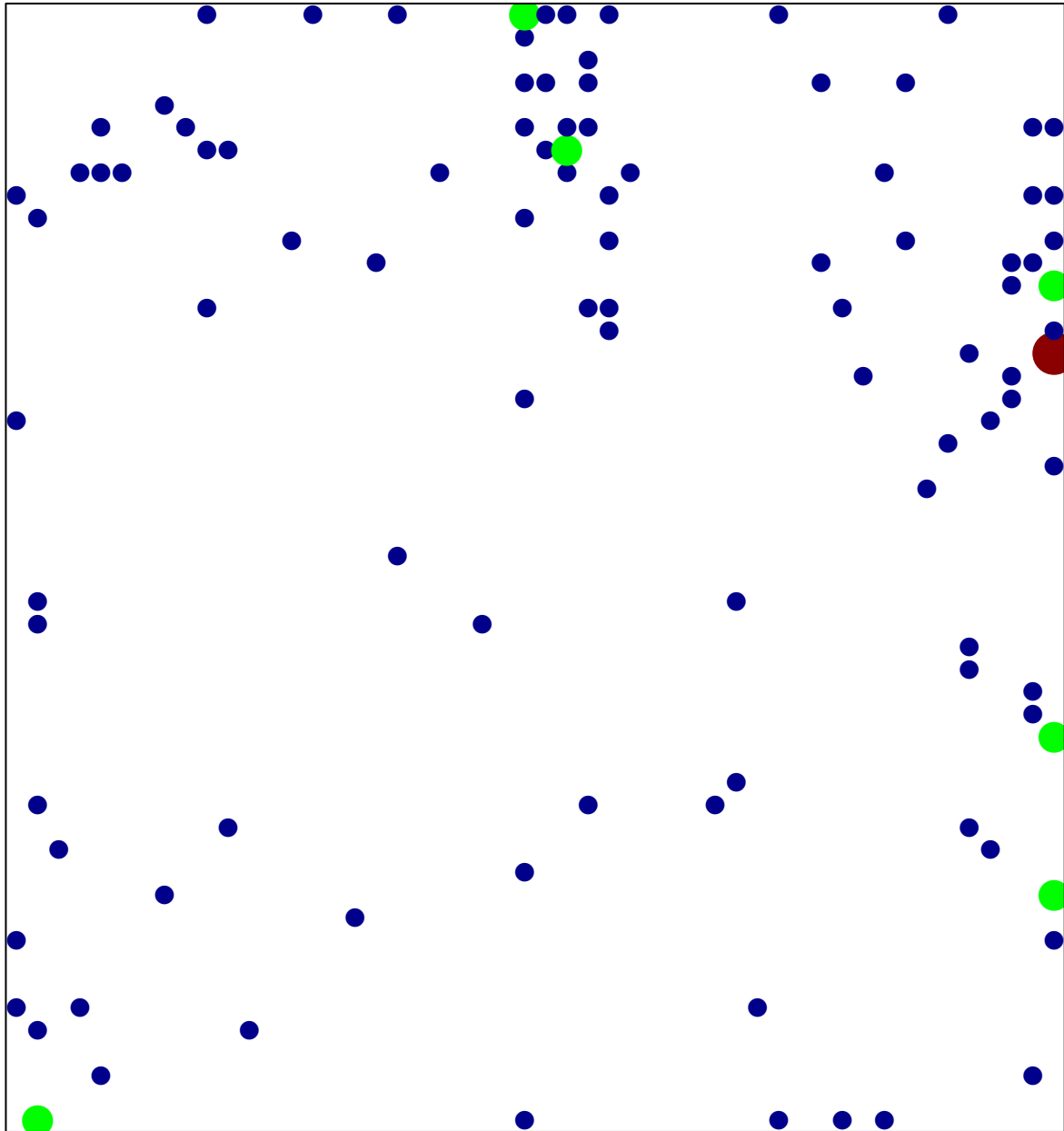
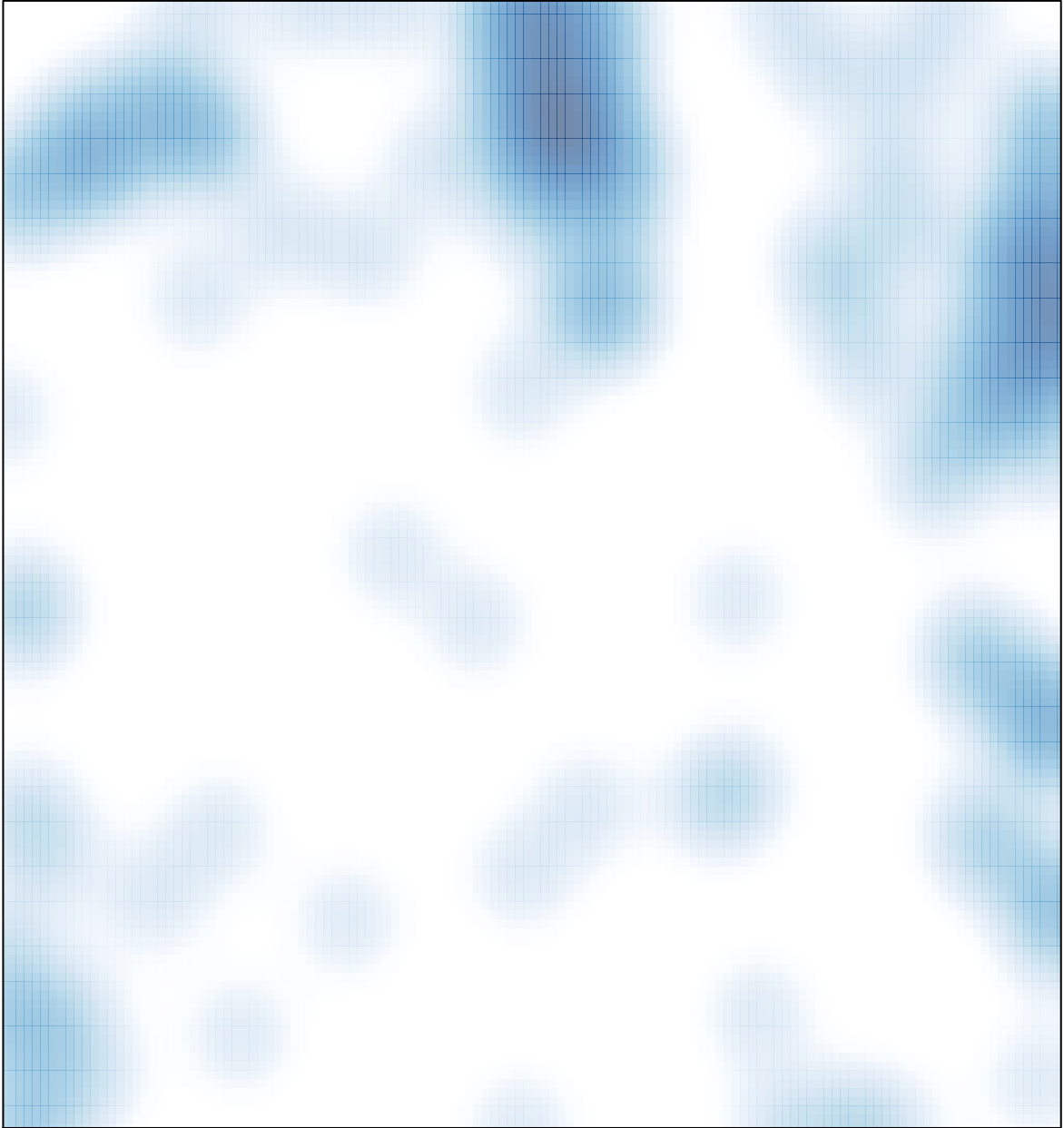


# hsa-miR-491-3p



# features = 112  
chi-square p = 0.75

# hsa-miR-491-3p



# features = 112 , max = 3