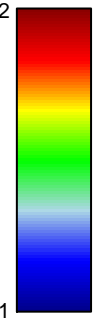
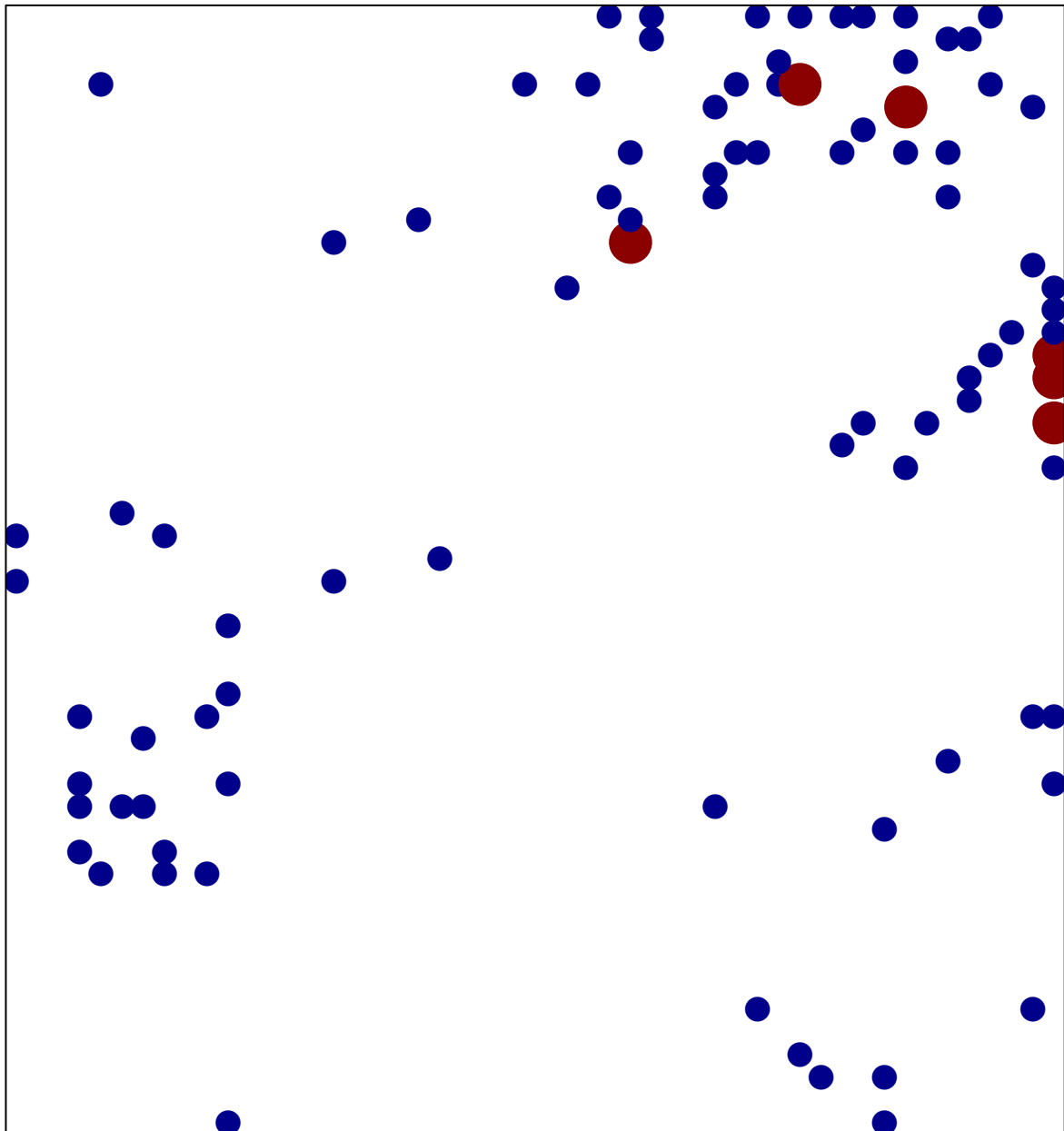
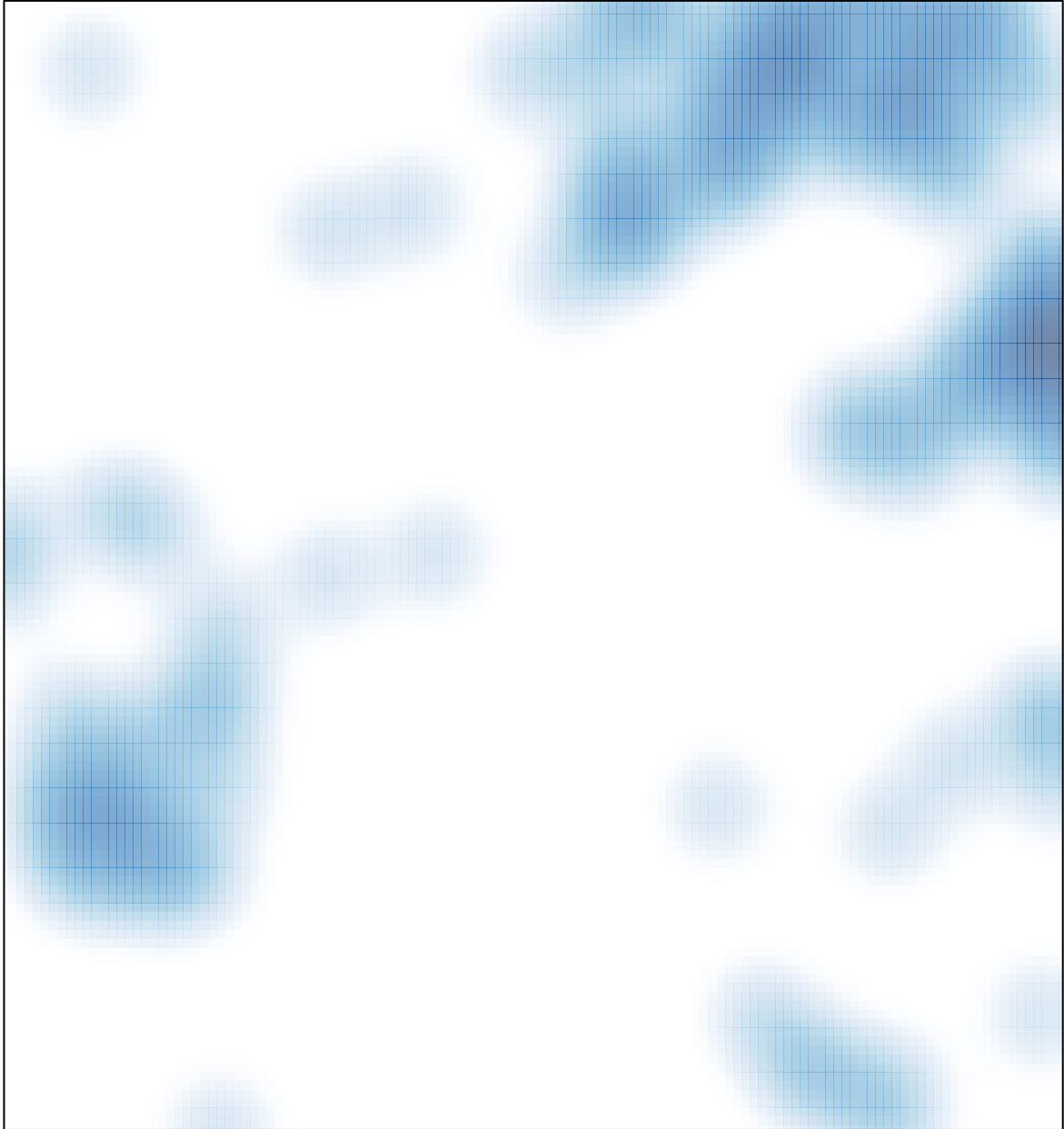


# spliceosomal complex



# features = 95  
chi-square p = 0.77

# spliceosomal complex



# features = 95 , max = 2