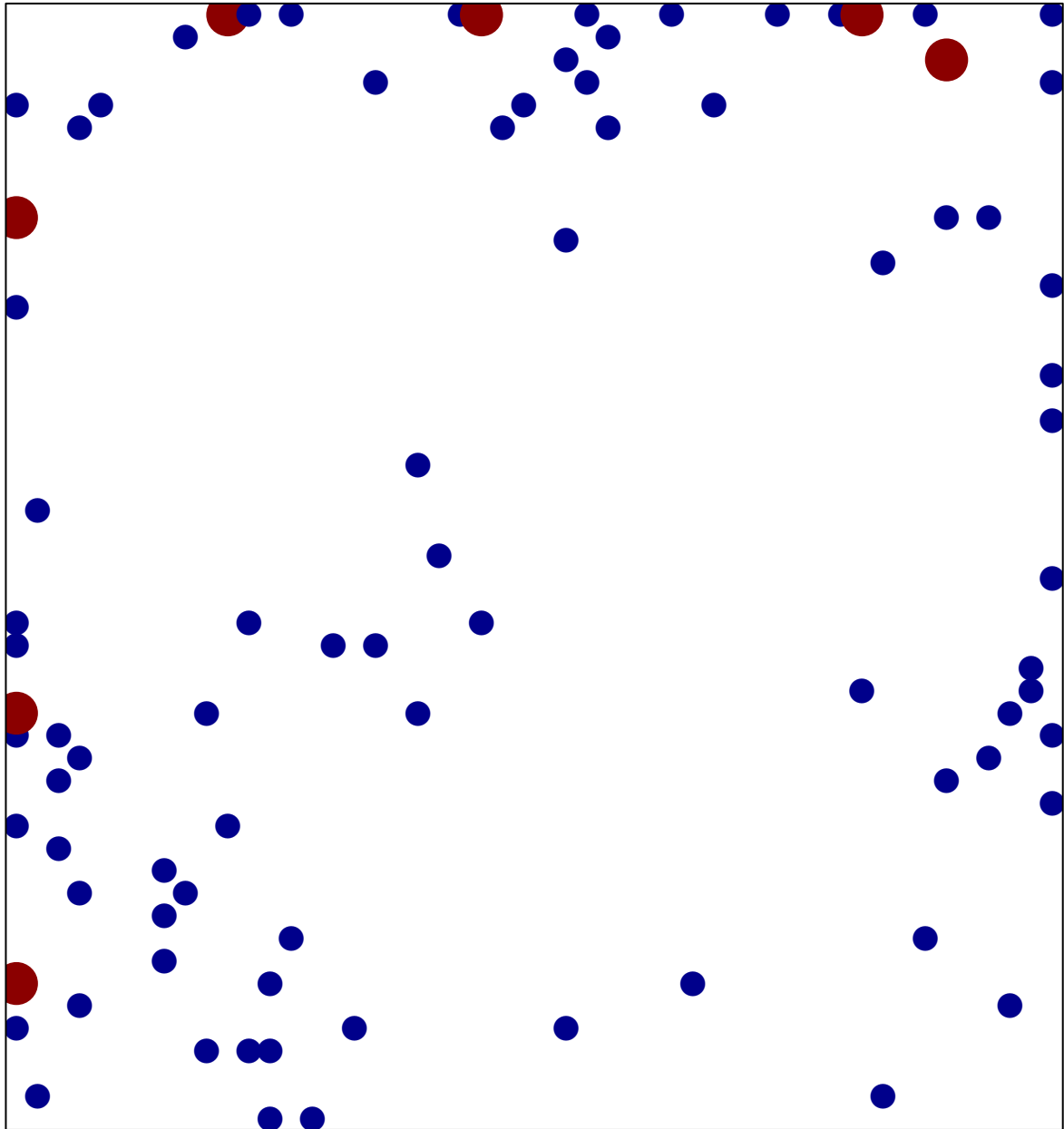
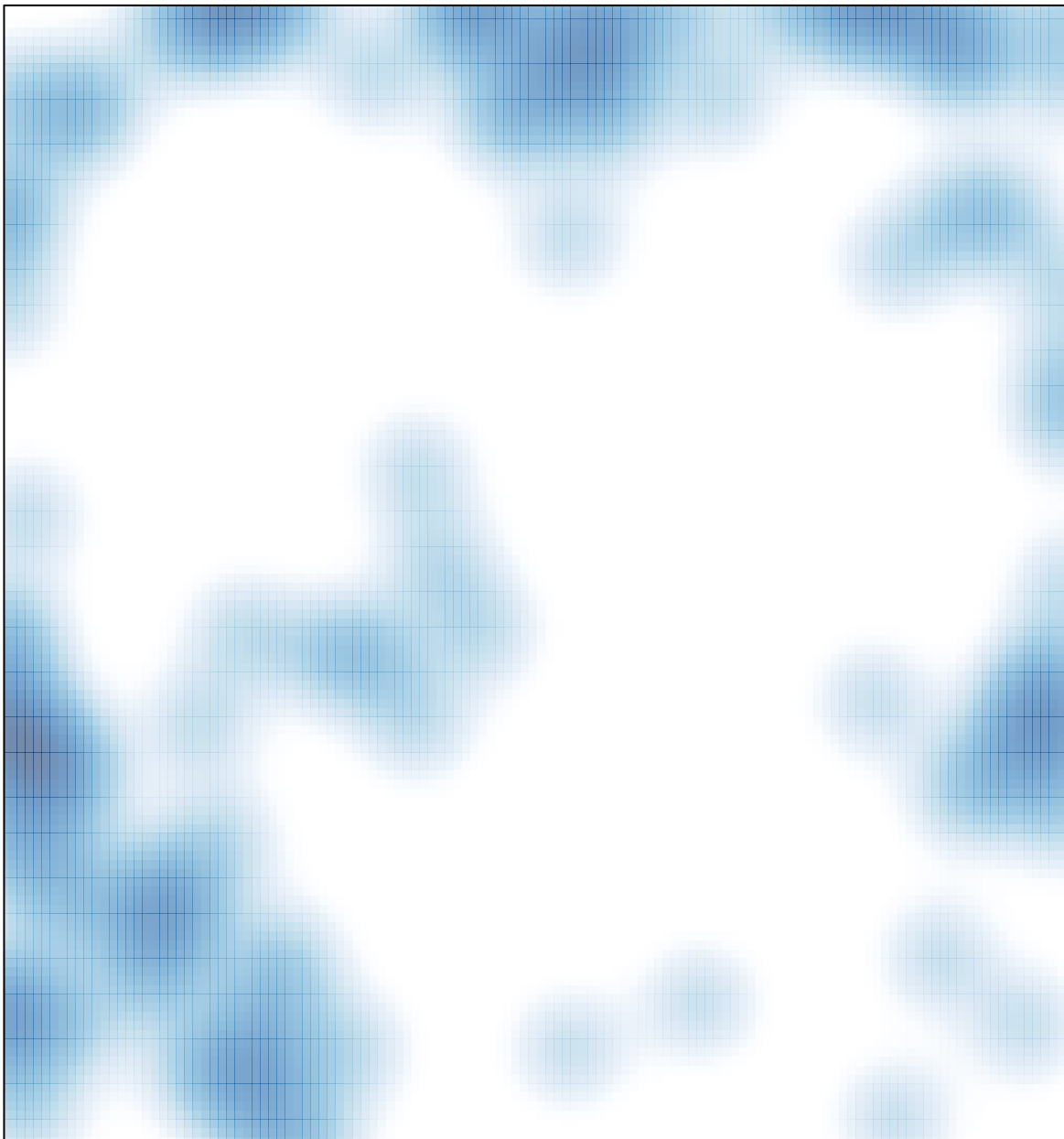


# hsa-miR-378



# features = 92  
chi-square p = 0.74

# hsa-miR-378



# features = 92 , max = 2