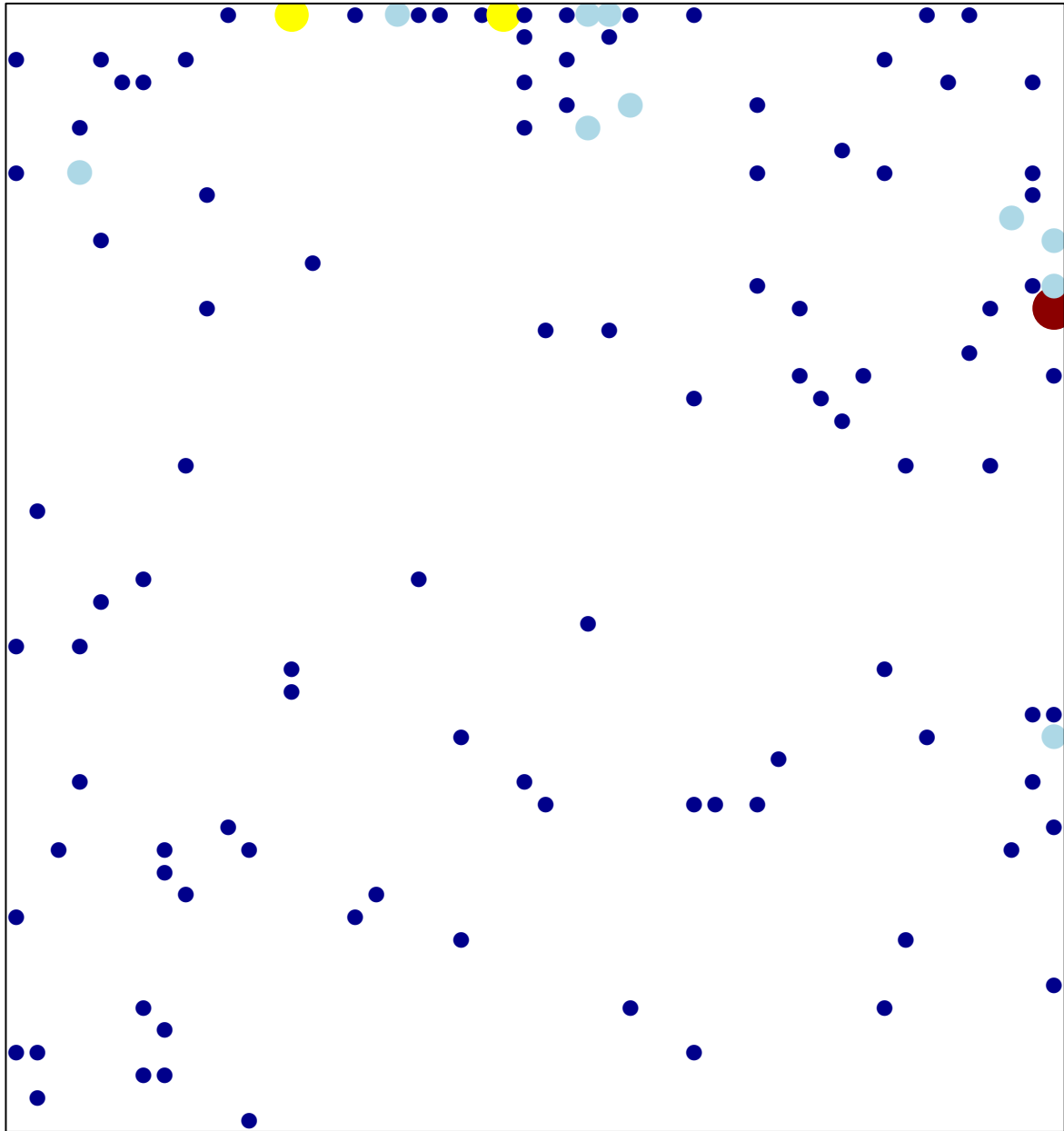
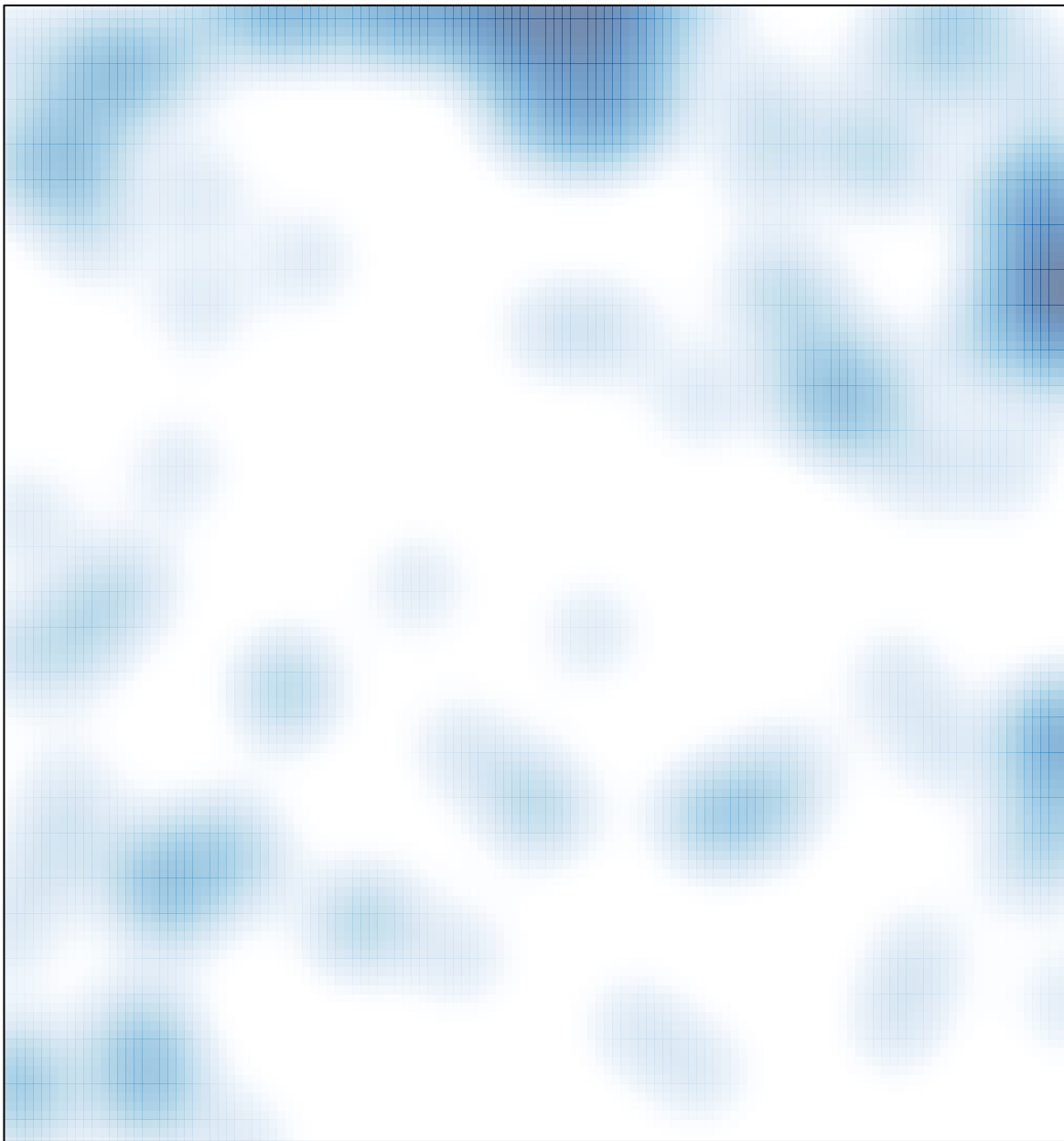


# hsa-miR-194



# features = 130  
chi-square p = 0.73

# hsa-miR-194



# features = 130 , max = 4