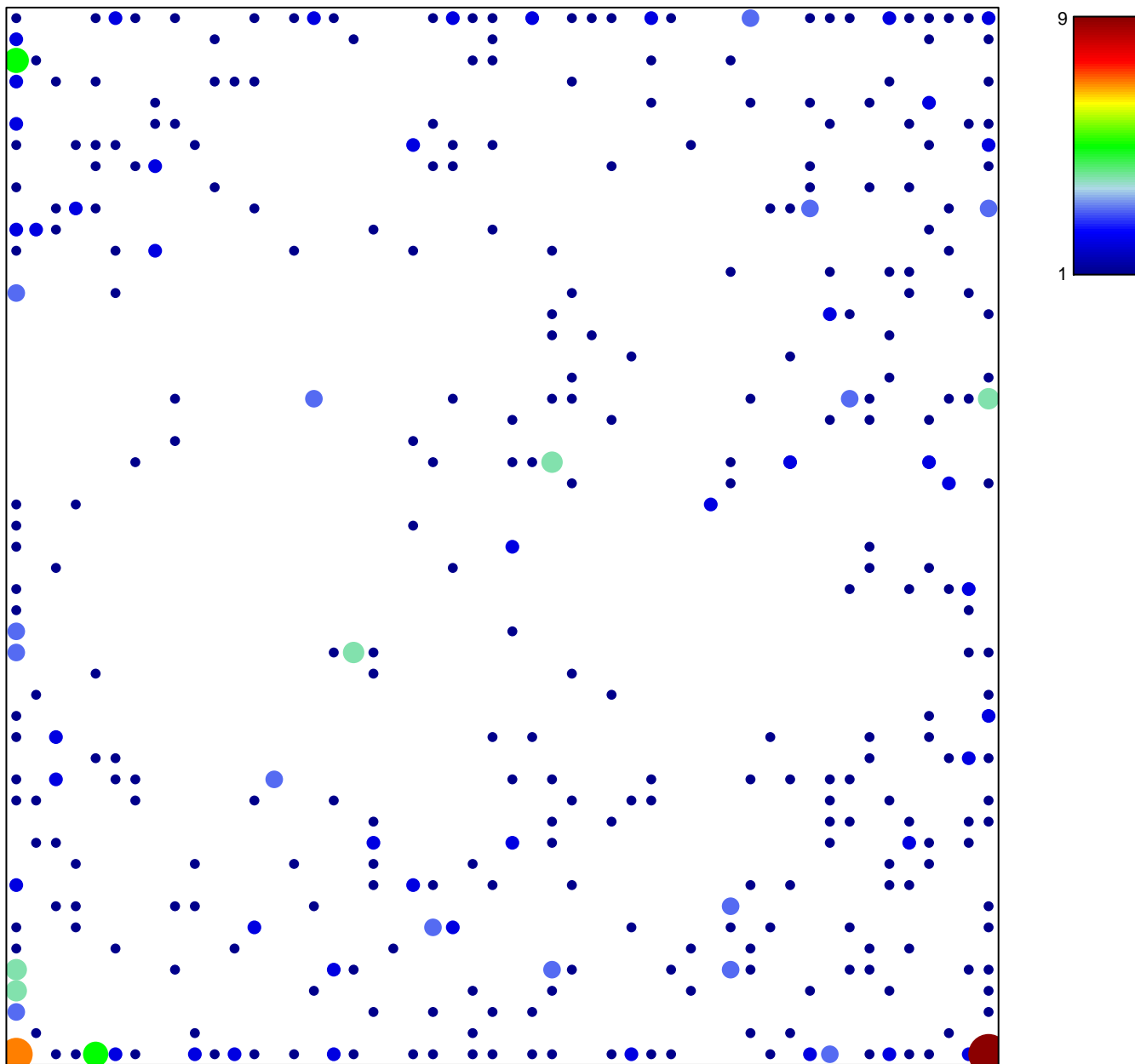
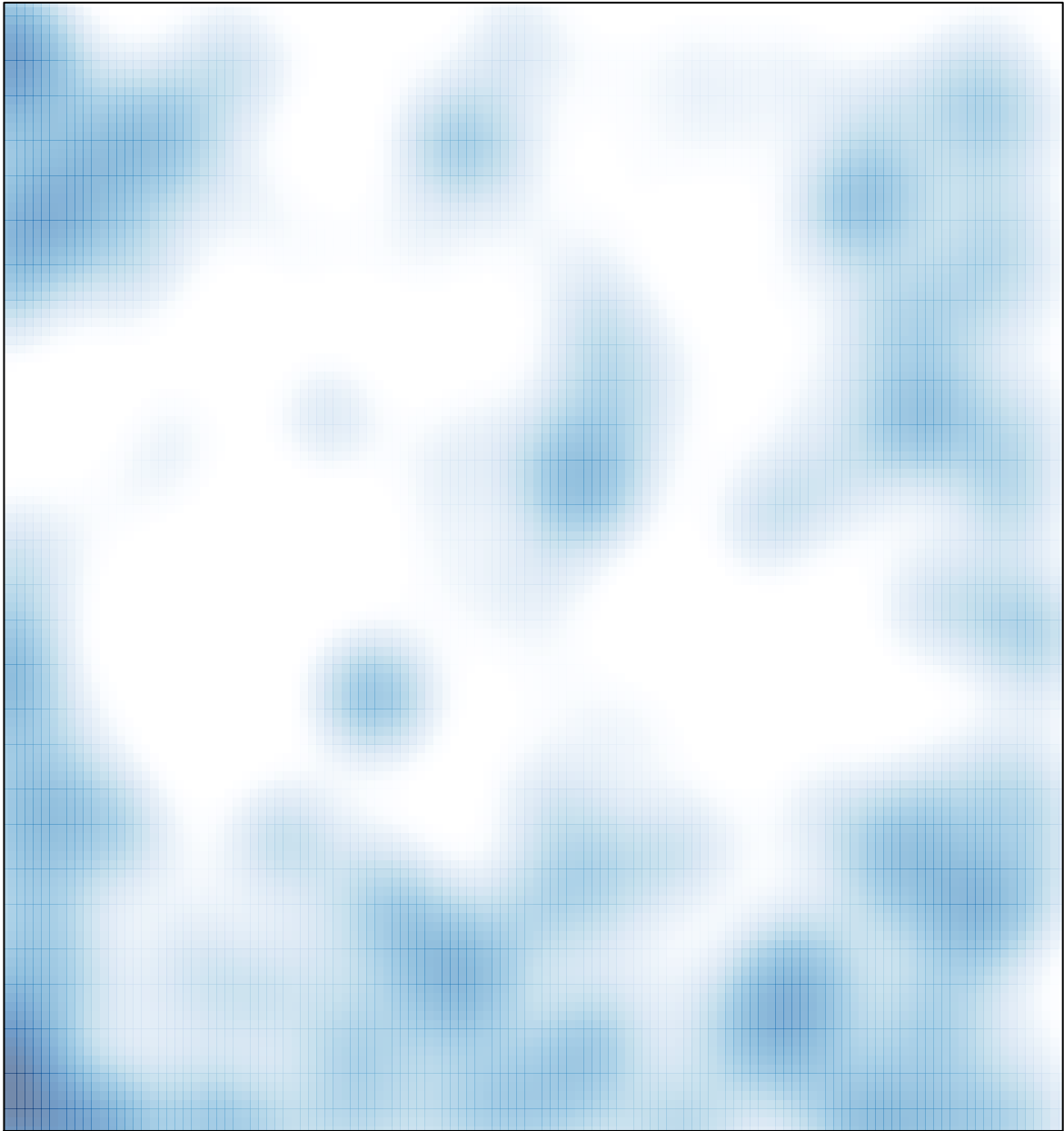


# GRAESSMANN\_APOPTOSIS\_BY\_SERUM\_DEPRIVATION\_UP



# features = 469  
chi-square p = 0.55

# AEISSMANN\_APOPTOSIS\_BY\_SERUM\_DEPRIVATION



# features = 469 , max = 9