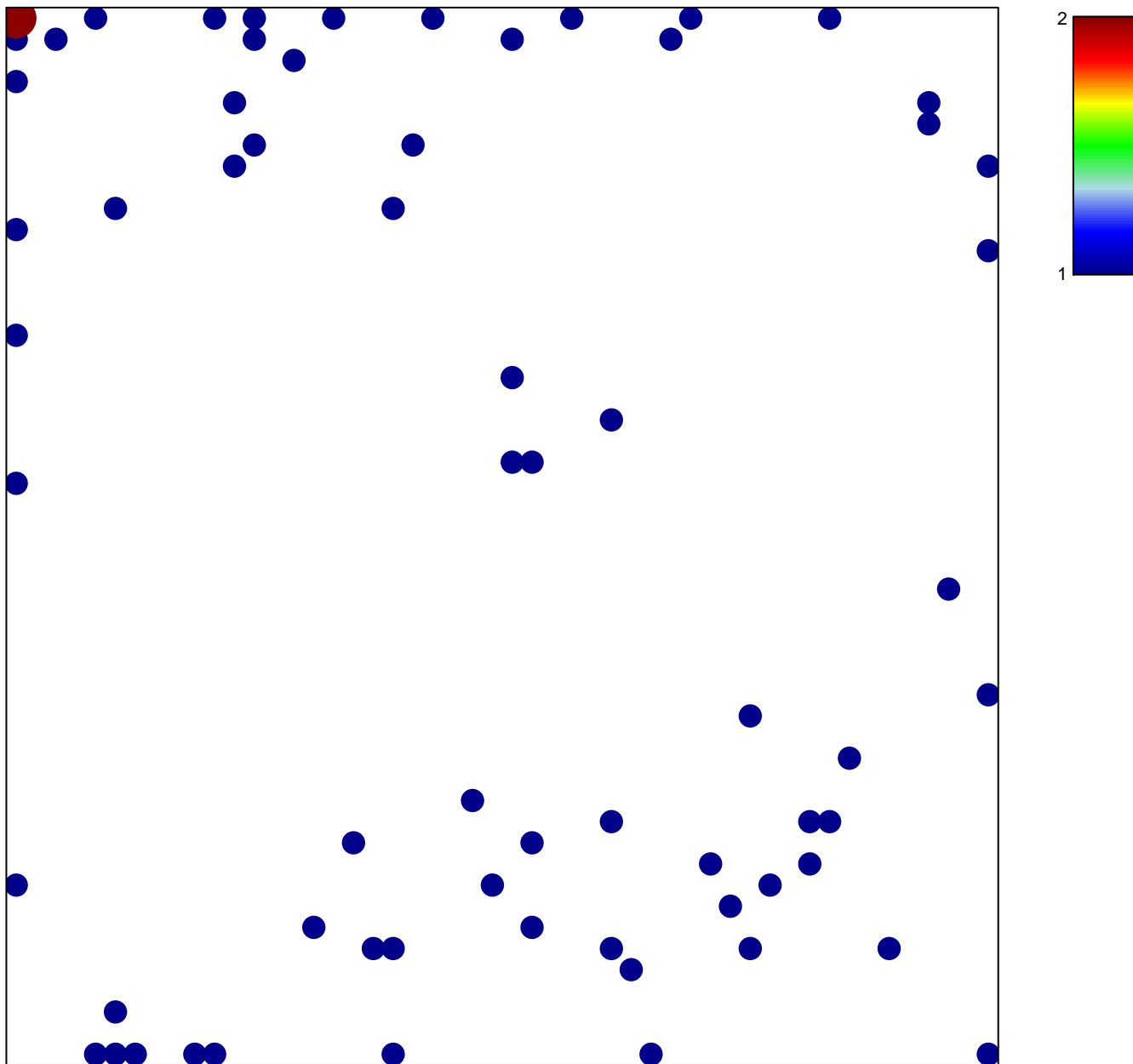
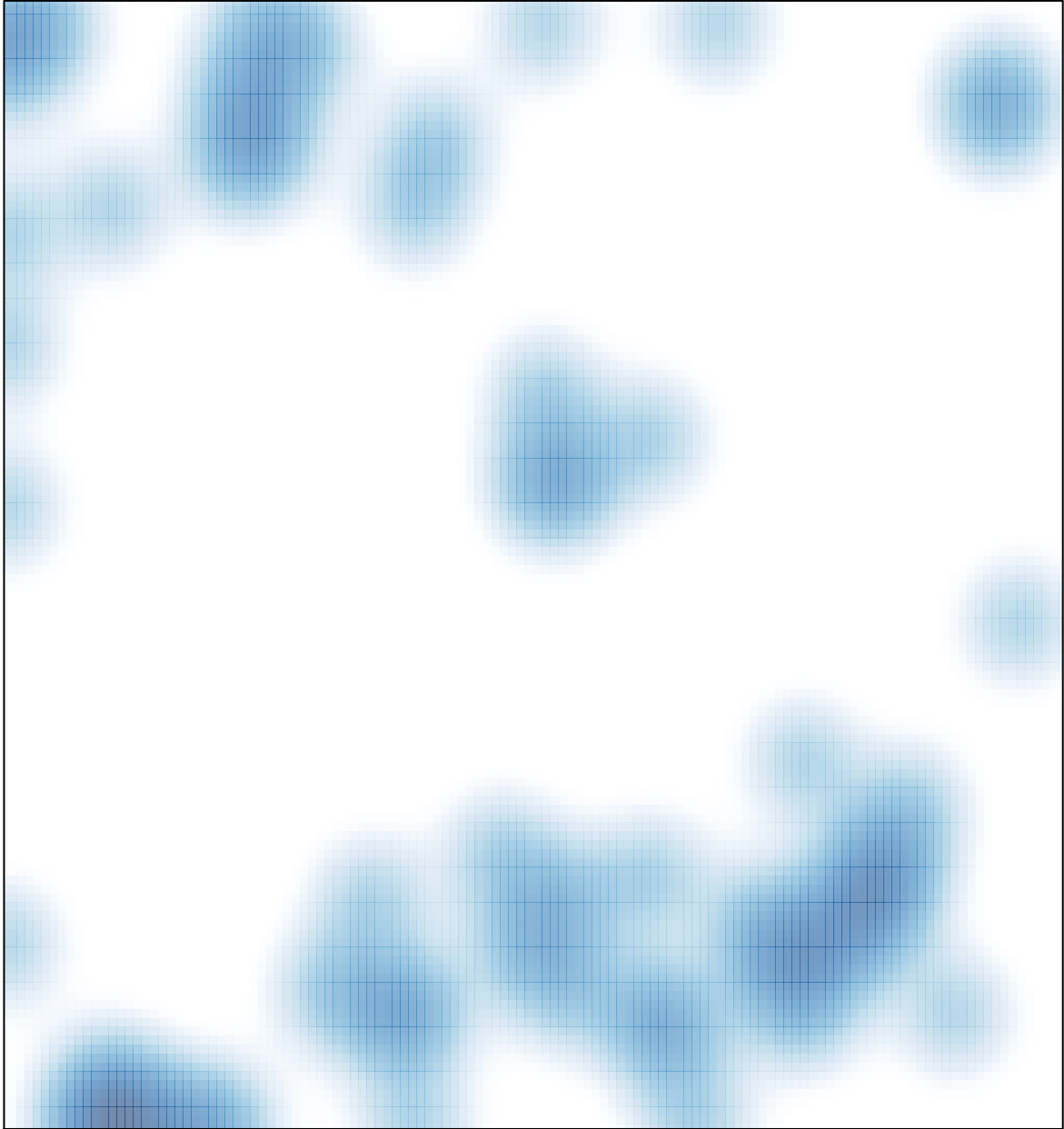


nucleic acid phosphodiester bond hydrolysis



features = 67
chi-square p = 0.77

nucleic acid phosphodiester bond hydrolysis



features = 67 , max = 2