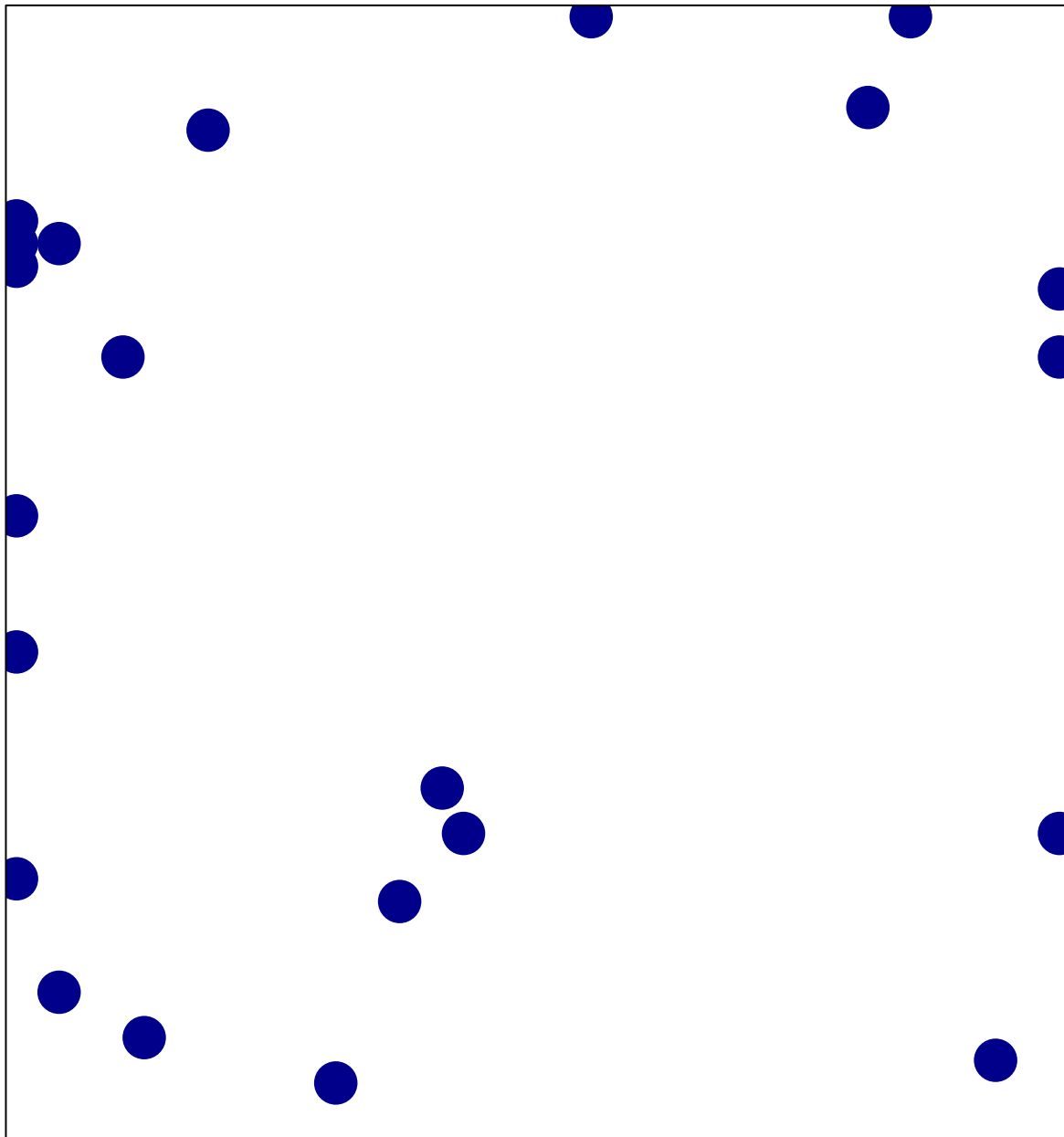
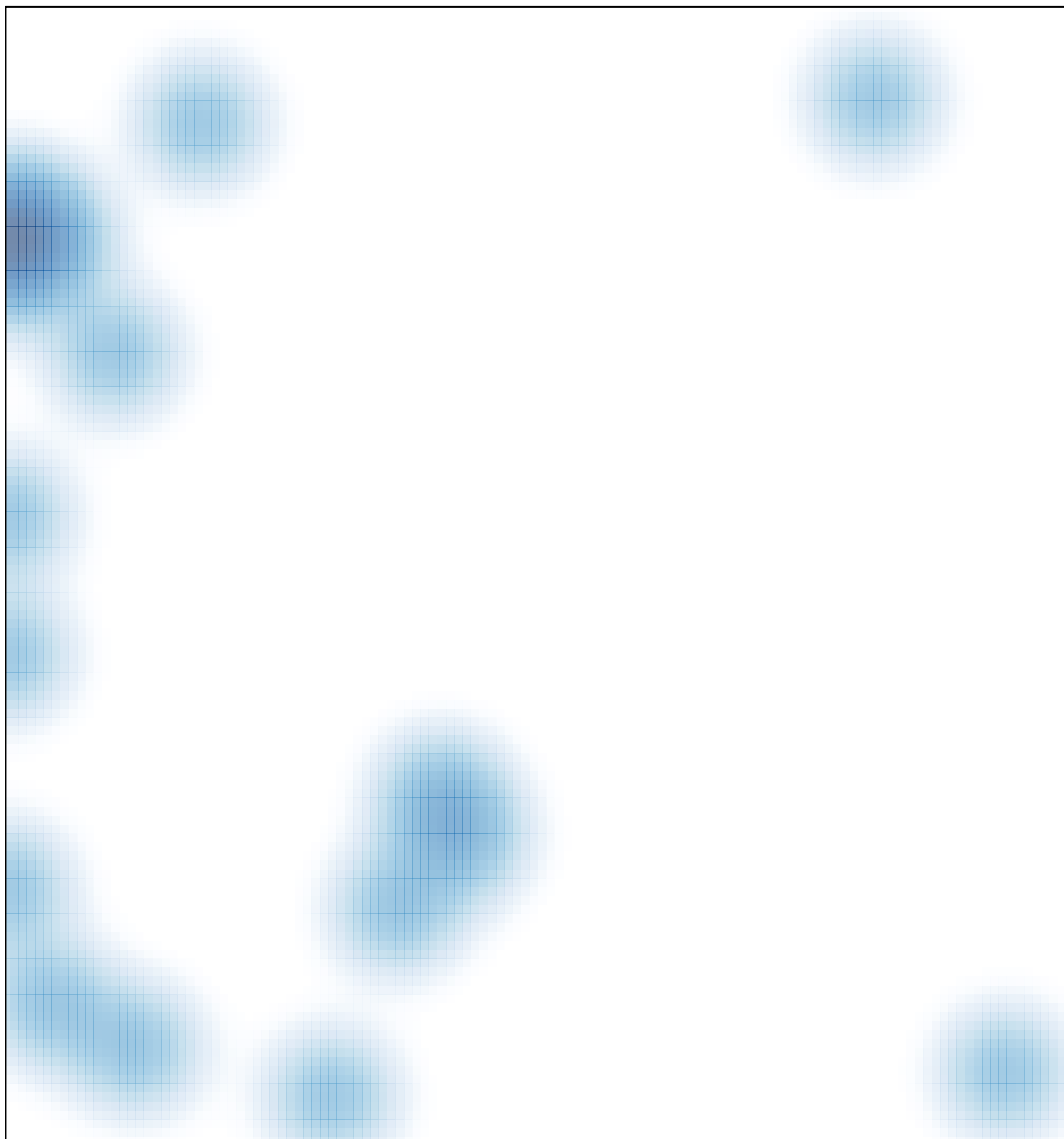


REACTOME_SULFUR_AMINO_ACID_METABOLISM



features = 22
chi-square p = 0.8

REACTOME_SULFUR_AMINO_ACID_METABOLISM



features = 22 , max = 1