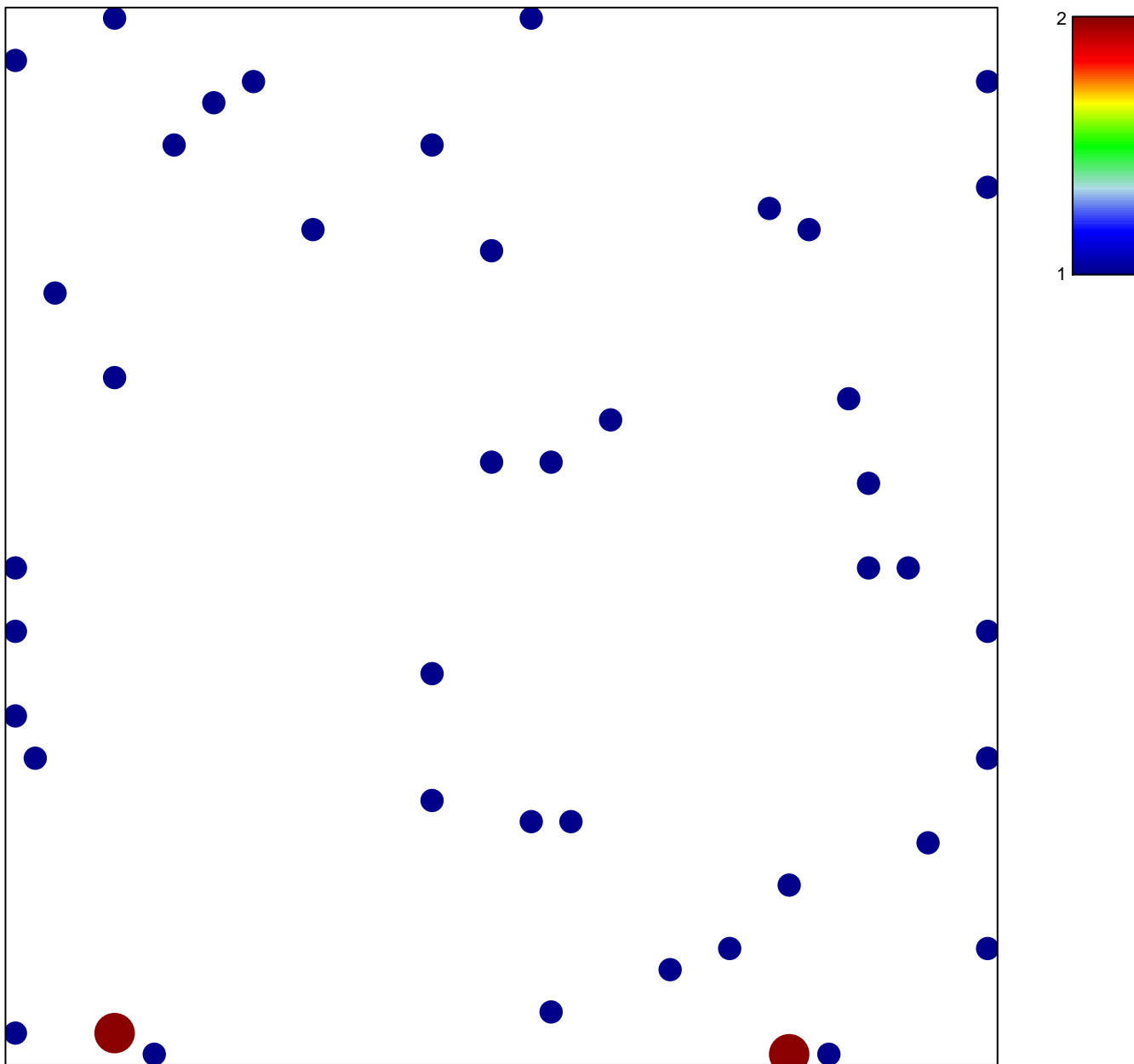
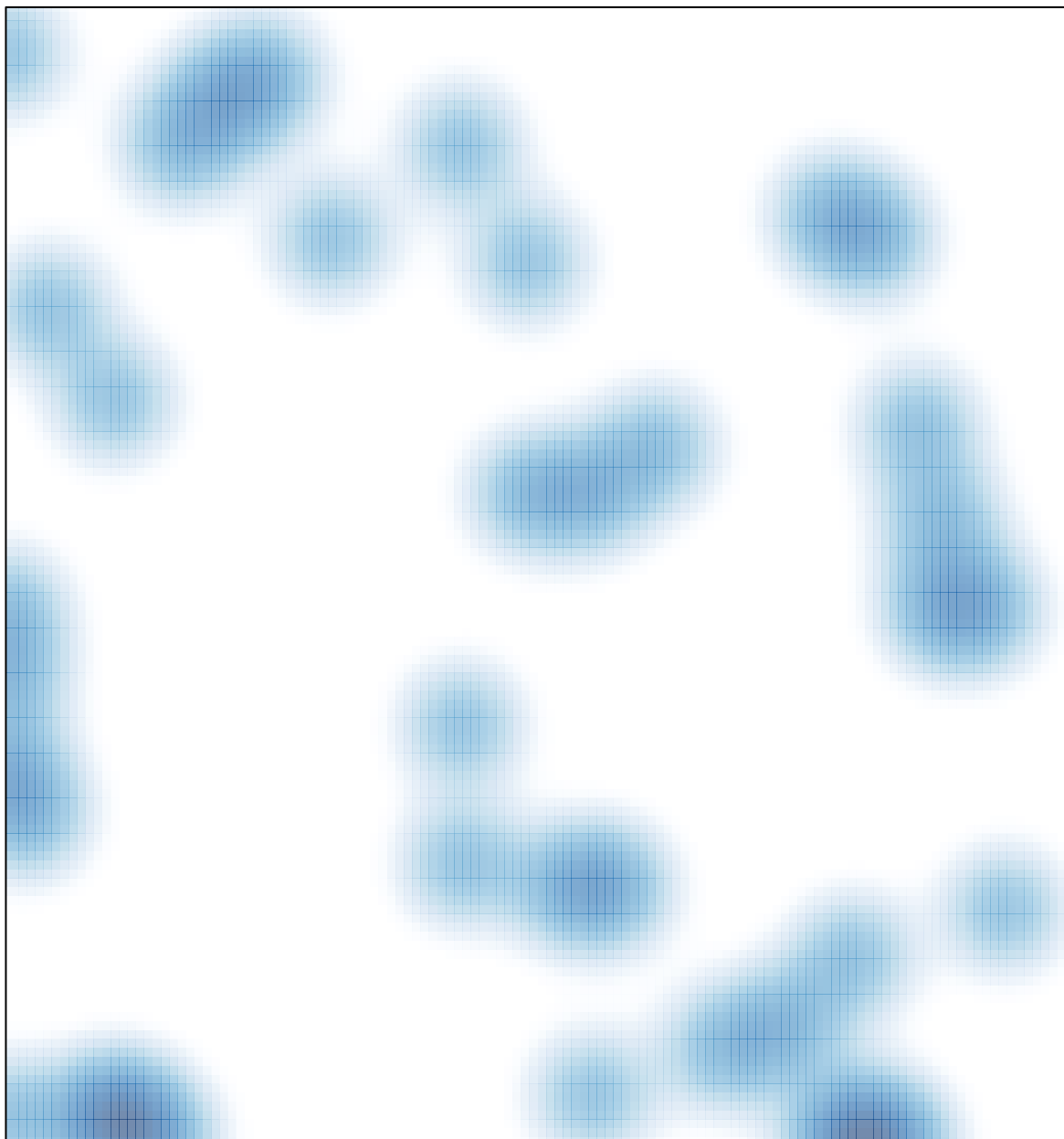


ROVERSI_GLIOMA_COPY_NUMBER_DN



features = 45
chi-square p = 0.82

ROVERSI_GLIOMA_COPY_NUMBER_DN



features = 45 , max = 2