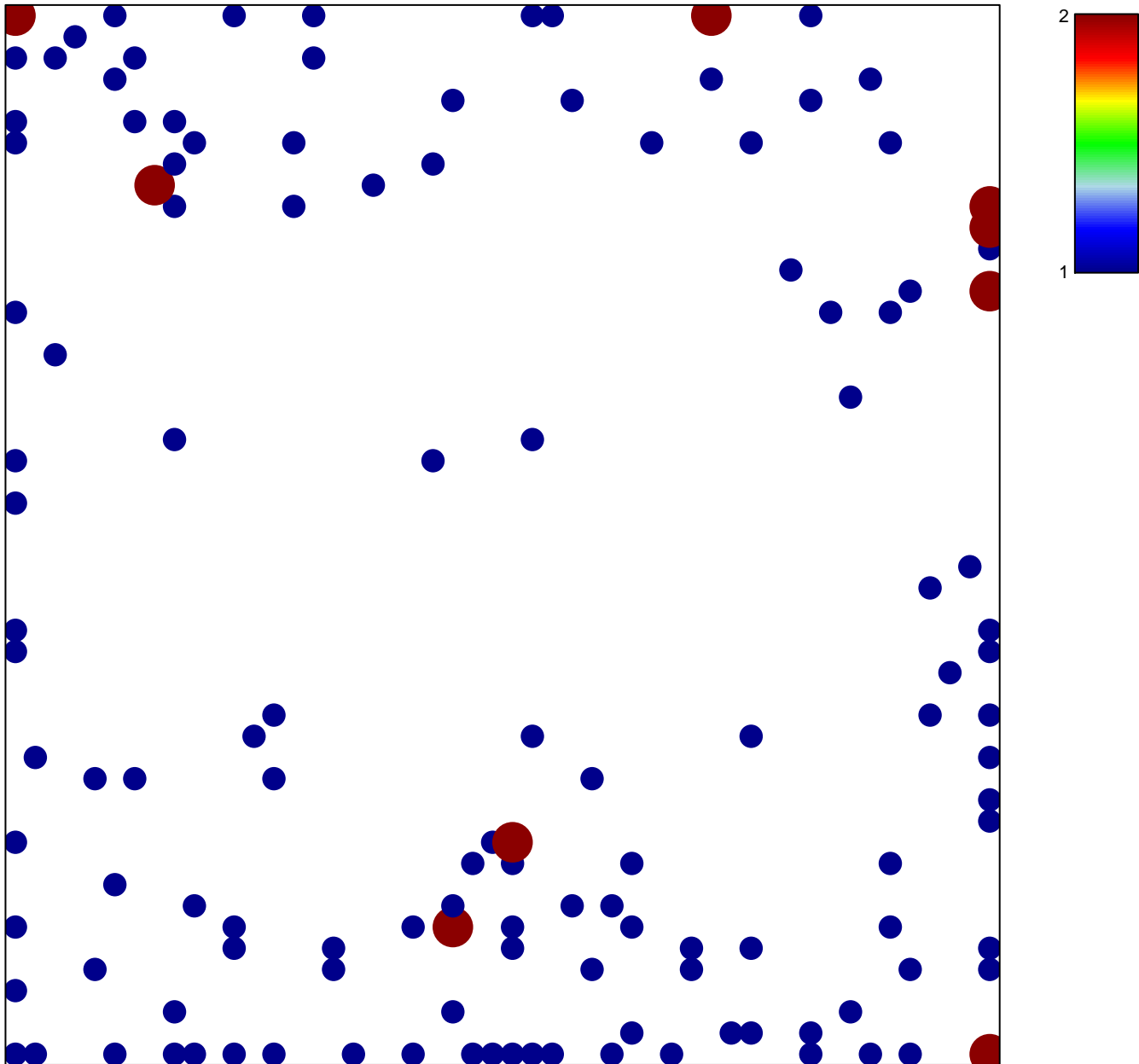
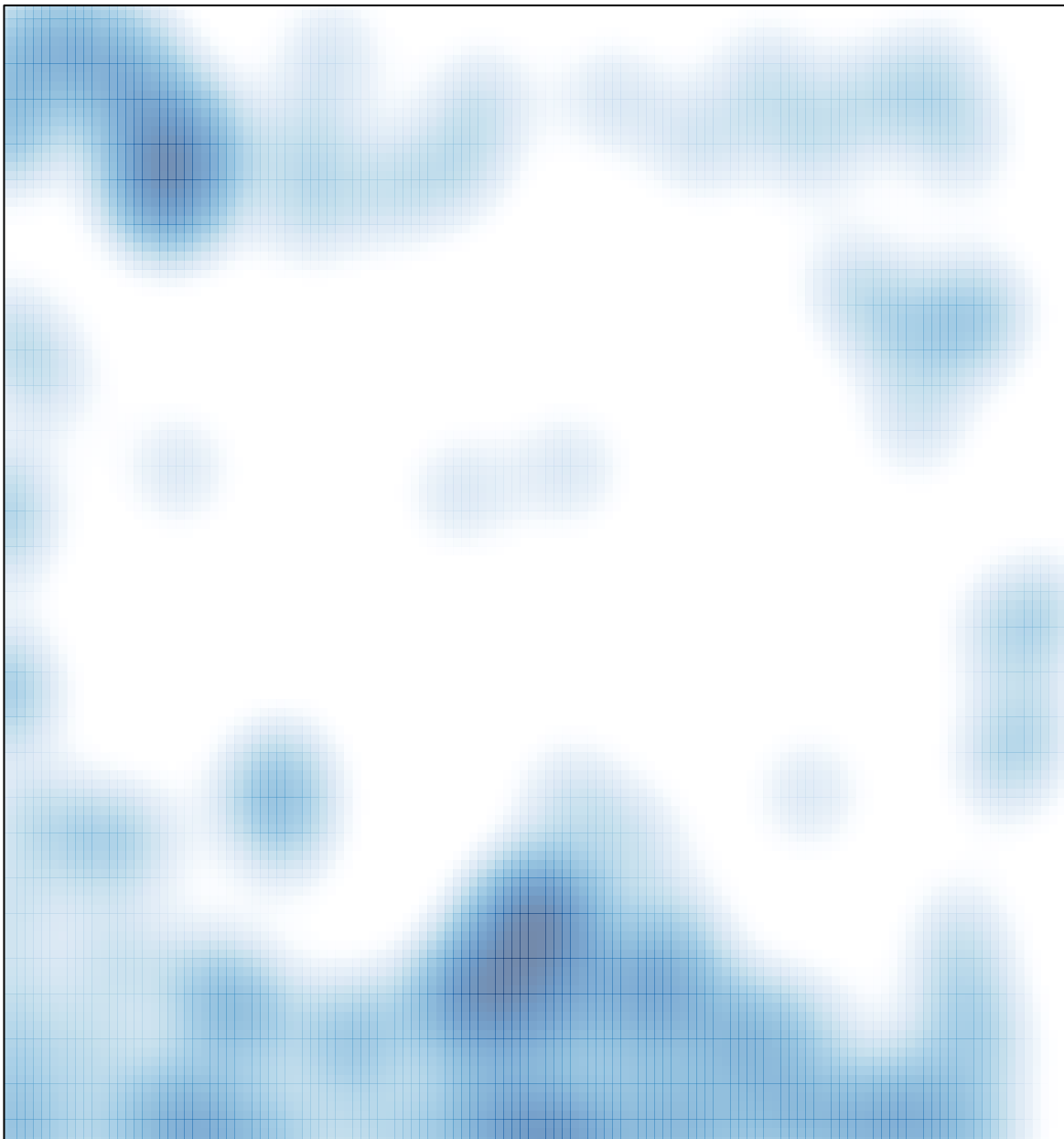


# hsa-miR-132



# features = 139  
chi-square p = 0.71

# hsa-miR-132



# features = 139 , max = 2