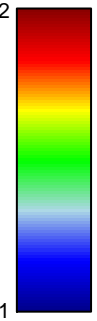
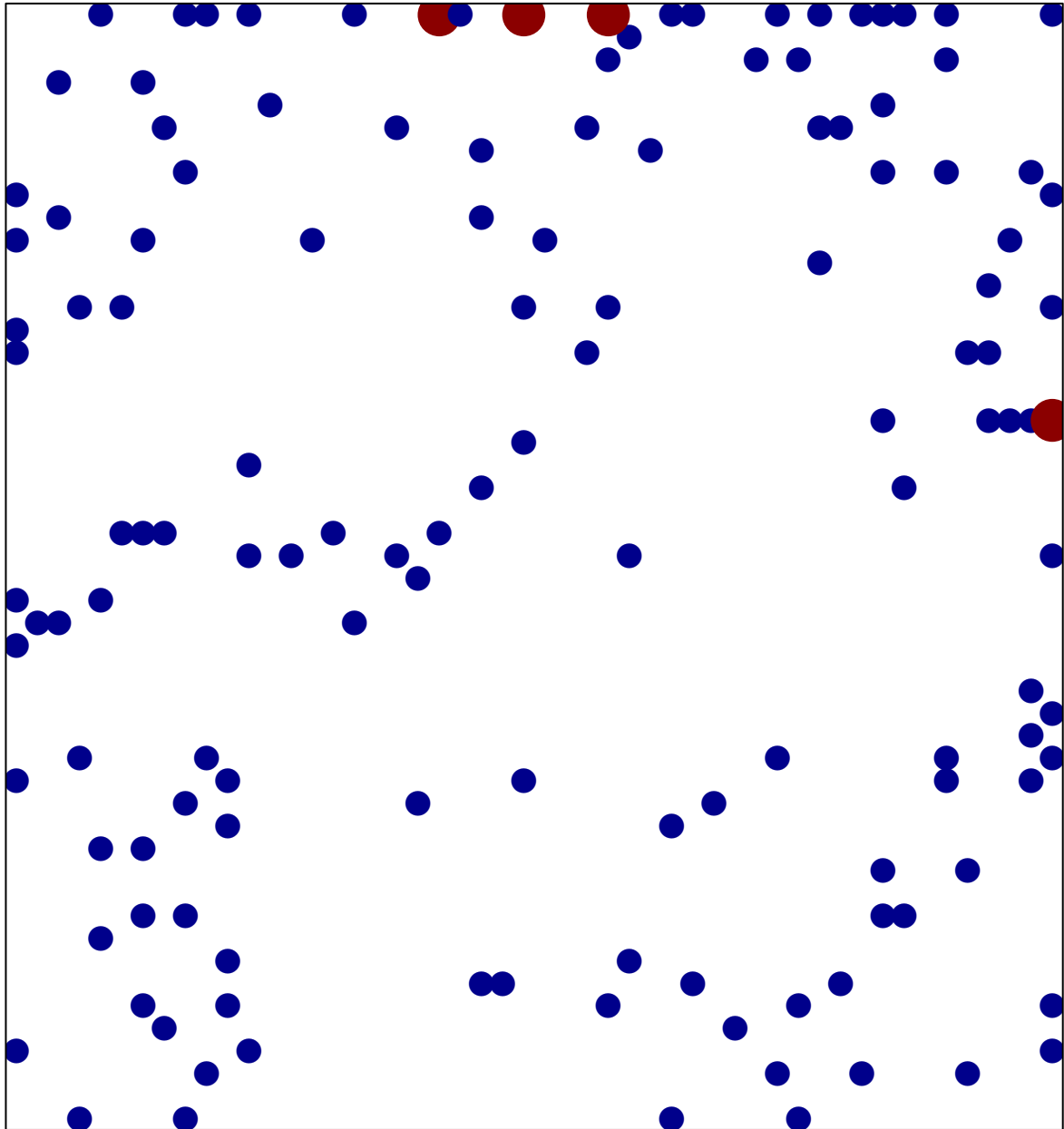
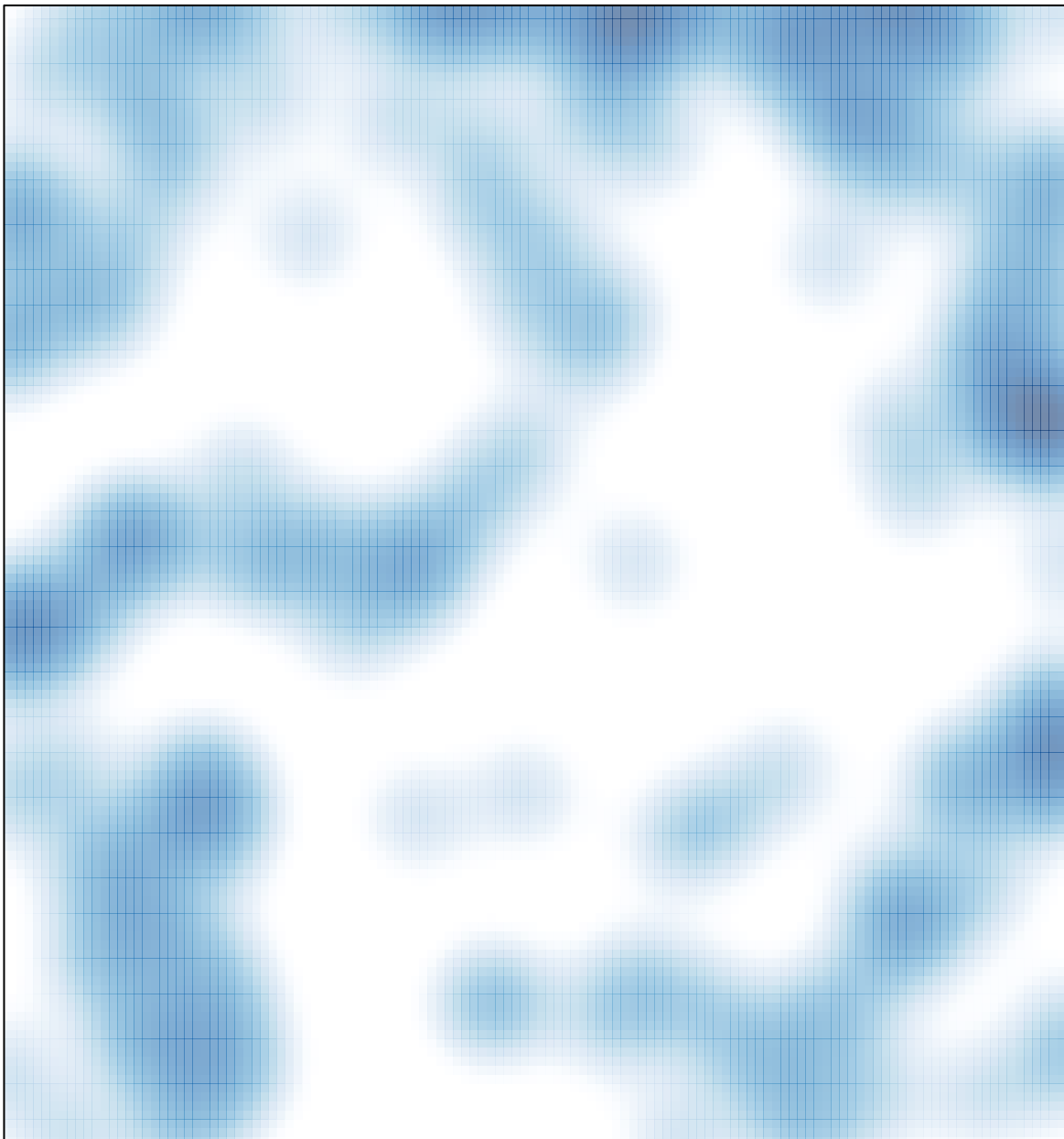


# hsa-miR-198



# features = 140  
chi-square p = 0.75

# hsa-miR-198



# features = 140 , max = 2