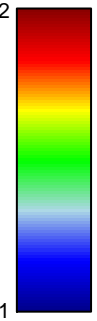
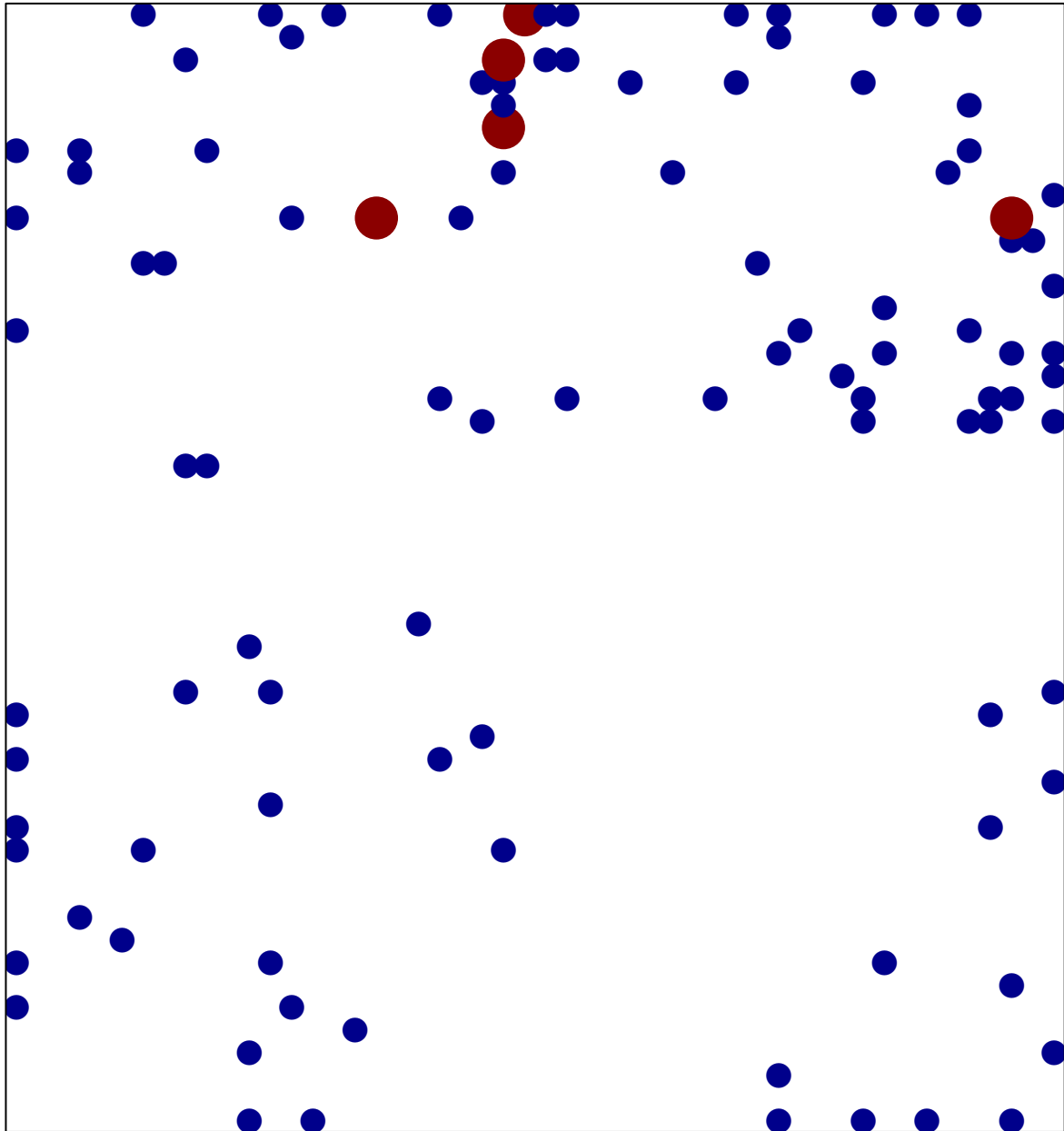
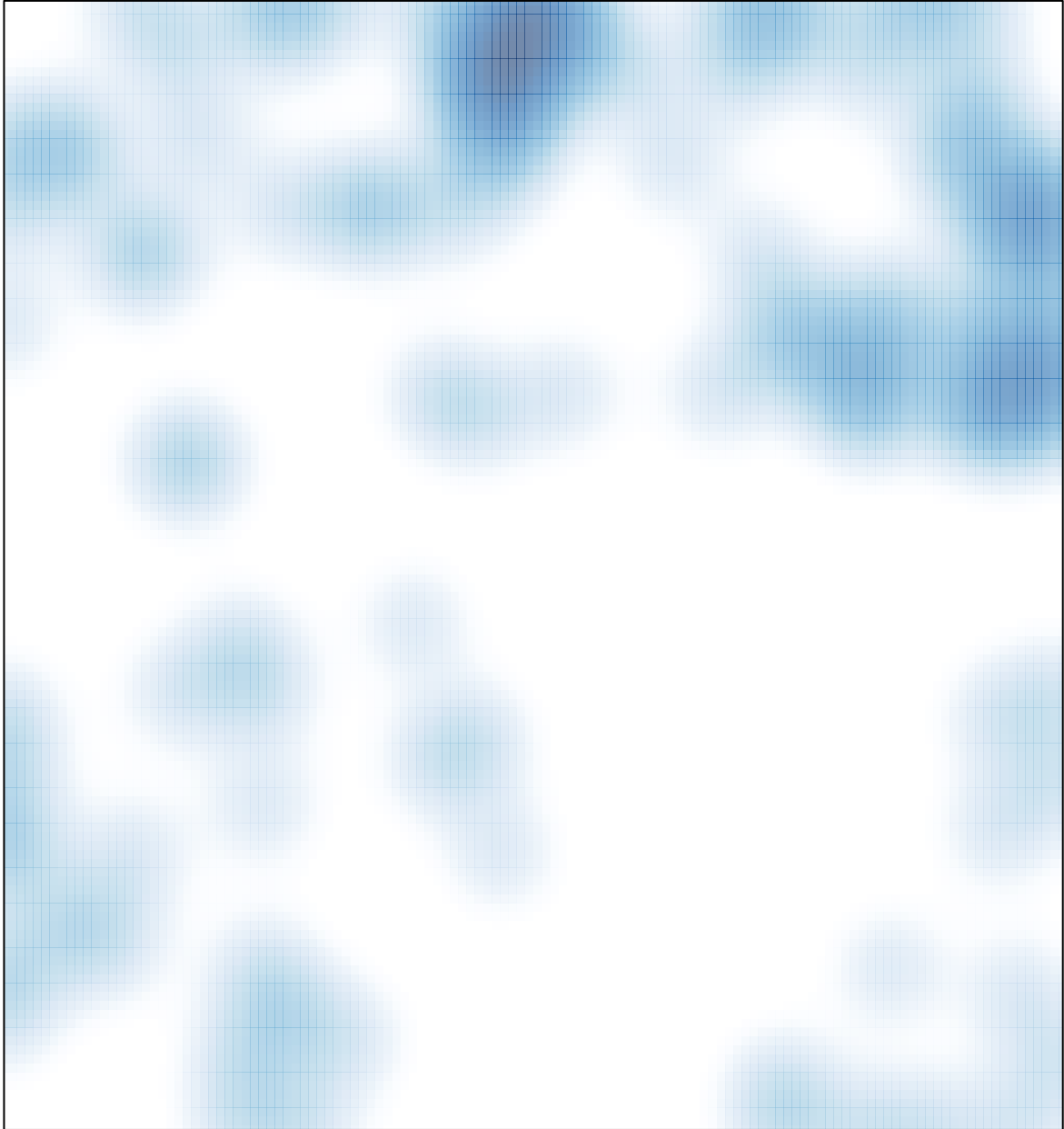


# hsa-miR-146a



# features = 109  
chi-square p = 0.76

# hsa-miR-146a



# features = 109 , max = 2