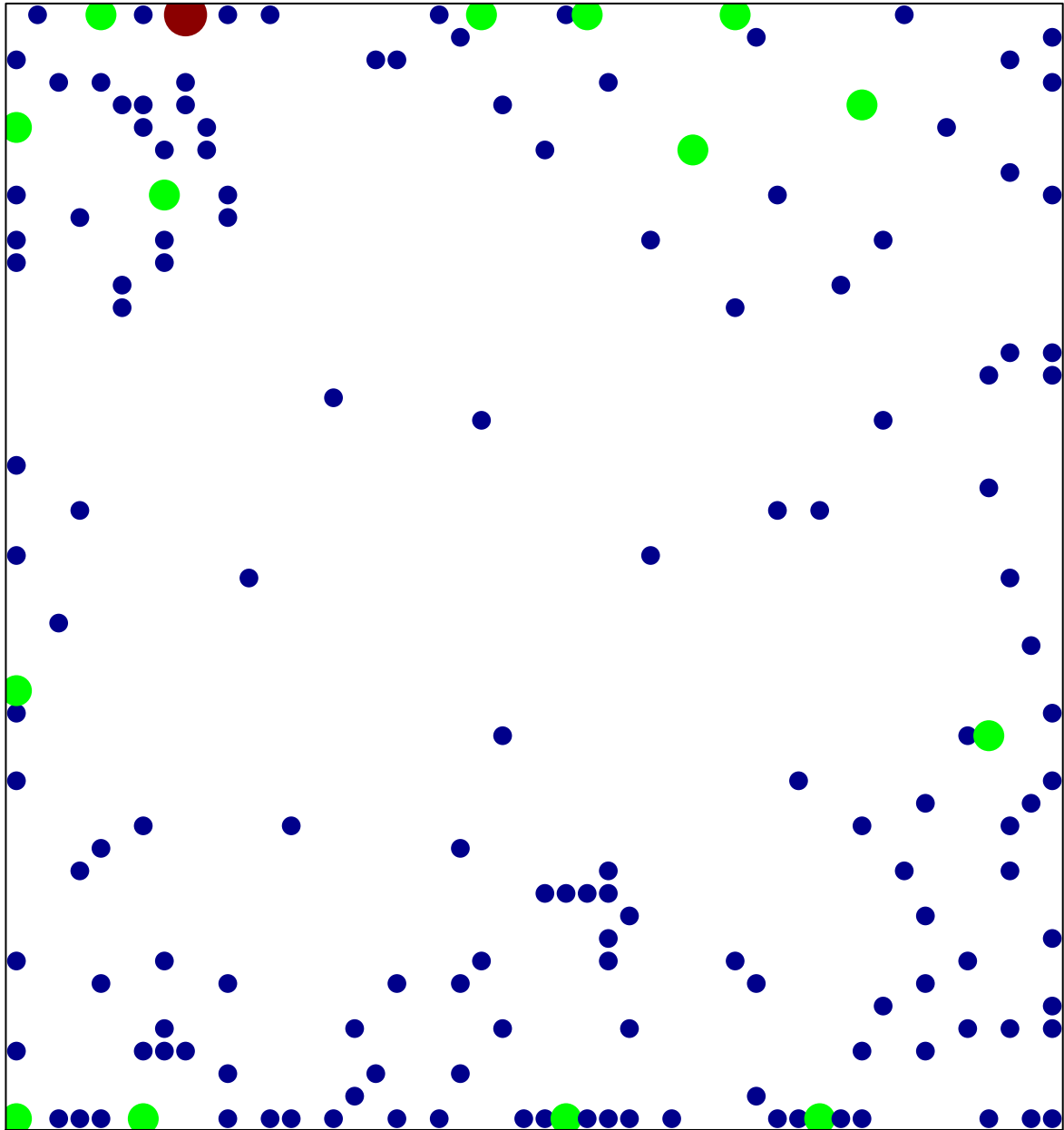
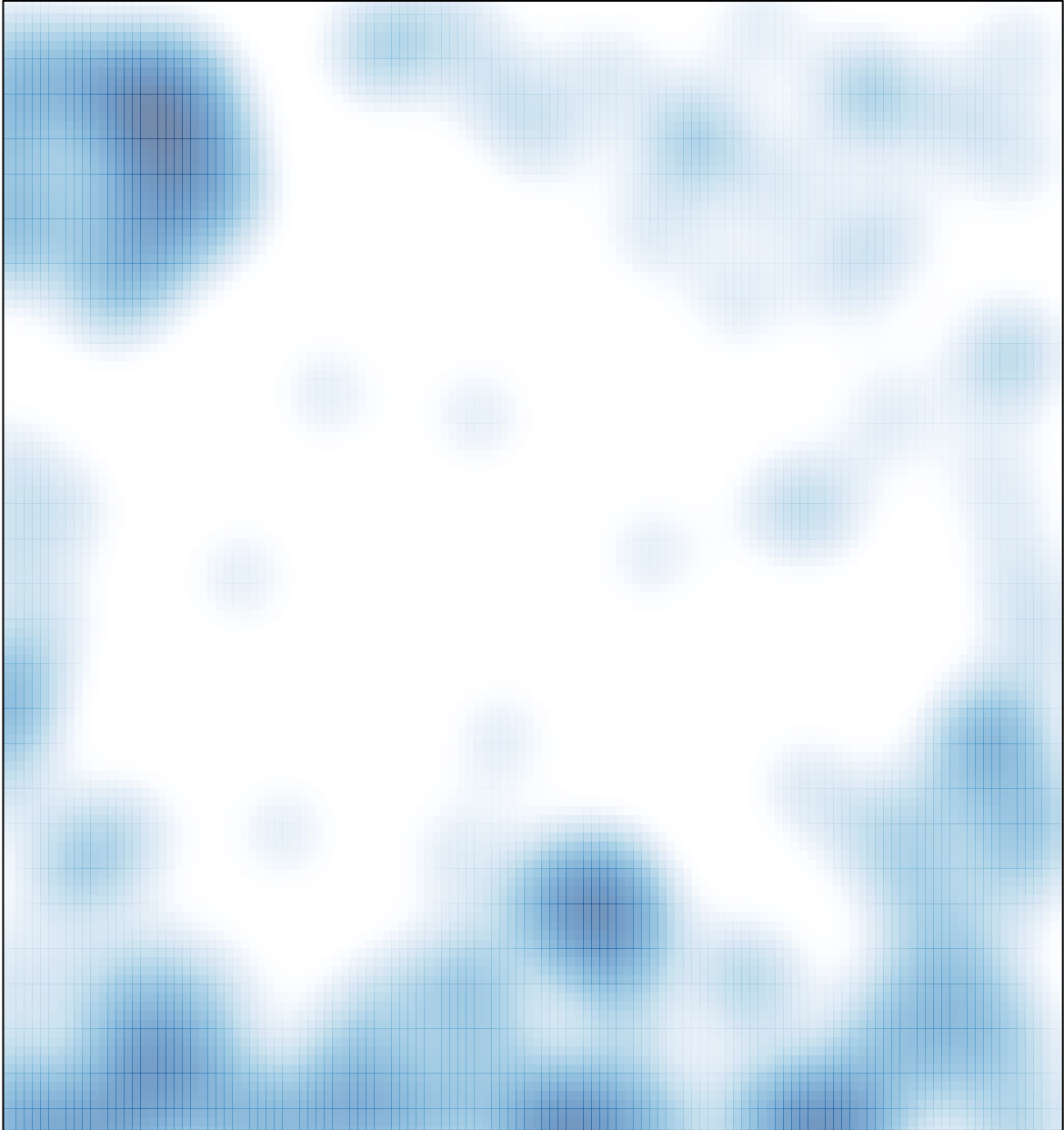


# hsa-miR-23b



# features = 176  
chi-square p = 0.68

# hsa-miR-23b



# features = 176 , max = 3