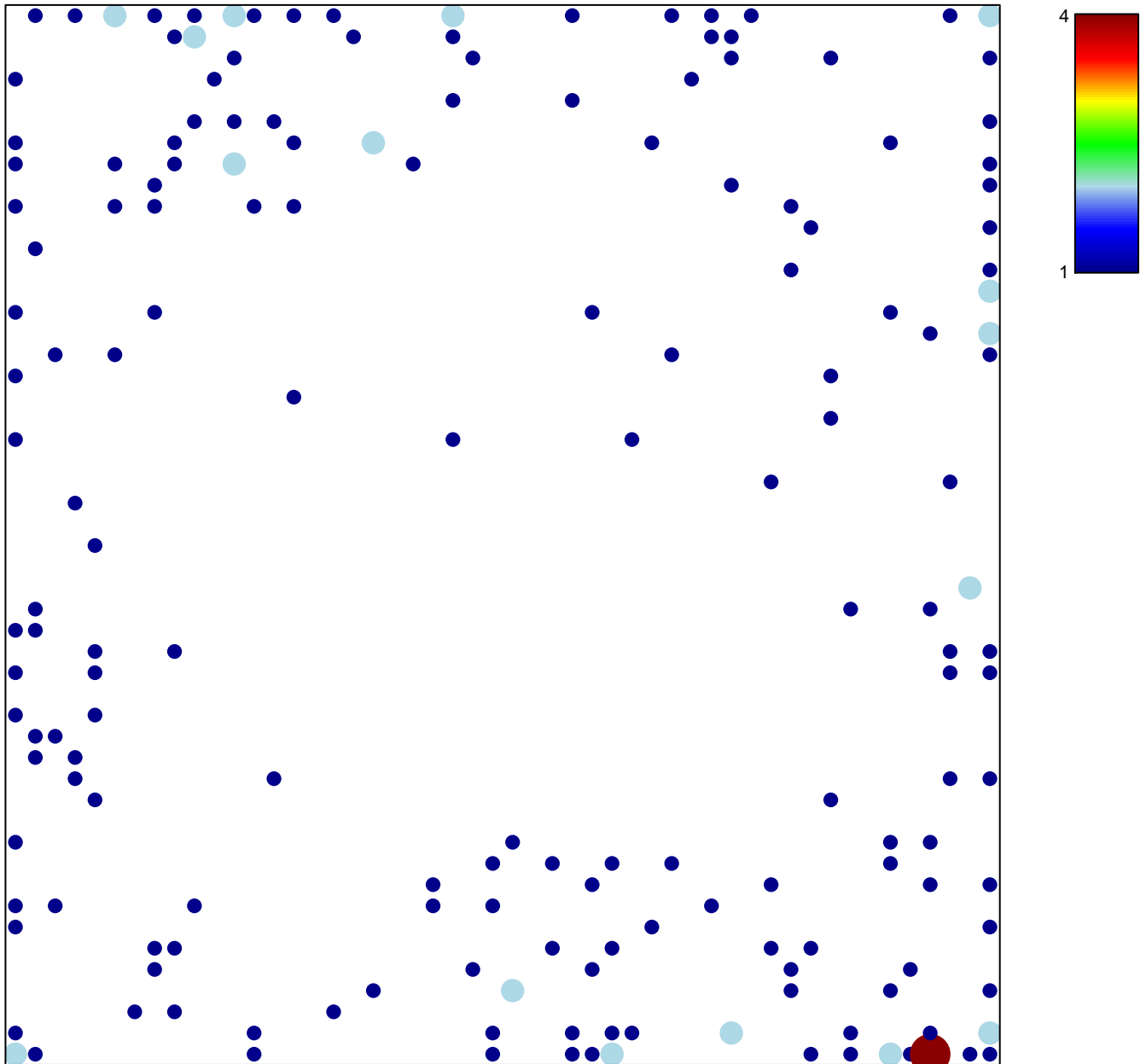
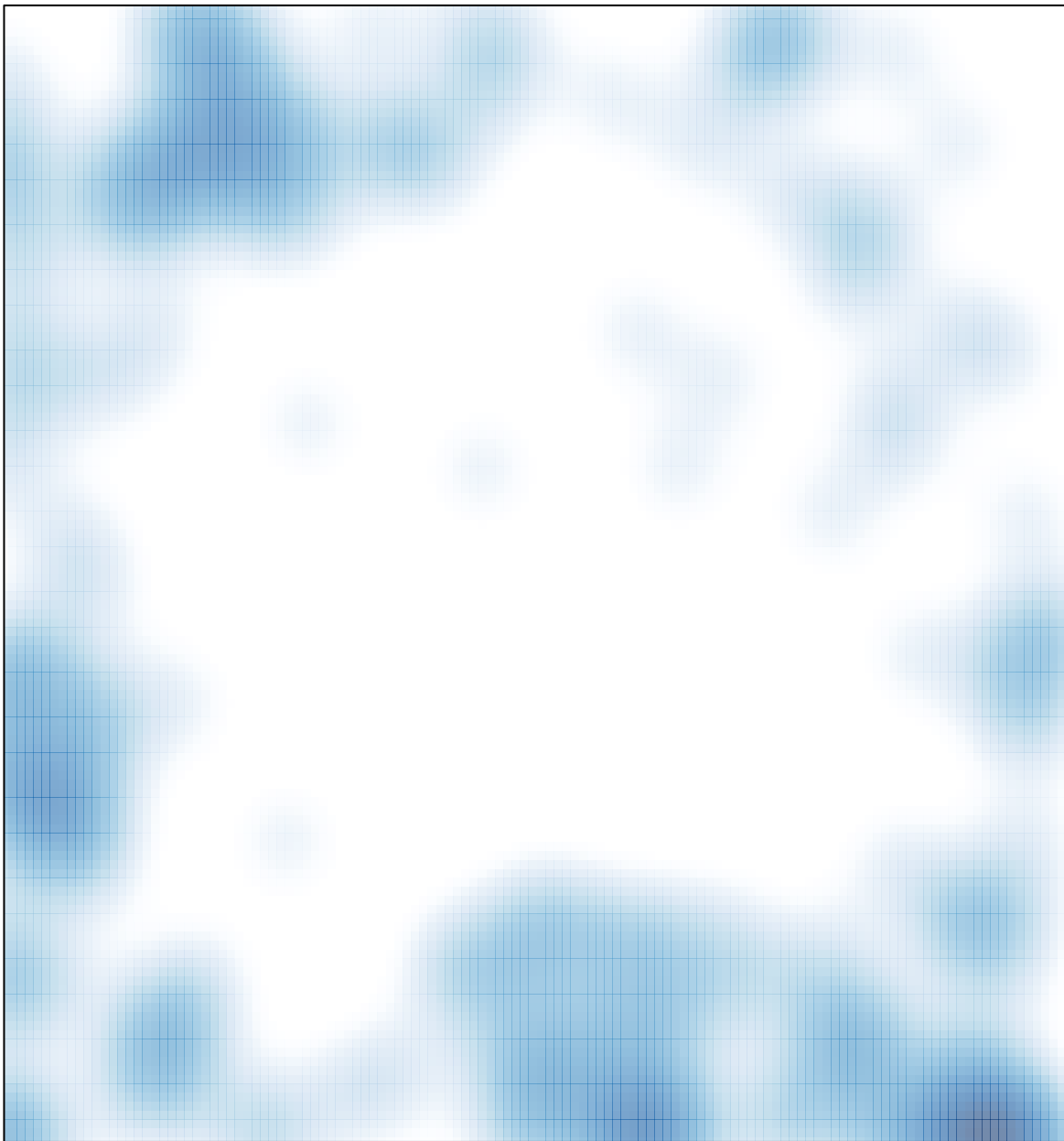


# hsa-miR-206



# features = 195  
chi-square p = 0.7

# hsa-miR-206



# features = 195 , max = 4