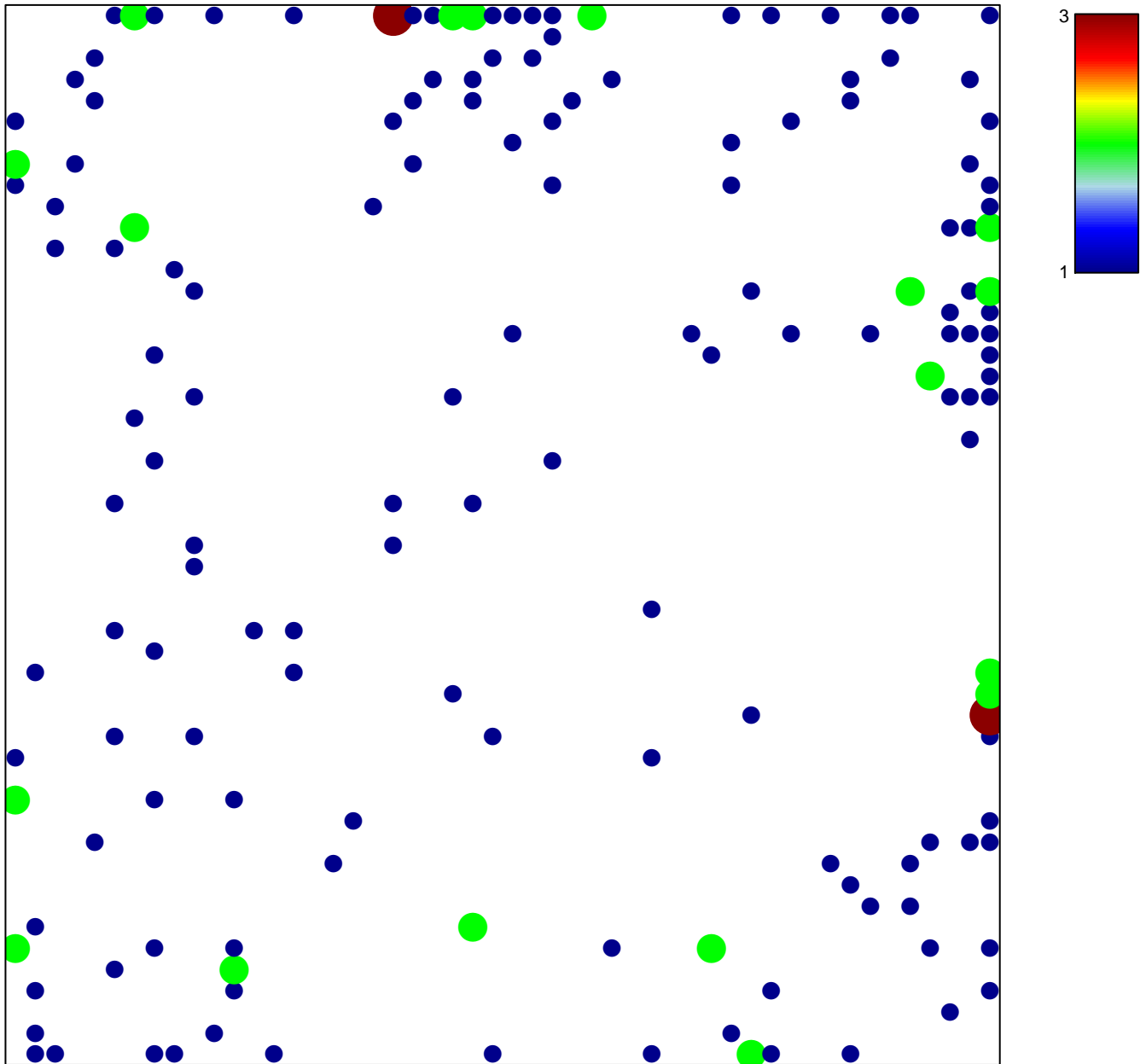
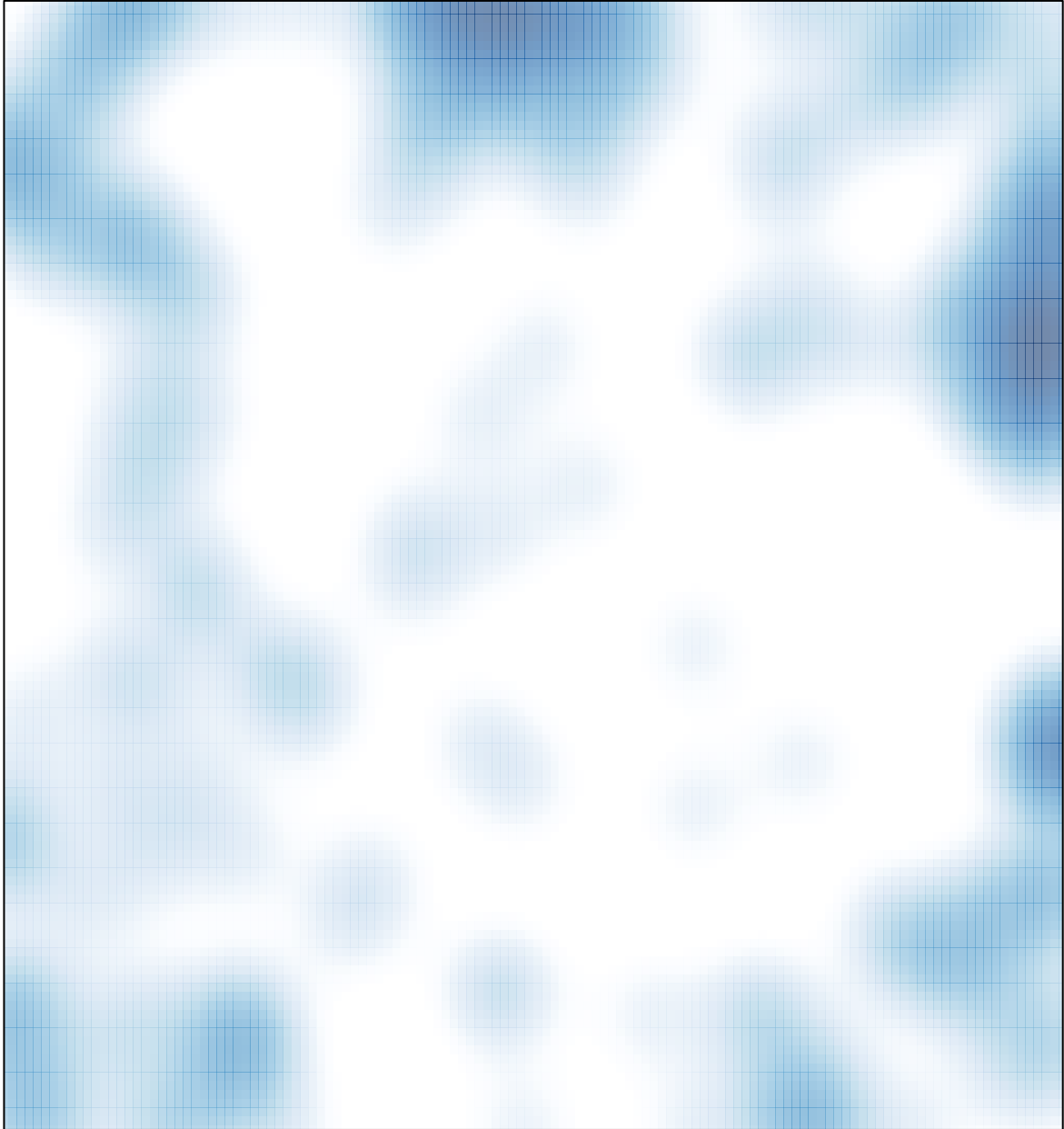


# hsa-miR-1



# features = 180  
chi-square p = 0.68

# hsa-miR-1



# features = 180 , max = 3