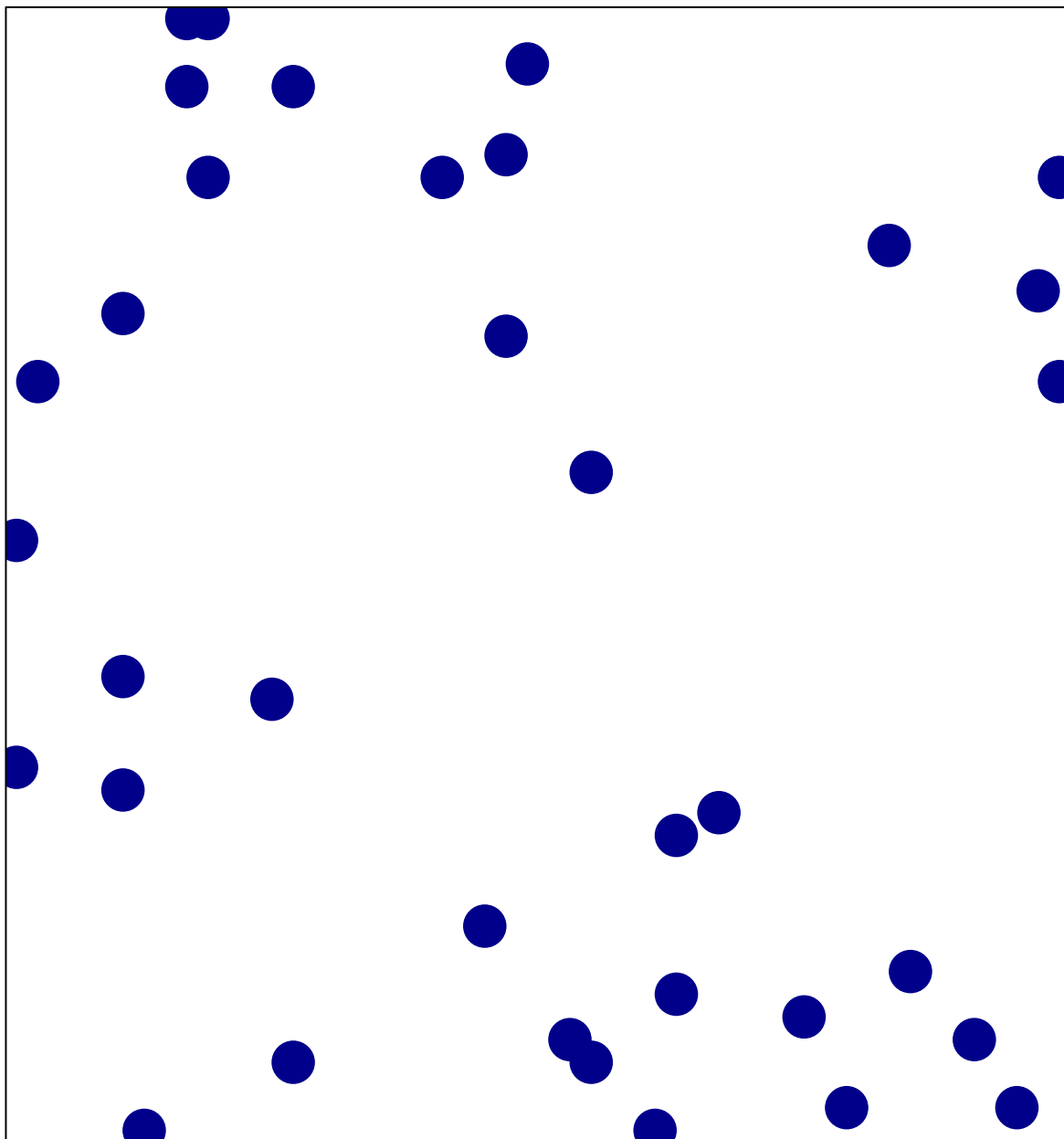
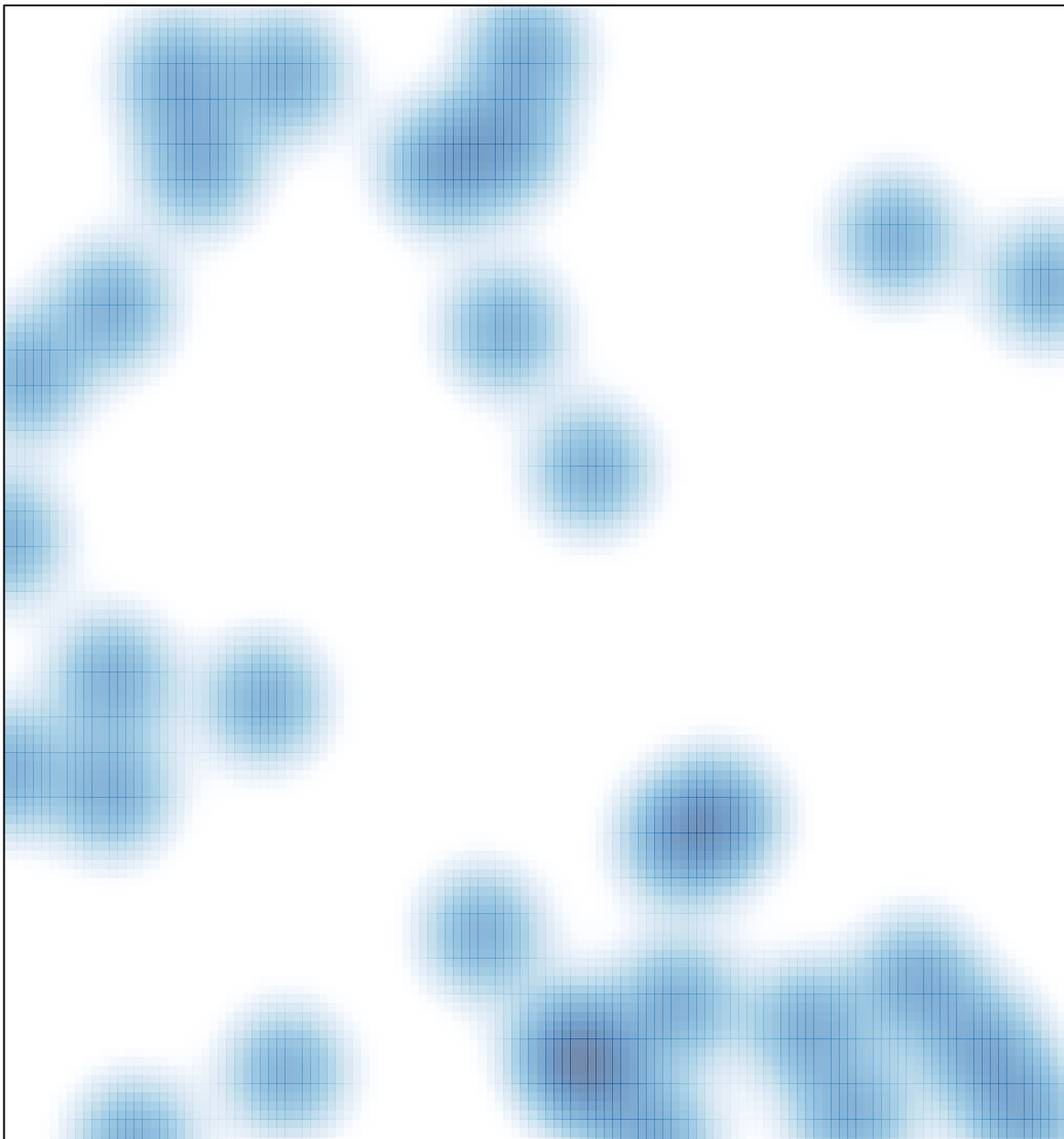


ROVERSI_GLIOMA_LOH_REGIONS



features = 35
chi-square p = 0.85

ROVERSI_GLIOMA_LOH_REGIONS



features = 35 , max = 1