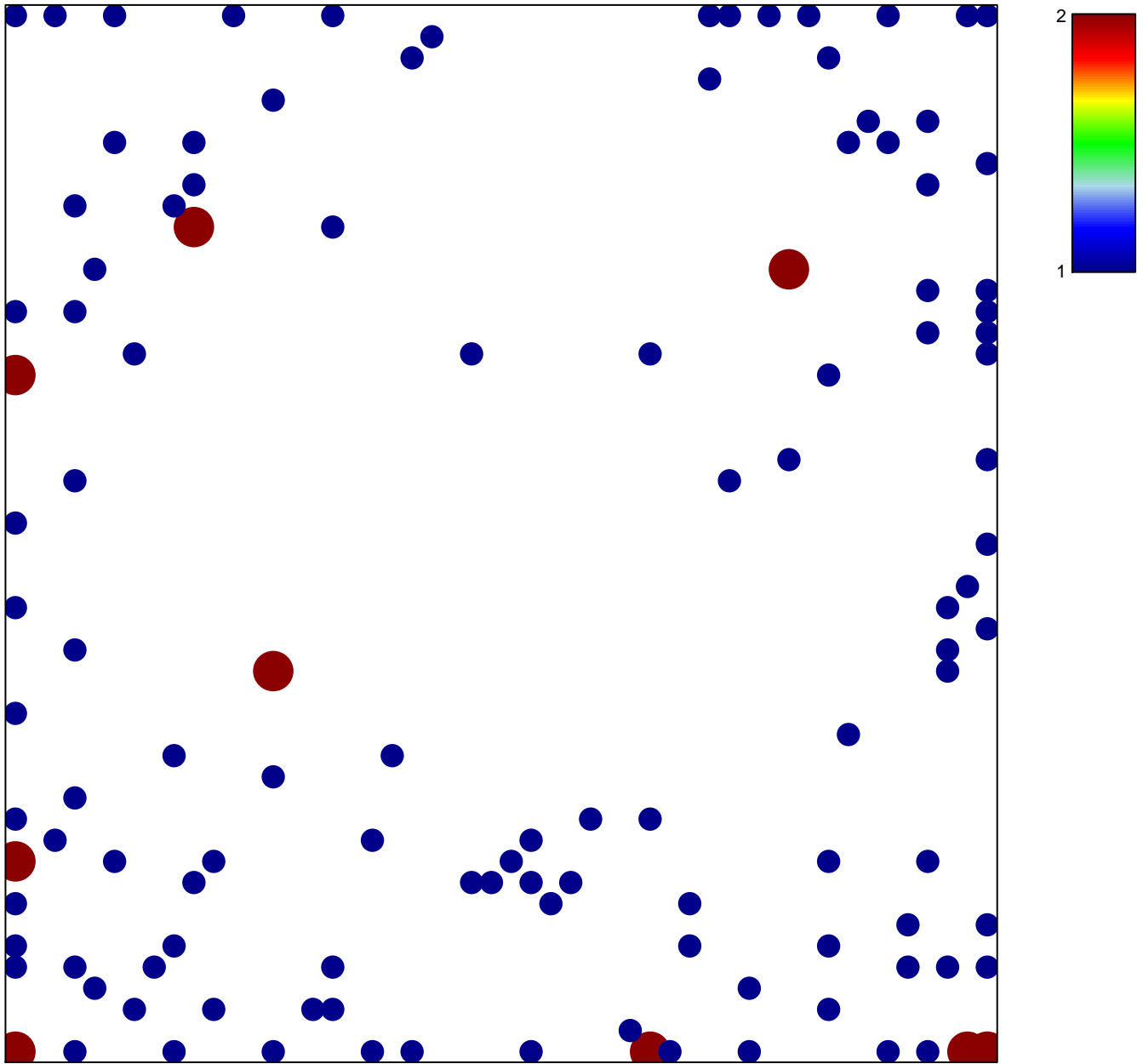
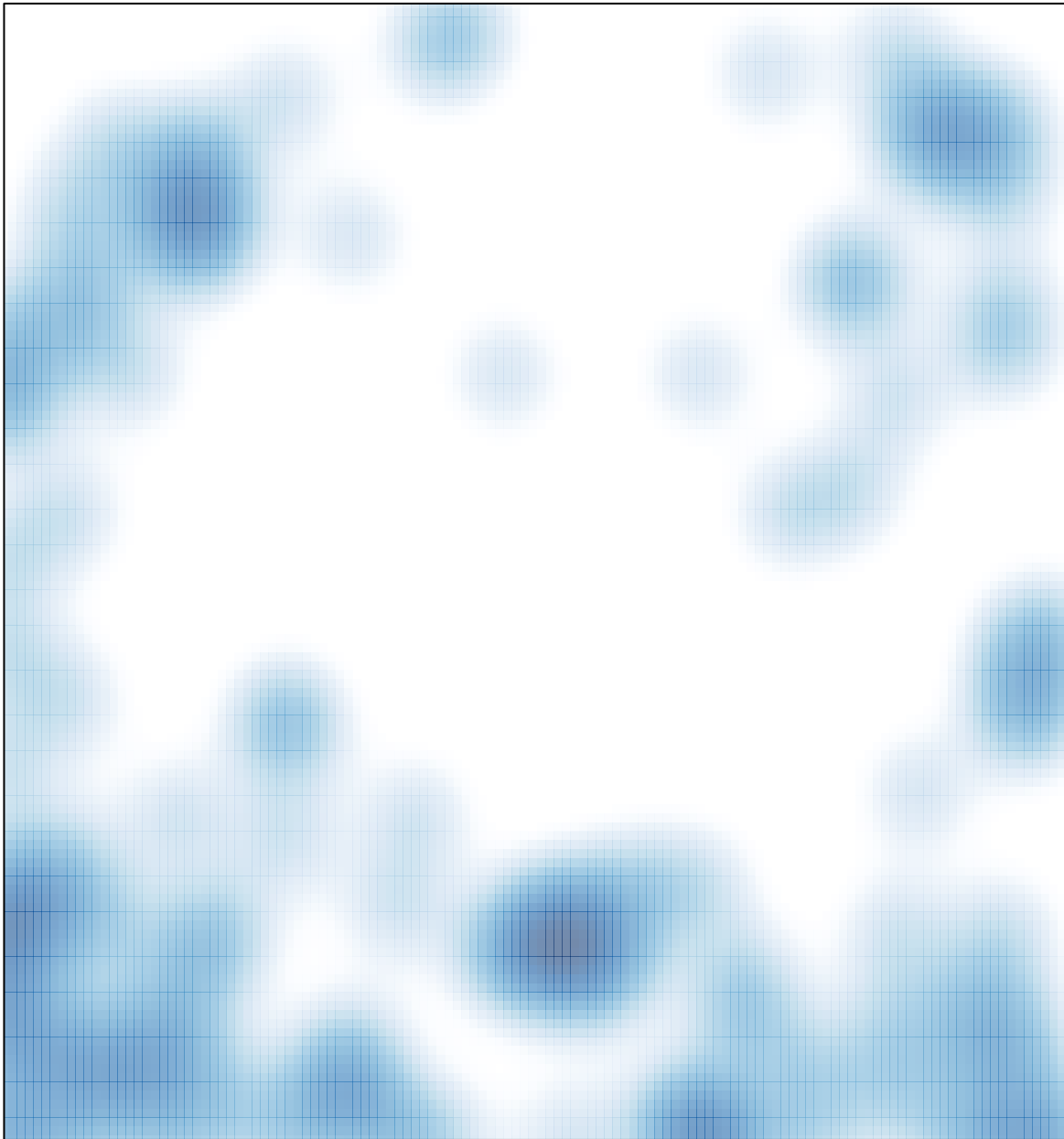


# hsa-miR-218



# features = 129  
chi-square p = 0.71

# hsa-miR-218



# features = 129 , max = 2