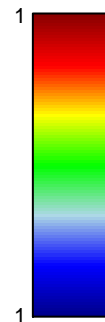
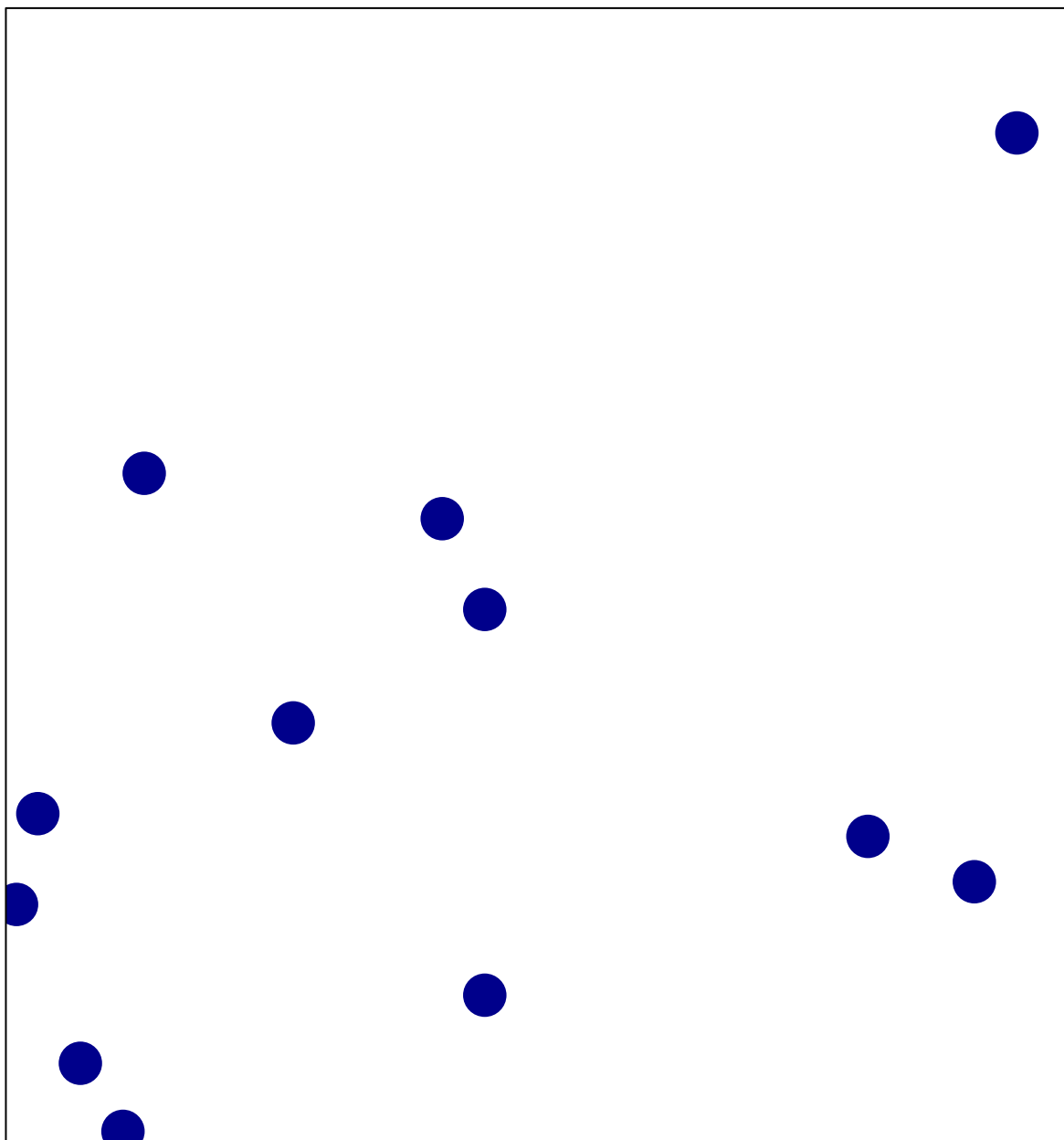
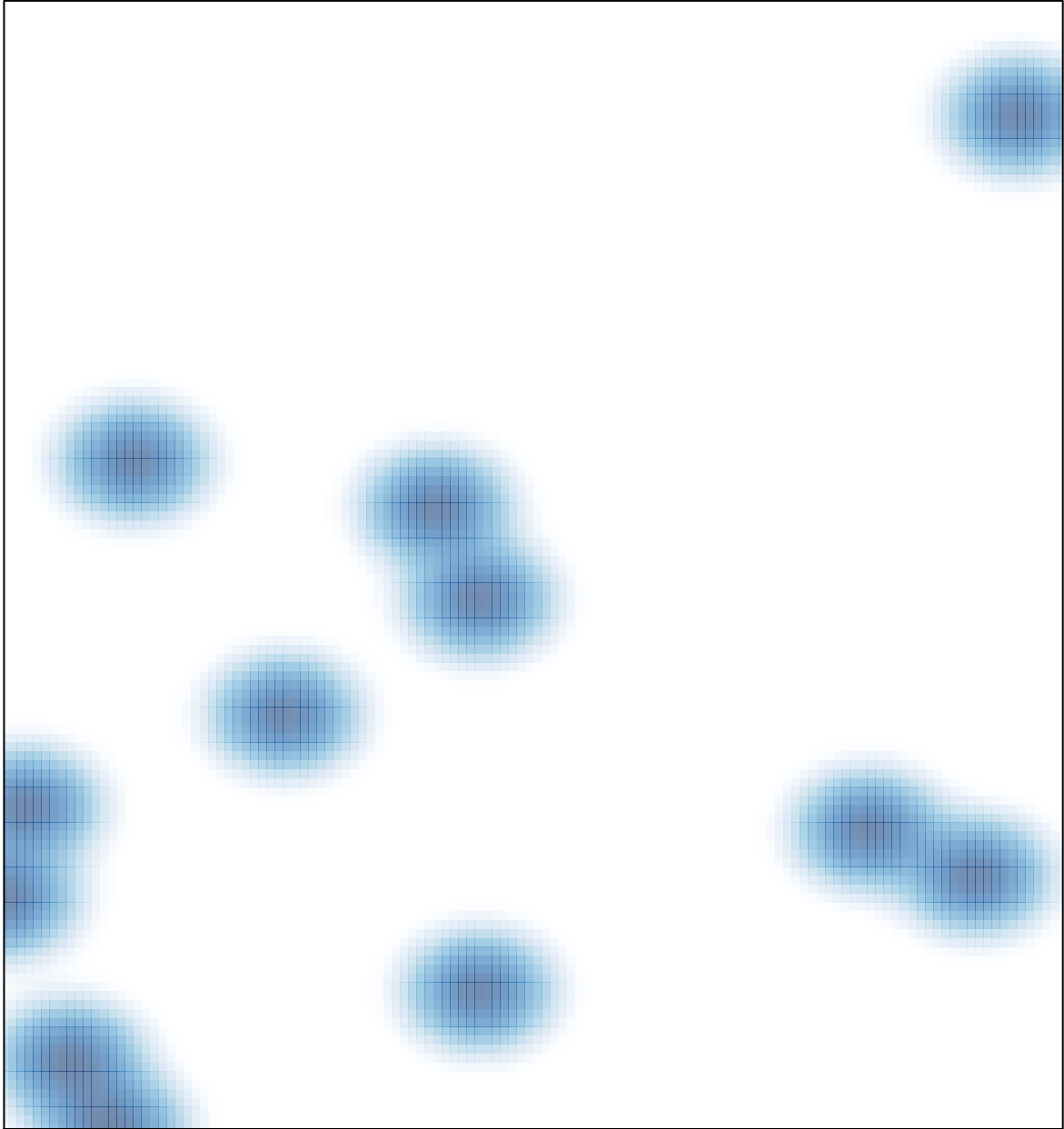


neutral amino acid transport



features = 12
chi-square p = 0.85

neutral amino acid transport



features = 12 , max = 1