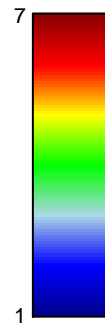
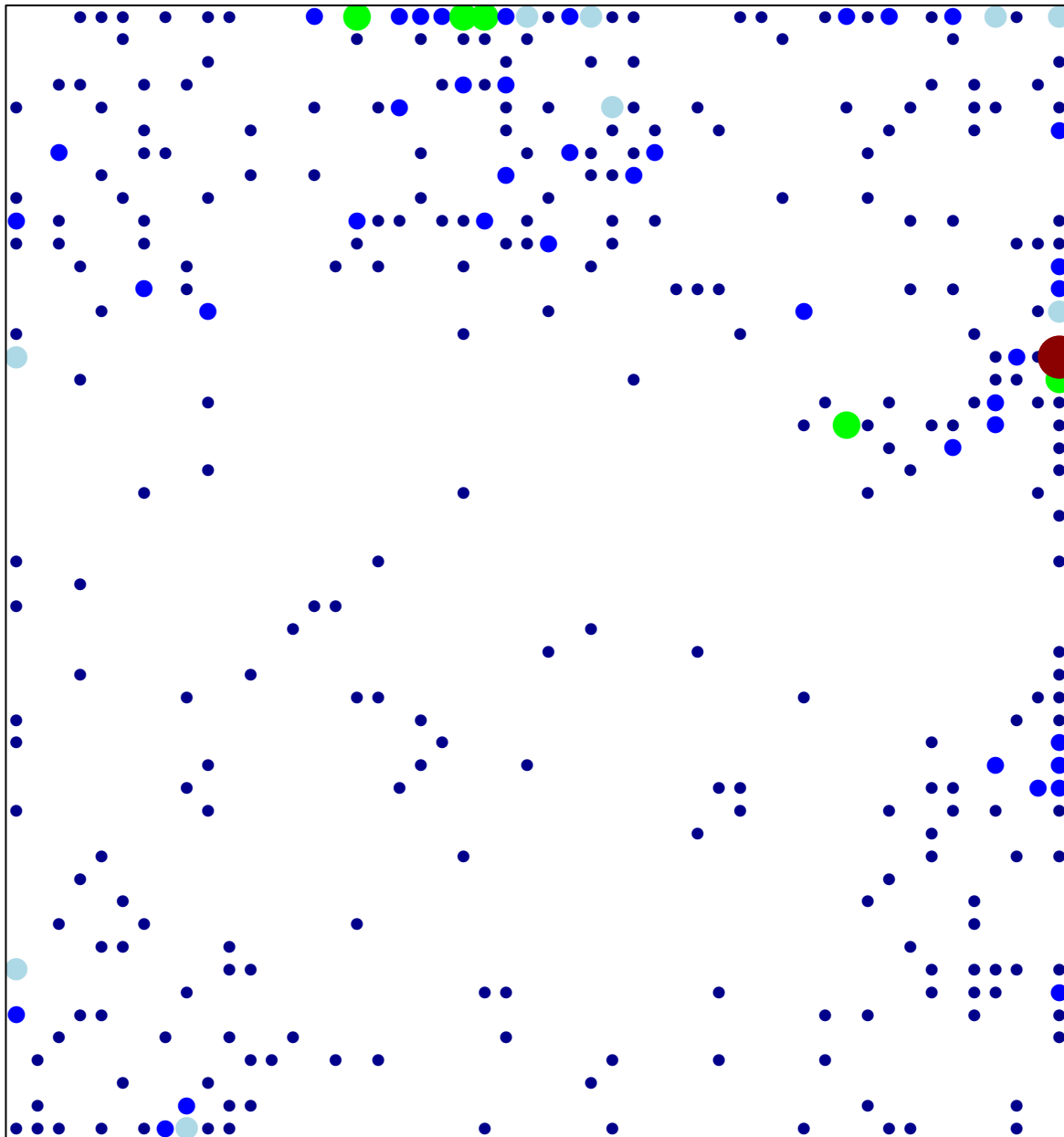
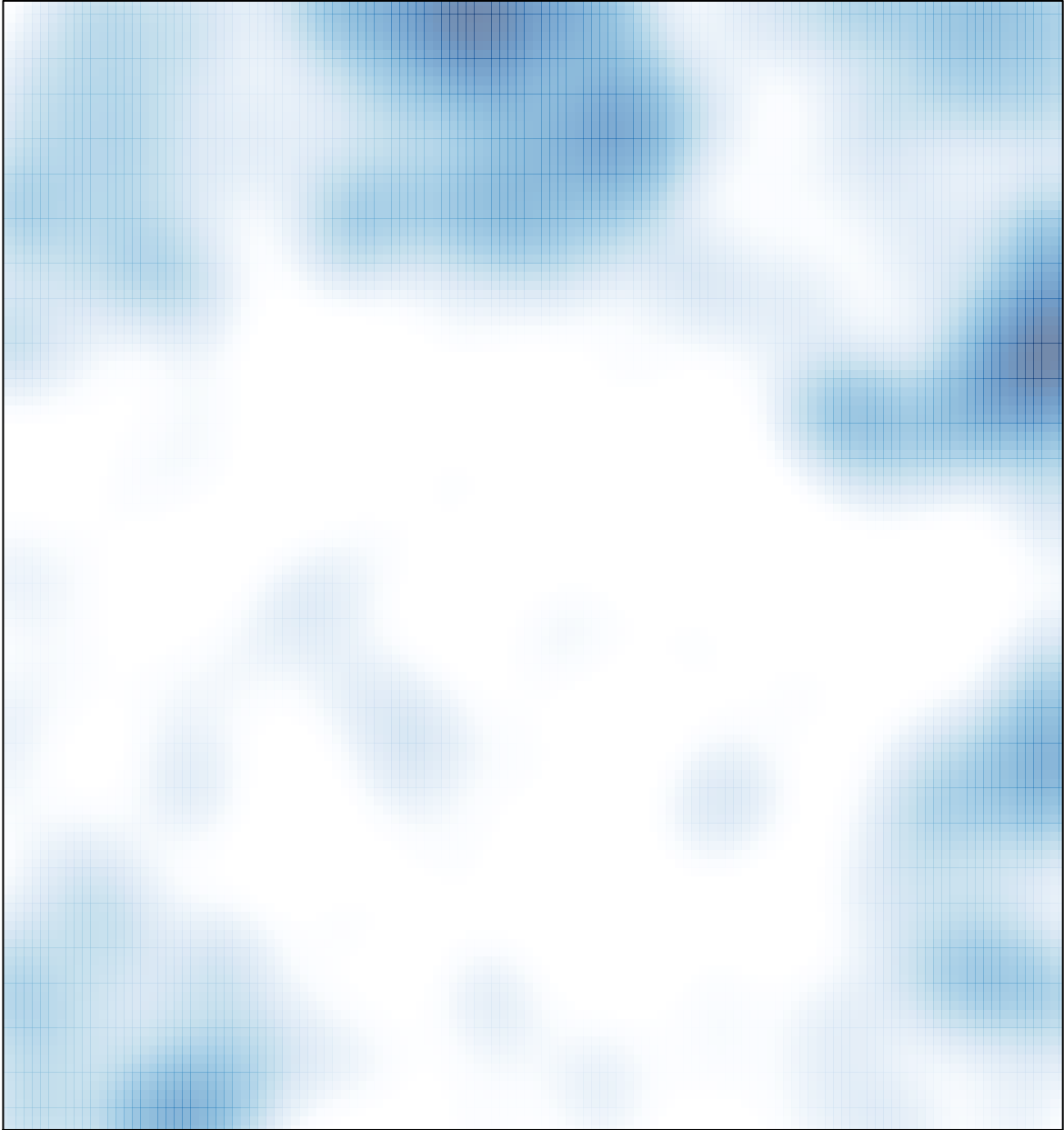


# hsa-miR-301b



# features = 398  
chi-square p = 0.59

# hsa-miR-301b



# features = 398 , max = 7