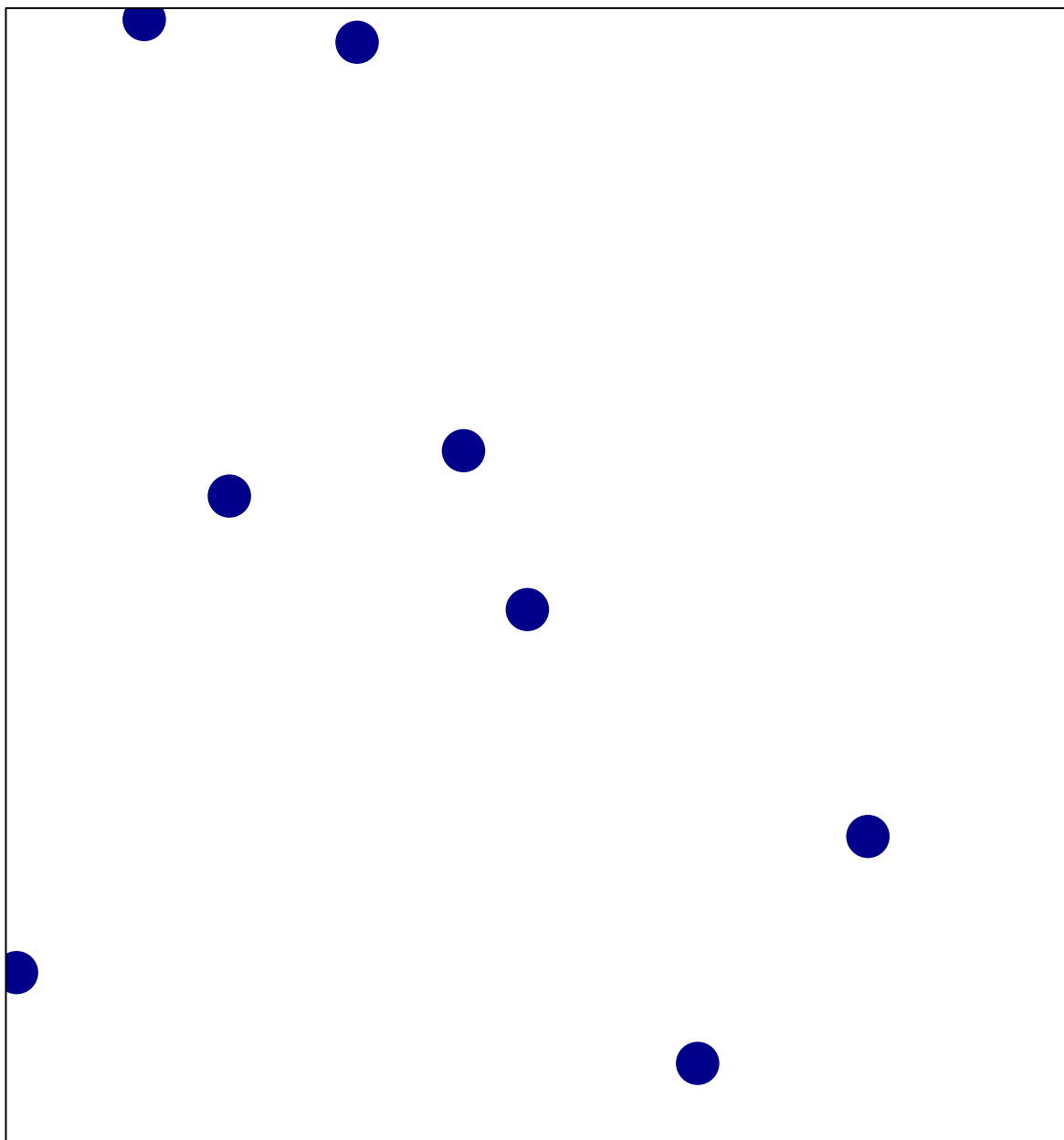
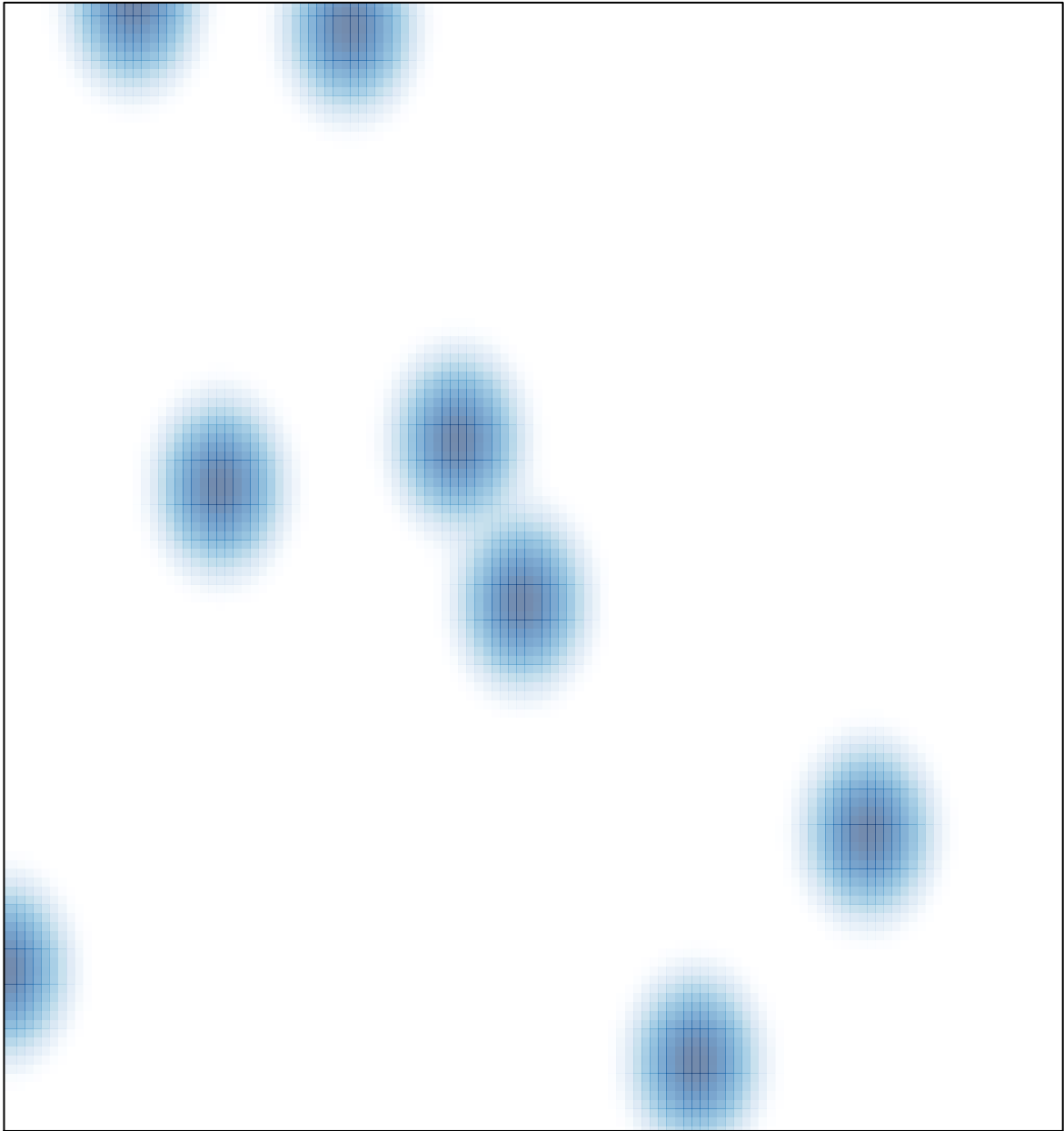


ME_GAMMA_CARBOXYLATION_TRANSPORT_AND_AMINO_TERMINAL_CLEAVAGE_OF_



features = 8
chi-square p = 0.85

BOXYLATION_TRANSPORT_AND_AMINO_TERMINAL



features = 8 , max = 1