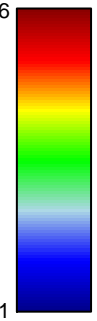
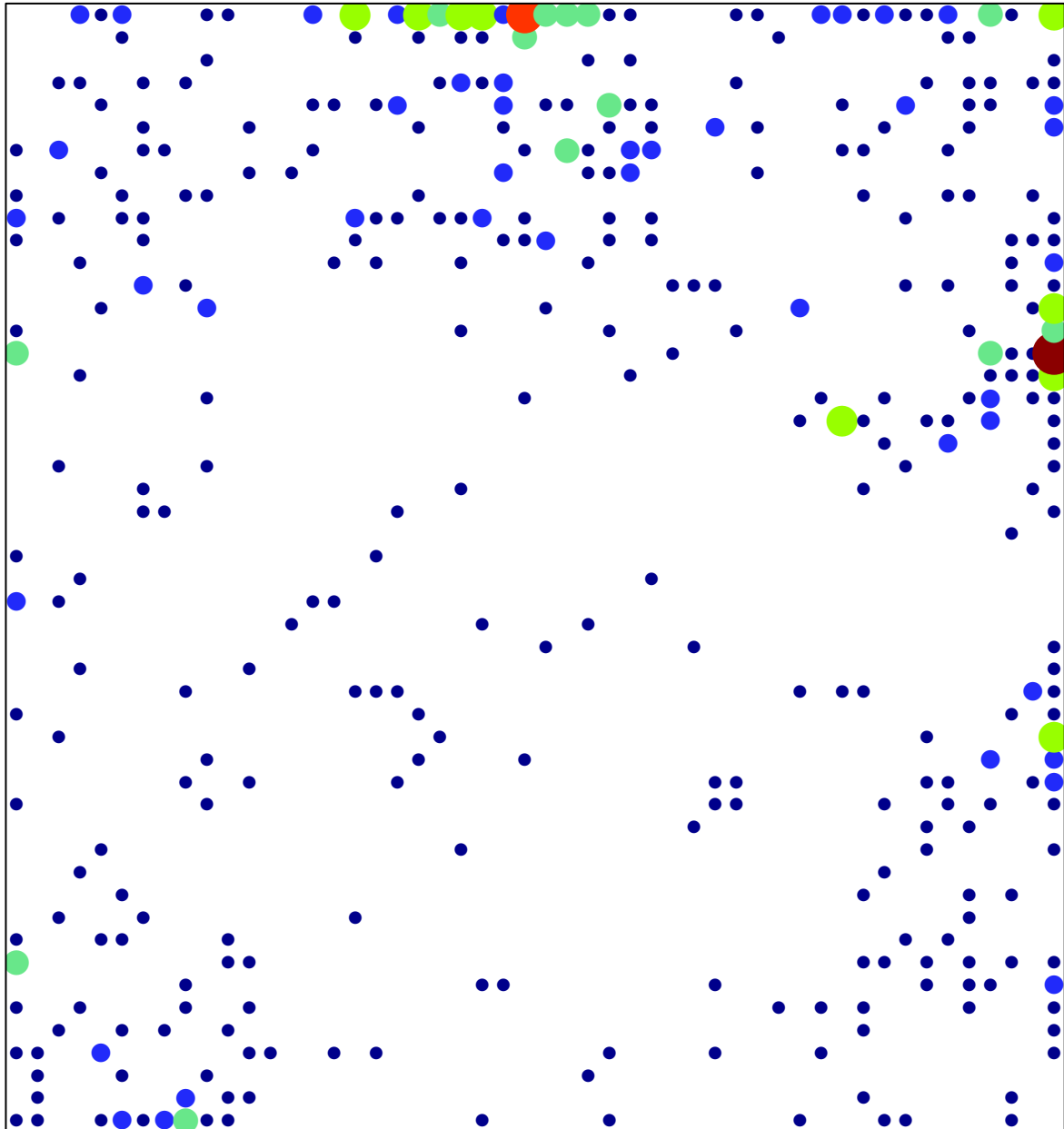
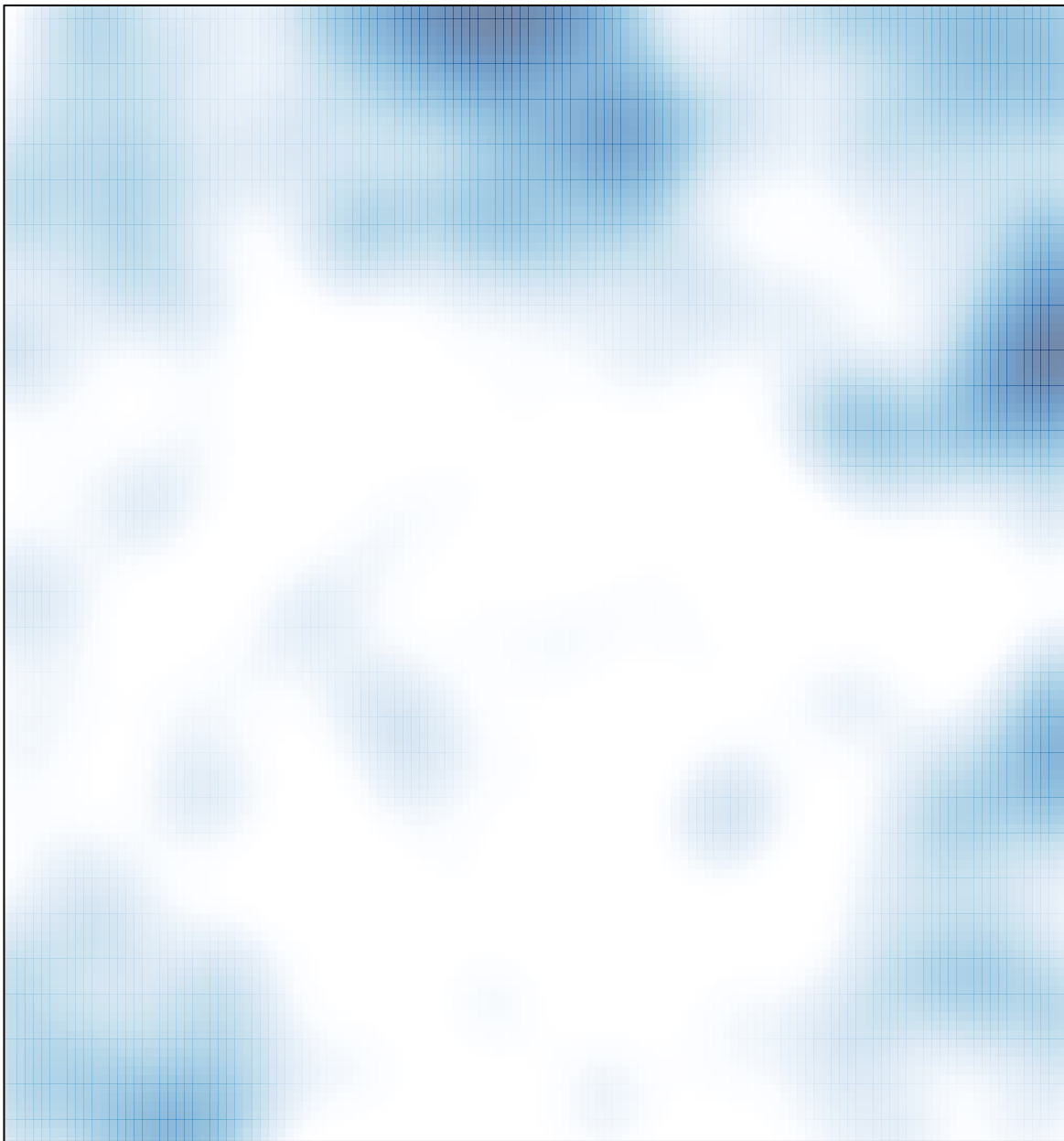


# hsa-miR-301a



# features = 464  
chi-square p = 0.56

# hsa-miR-301a



# features = 464 , max = 6