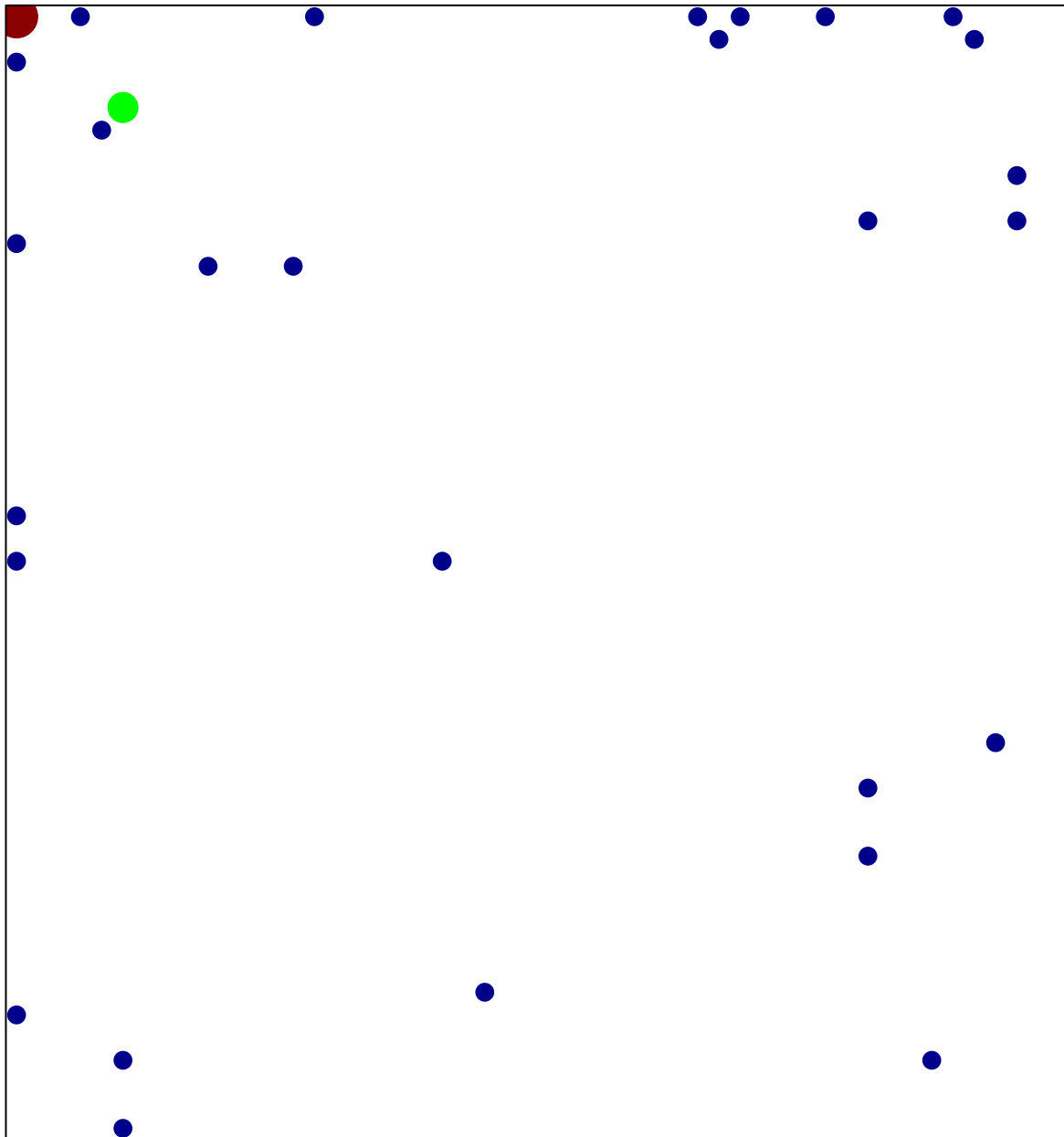
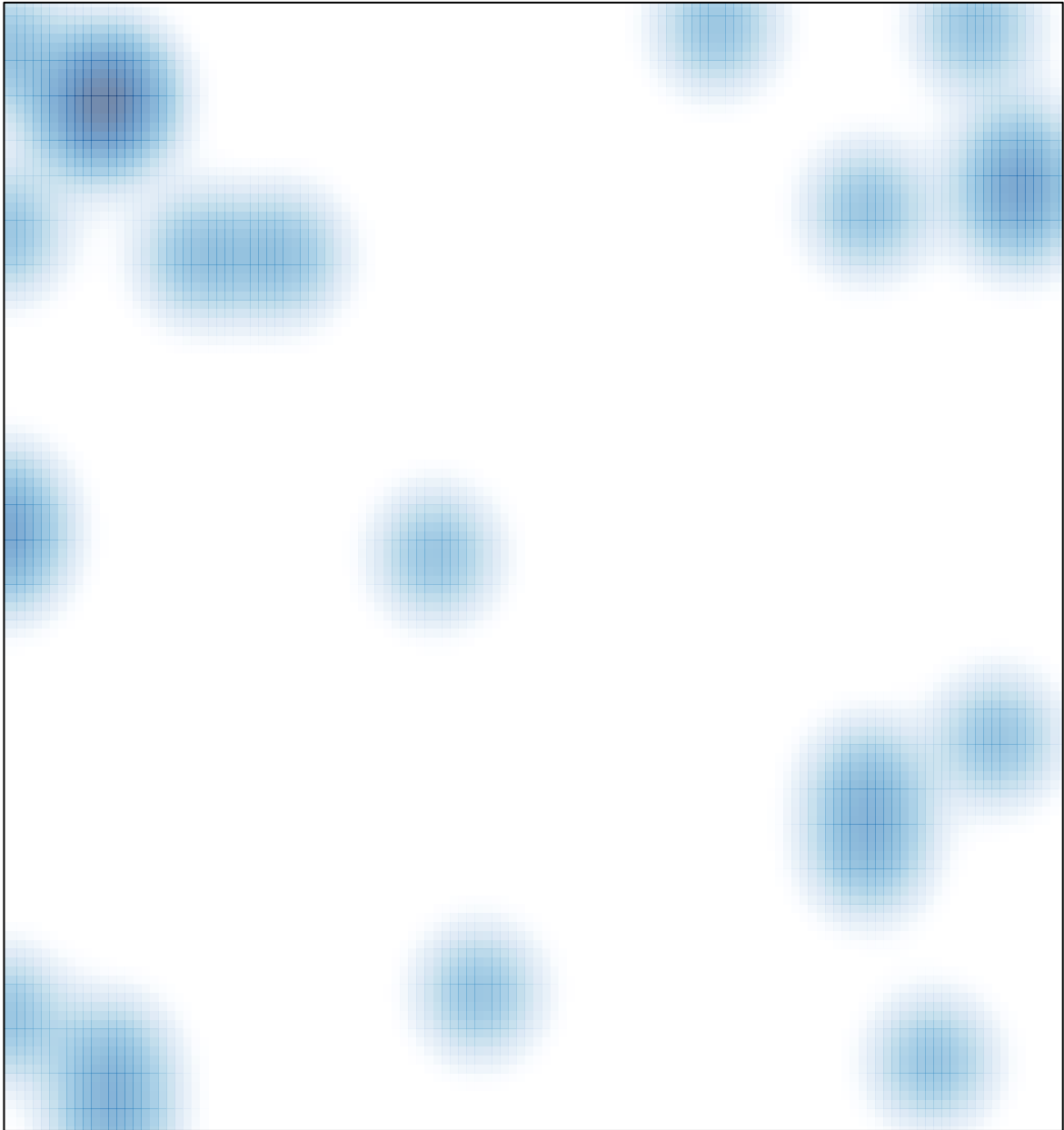


RNA phosphodiester bond hydrolysis, endonucleolytic



features = 32
chi-square p = 0.74

RNA phosphodiester bond hydrolysis, endonucleolyt



features = 32 , max = 3