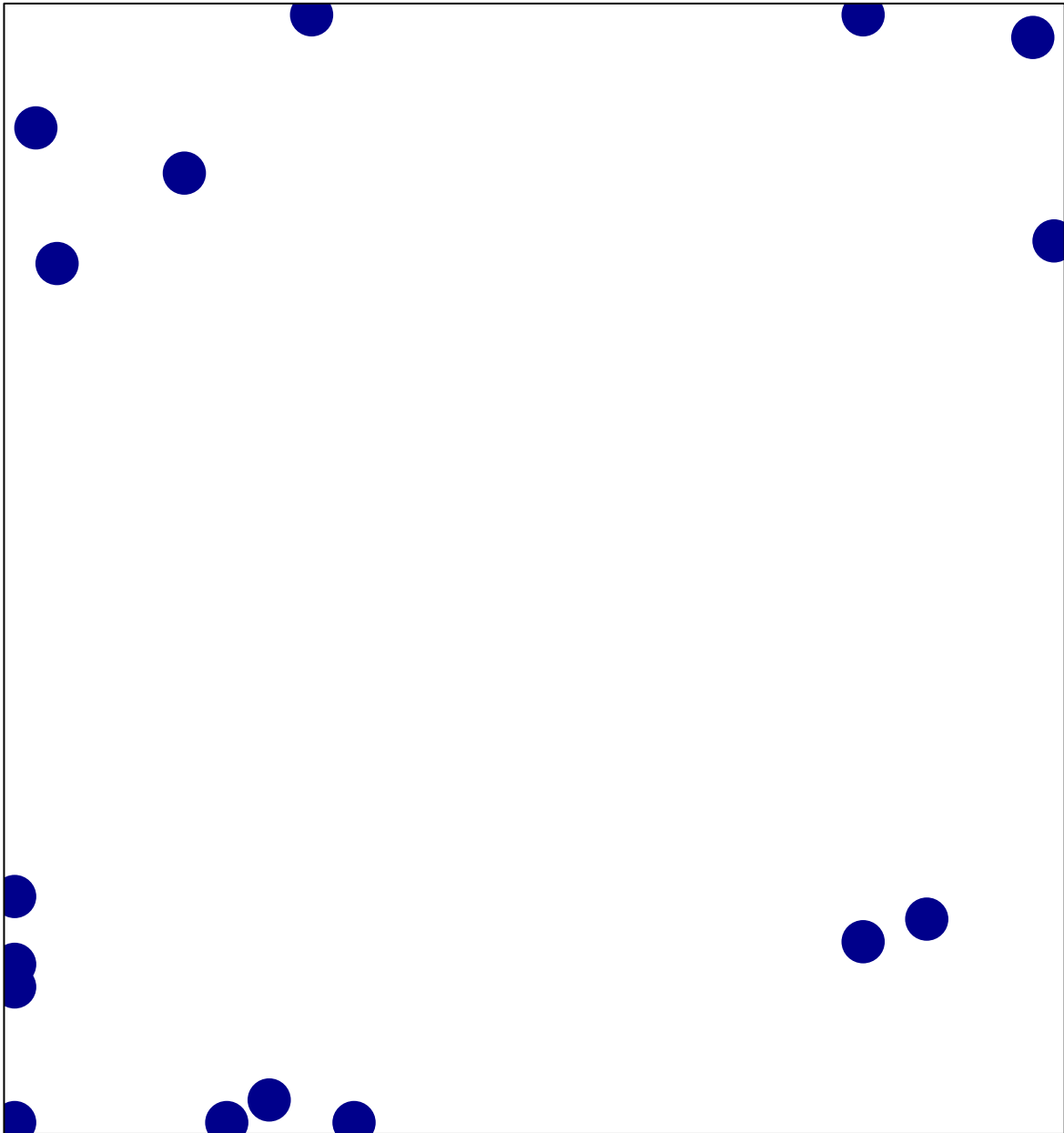
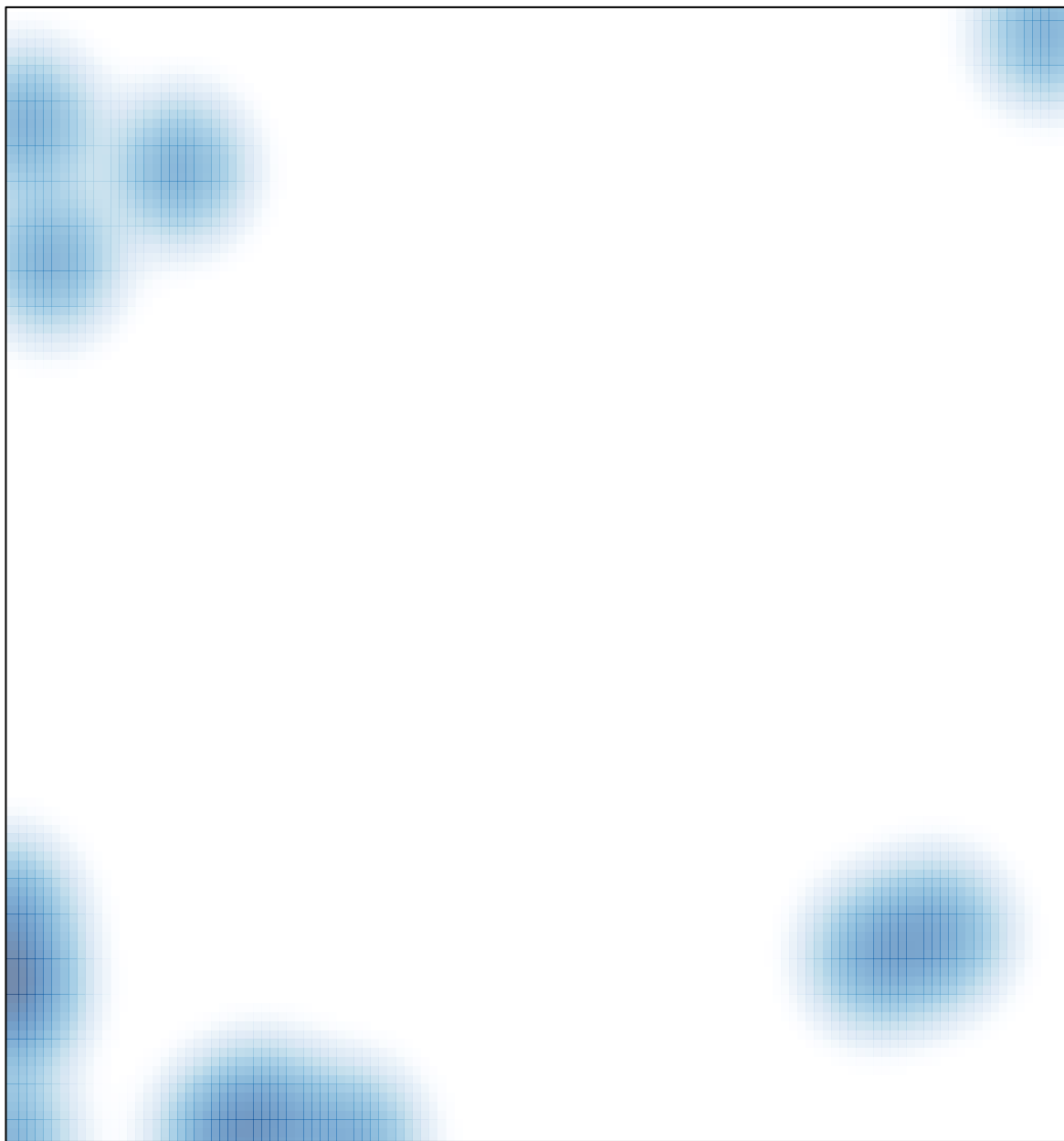


REACTOME_BRANCHED_CHAIN_AMINO_ACID_CATABOLISM



features = 16
chi-square p = 0.82

CTOME_BRANCHED_CHAIN_AMINO_ACID_CATABO



features = 16 , max = 1