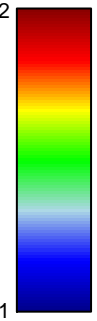
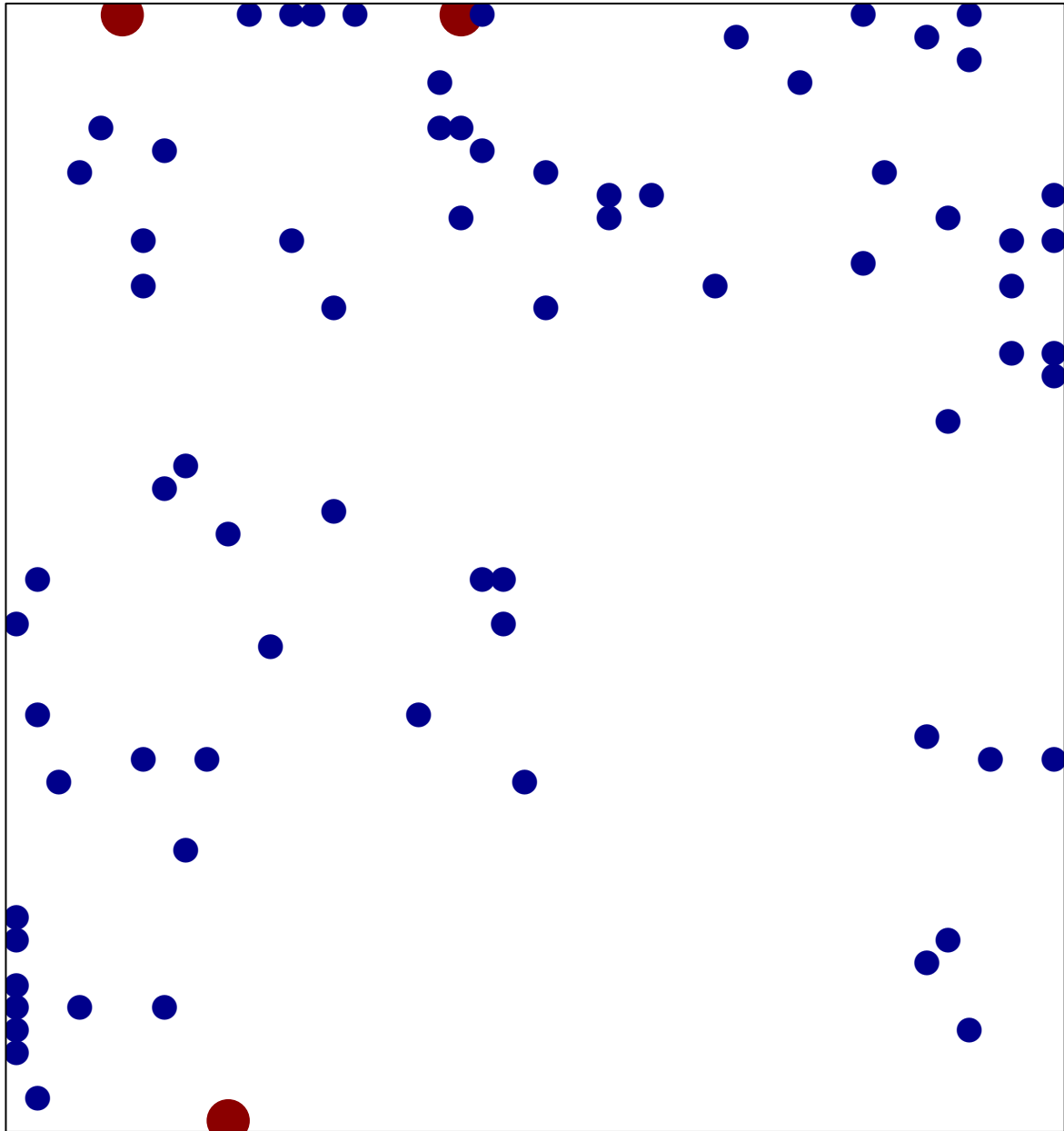
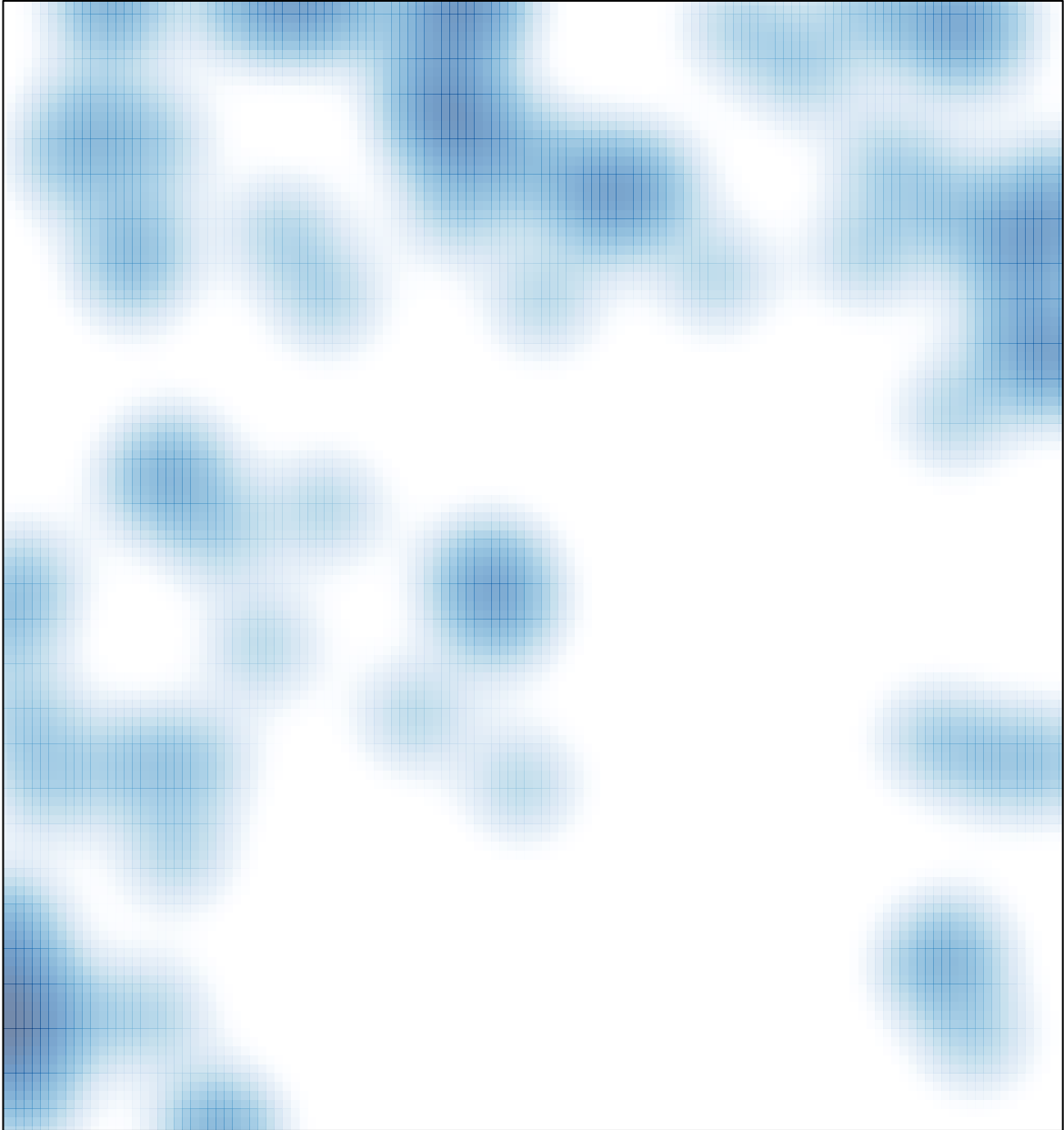


# hsa-miR-205



# features = 78  
chi-square p = 0.8

# hsa-miR-205



# features = 78 , max = 2