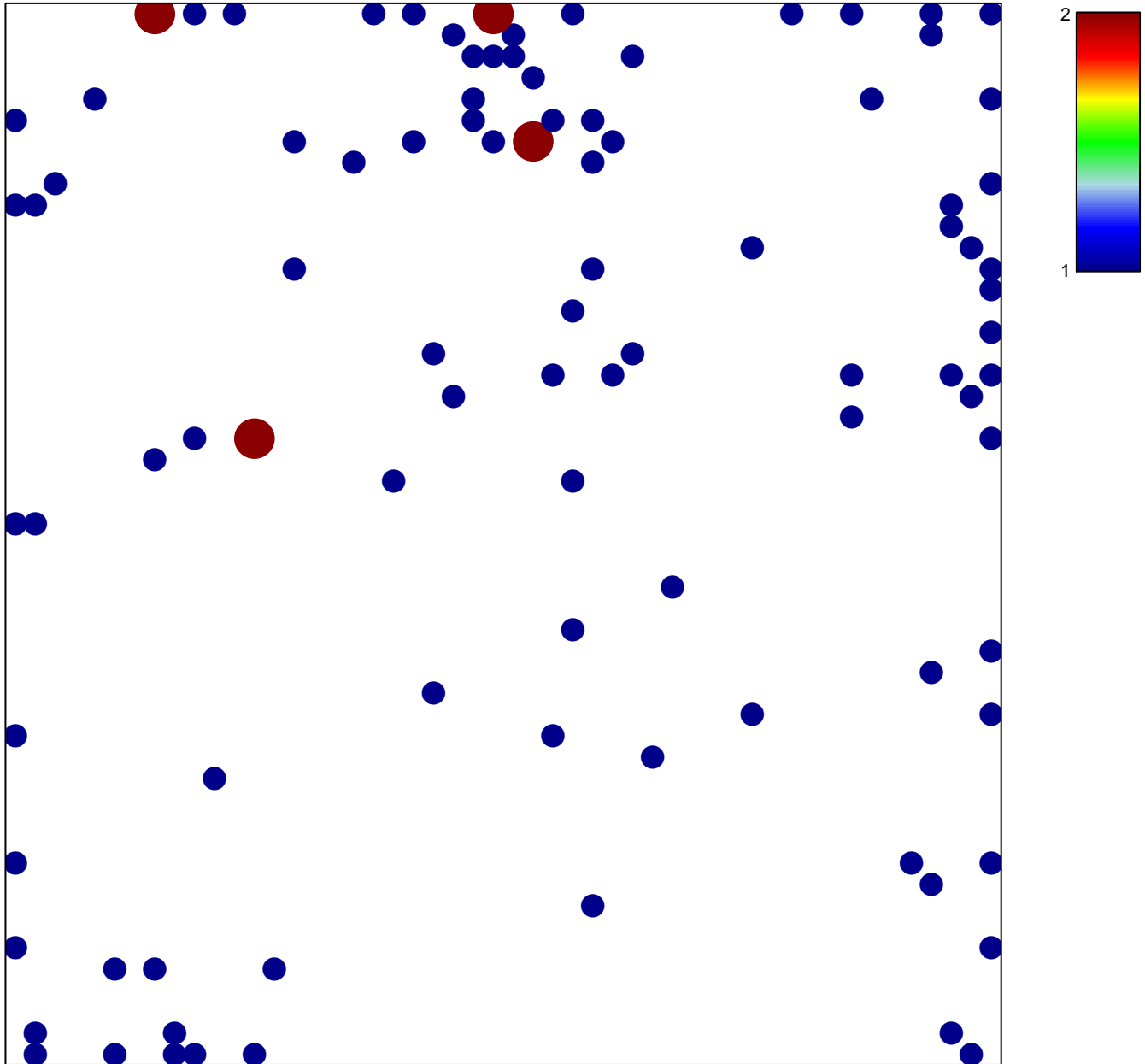
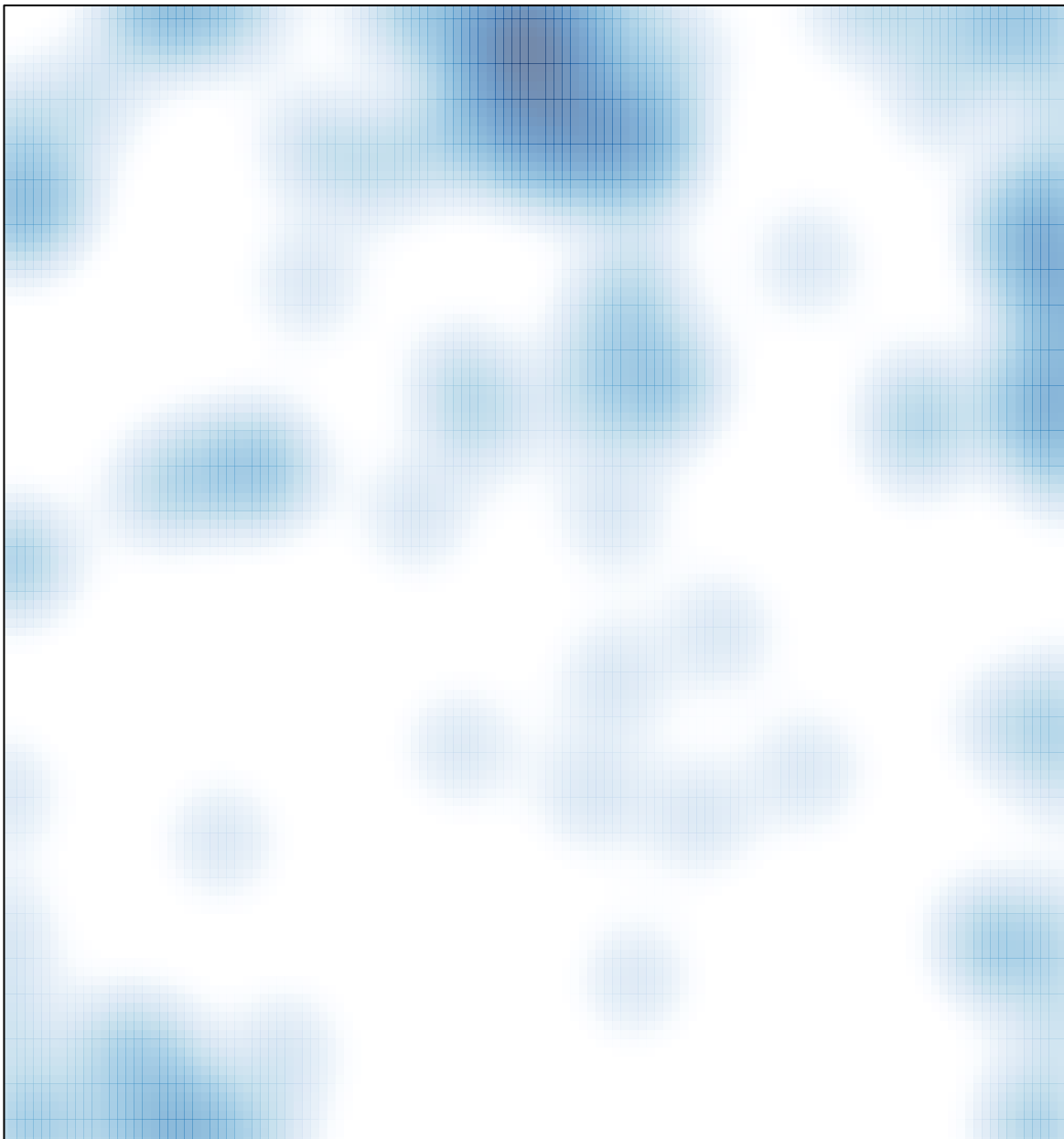


# hsa-miR-33b



# features = 100  
chi-square p = 0.76

# hsa-miR-33b



# features = 100 , max = 2