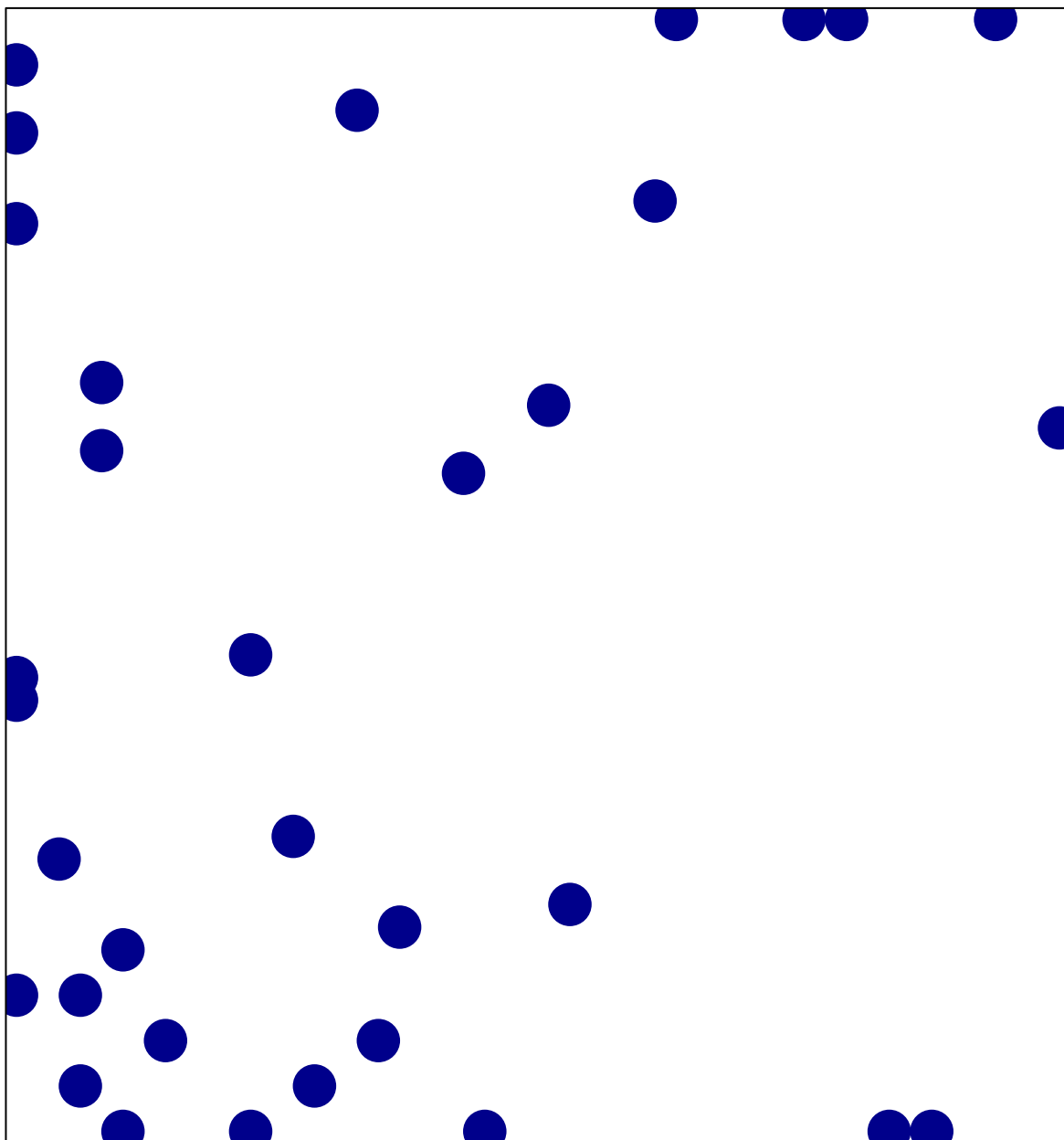
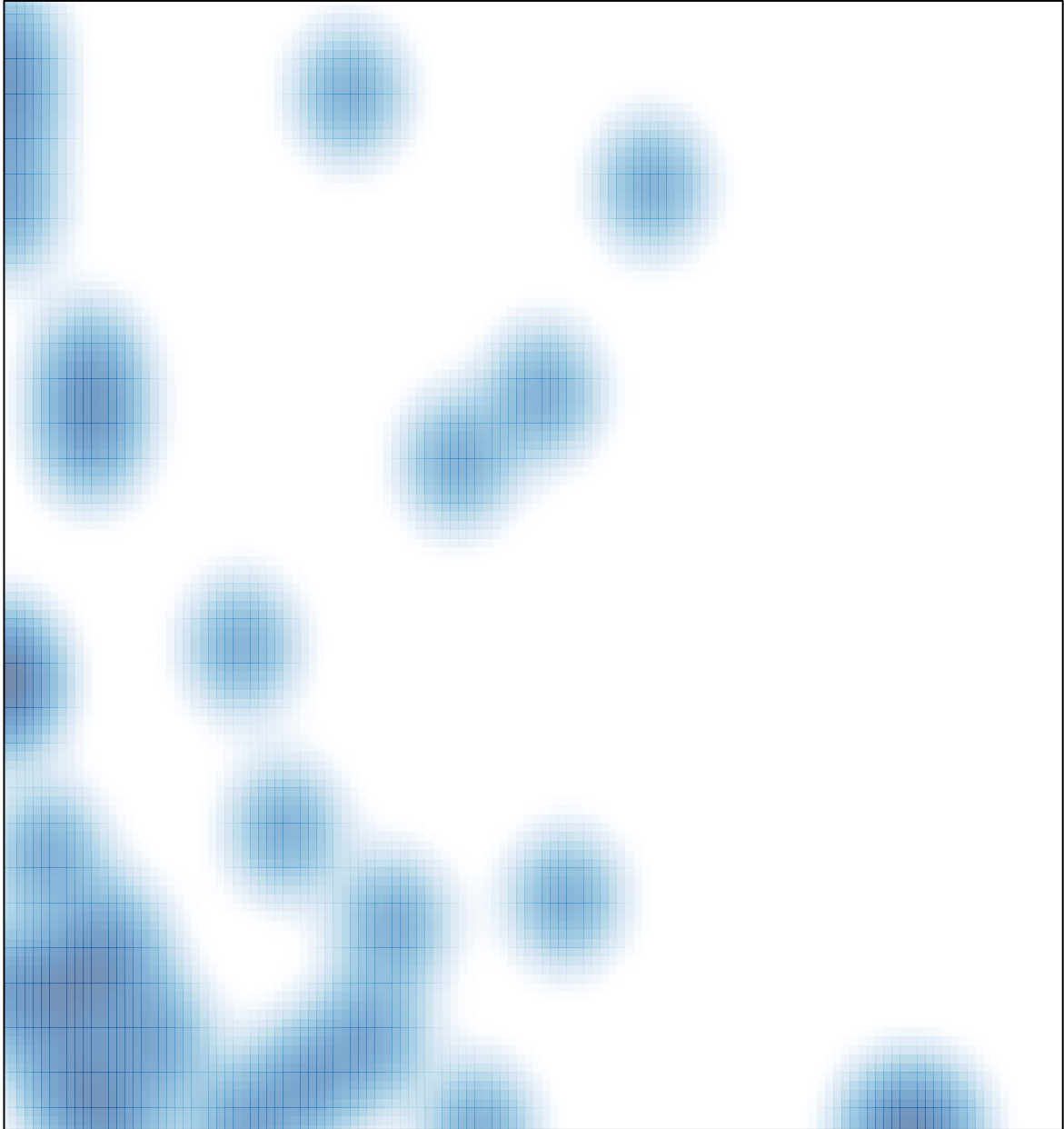


NUTT_GBM_VS_AO_GLIOMA_DN



features = 33
chi-square p = 0.81

NUTT_GBM_VS_AO_GLIOMA_DN



features = 33 , max = 1