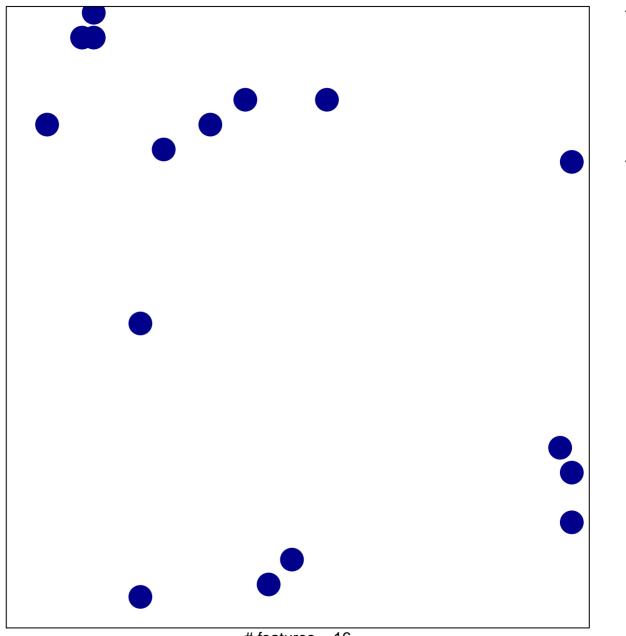
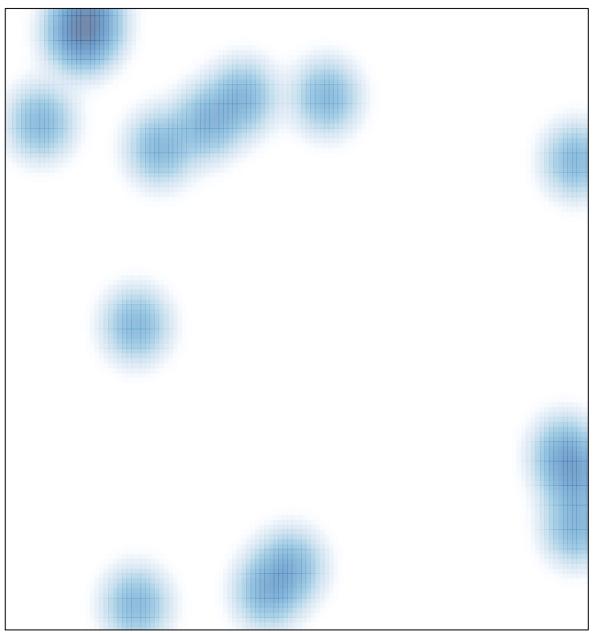
REACTOME_PEROXISOMAL_LIPID_METABOLISM



features = 16 chi-square p = 0.83

REACTOME_PEROXISOMAL_LIPID_METABOLISM



features = 16, max = 1