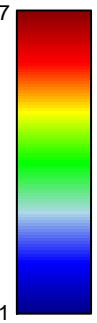
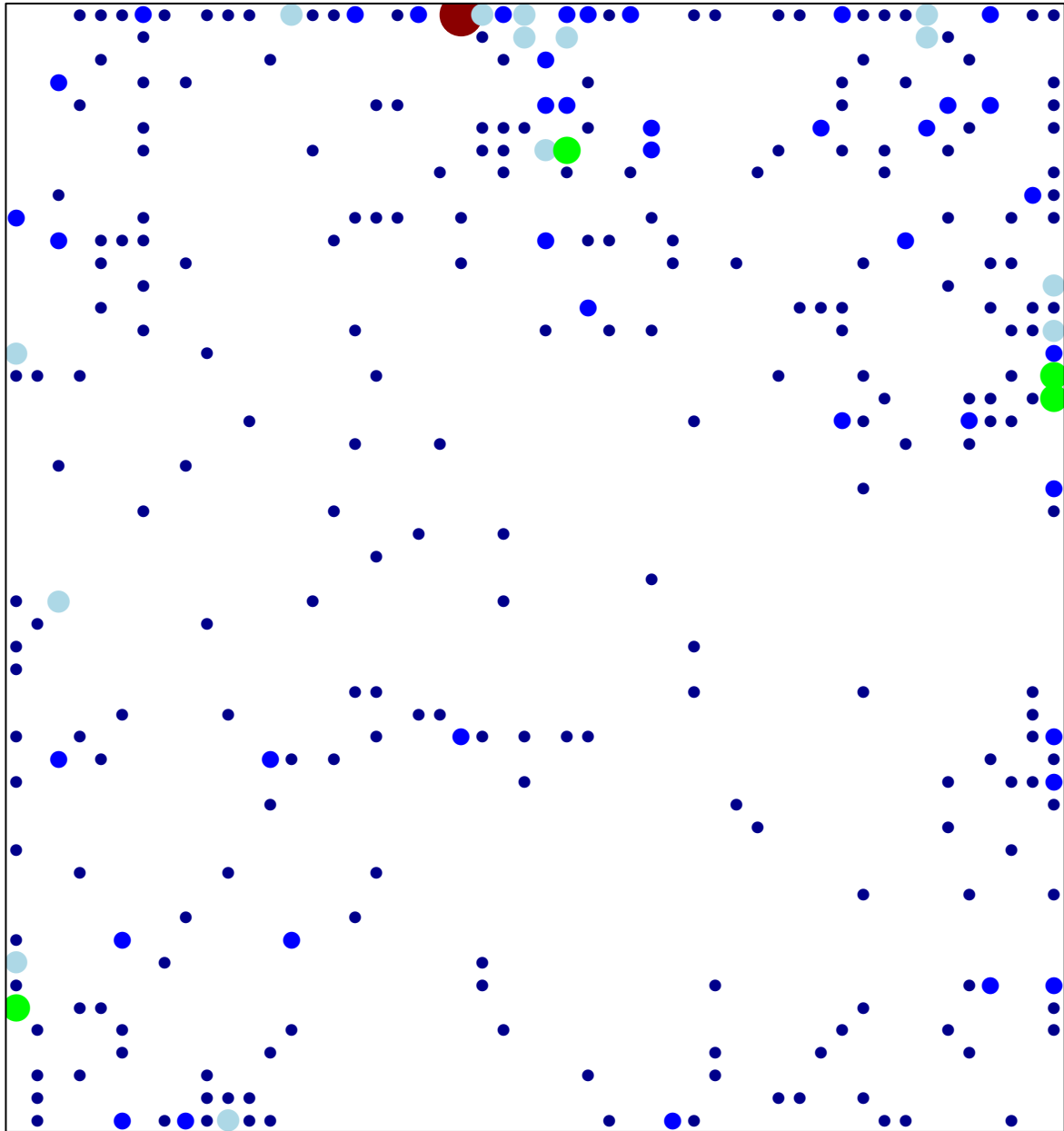
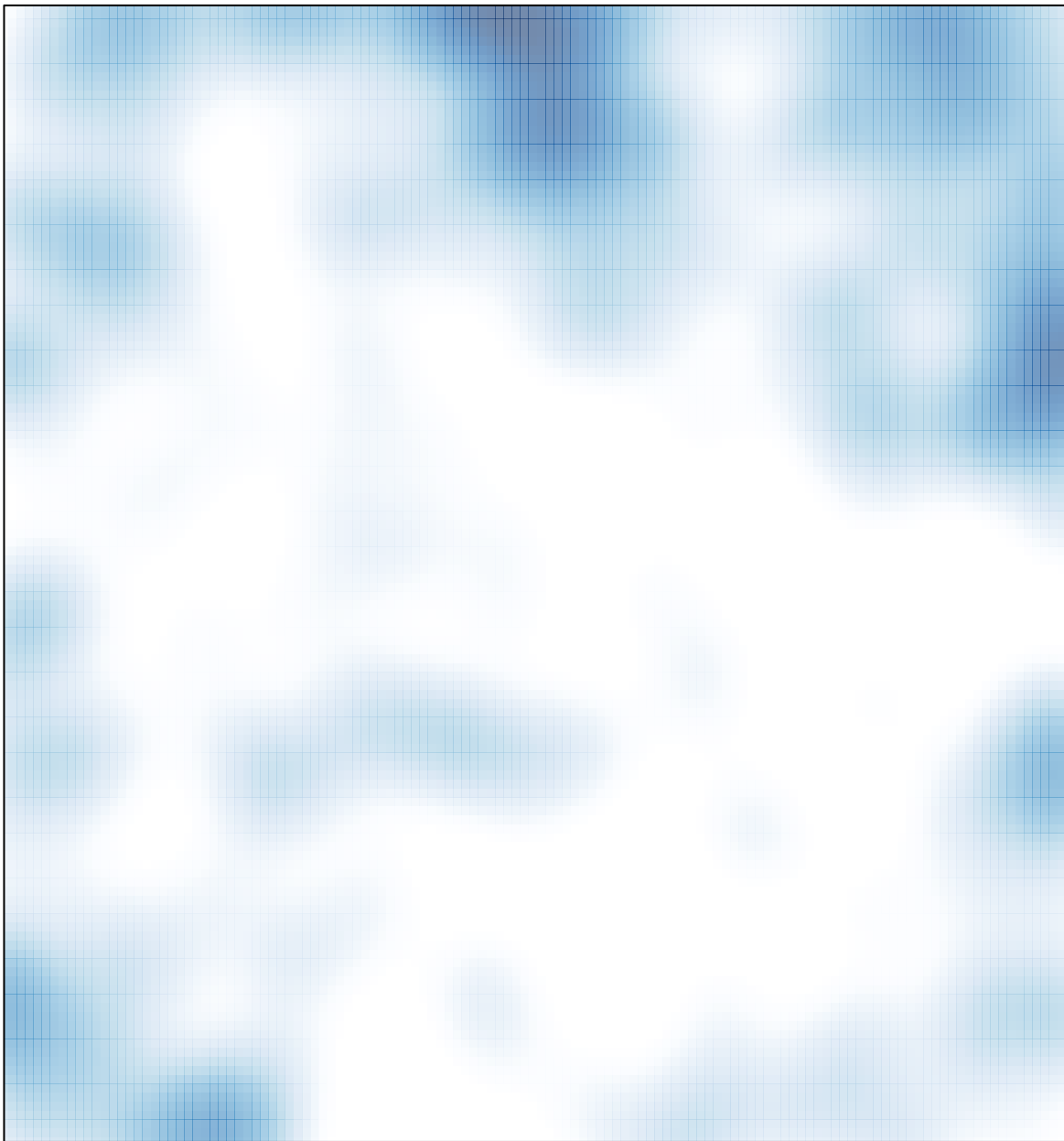


# hsa-miR-373



# features = 378  
chi-square p = 0.61

# hsa-miR-373



# features = 378 , max = 7