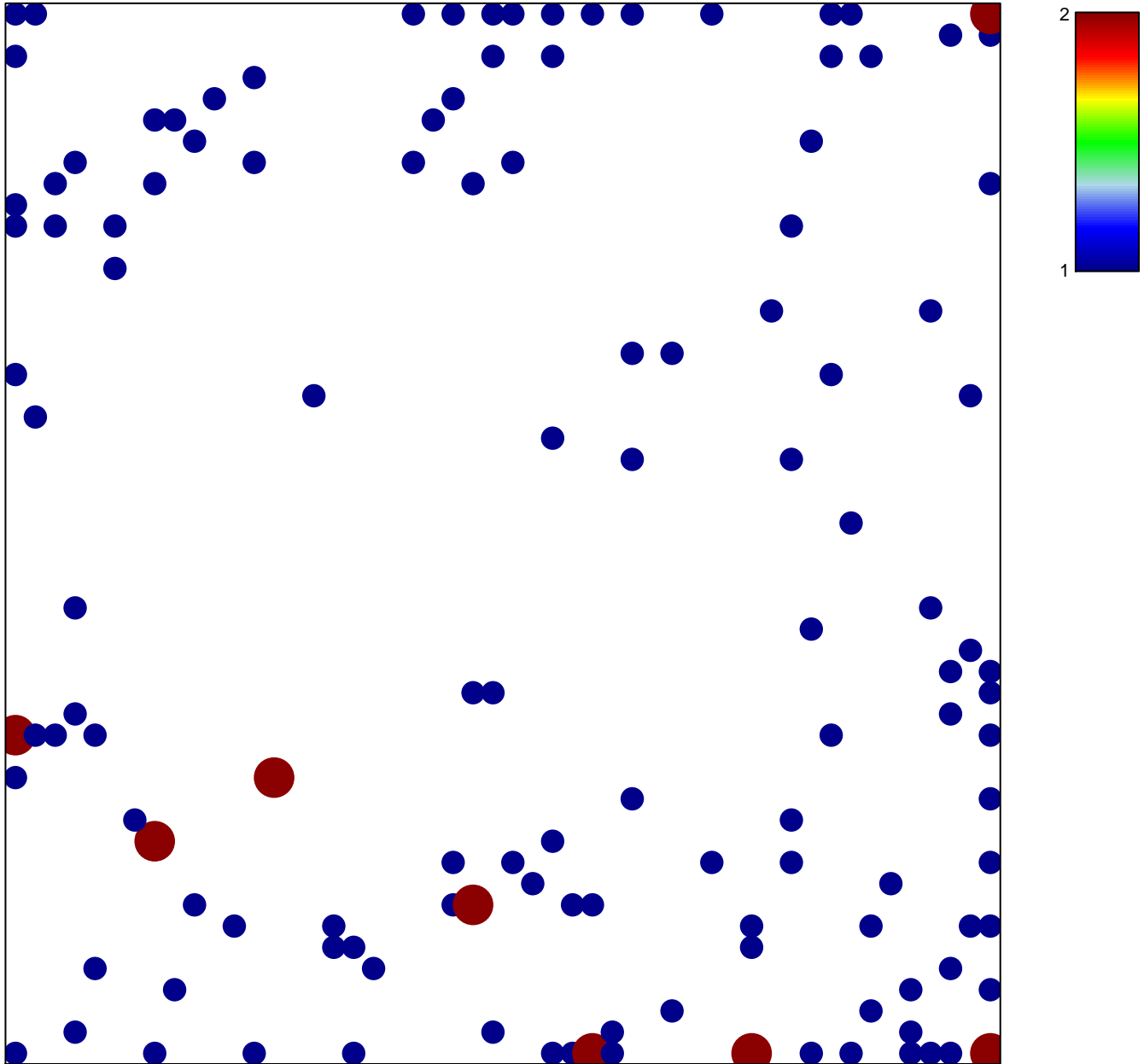
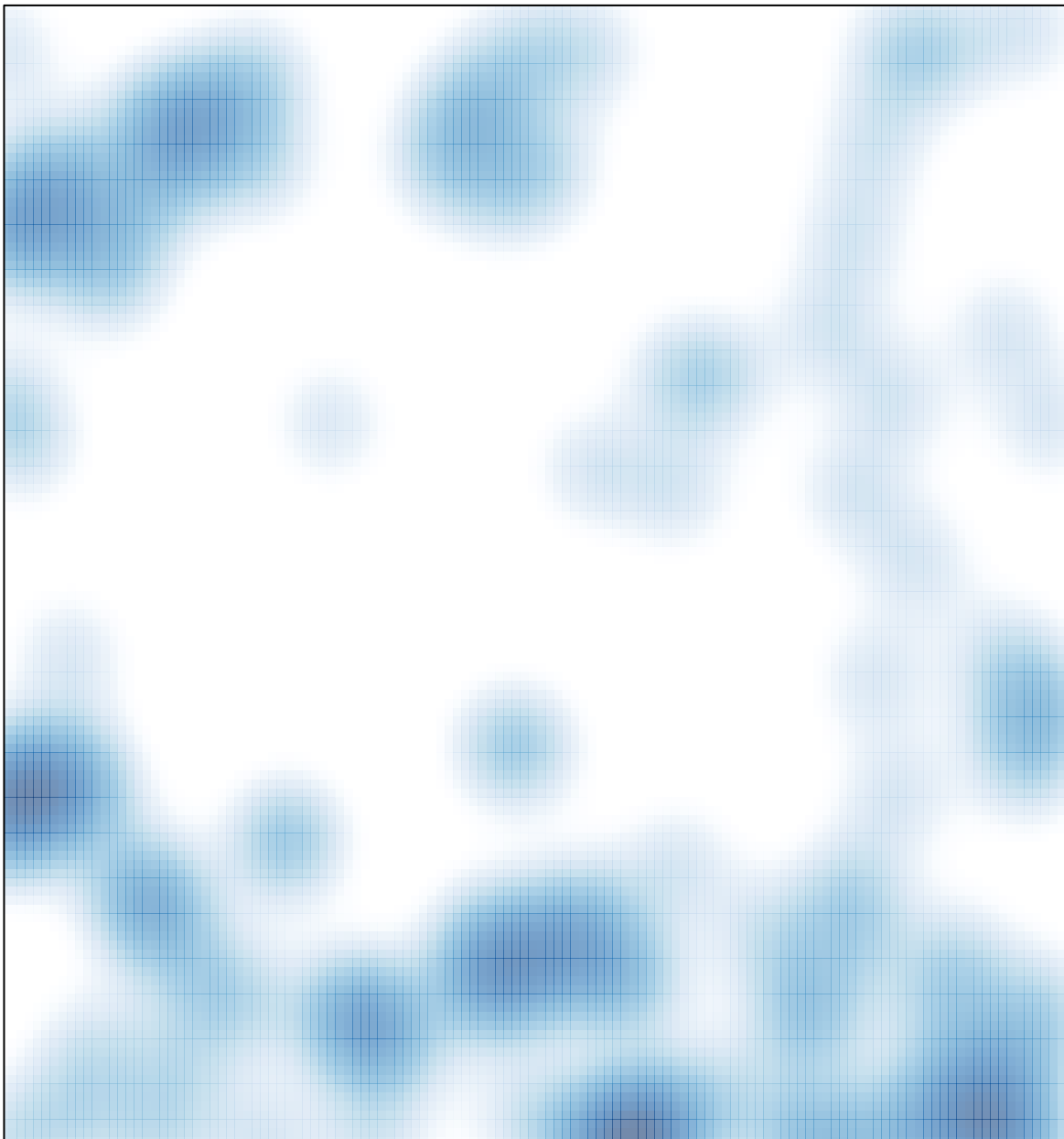


# hsa-miR-155



# features = 136  
chi-square p = 0.74

# hsa-miR-155



# features = 136 , max = 2