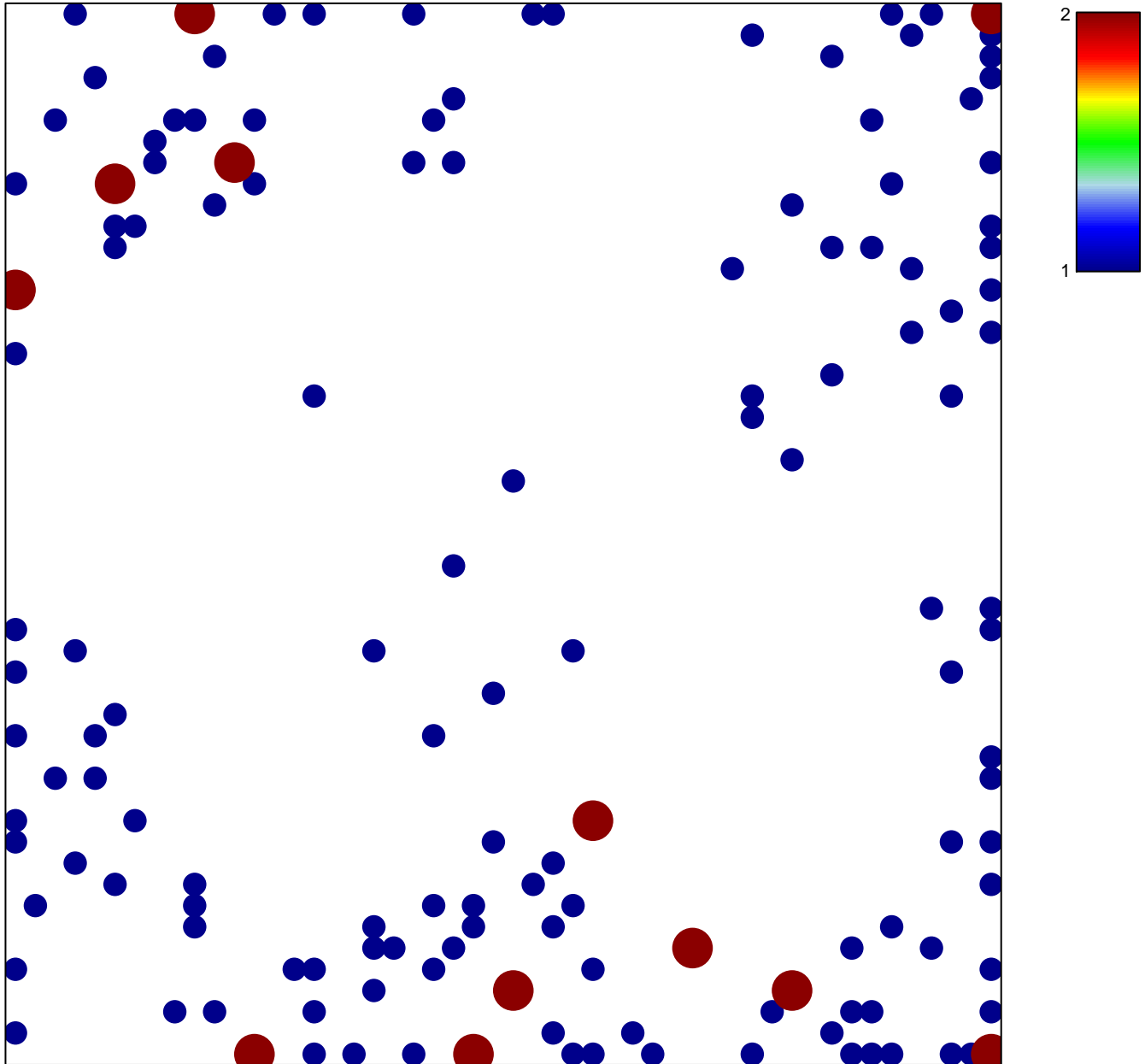
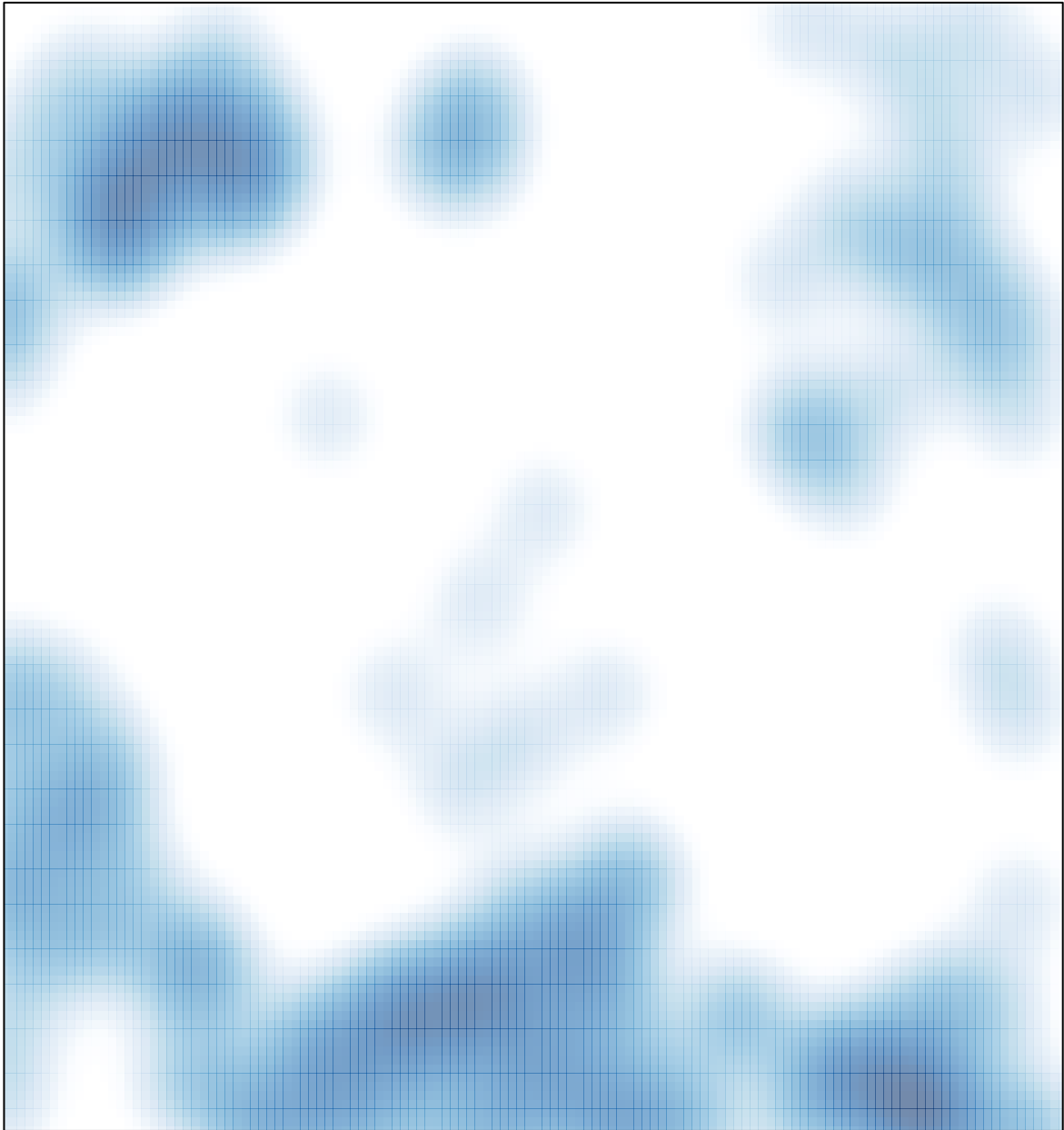


# hsa-miR-31



# features = 155  
chi-square p = 0.72

# hsa-miR-31



# features = 155 , max = 2