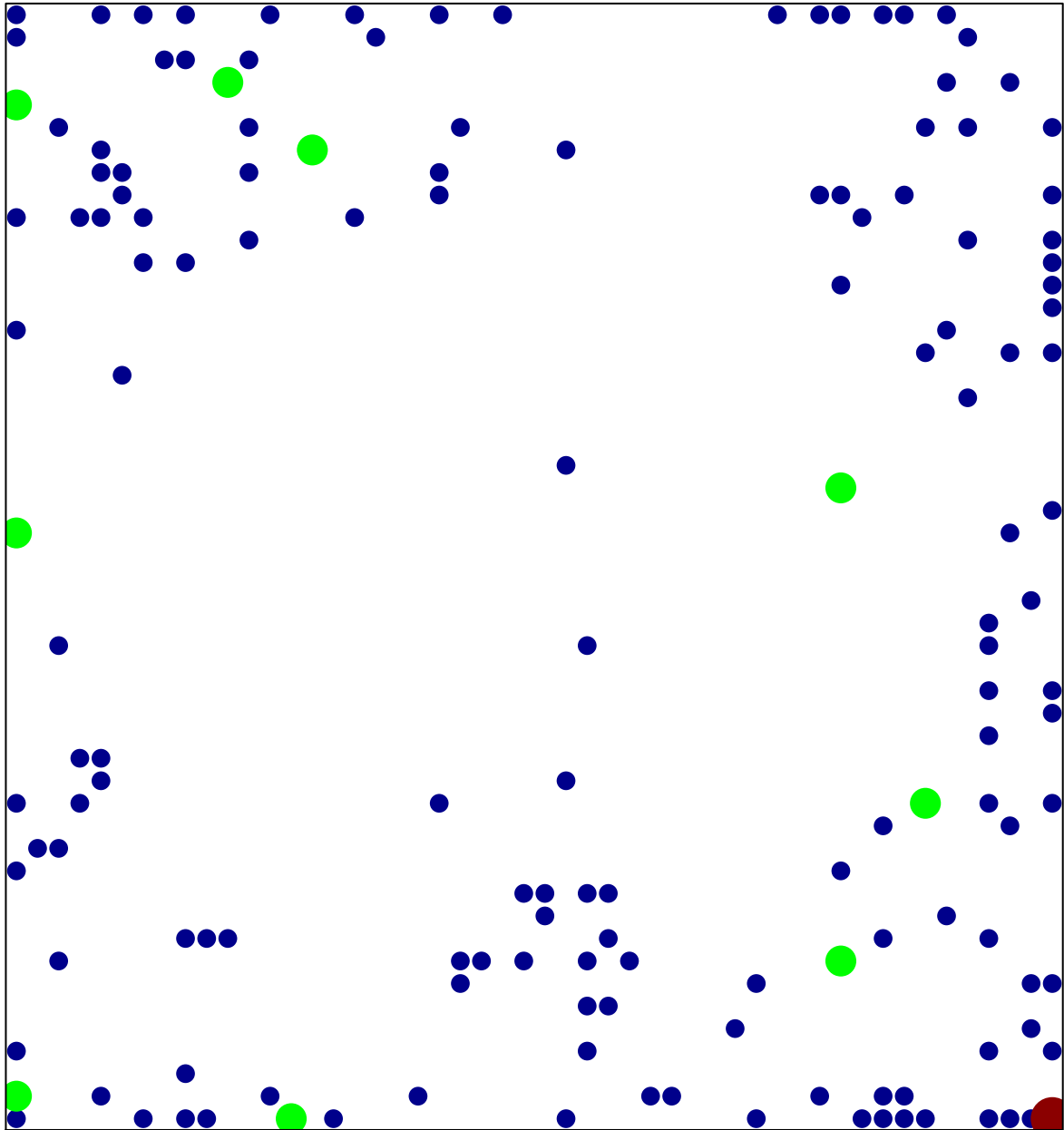
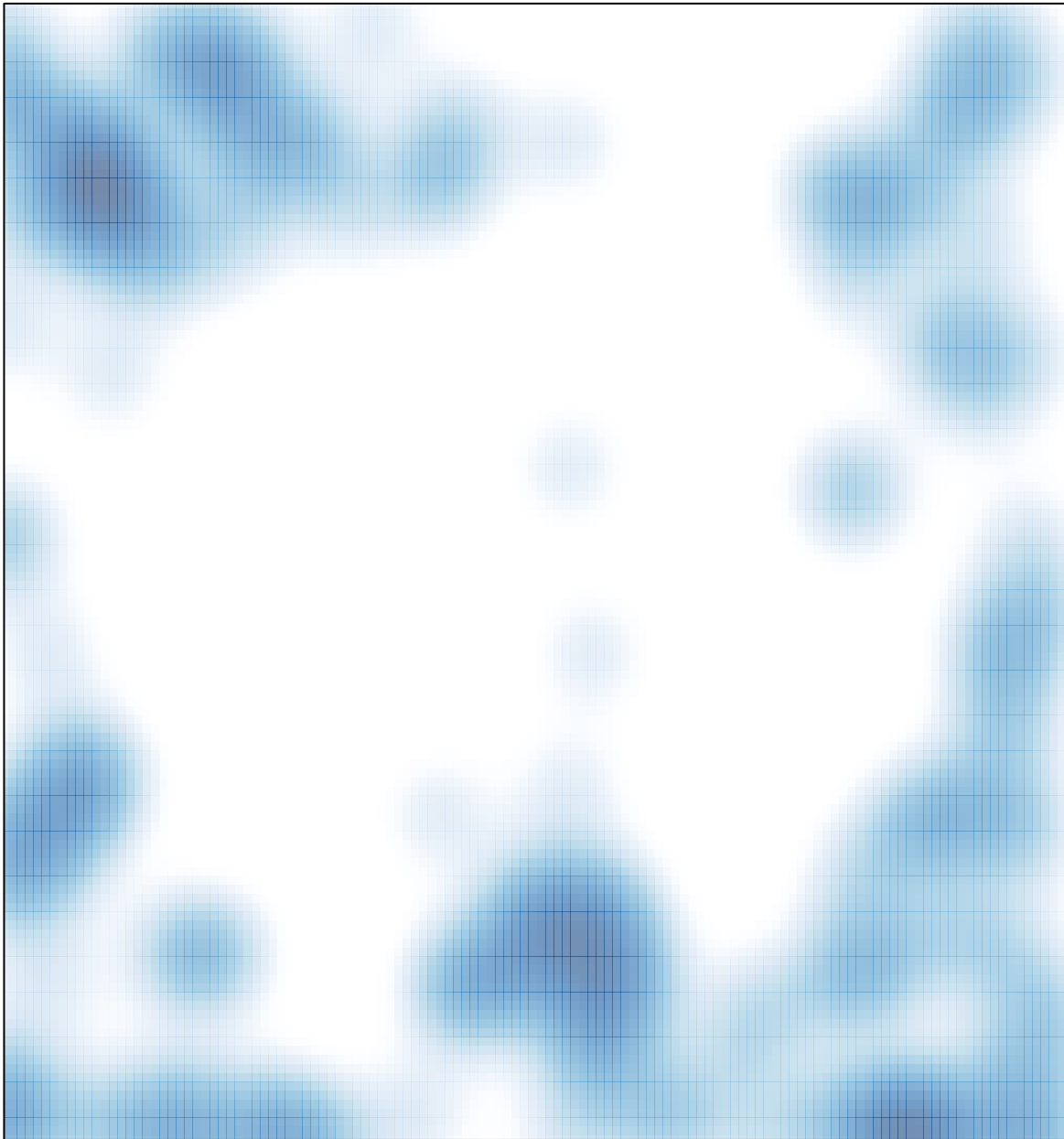


# hsa-miR-495



# features = 163  
chi-square p = 0.72

# hsa-miR-495



# features = 163 , max = 3