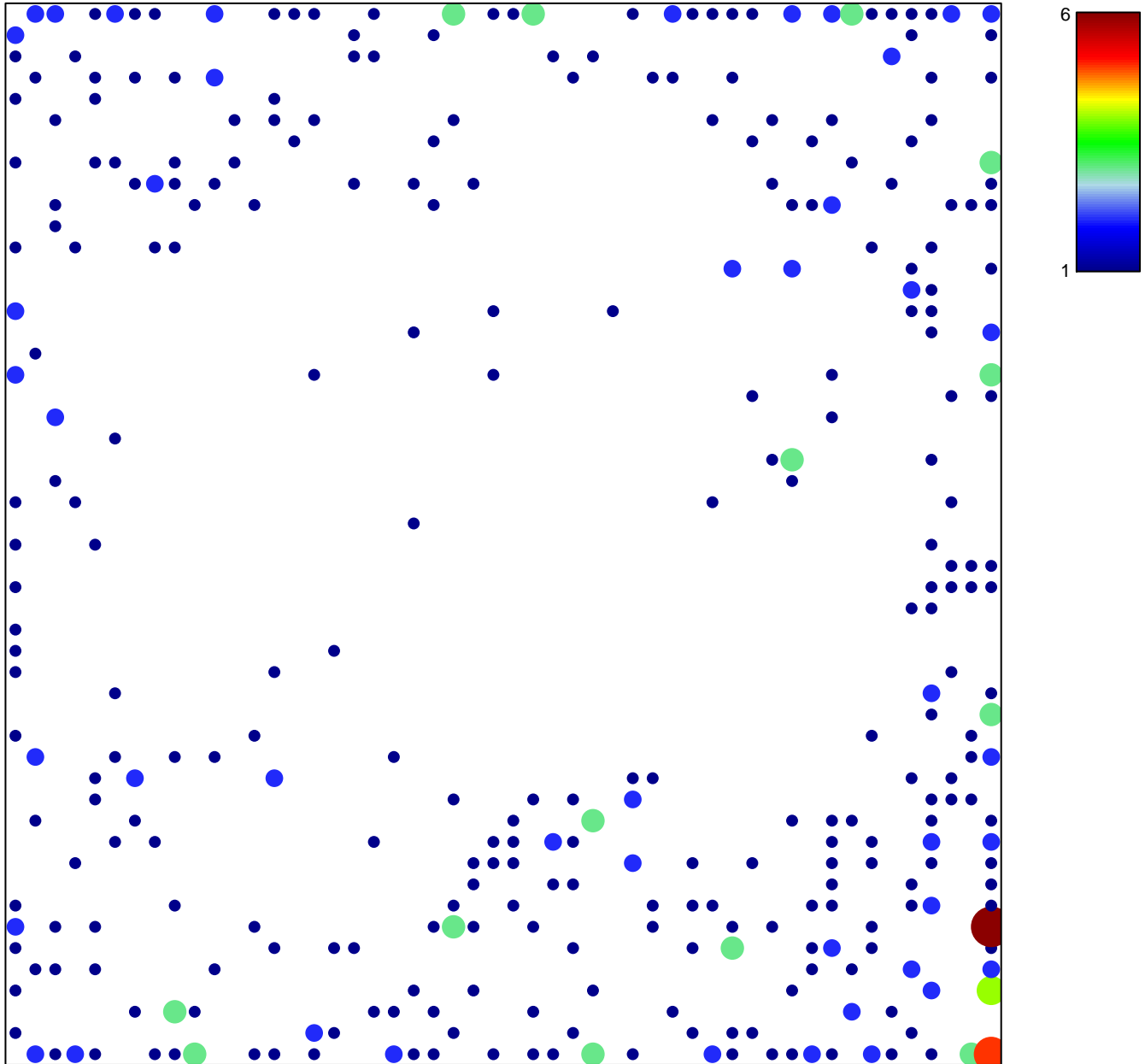
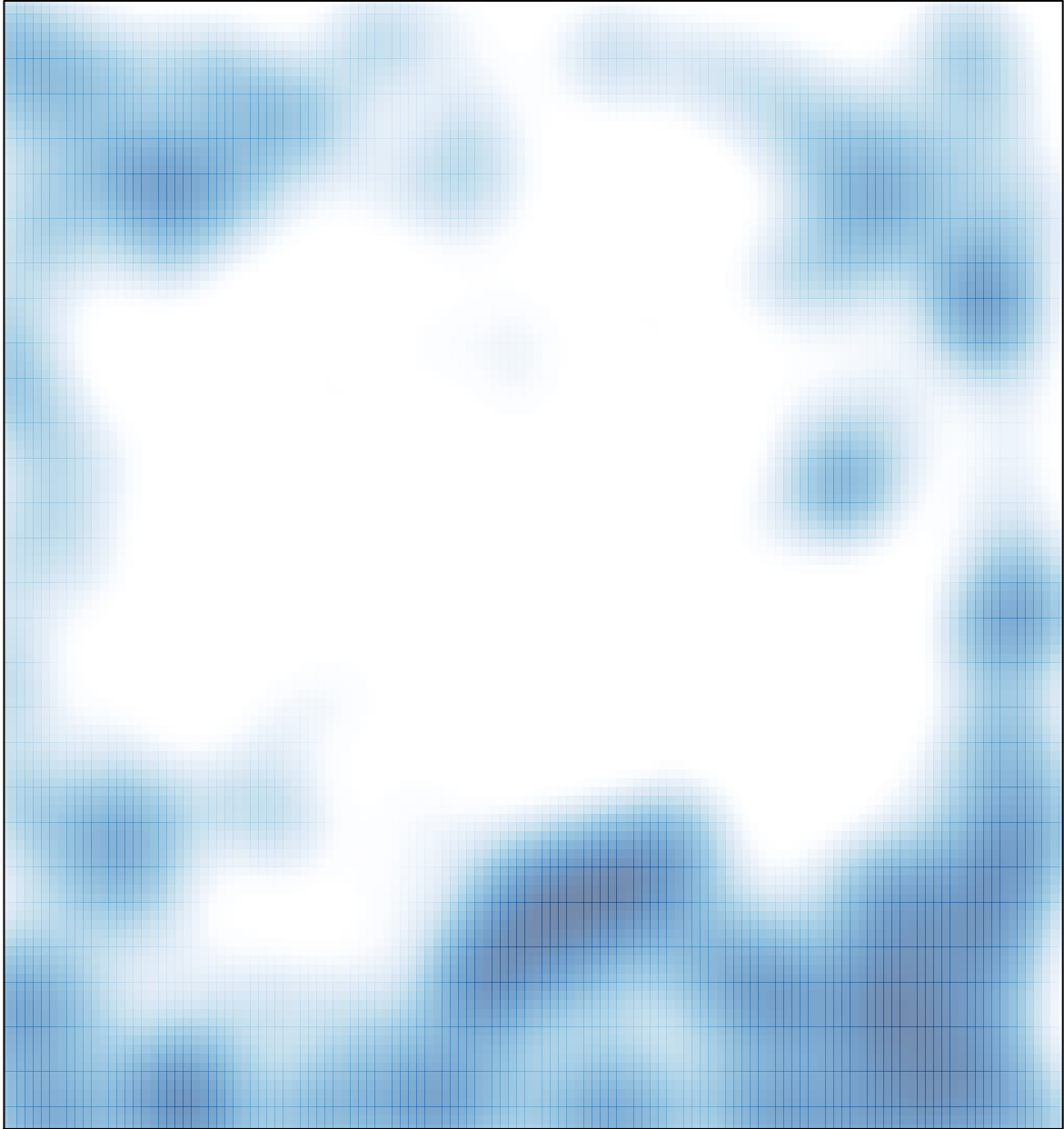


# hsa-miR-372



# features = 411  
chi-square p = 0.58

# hsa-miR-372



# features = 411 , max = 6