REACTOME_TRNA_AMINOACYLATION
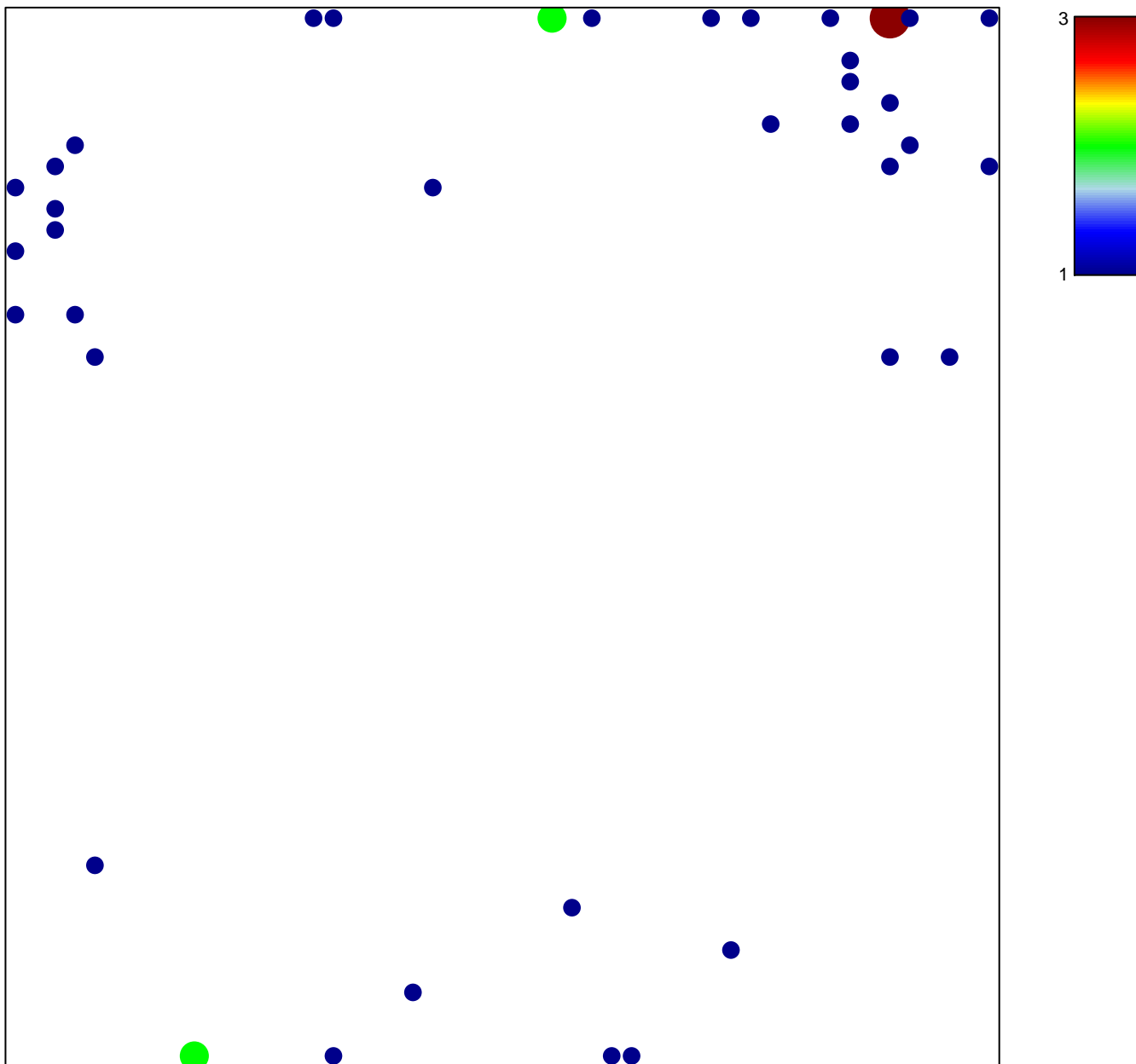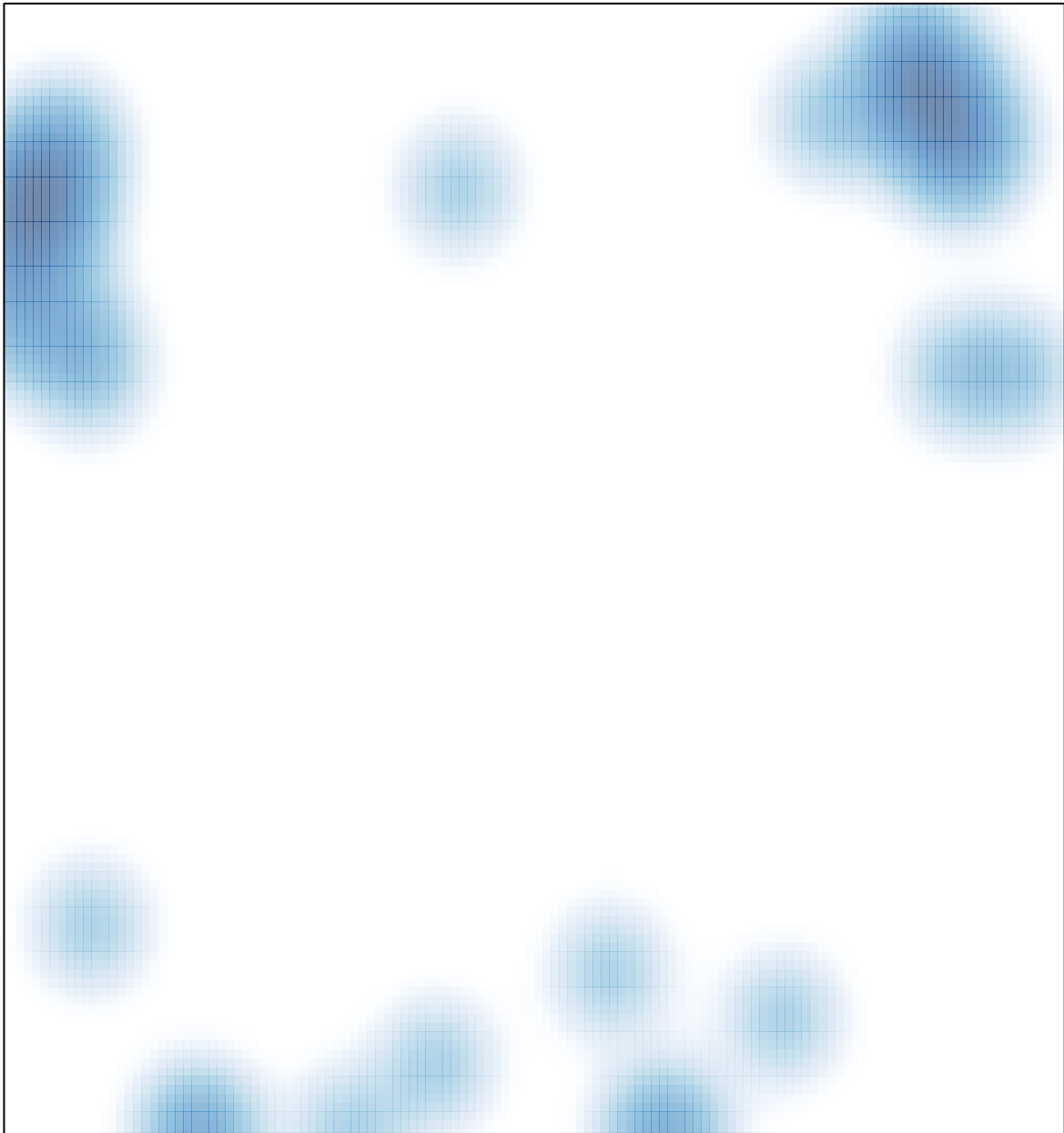

features = 42

chi−square p = 0.72

REACTOME_TRNA_AMINOACYLATION

features = 42 , max = 3