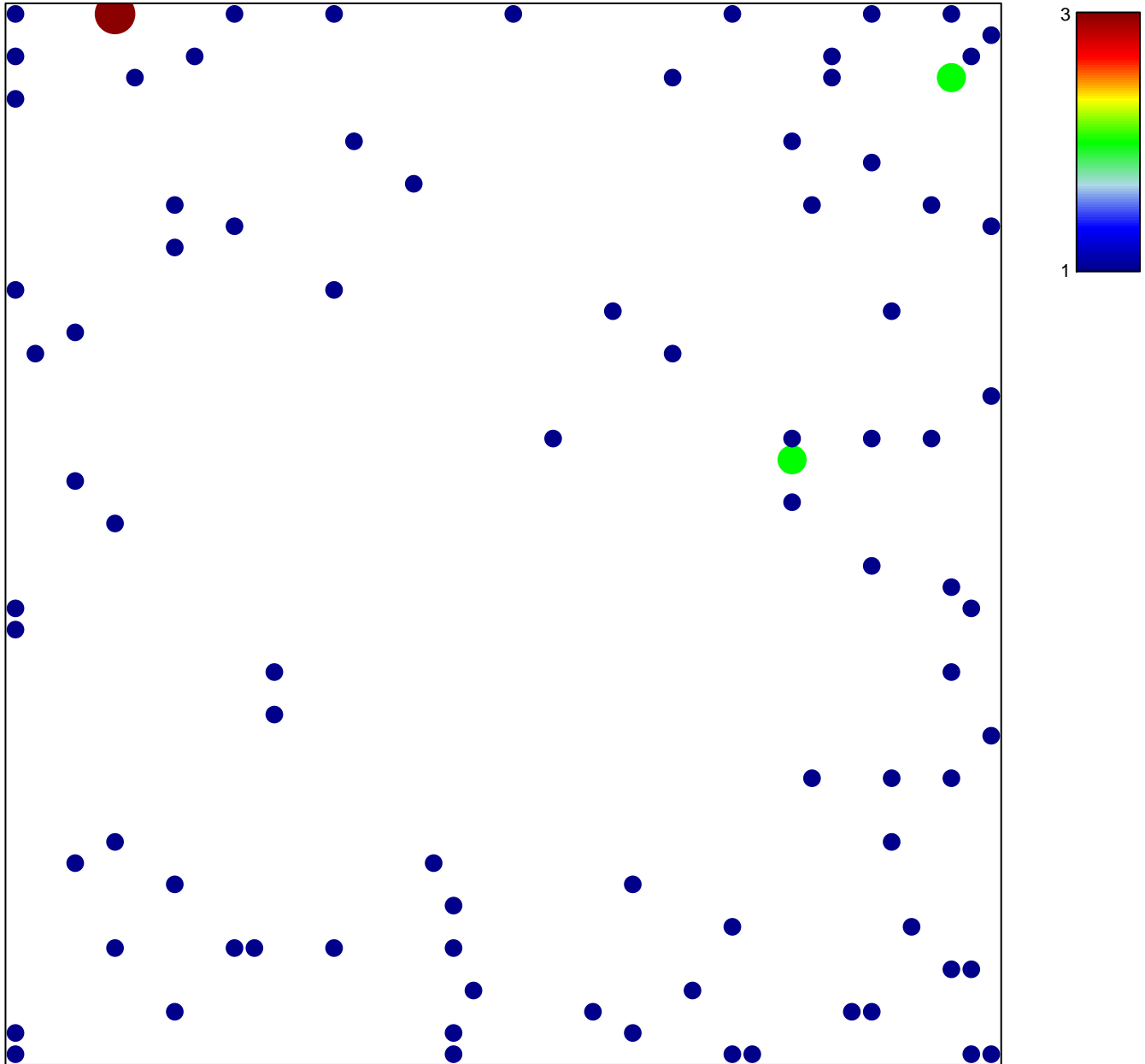
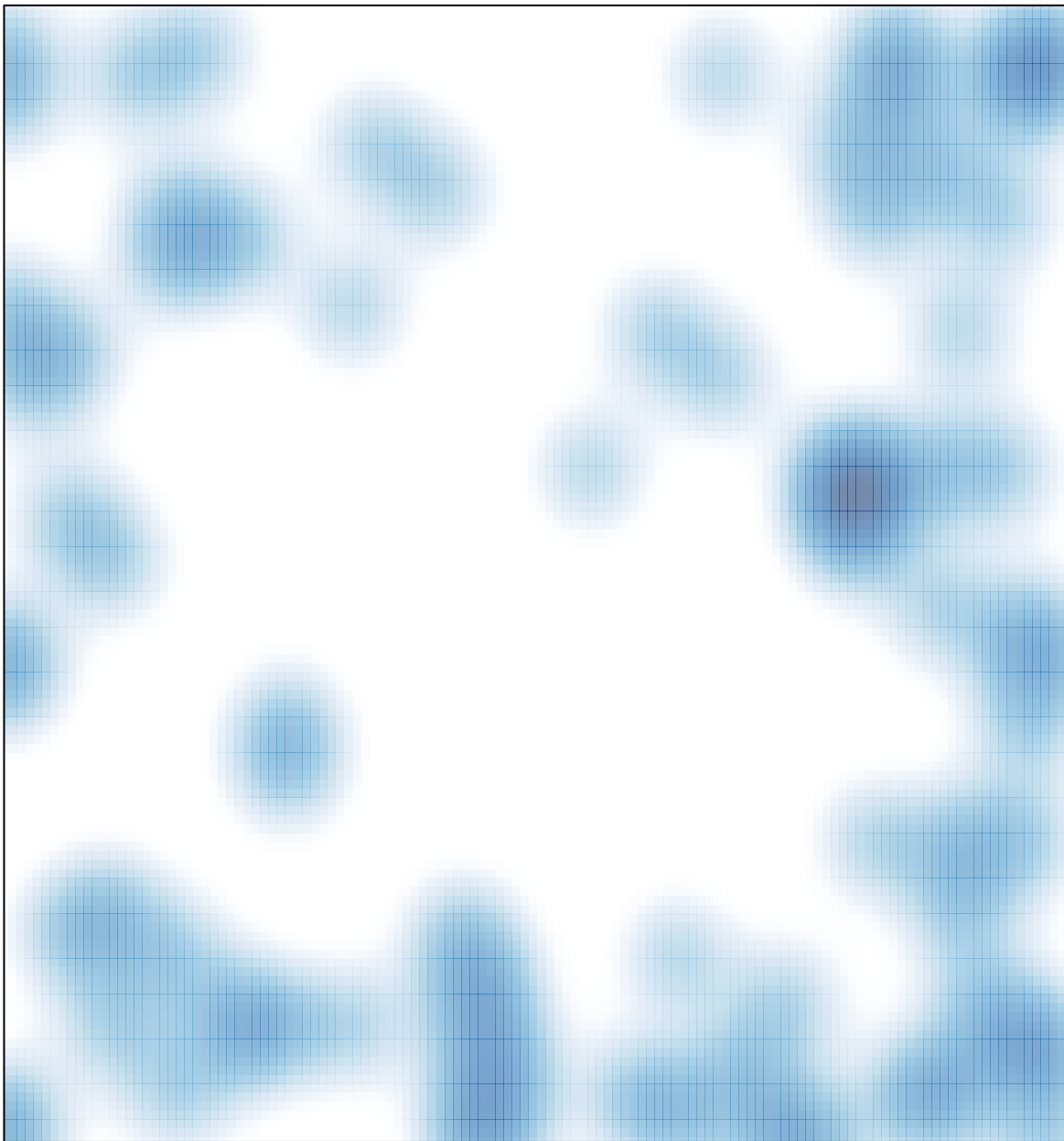


# hsa-miR-378



# features = 91  
chi-square p = 0.78

# hsa-miR-378



# features = 91 , max = 3