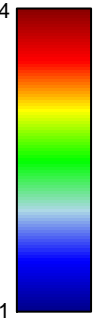
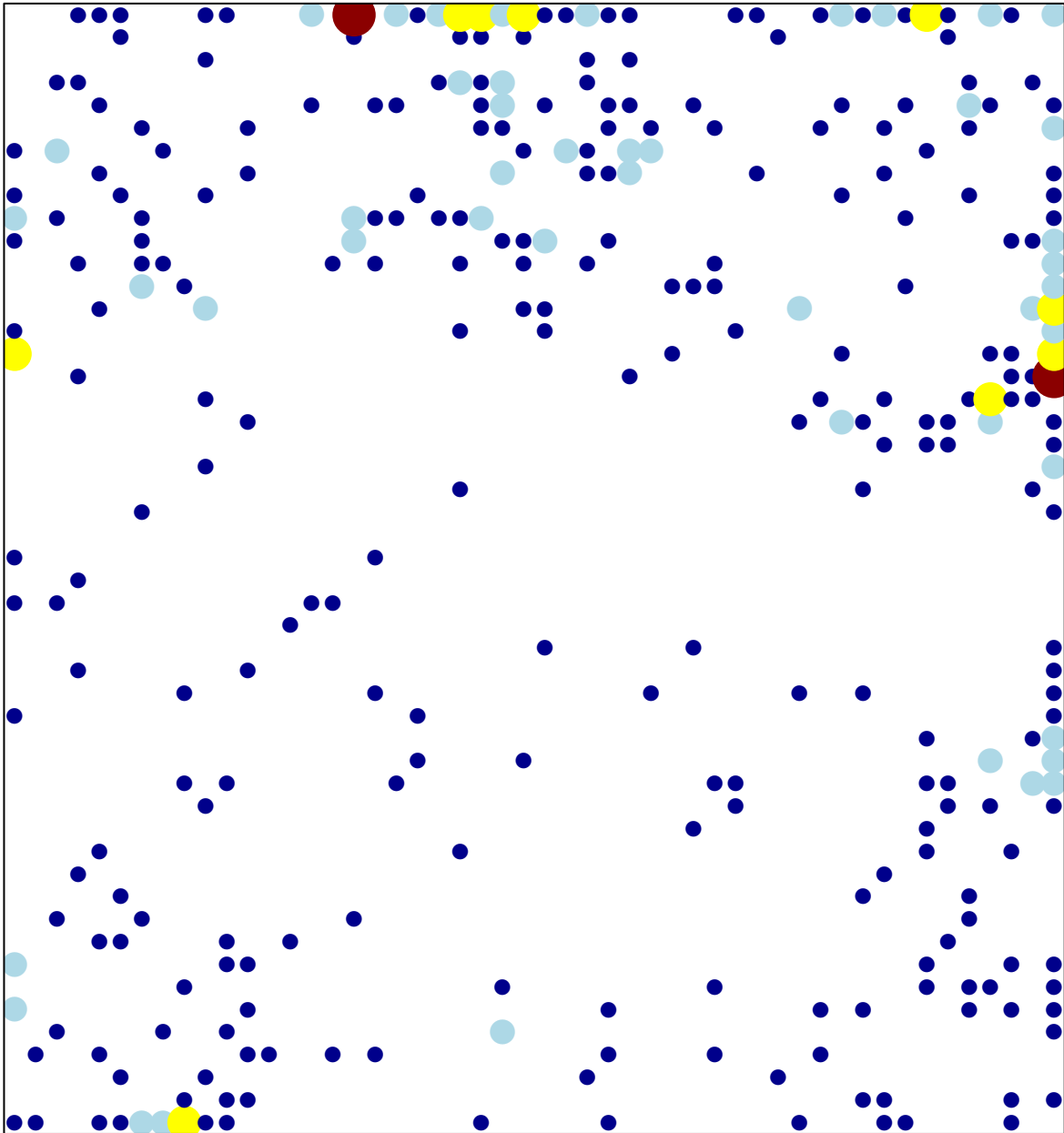
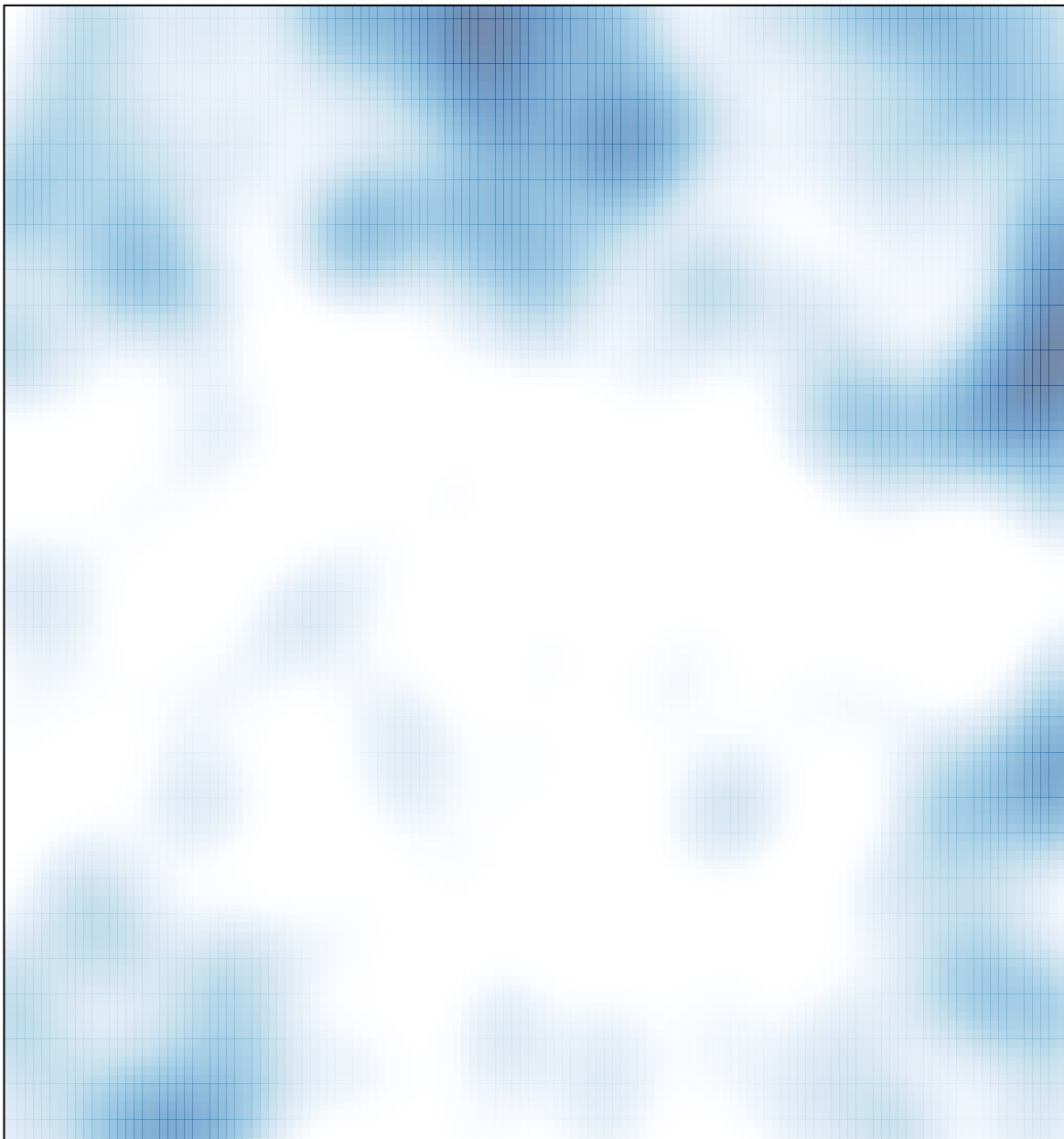


# hsa-miR-454



# features = 383  
chi-square p = 0.61

# hsa-miR-454



# features = 383 , max = 4