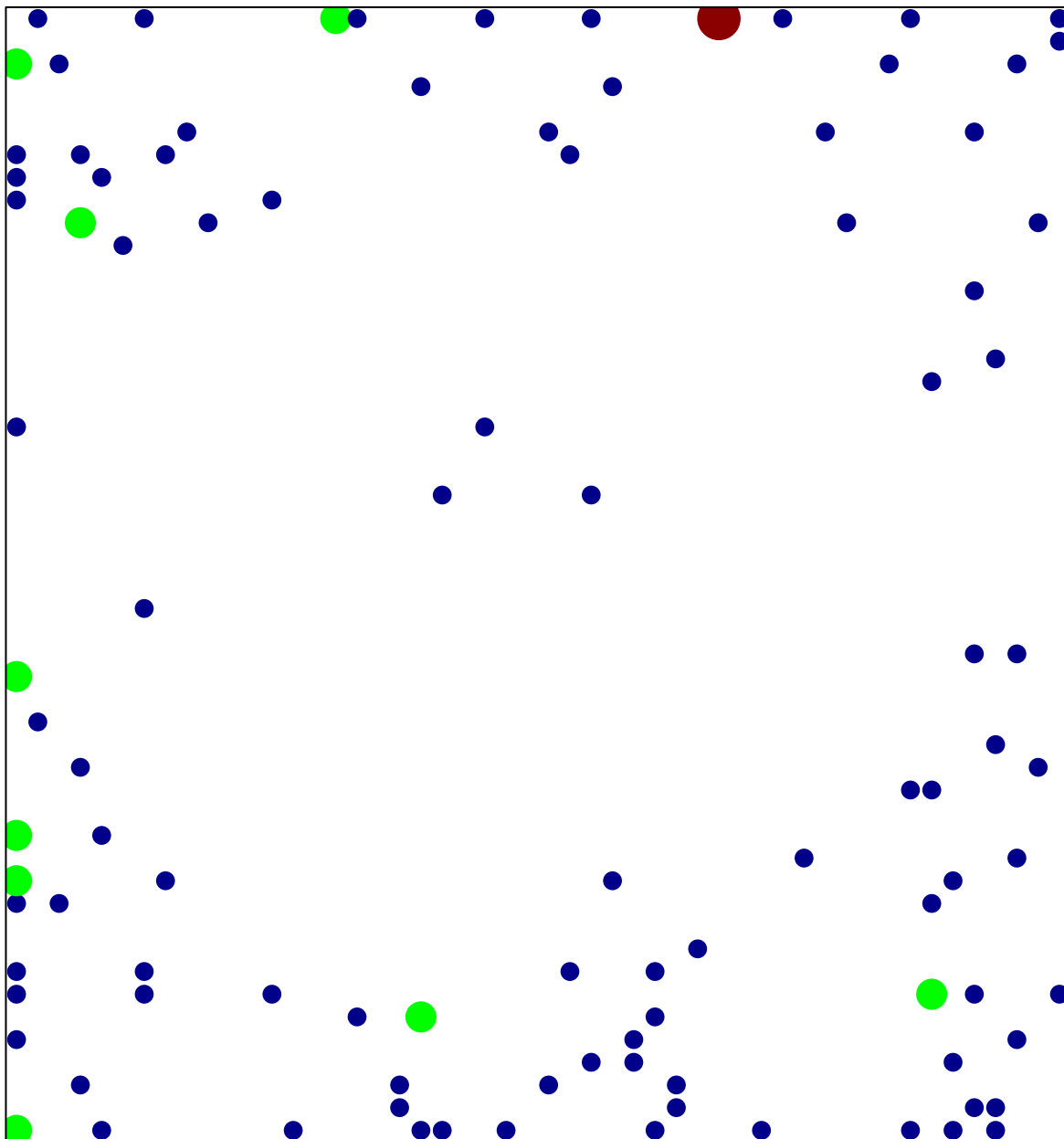
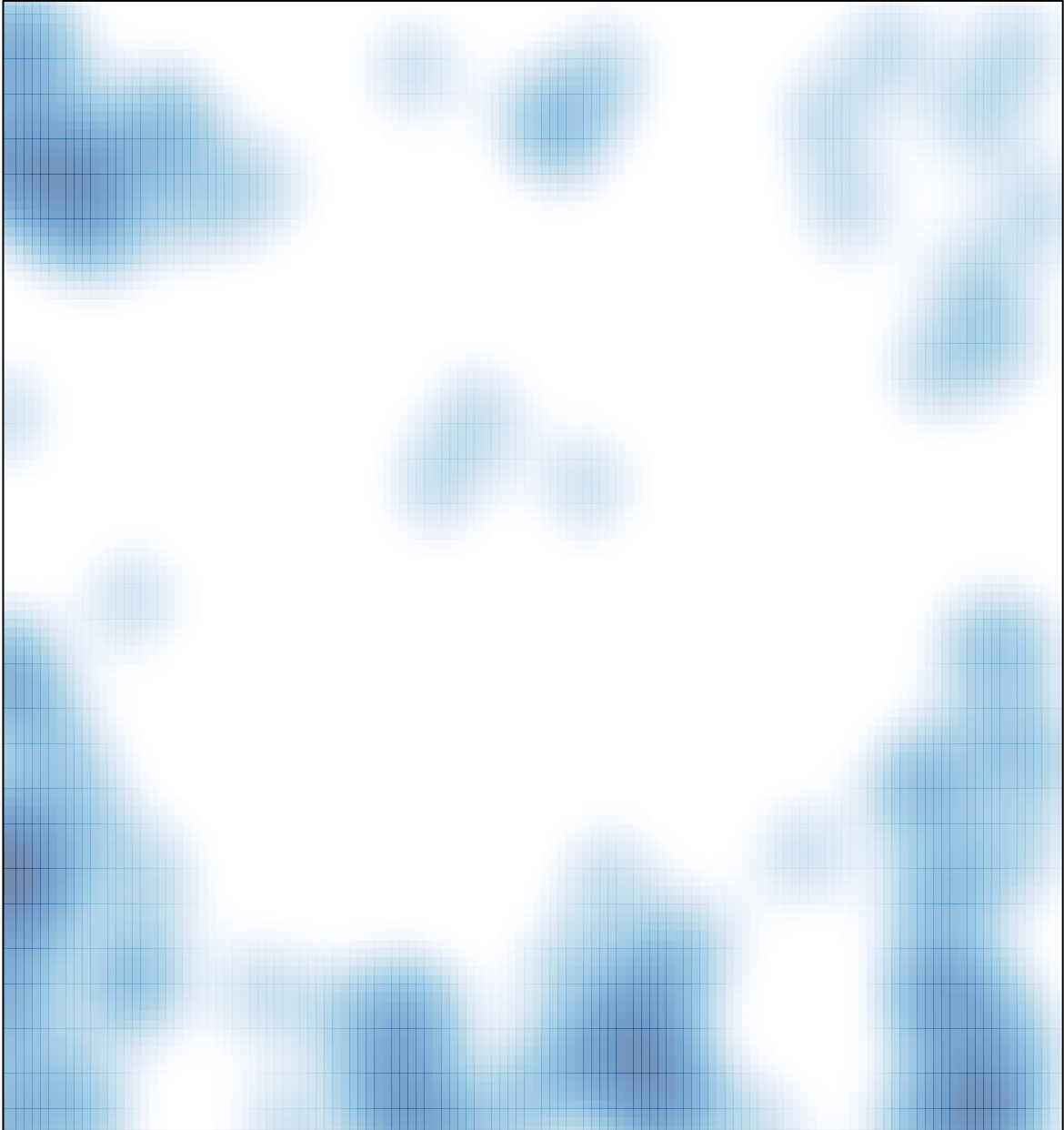


# hsa-miR-297



# features = 112  
chi-square p = 0.72

# hsa-miR-297



# features = 112 , max = 3