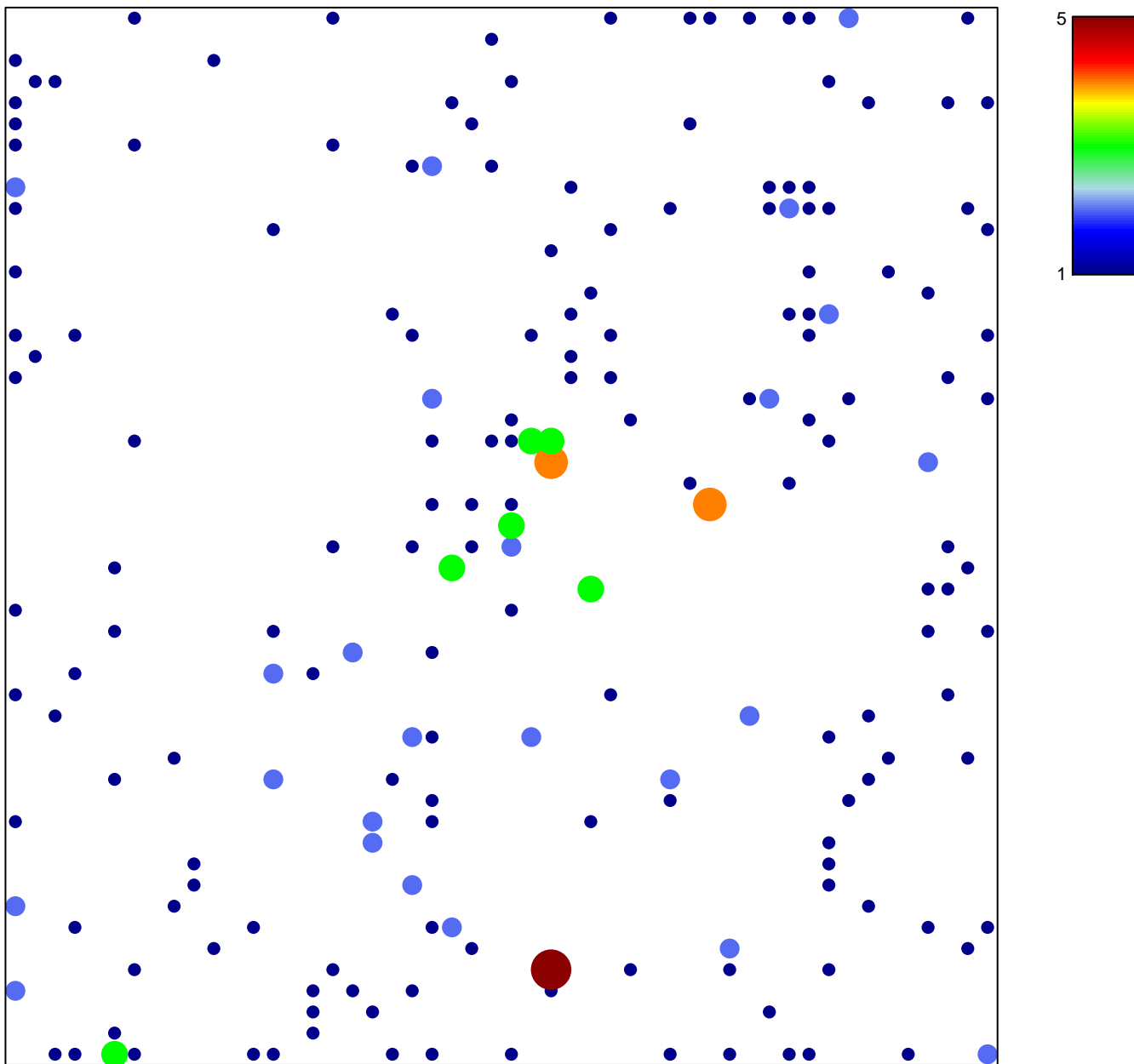
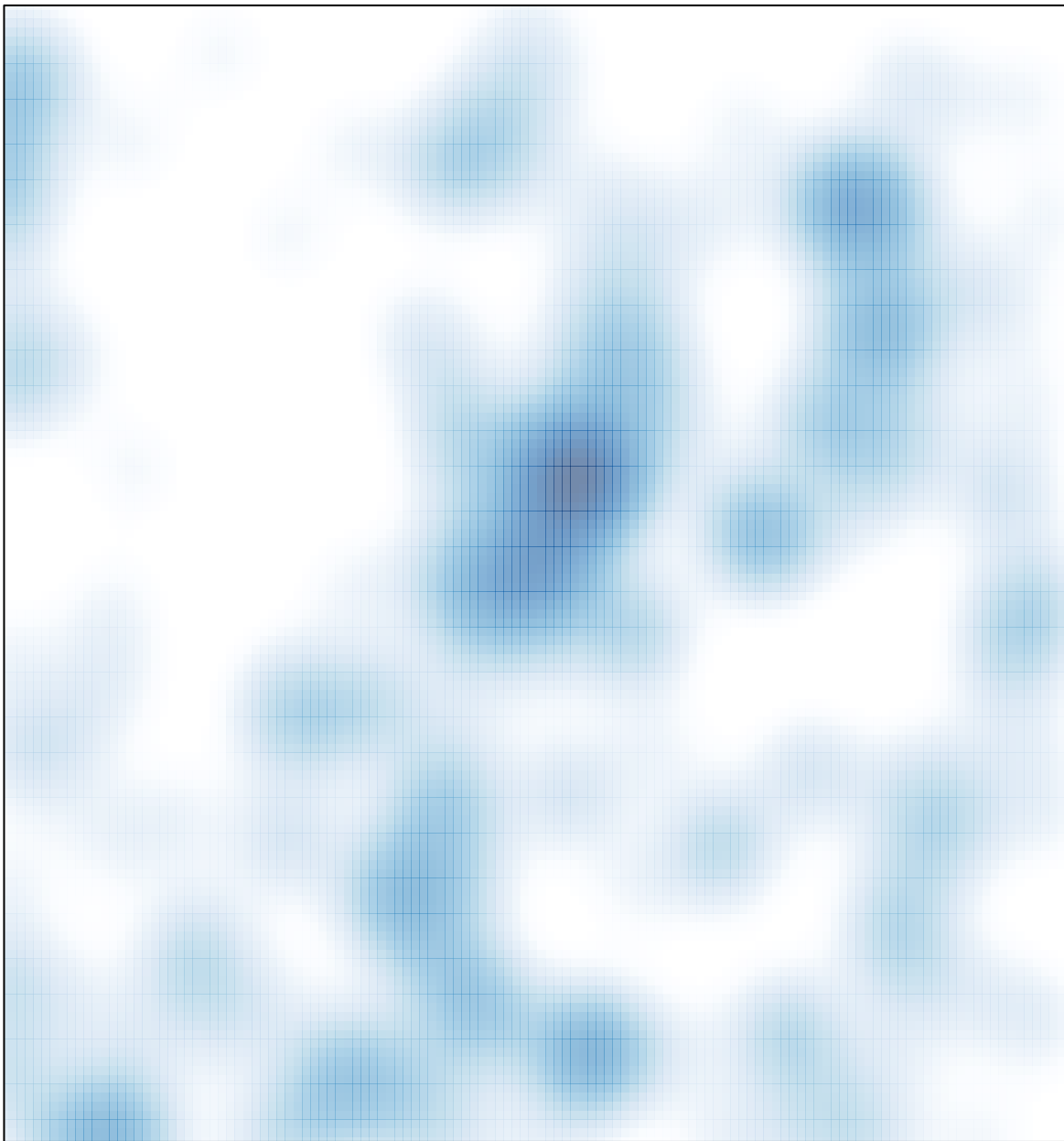


# SENGUPTA\_NASOPHARYNGEAL\_CARCINOMA\_DN



# features = 238  
chi-square p = 0.77

# SENGUPTA\_NASOPHARYNGEAL\_CARCINOMA\_DM



# features = 238 , max = 5