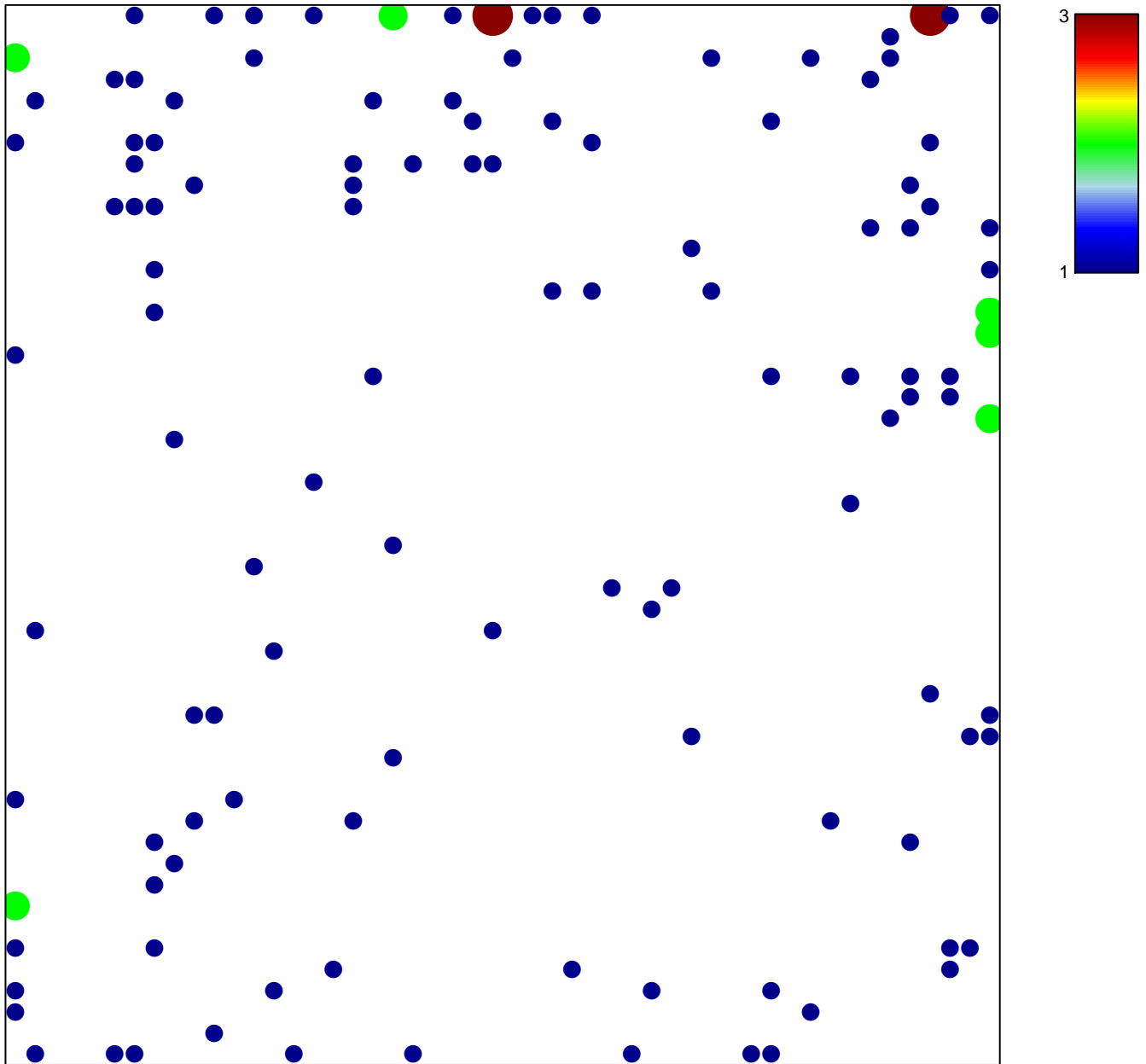
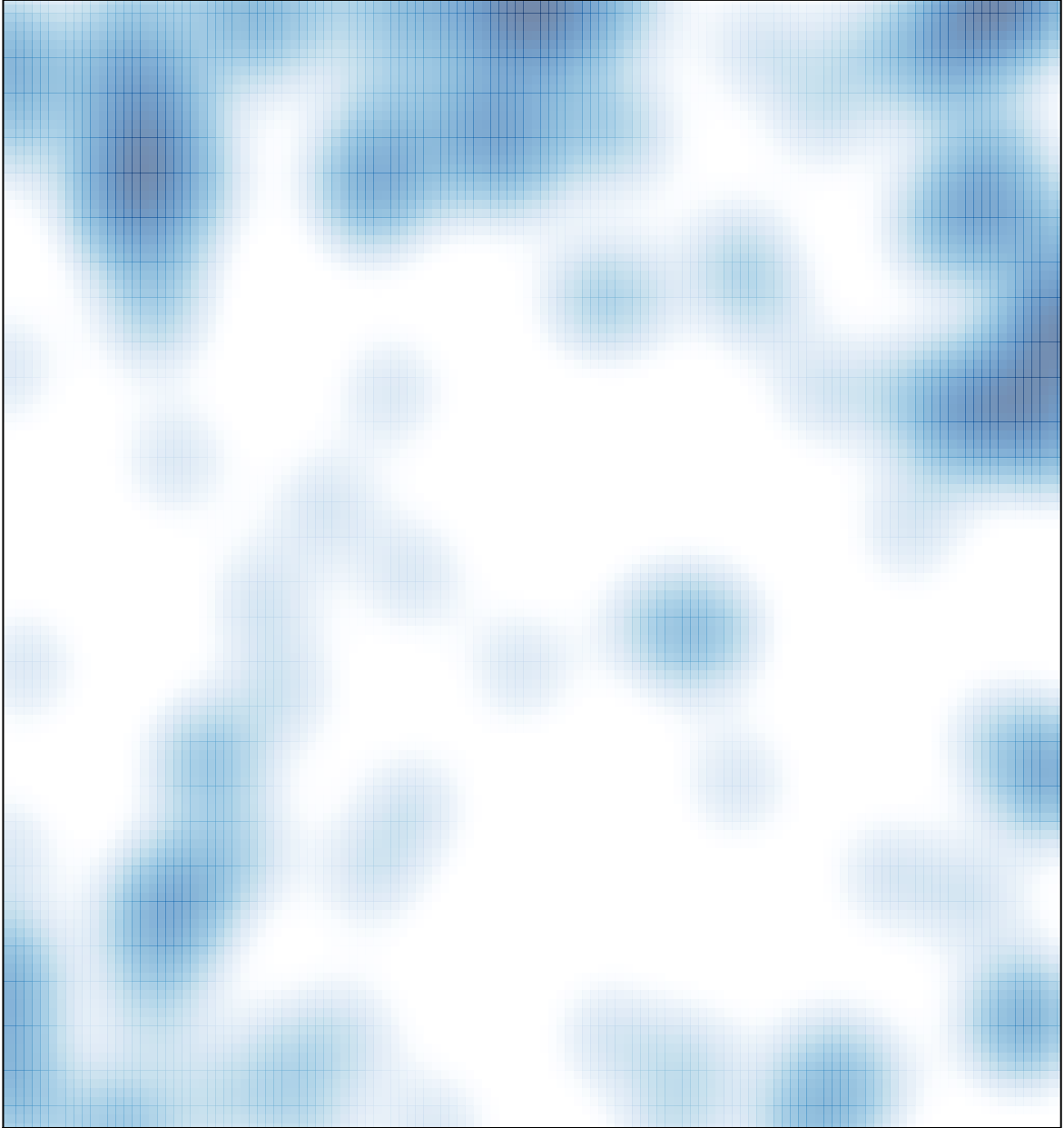


# hsa-miR-218



# features = 131  
chi-square p = 0.75

# hsa-miR-218



# features = 131 , max = 3