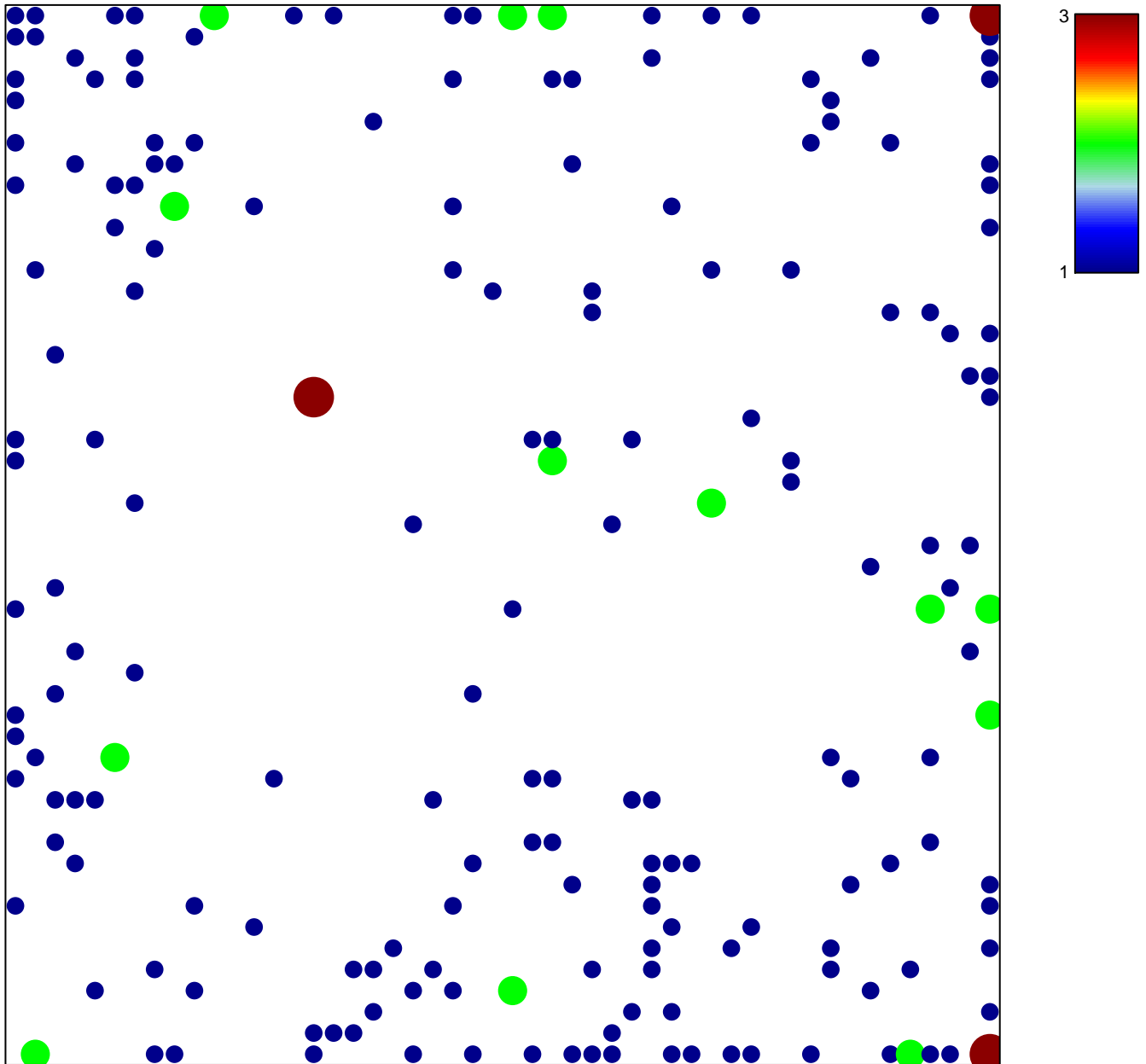
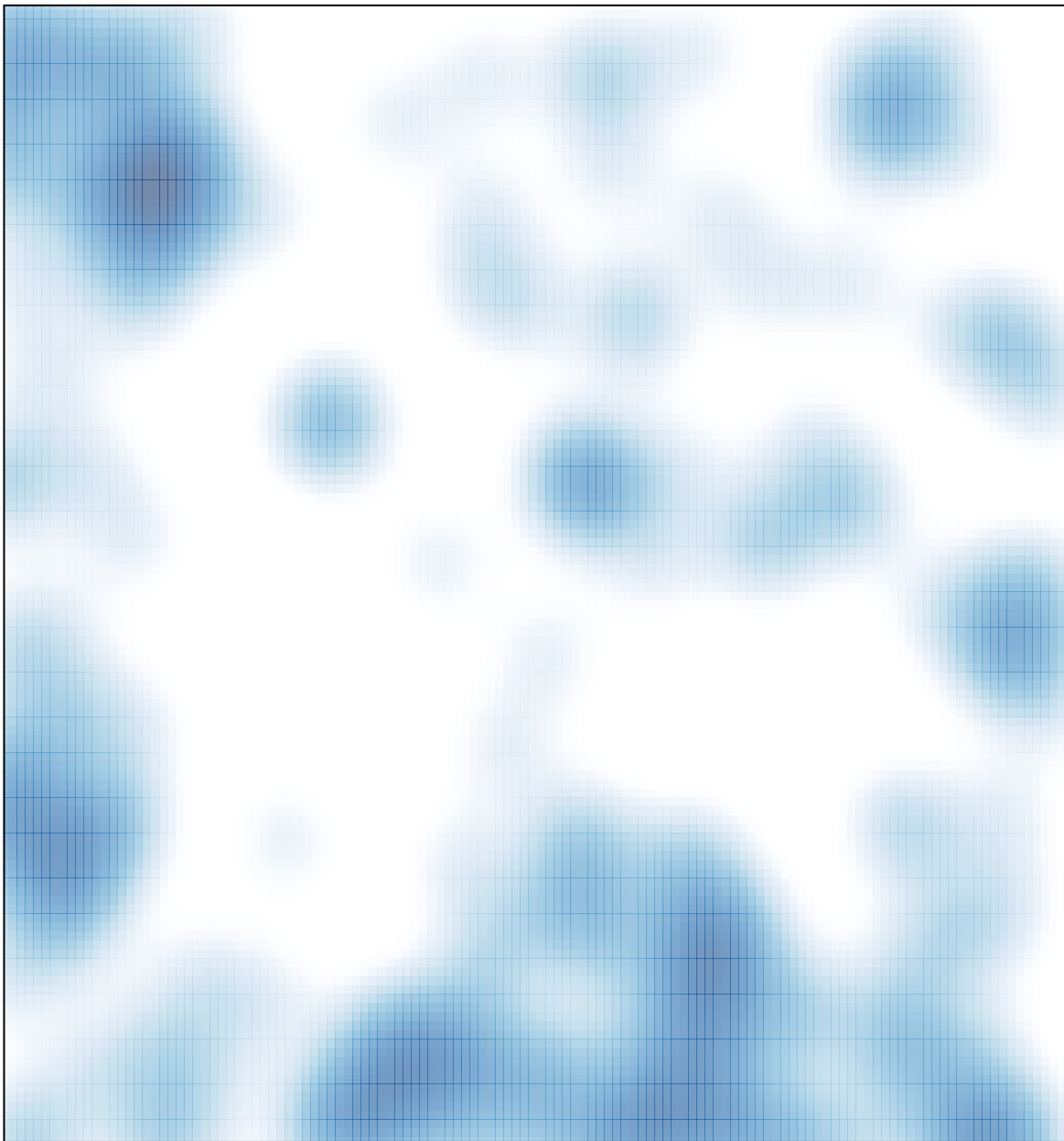


# hsa-miR-196a



# features = 209  
chi-square p = 0.7

# hsa-miR-196a



# features = 209 , max = 3