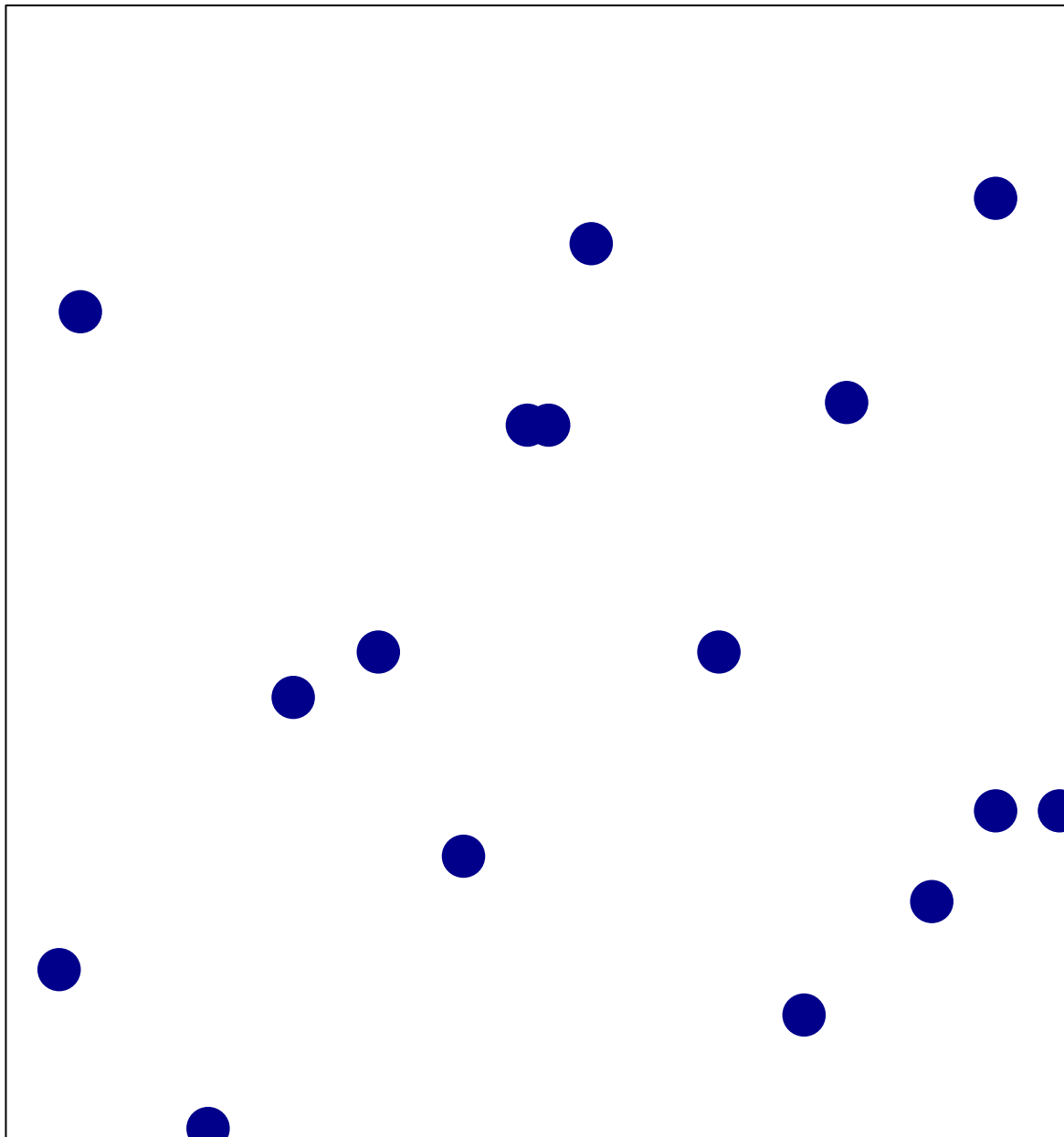
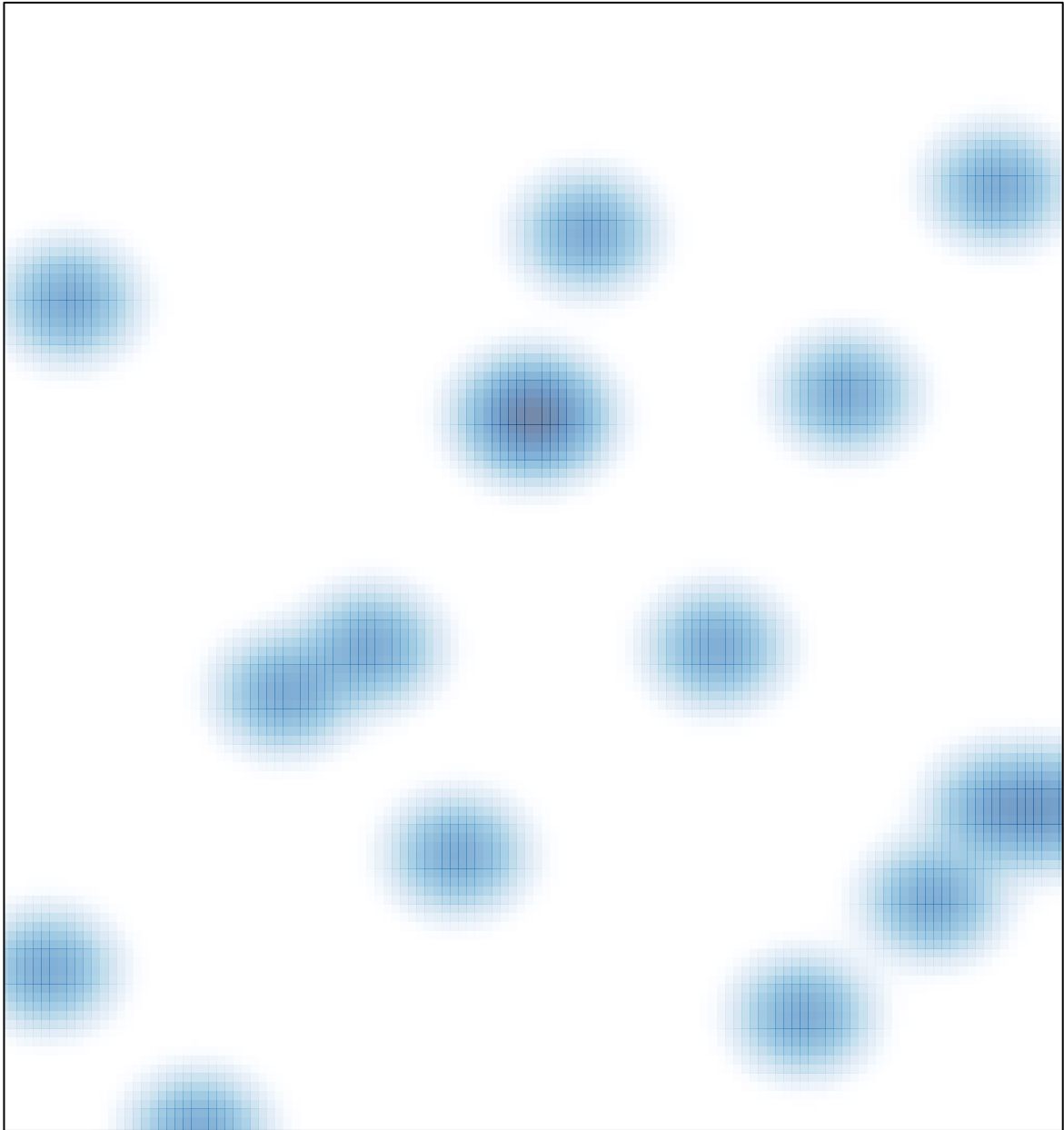


NUTT_GBM_VS_AO_GLIOMA_DN



features = 16
chi-square p = 0.85

NUTT_GBM_VS_AO_GLIOMA_DN



features = 16 , max = 1