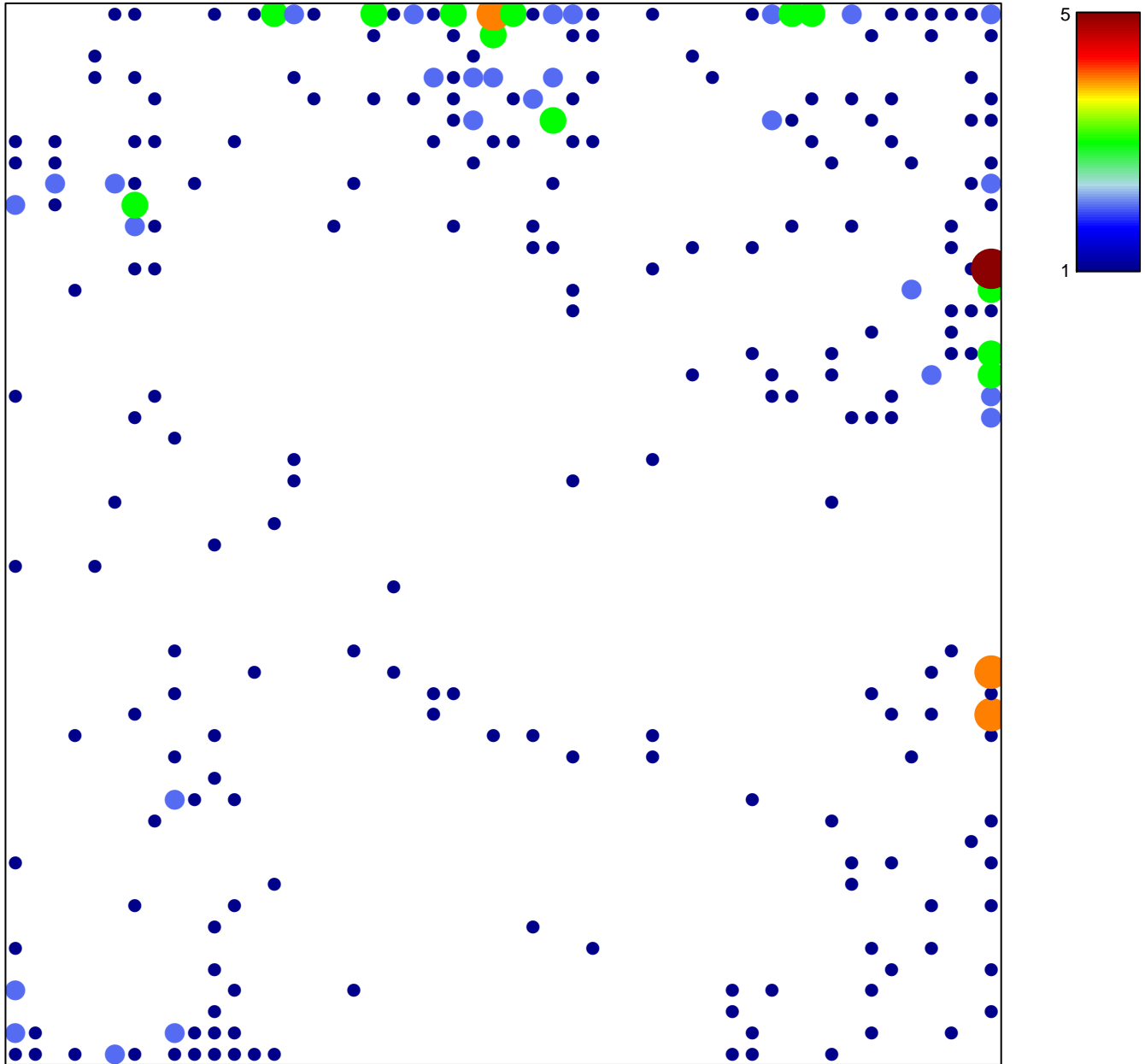
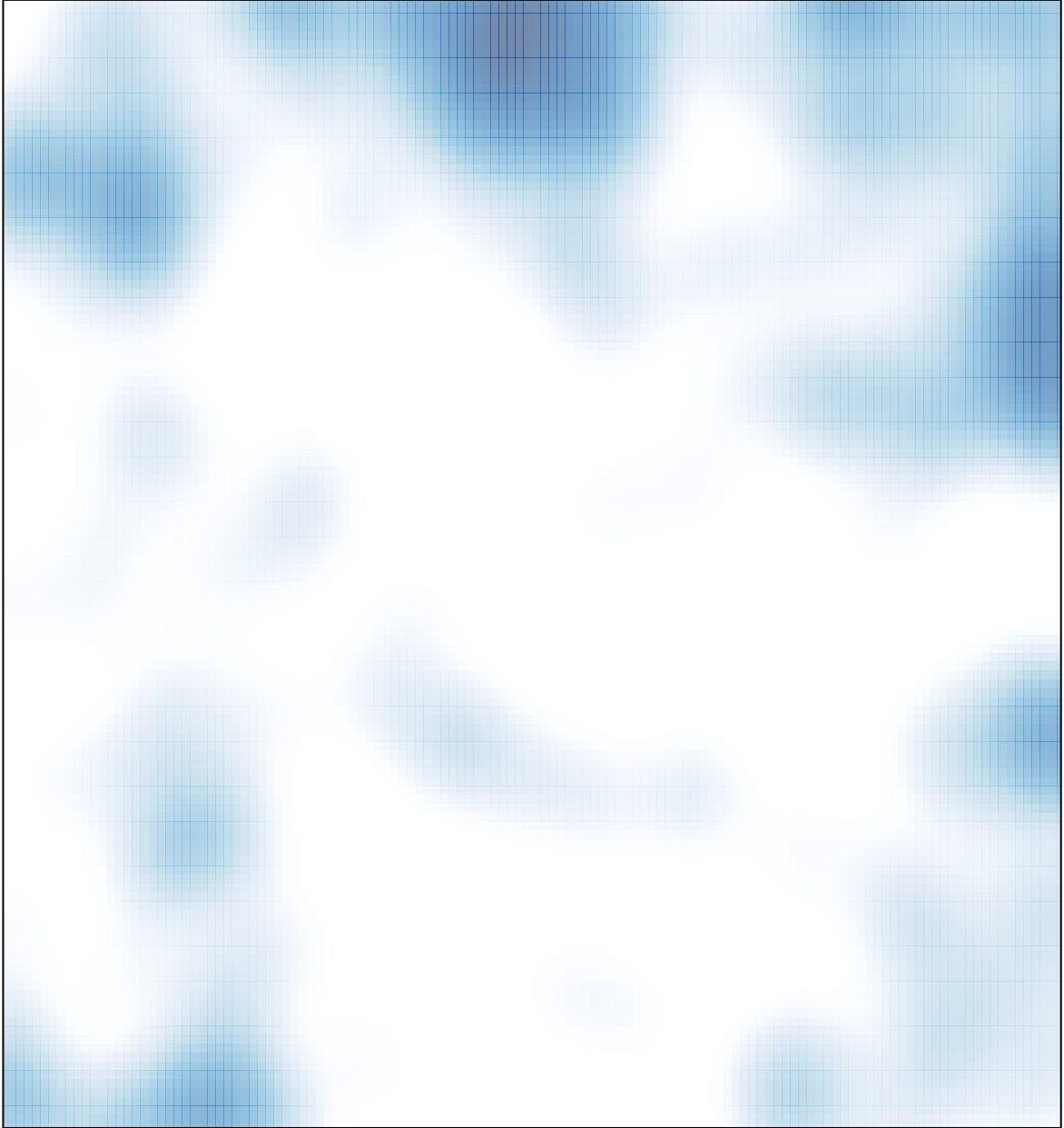


# hsa-miR-30e



# features = 315  
chi-square p = 0.58

# hsa-miR-30e



# features = 315 , max = 5