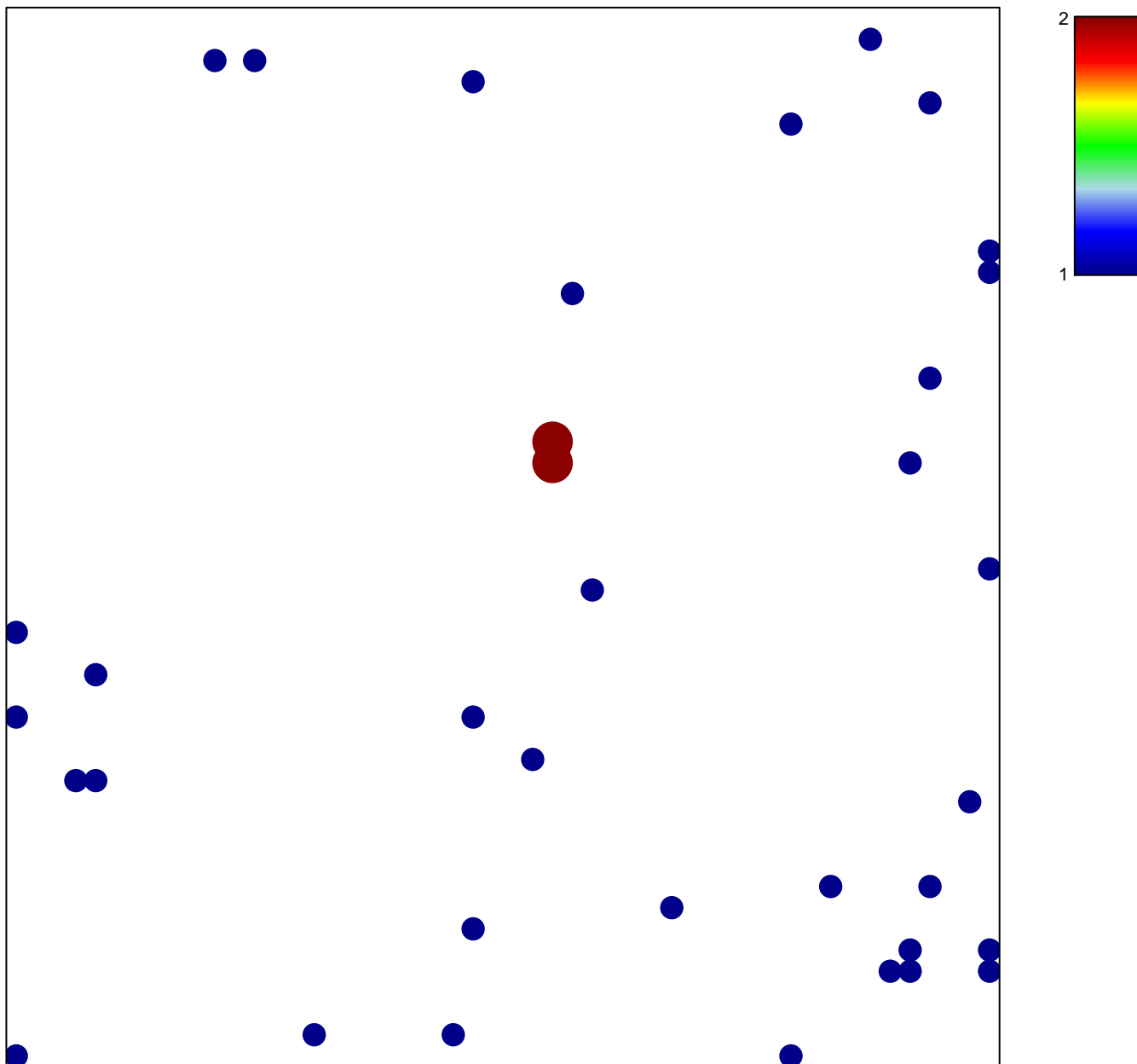
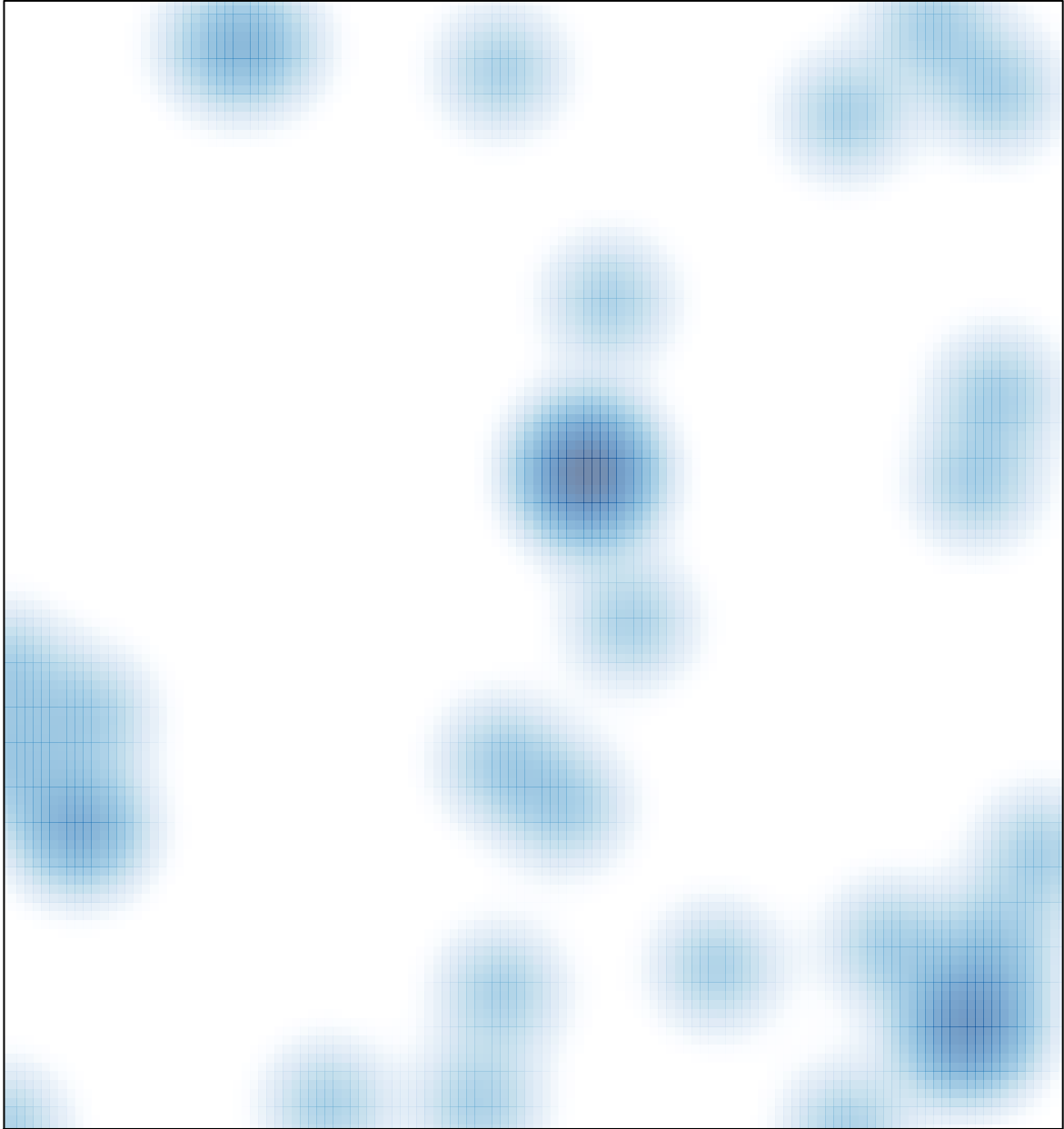


# actin cytoskeleton reorganization



# features = 38  
chi-square p = 0.81

# actin cytoskeleton reorganization



# features = 38 , max = 2