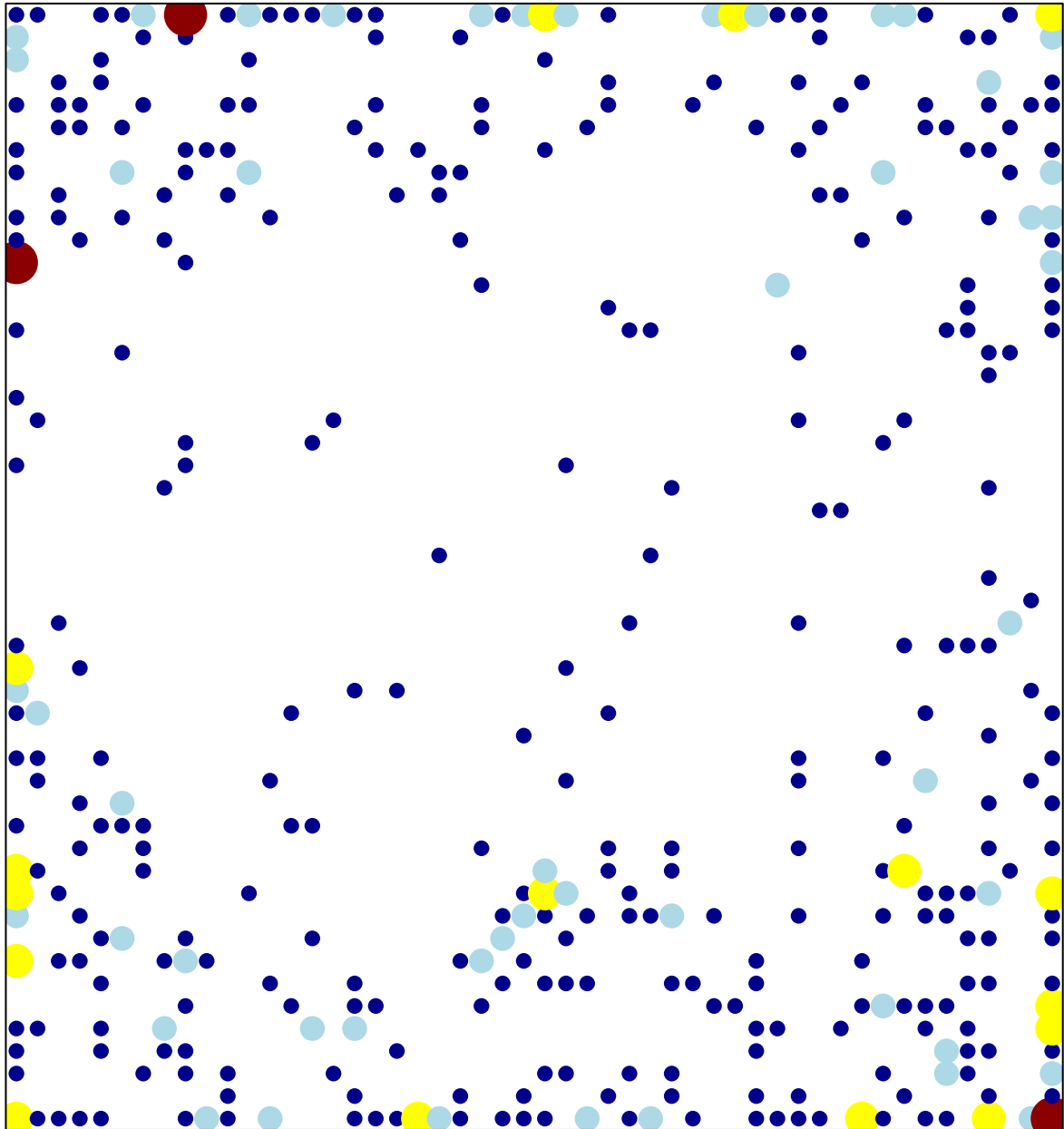
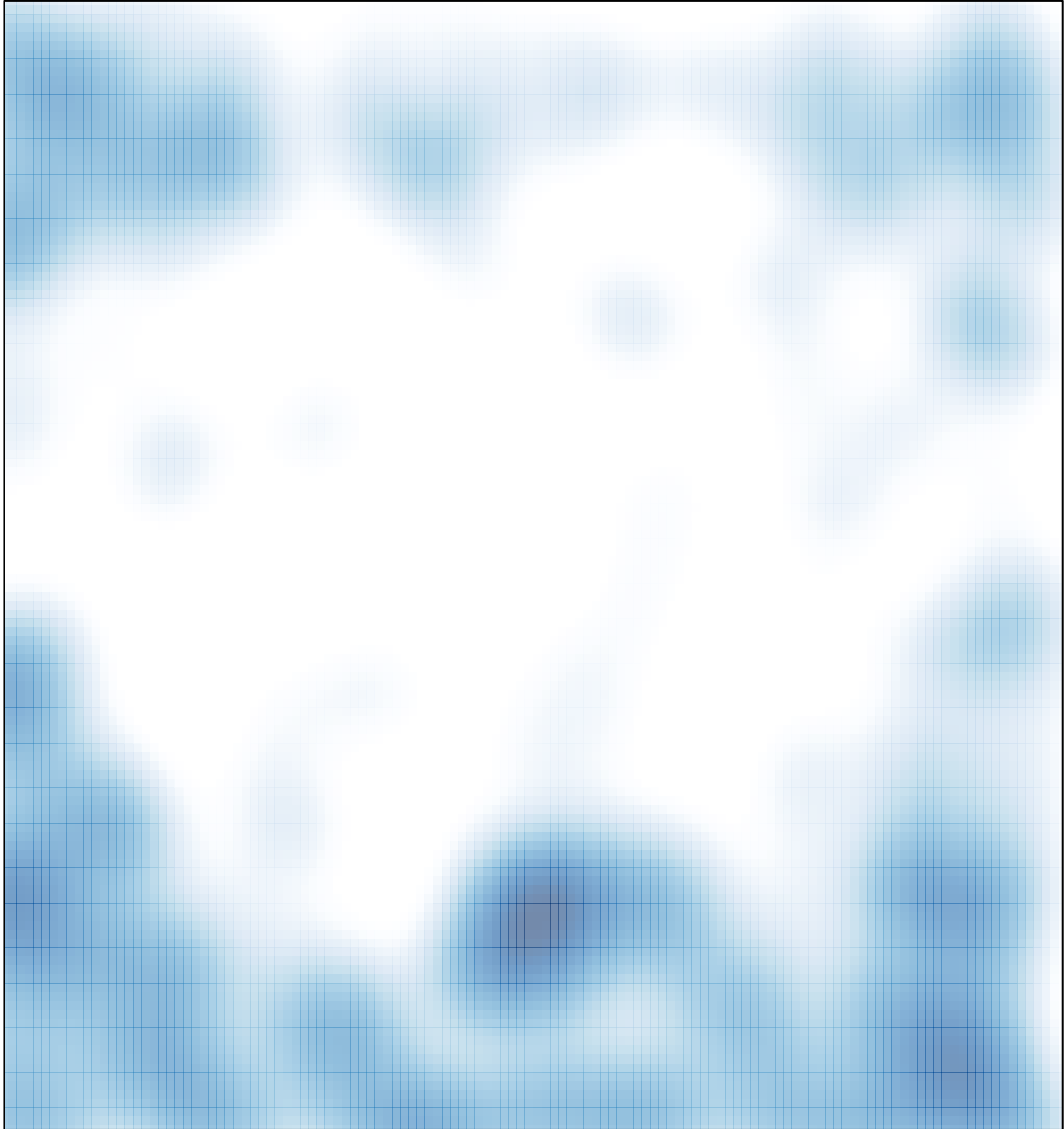


# hsa-miR-301a



# features = 467  
chi-square p = 0.57

# hsa-miR-301a



# features = 467 , max = 4