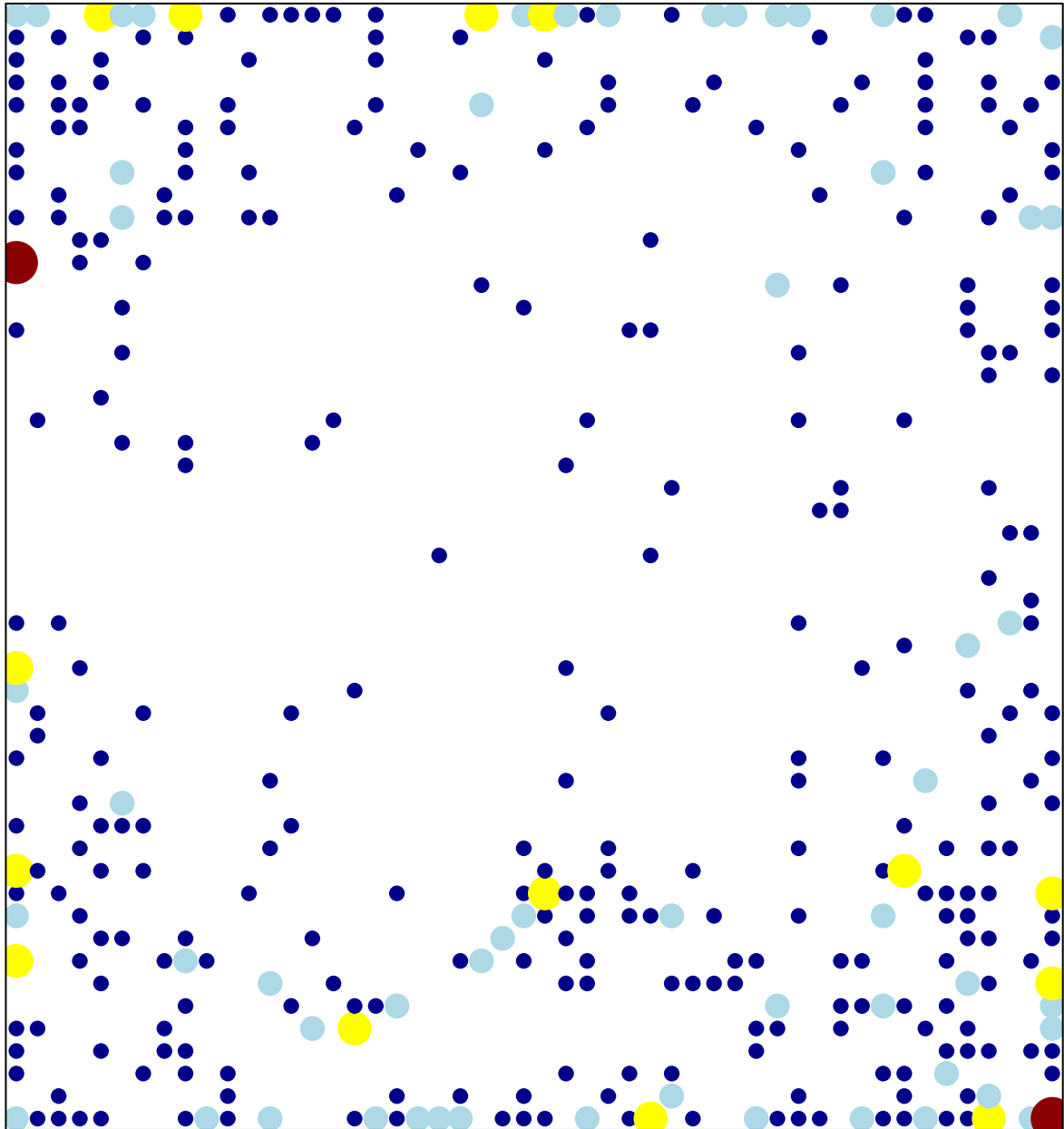
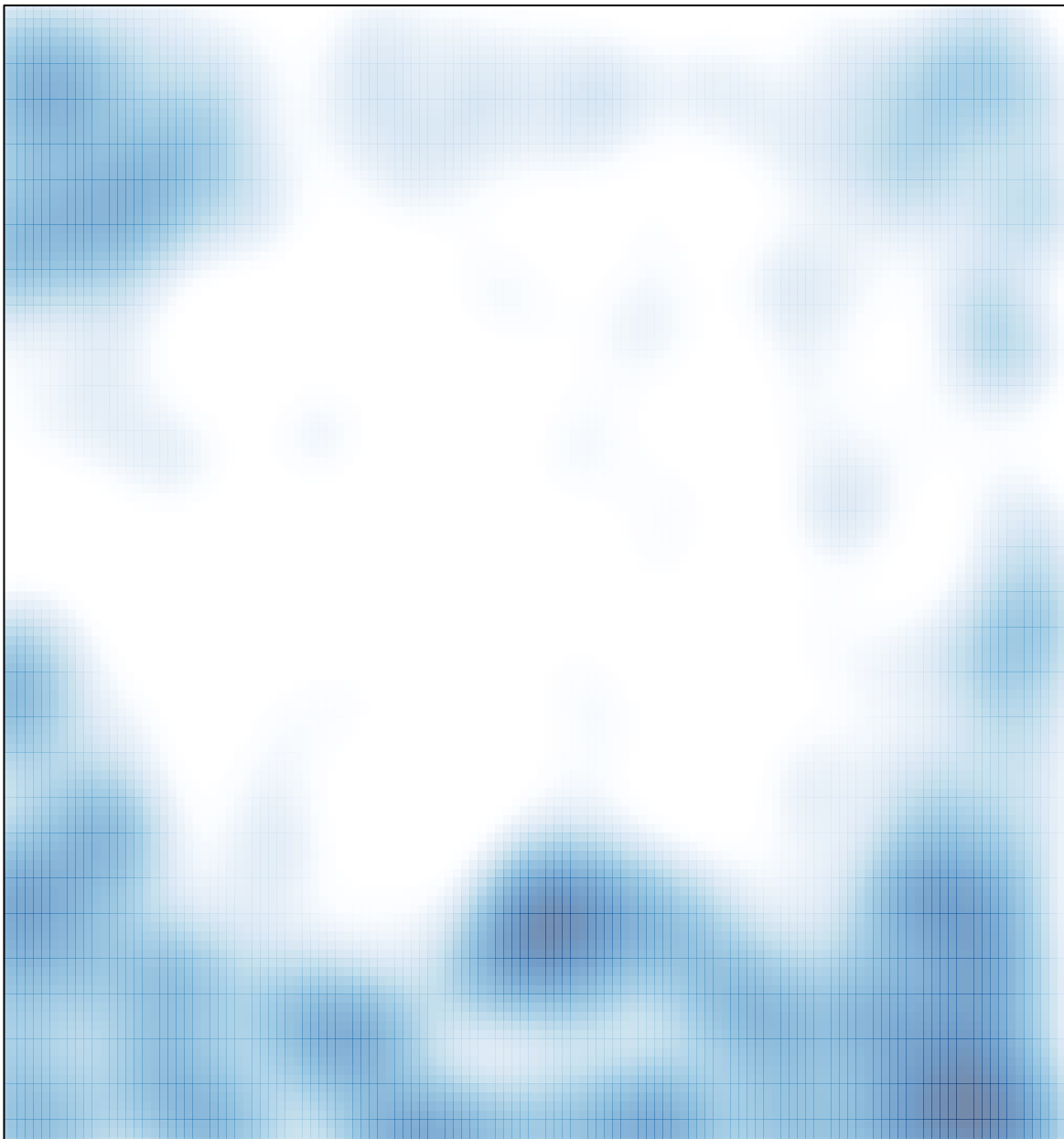


# hsa-miR-130a



# features = 456  
chi-square p = 0.58

# hsa-miR-130a



# features = 456 , max = 4