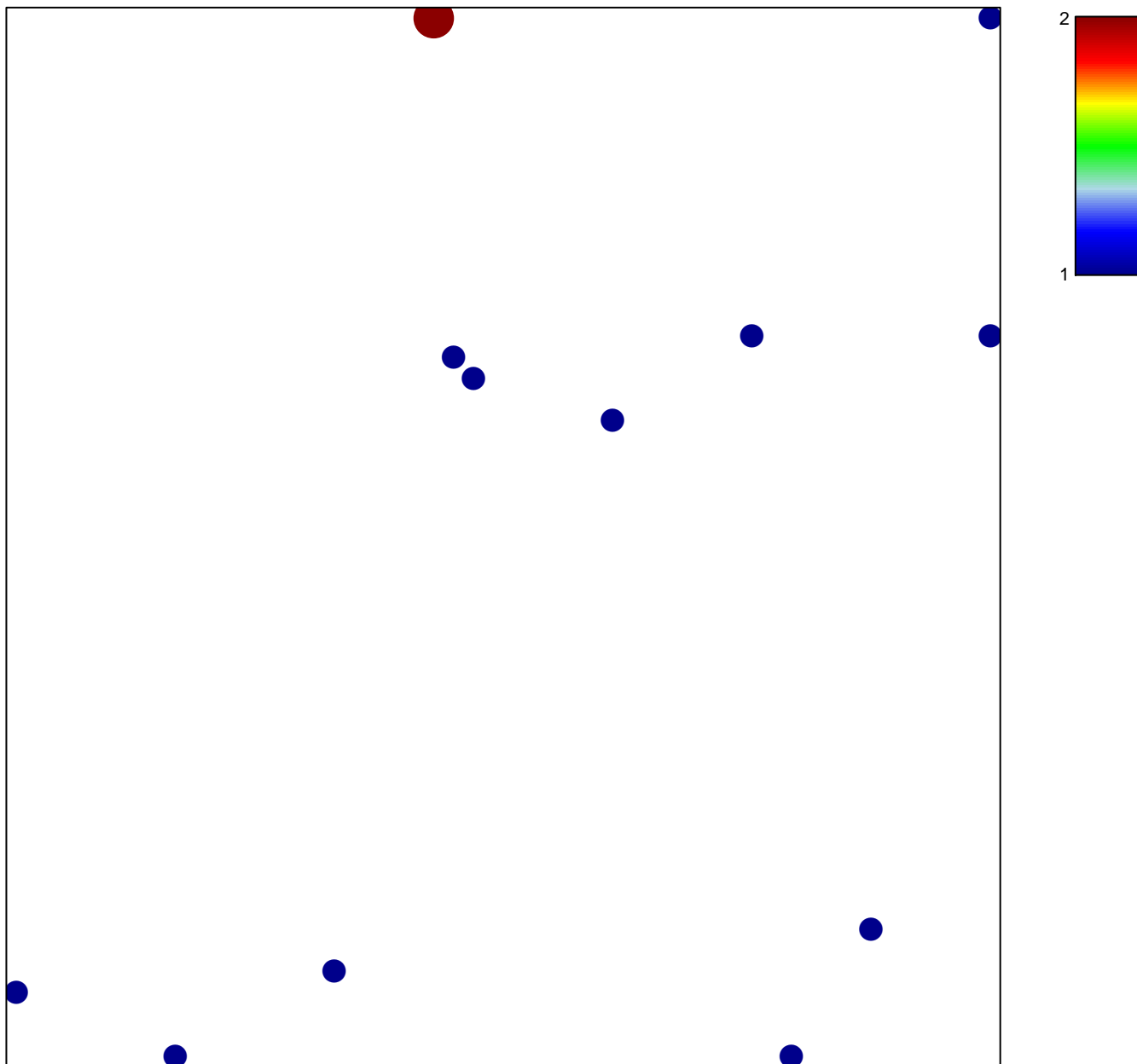
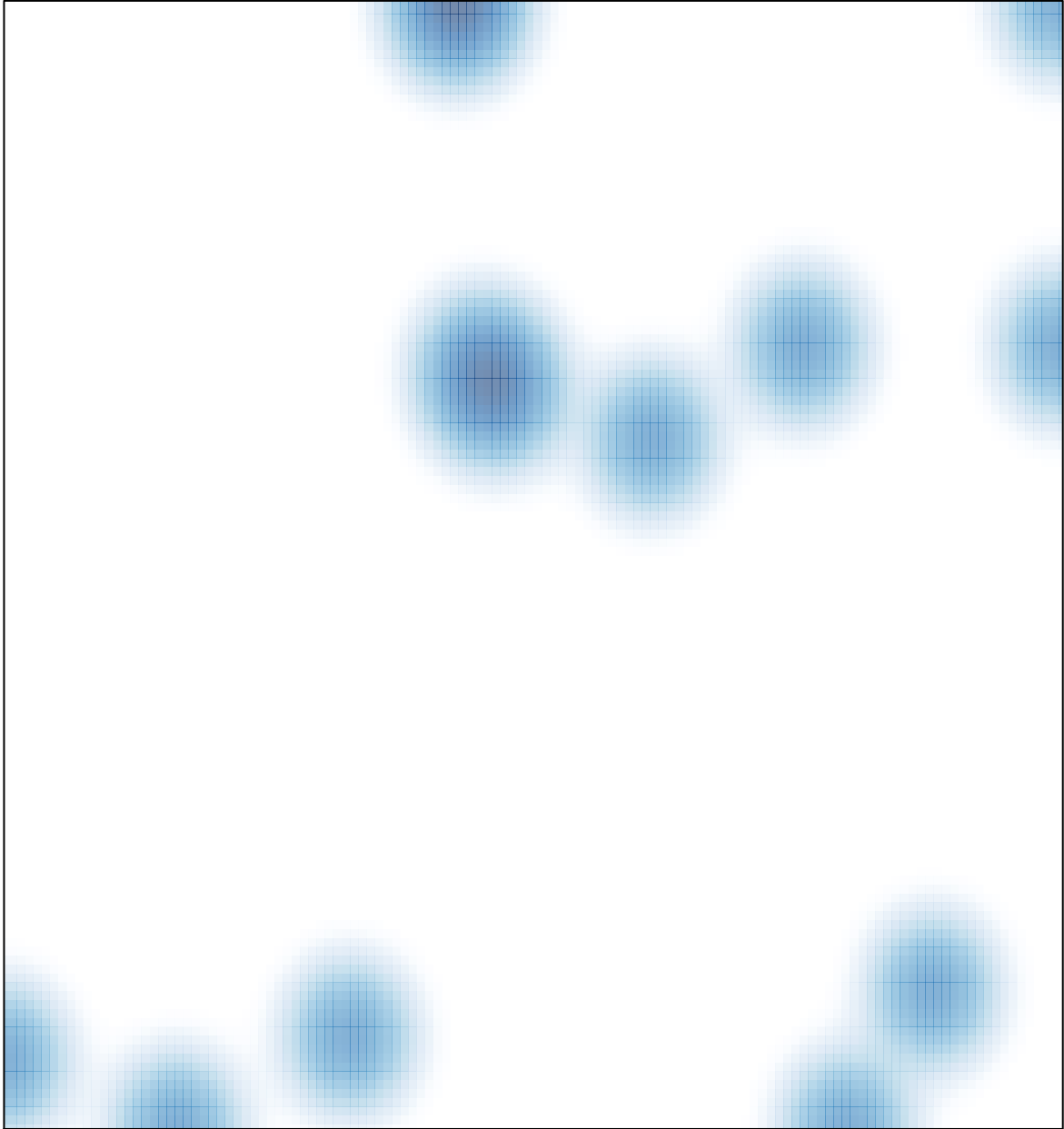


# SMAD protein signal transduction



# features = 13  
chi-square p = 0.82

# SMAD protein signal transduction



# features = 13 , max = 2