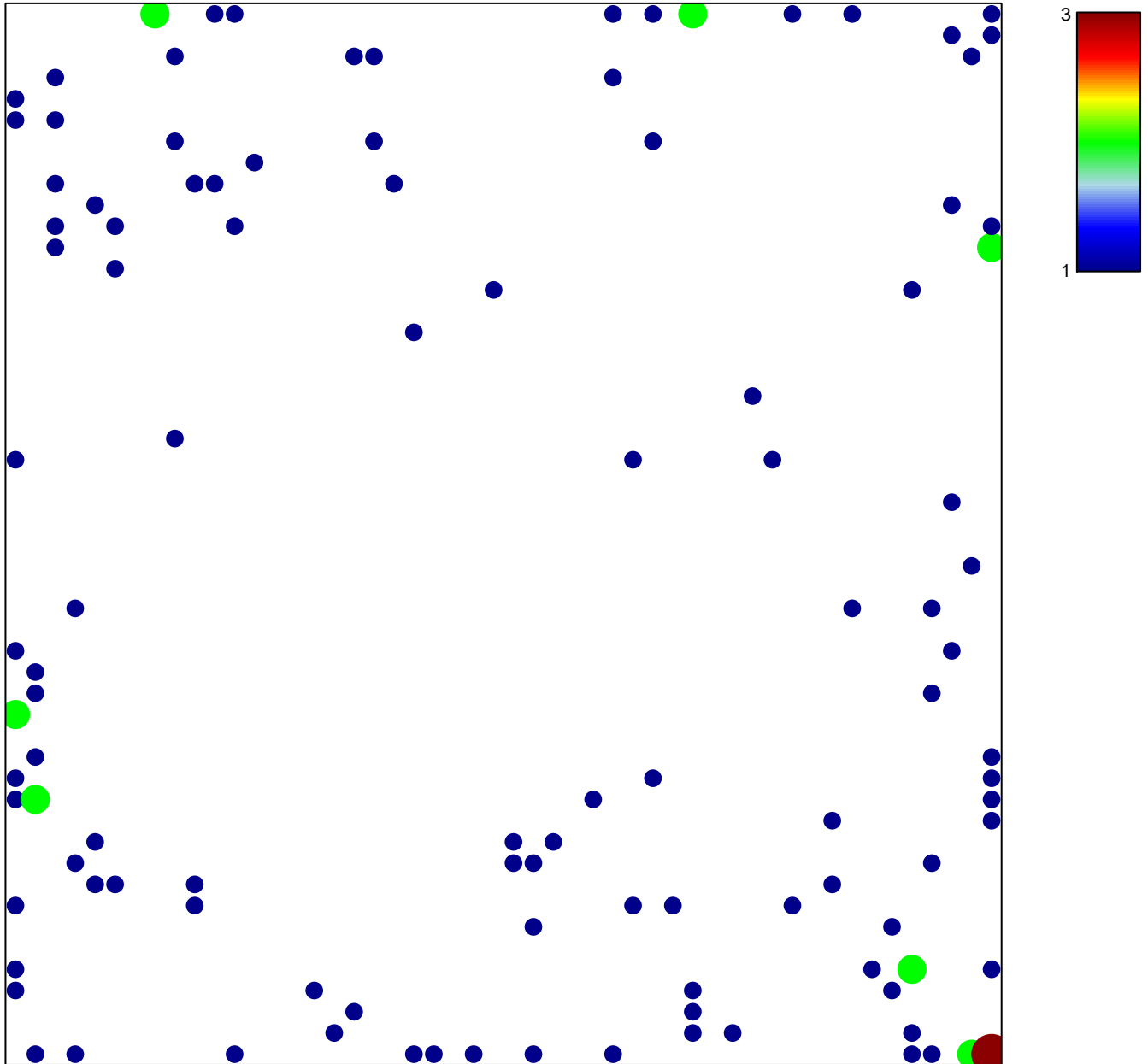
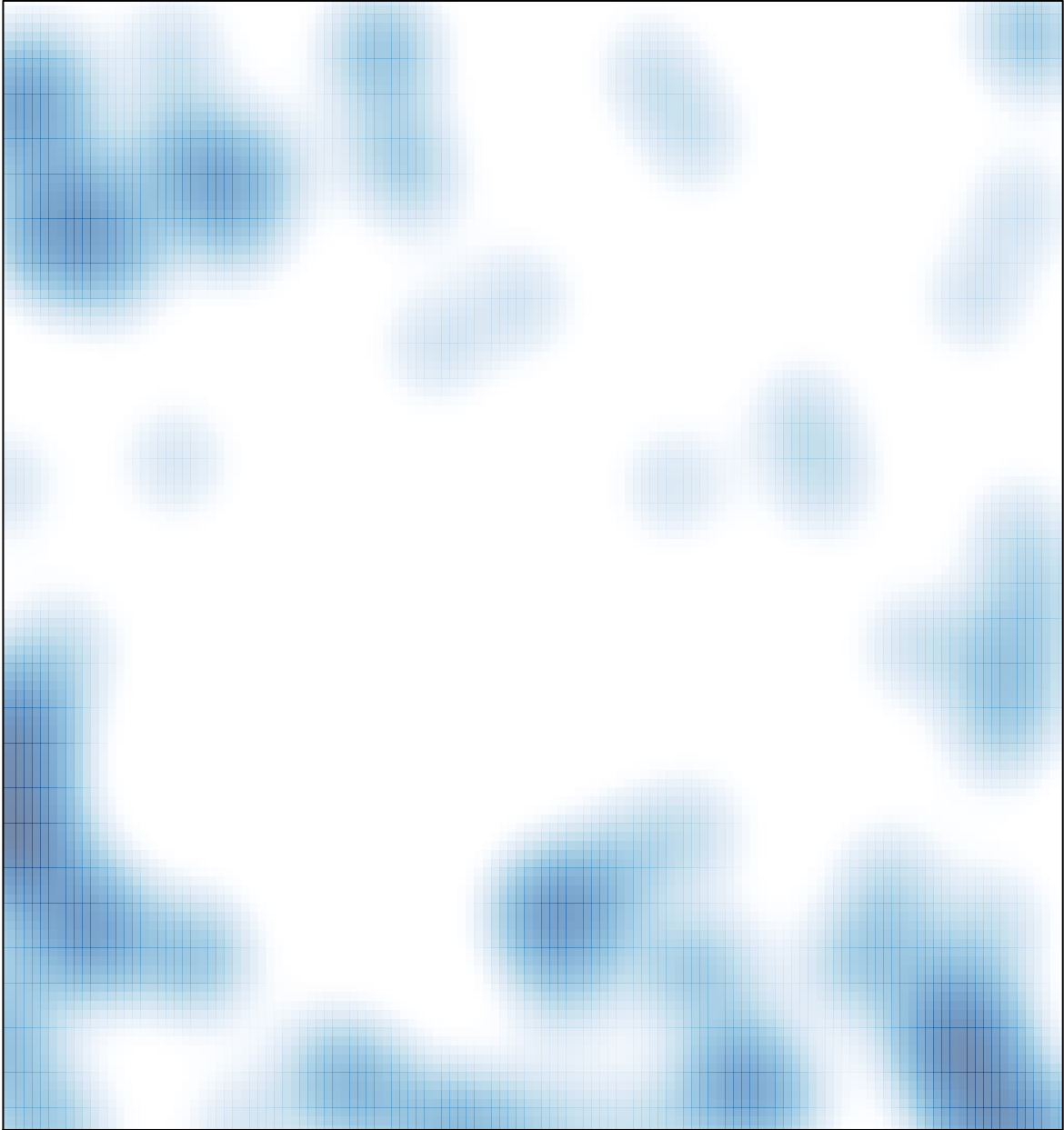


# hsa-miR-105



# features = 120  
chi-square p = 0.74

# hsa-miR-105



# features = 120 , max = 3