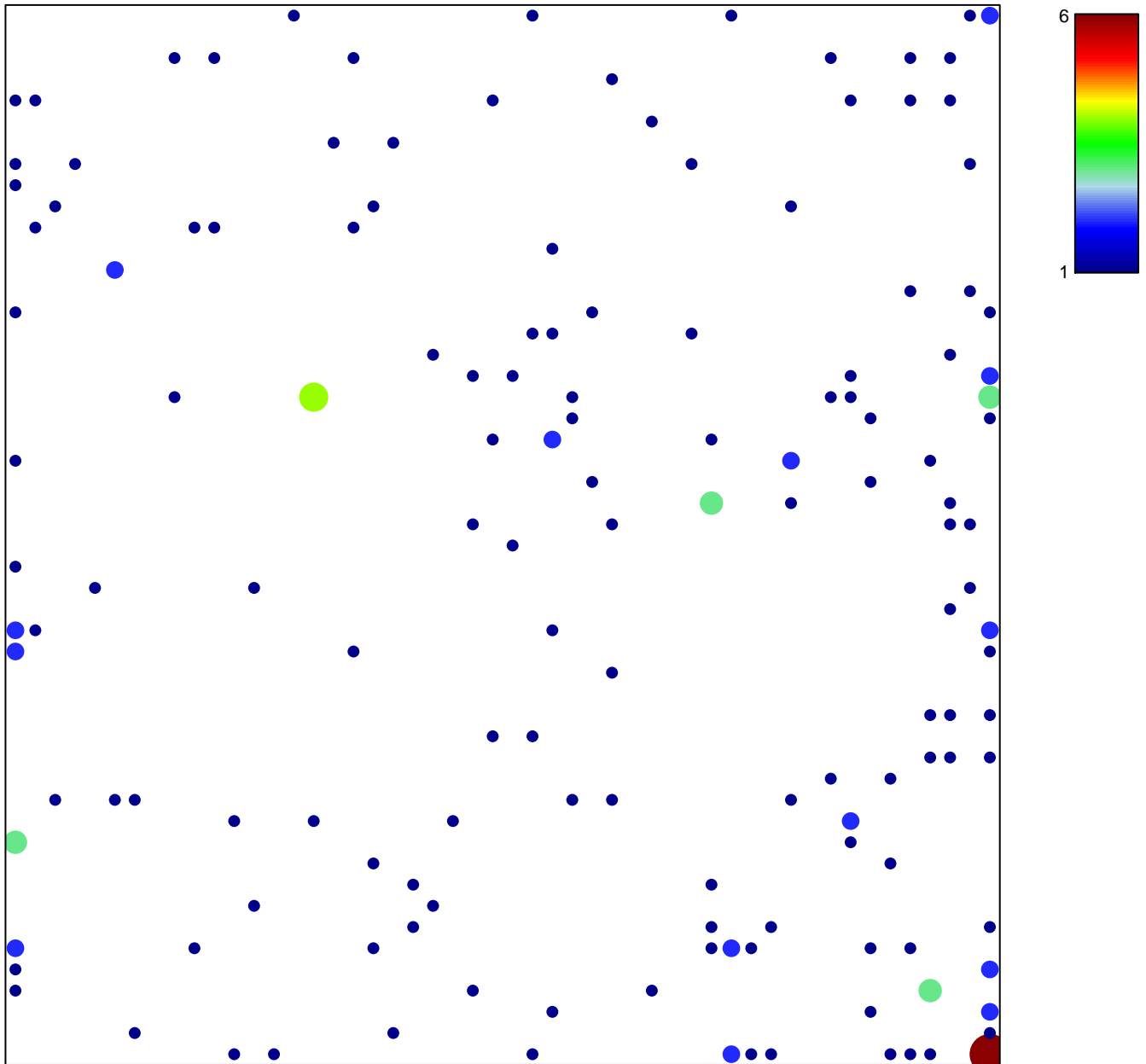
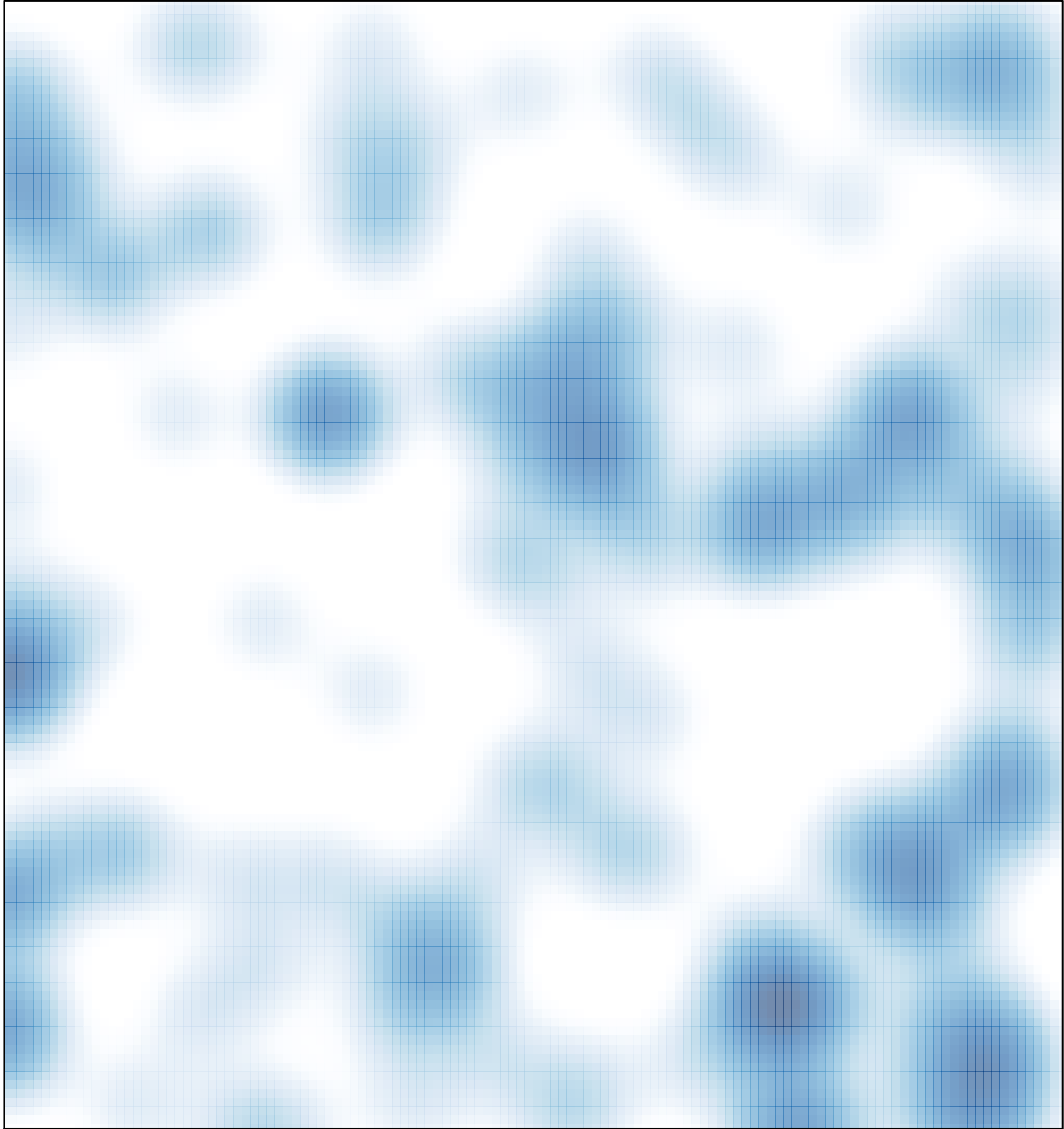


VERHAAK_GLIOMASTOMA_MESENCHYMAL



features = 179
chi-square p = 0.71

VERHAAK_GLIOLASTOMA_MESENCHYMAL



features = 179 , max = 6