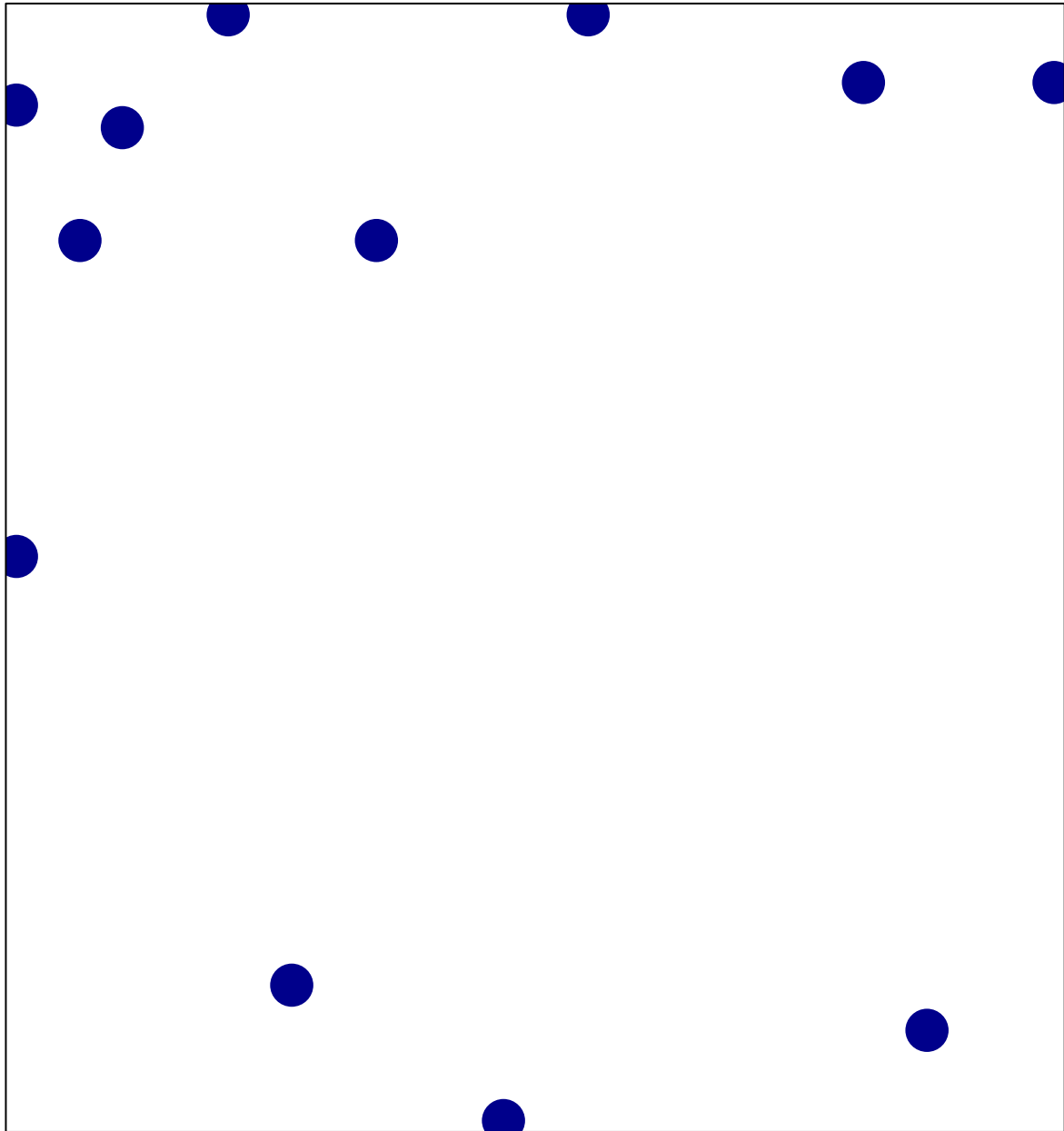
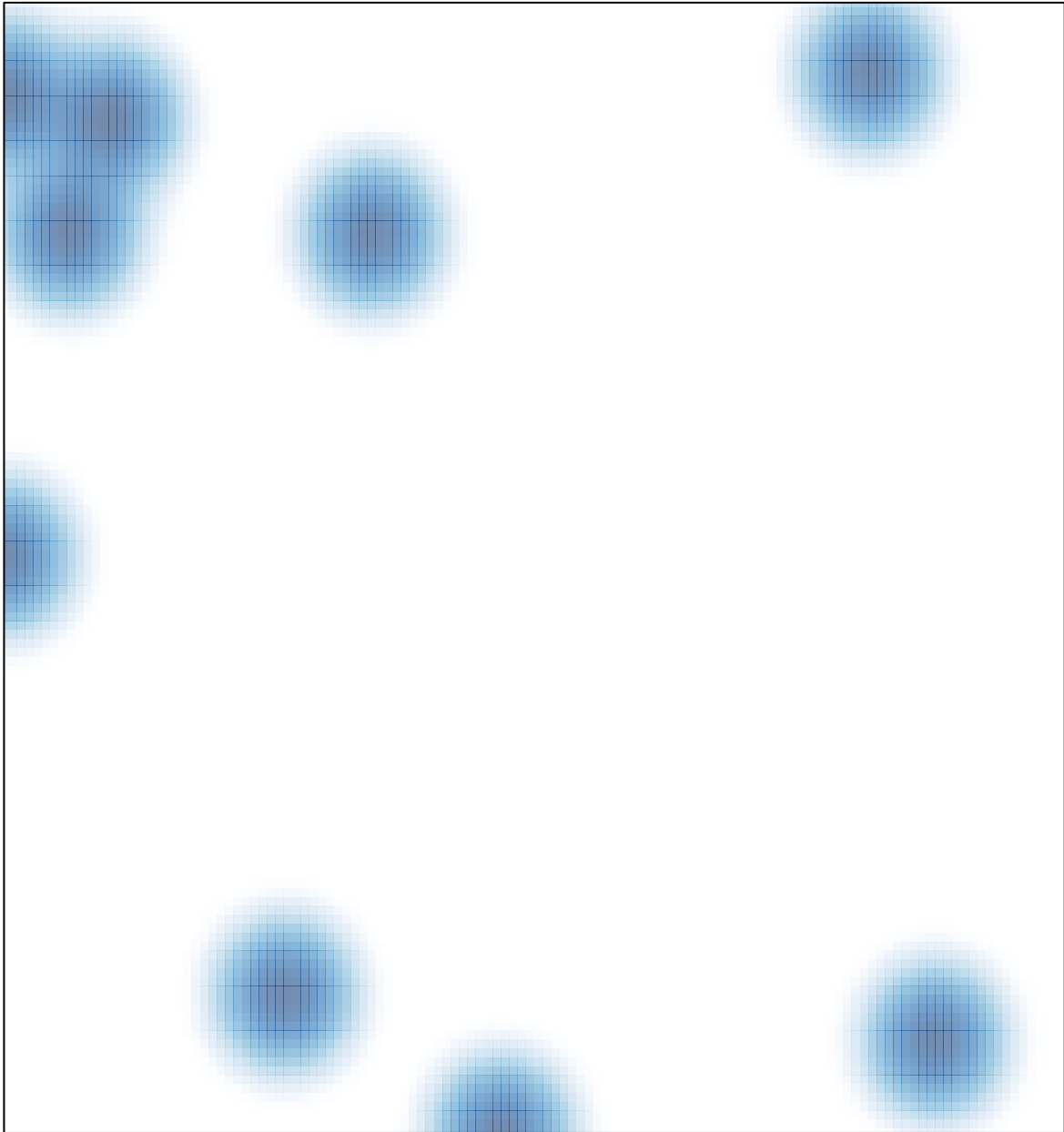


mitochondrial genome maintenance



features = 12
chi-square p = 0.83

mitochondrial genome maintenance



features = 12 , max = 1