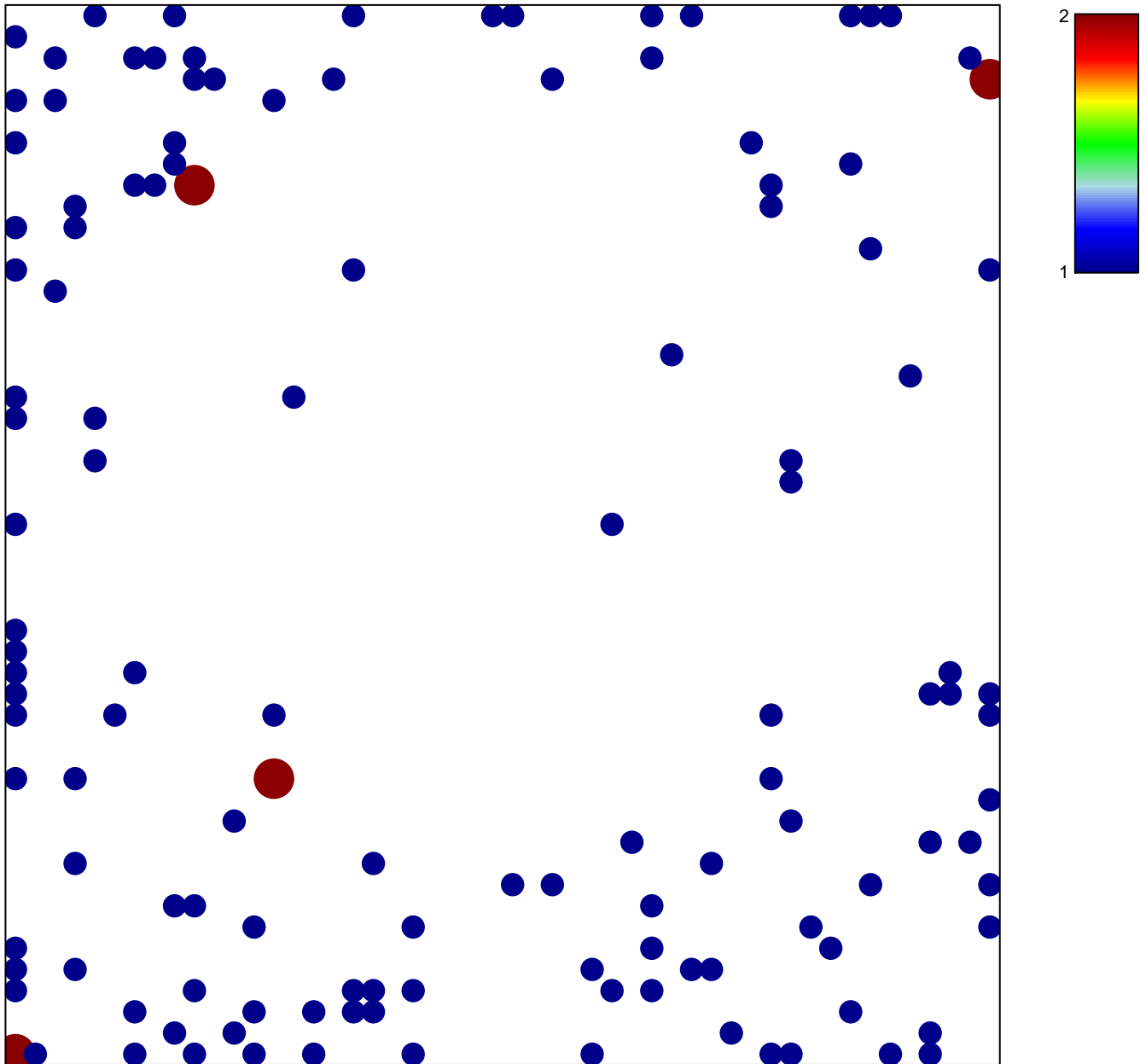
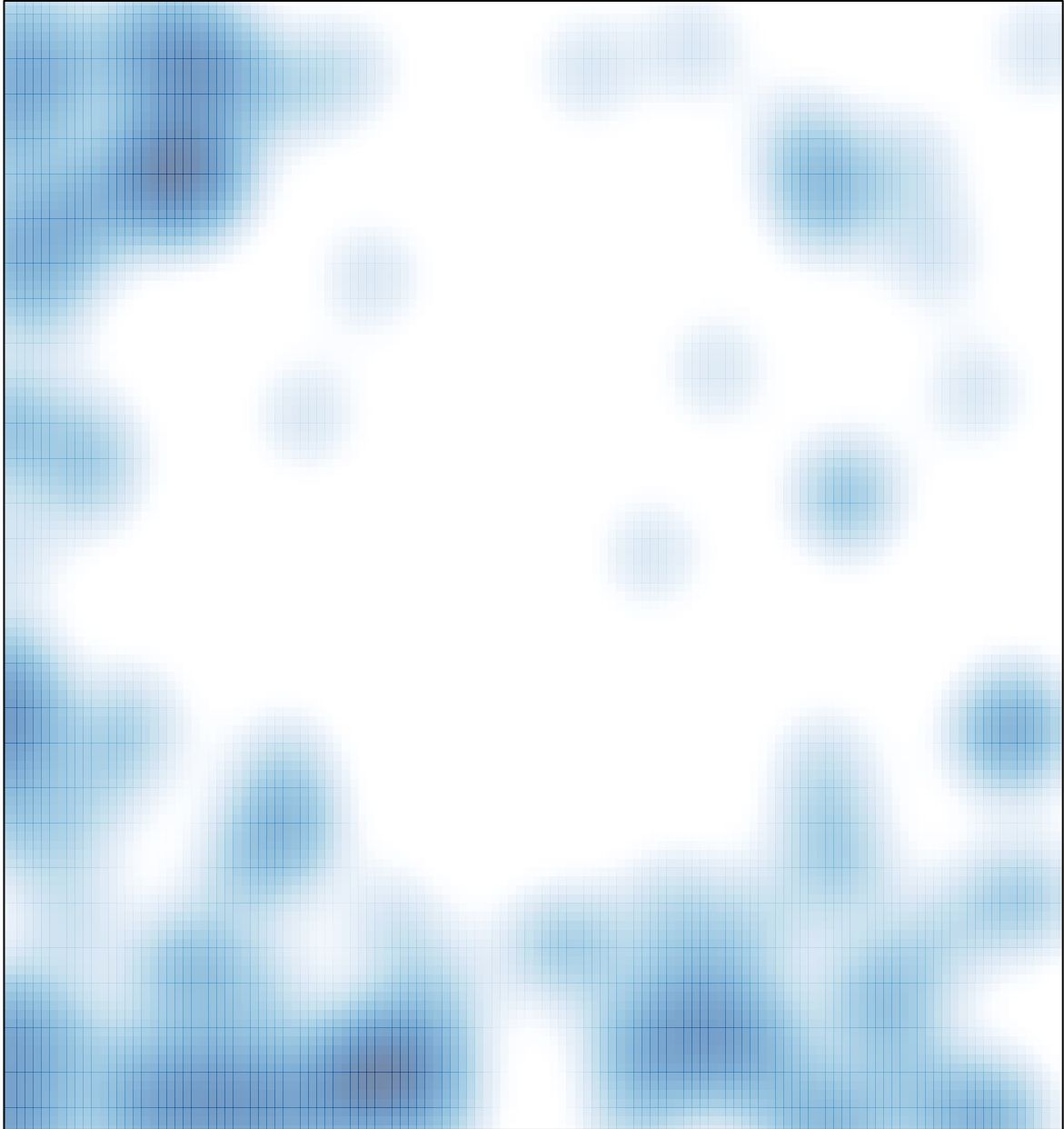


# hsa-miR-198



# features = 133  
chi-square p = 0.73

# hsa-miR-198



# features = 133 , max = 2