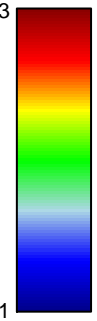
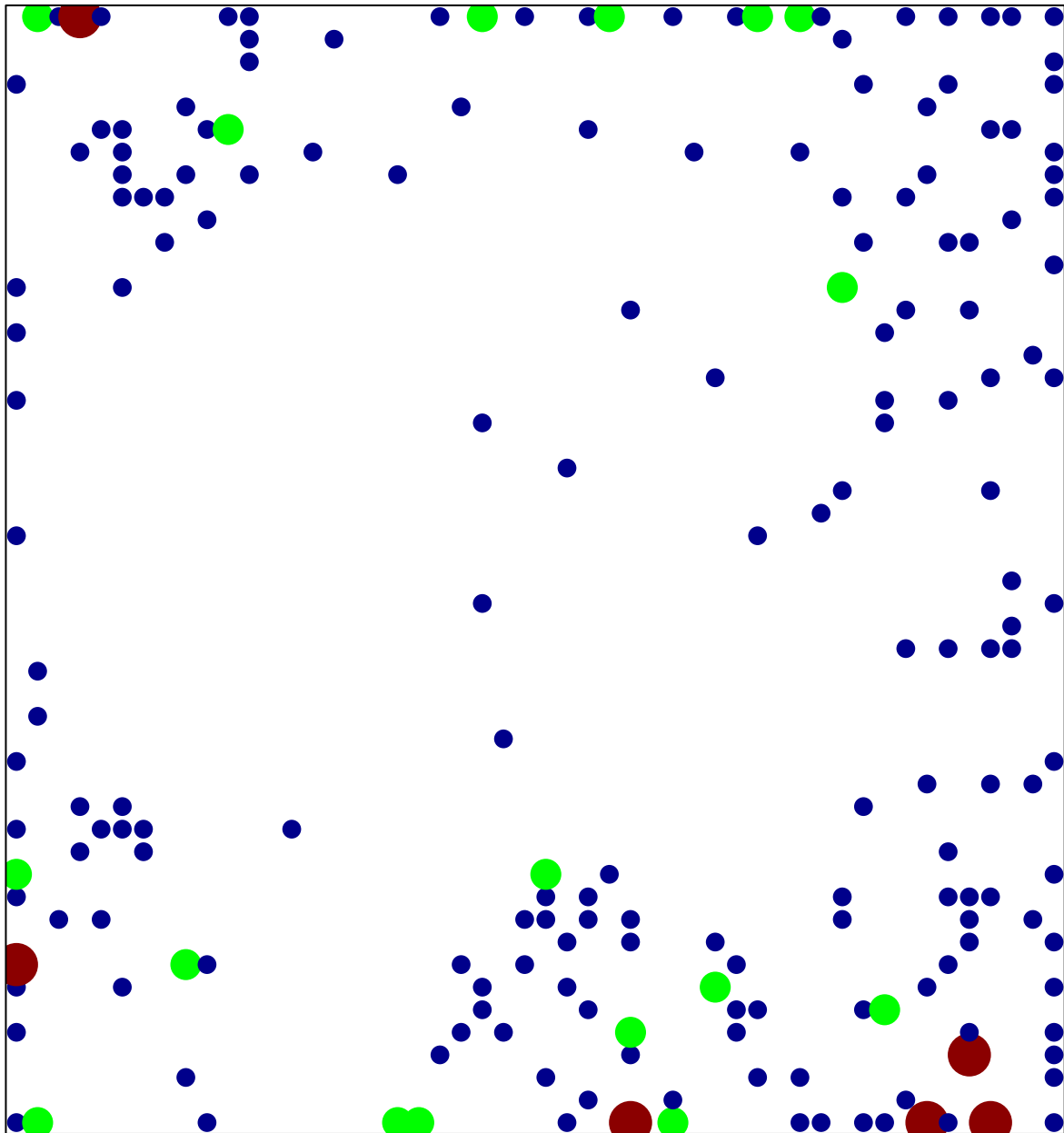
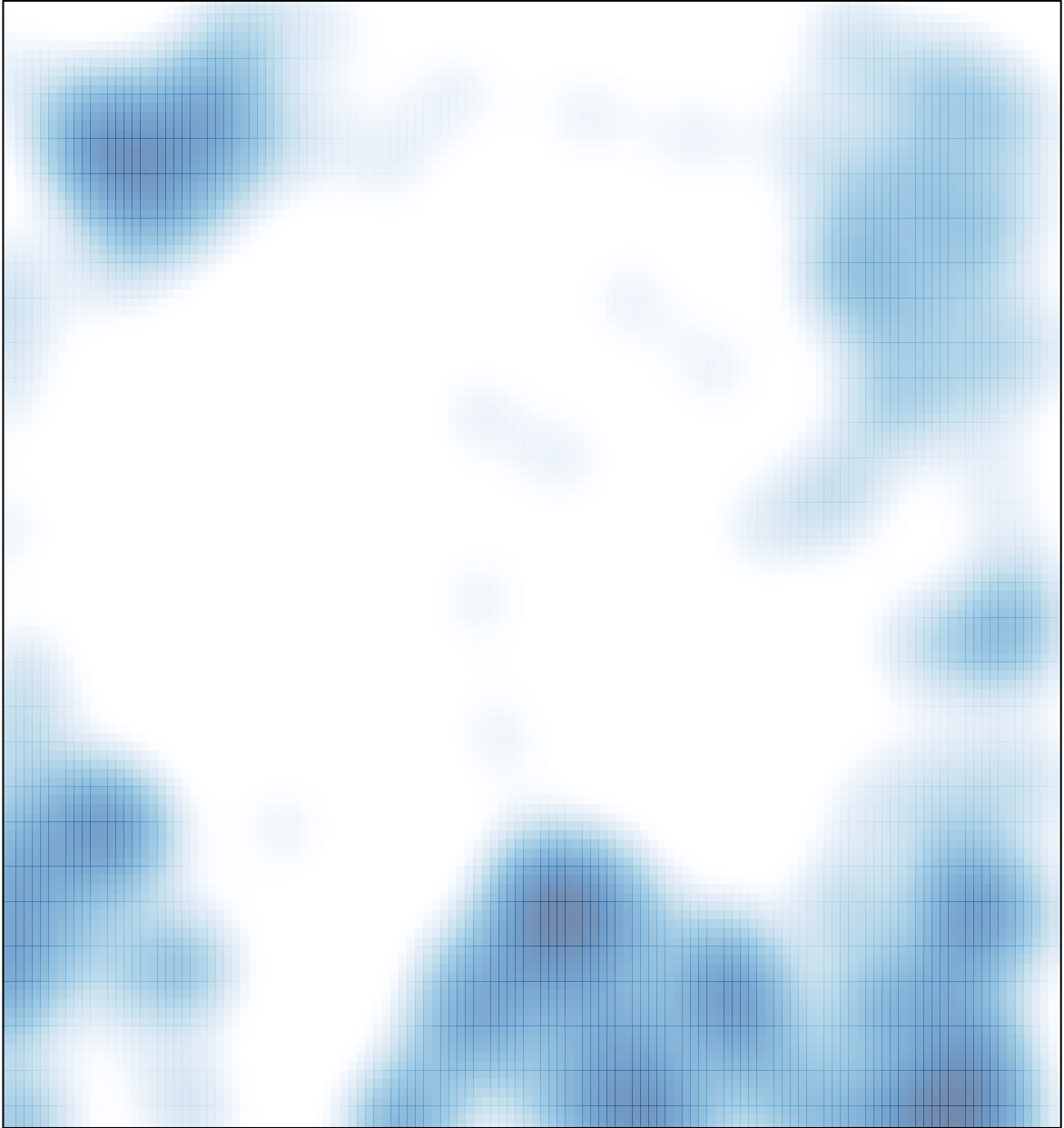


# hsa-miR-181b



# features = 224  
chi-square p = 0.67

# hsa-miR-181b



# features = 224 , max = 3