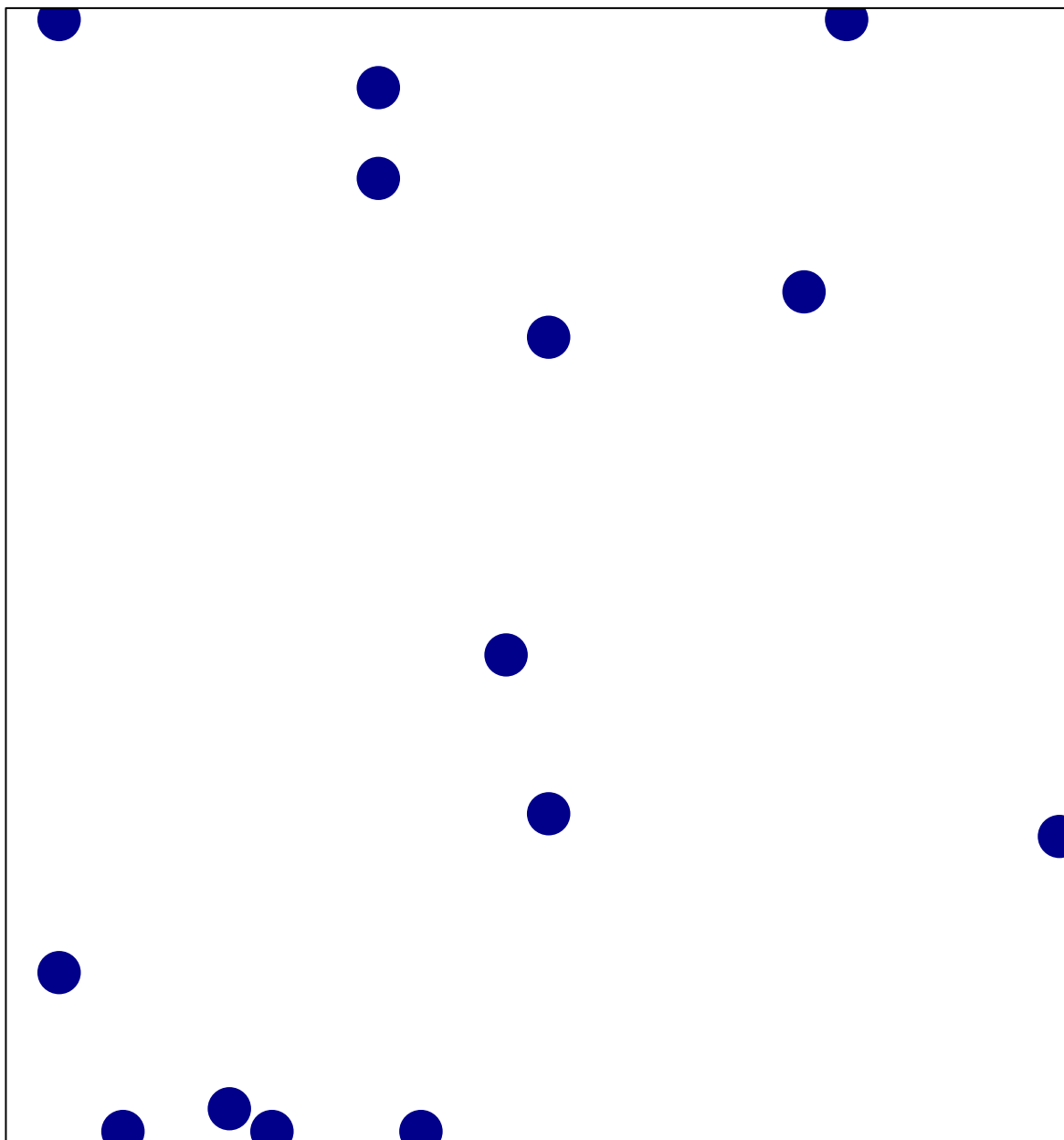


ROVERSI_GLIOMA_LOH_REGIONS



features = 14
chi-square p = 0.83

ROVERSI_GLIOMA_LOH_REGIONS



features = 14 , max = 1