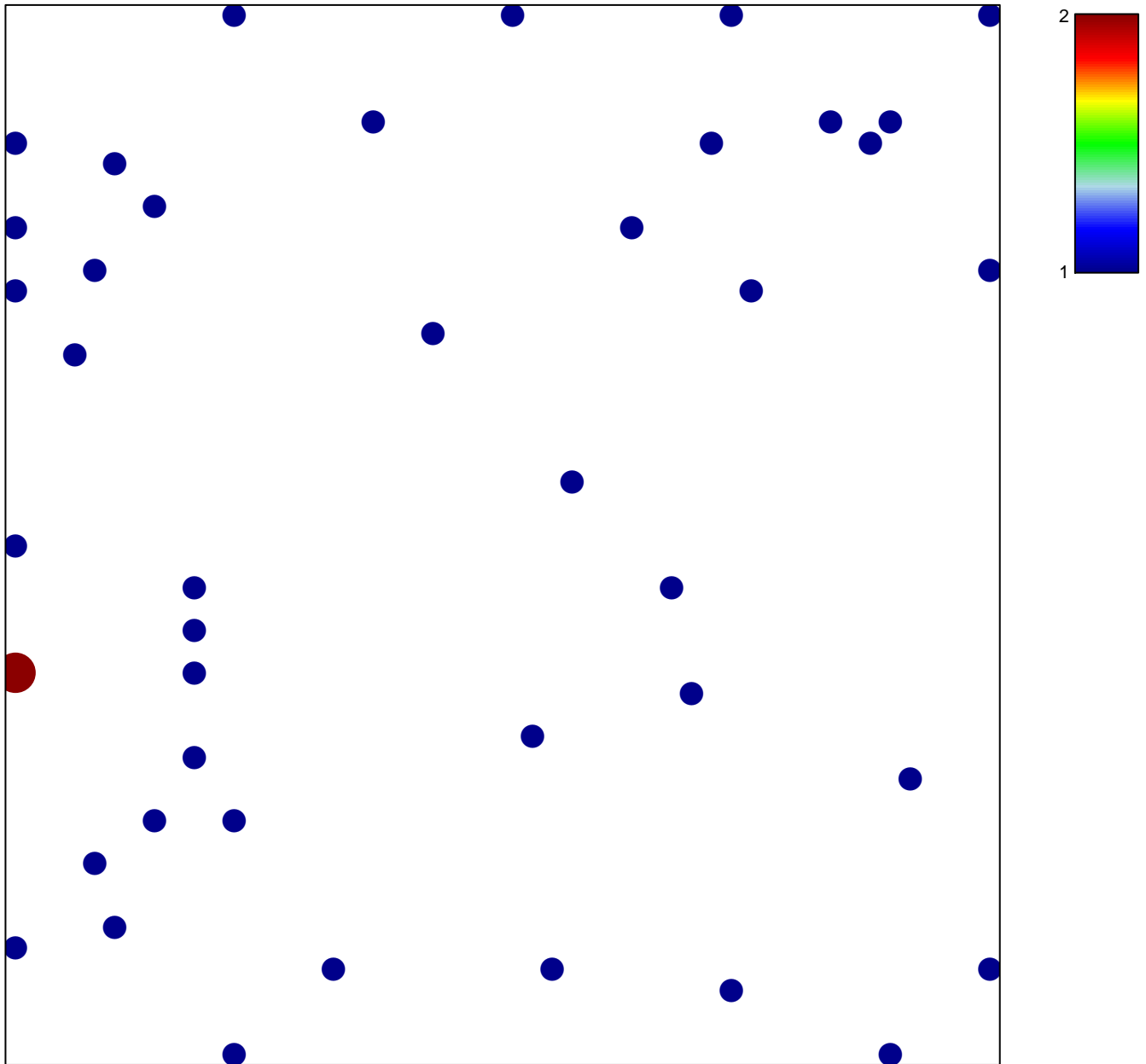
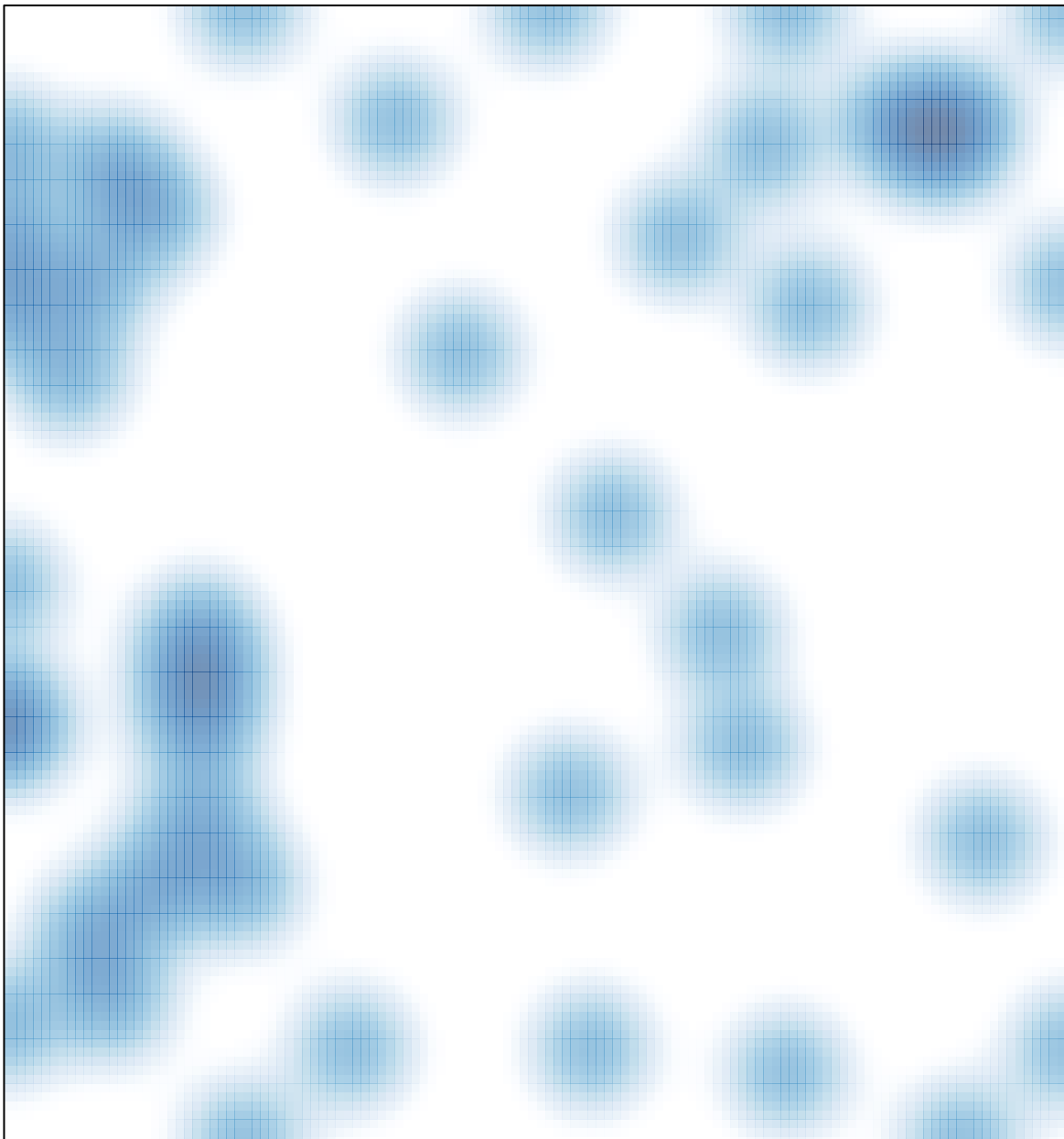


# hsa-miR-661



# features = 43  
chi-square p = 0.8

# hsa-miR-661



# features = 43 , max = 2