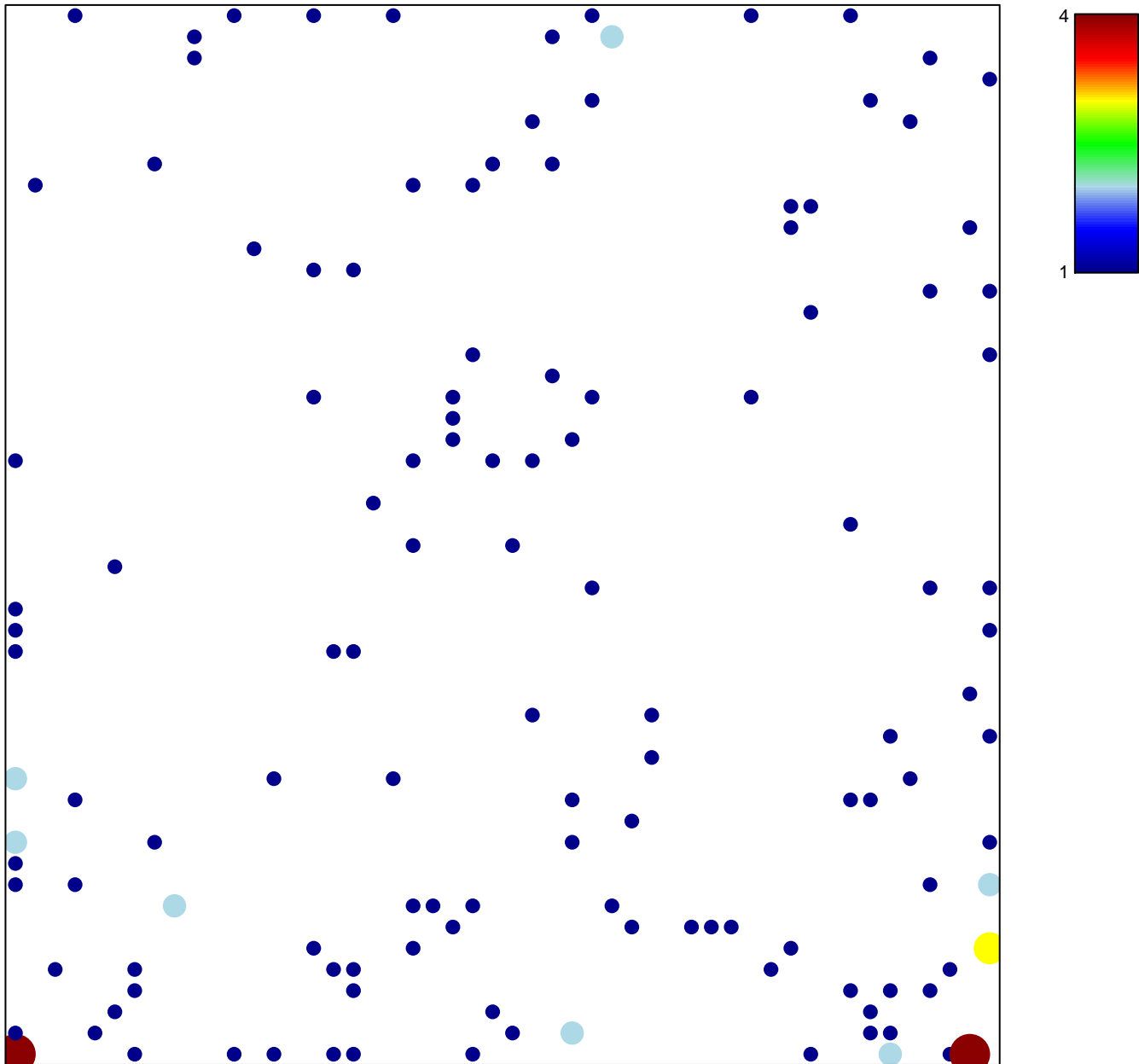
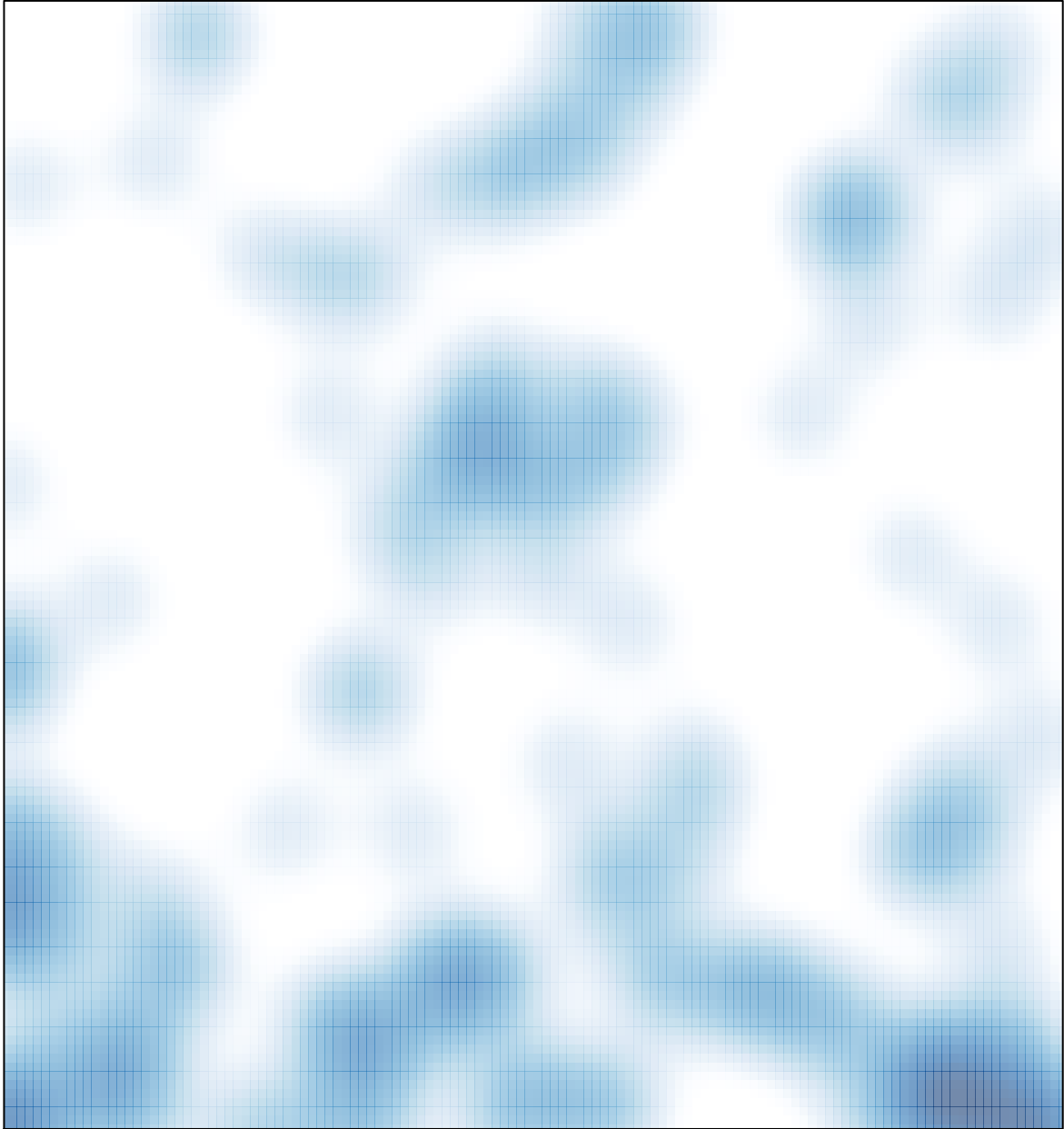


VERHAAK_GLIOMASTOMA_CLASSICAL



features = 145
chi-square p = 0.75

VERHAAK_GLIOMASTOMA_CLASSICAL



features = 145 , max = 4