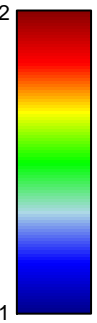
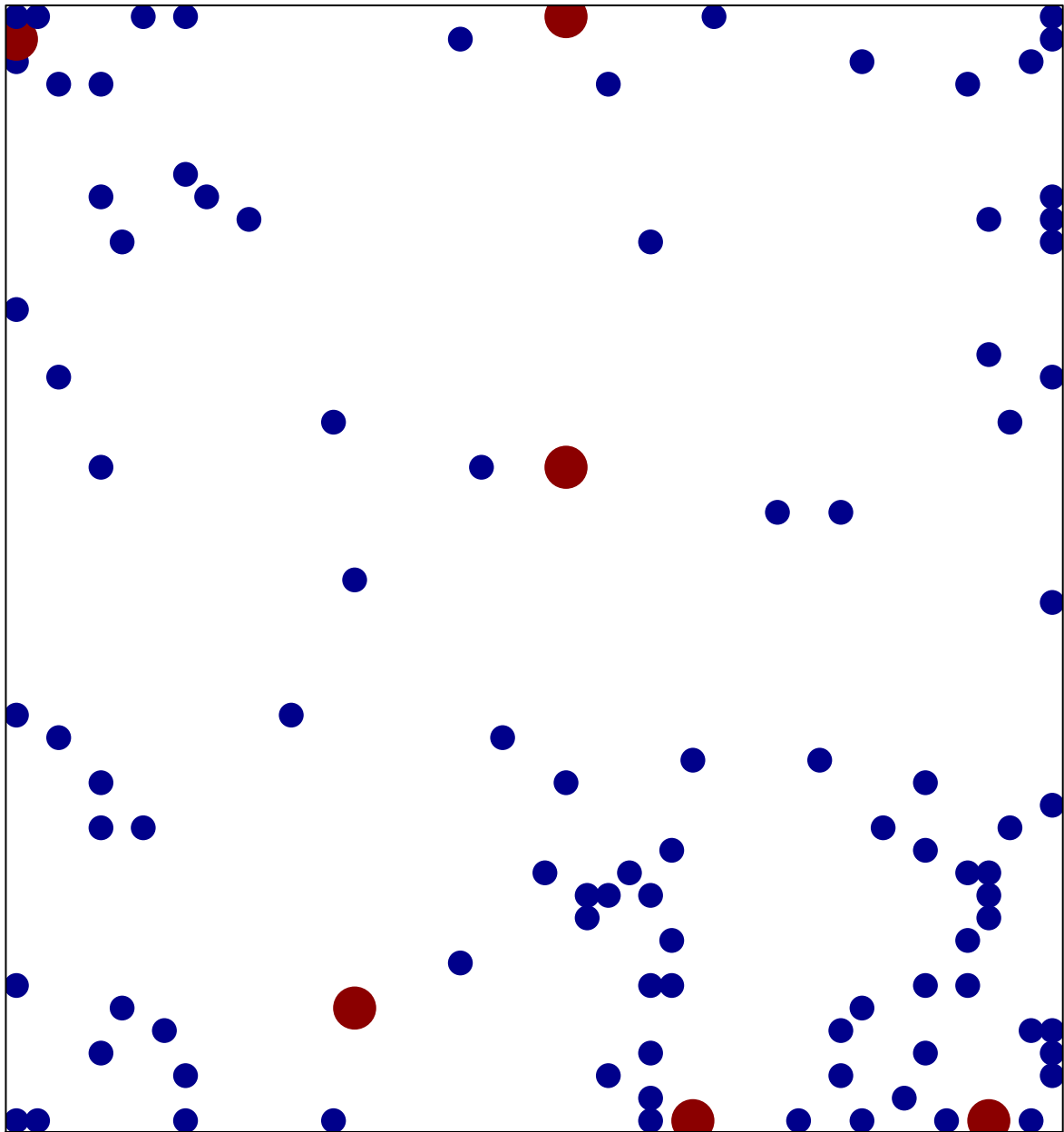
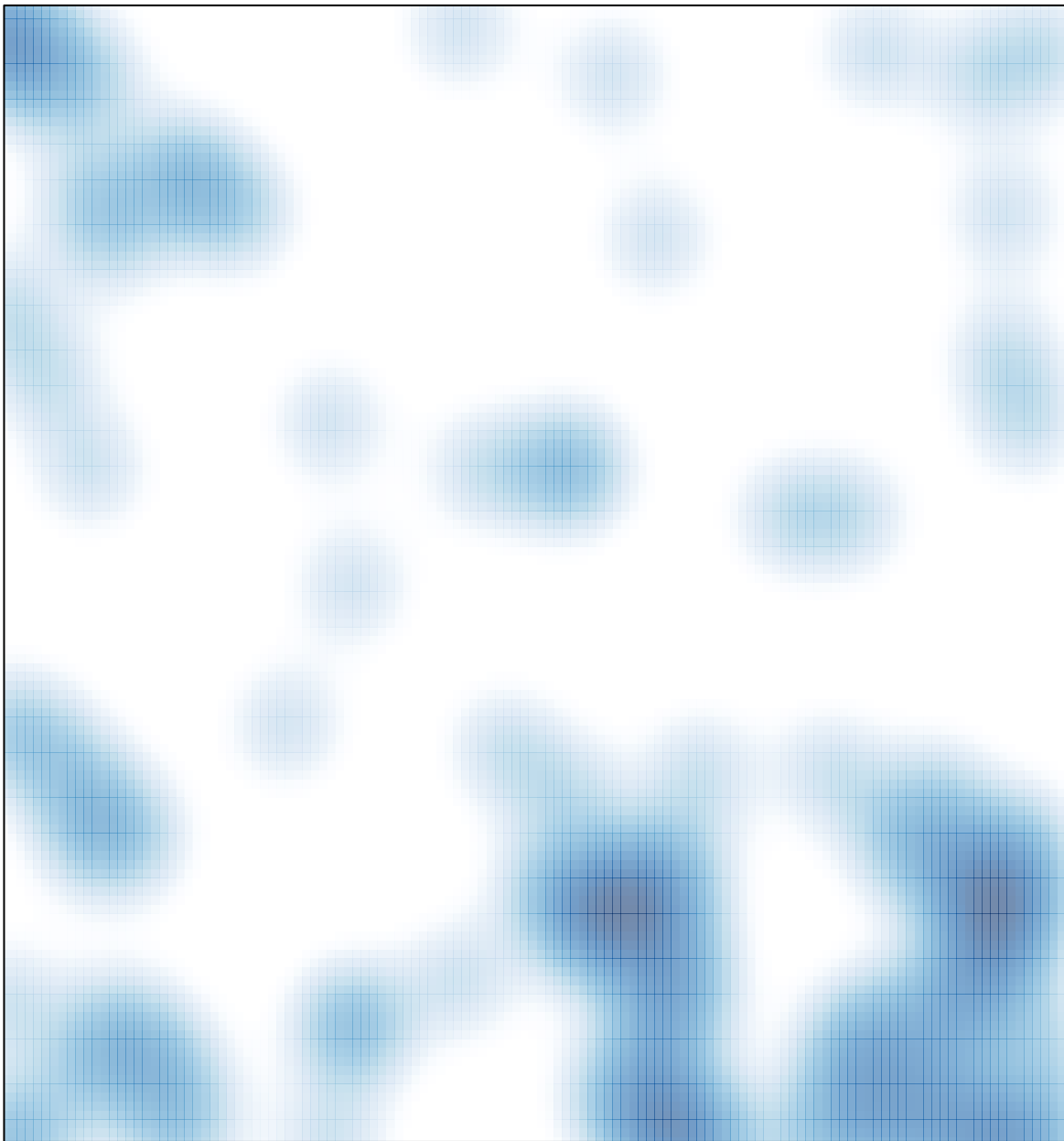


# hsa-miR-211



# features = 108  
chi-square p = 0.75

# hsa-miR-211



# features = 108 , max = 2