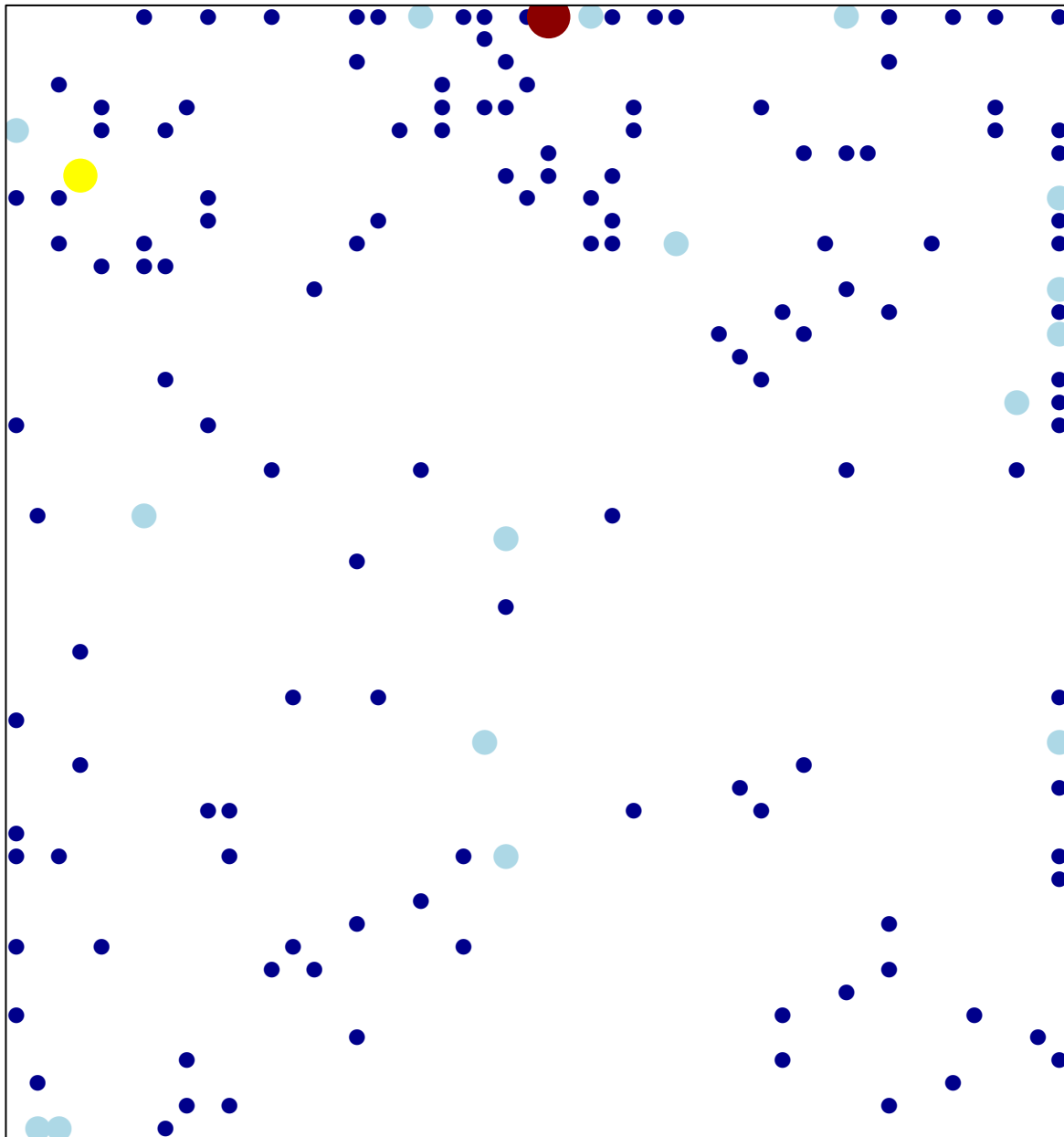
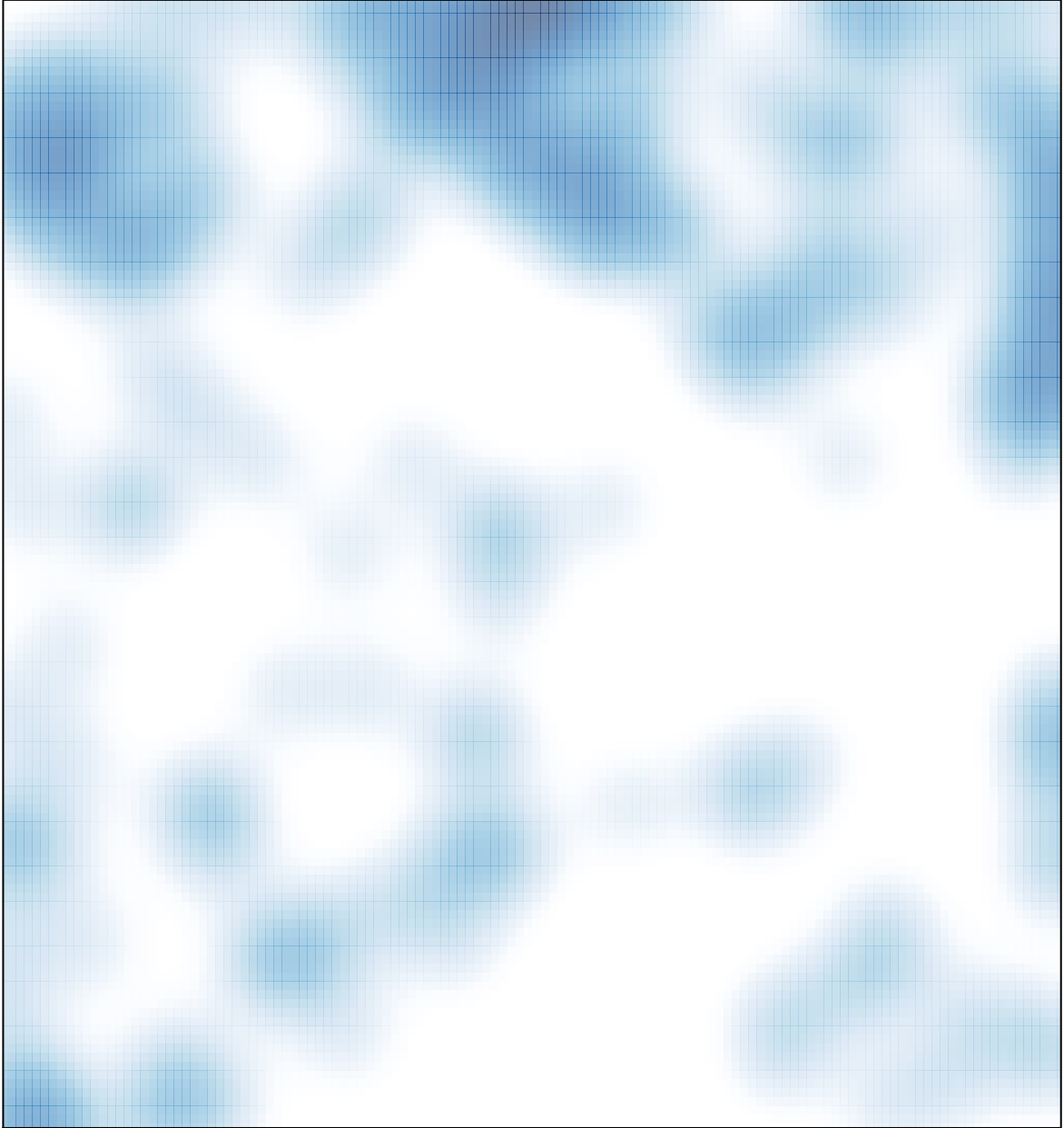


# hsa-miR-29a



# features = 172  
chi-square p = 0.7

# hsa-miR-29a



# features = 172 , max = 4