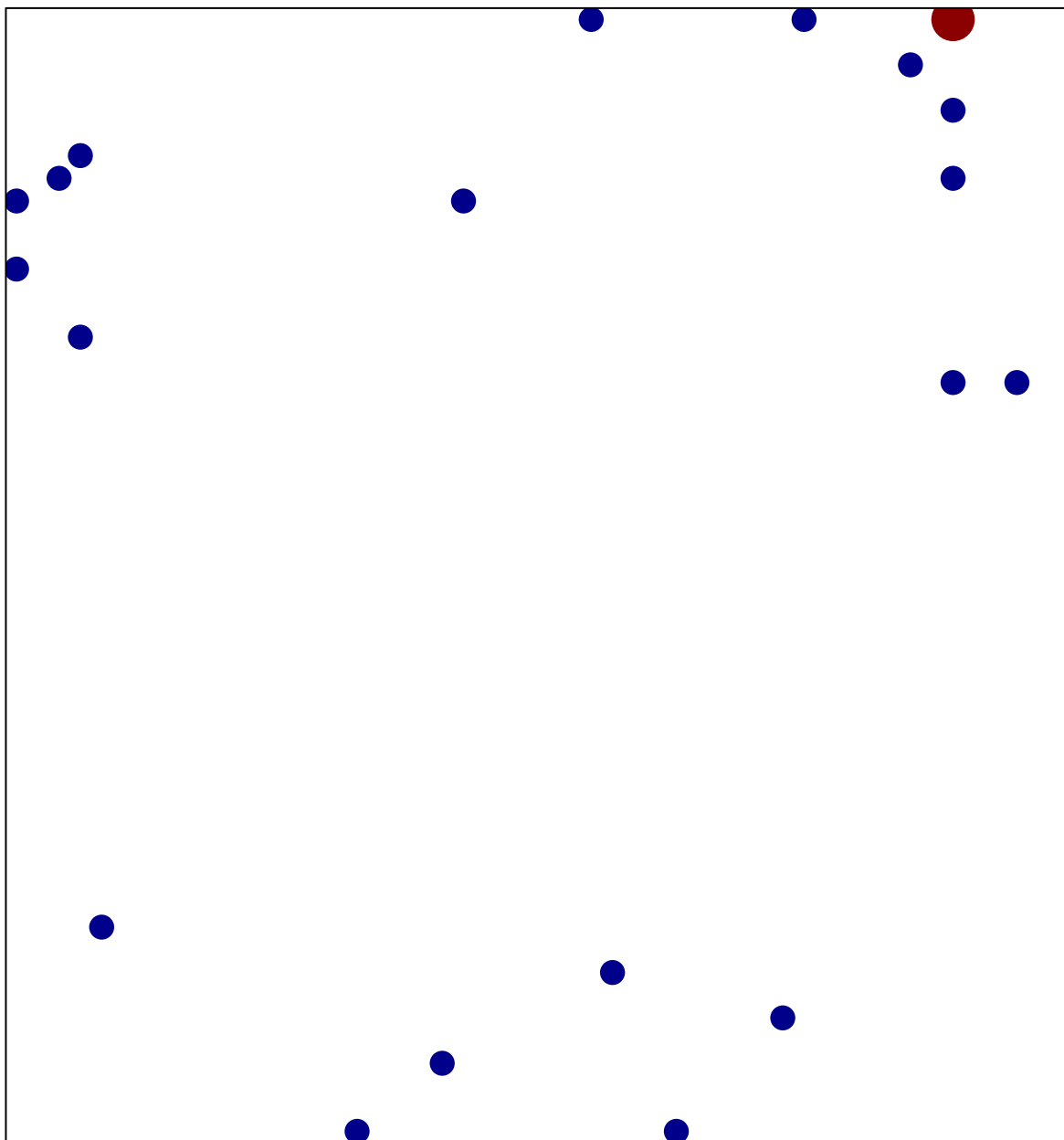
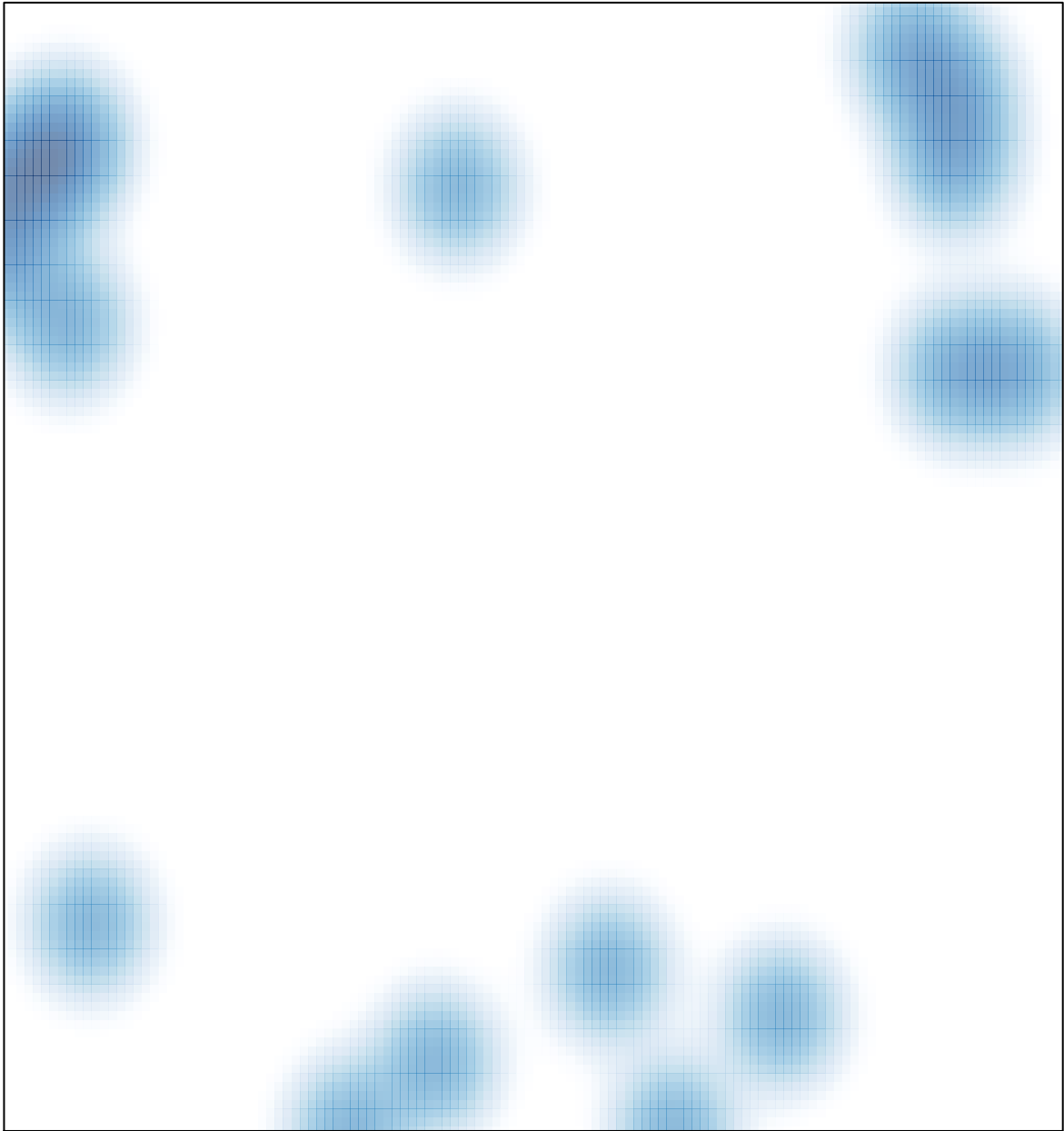


REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATION



features = 21
chi-square p = 0.8

REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATI



features = 21 , max = 2