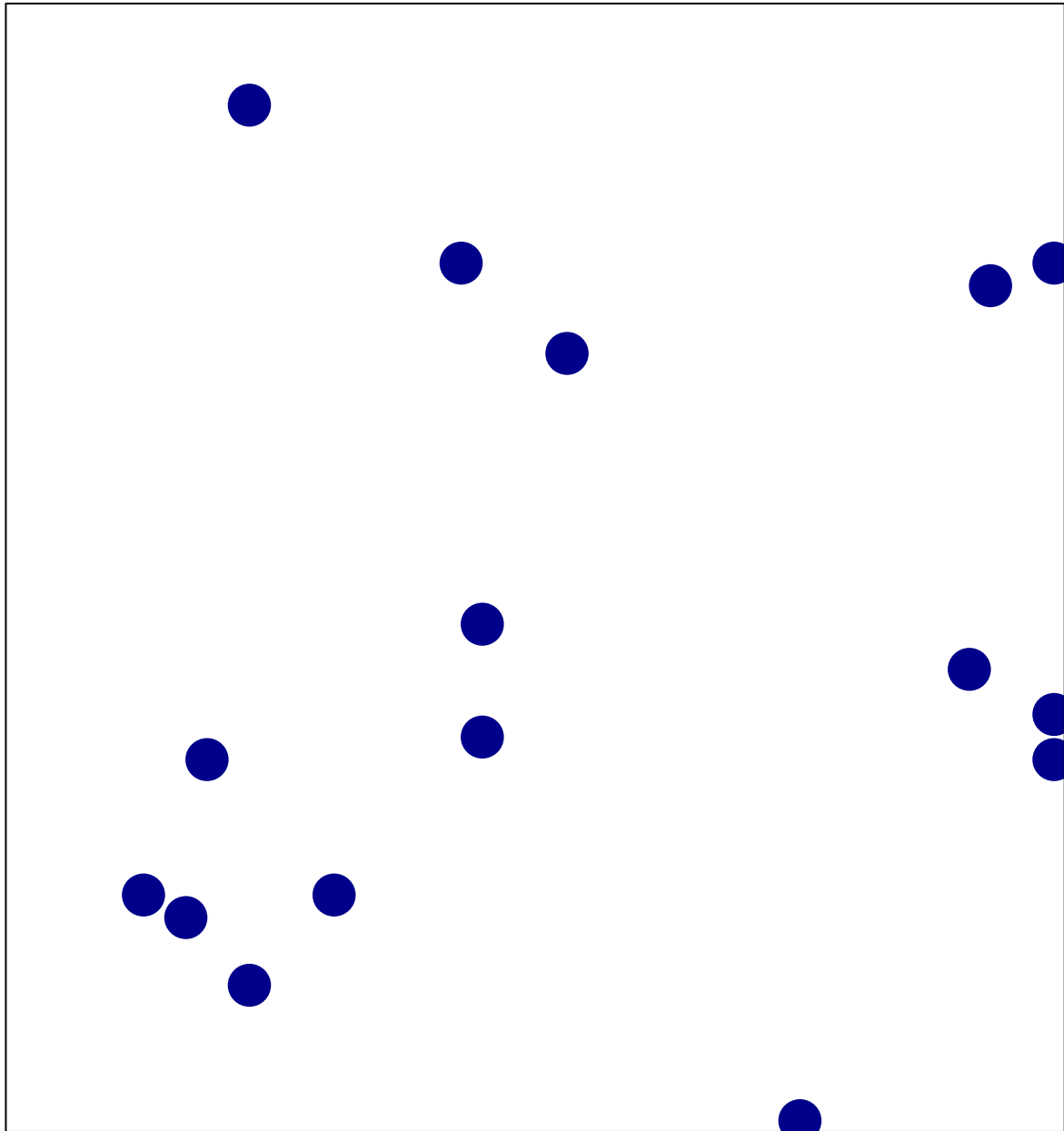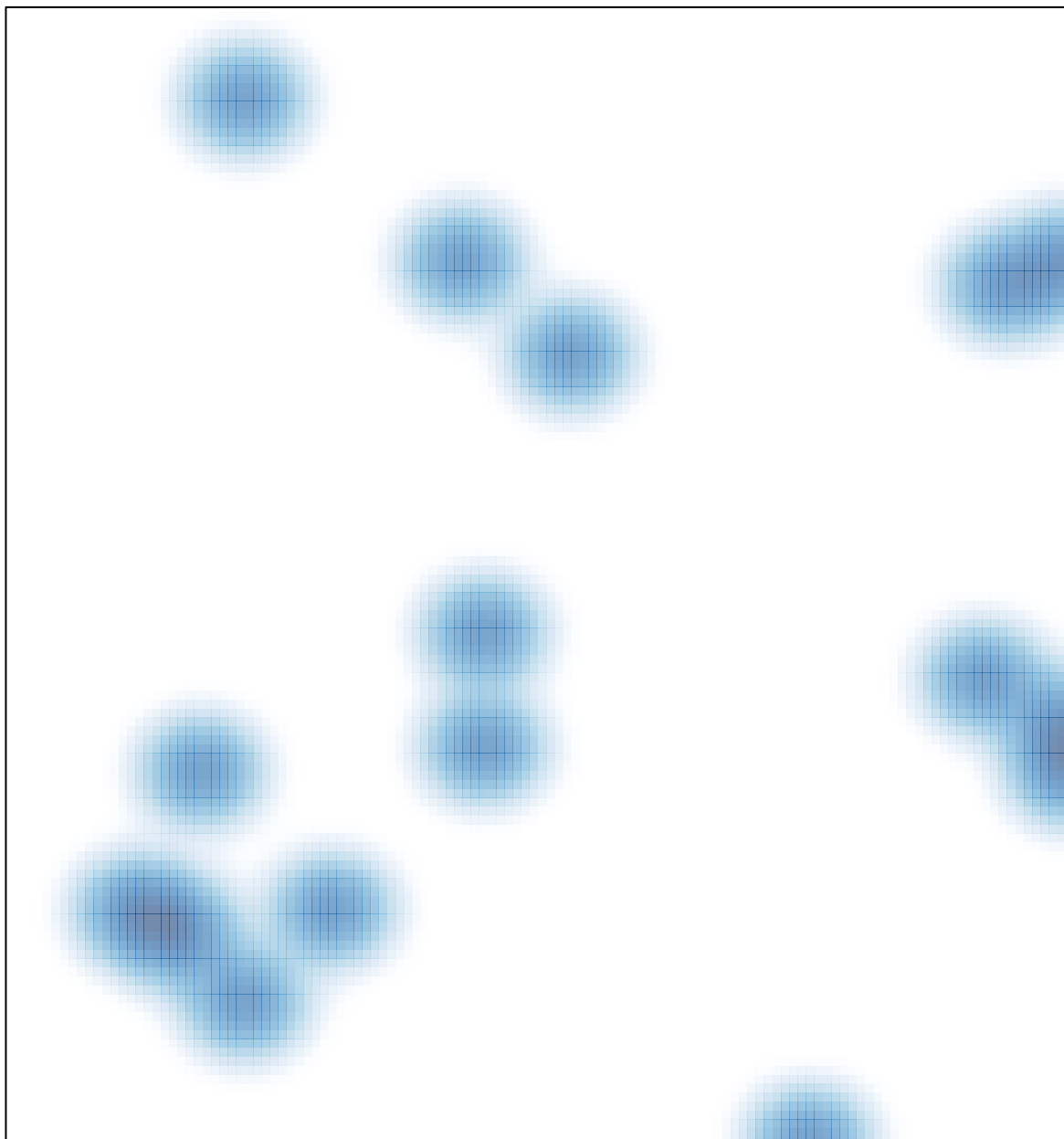
features = 16
chi−square p = 0.83

histone acetyl–lysine binding

features = 16 , max = 1