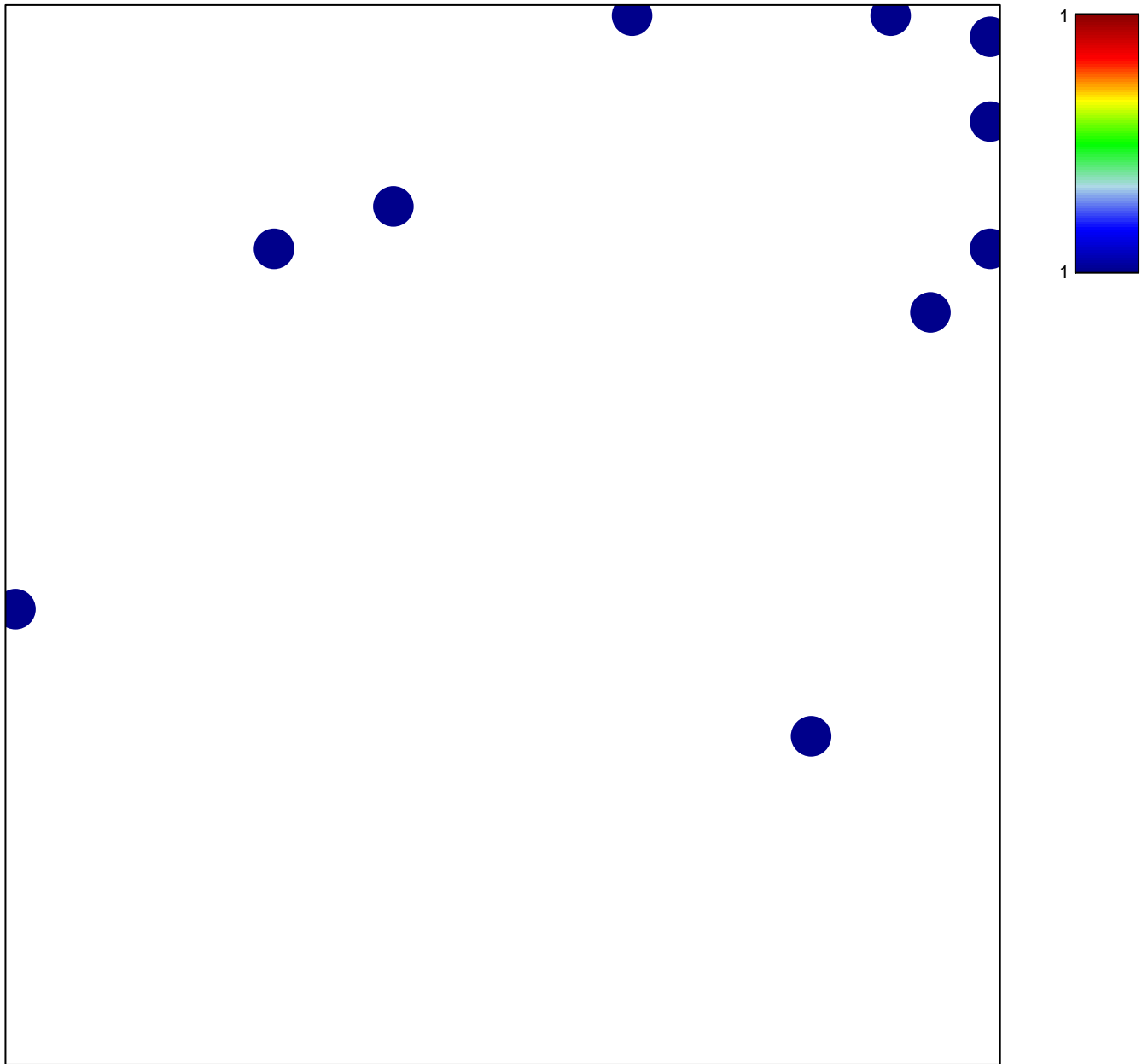
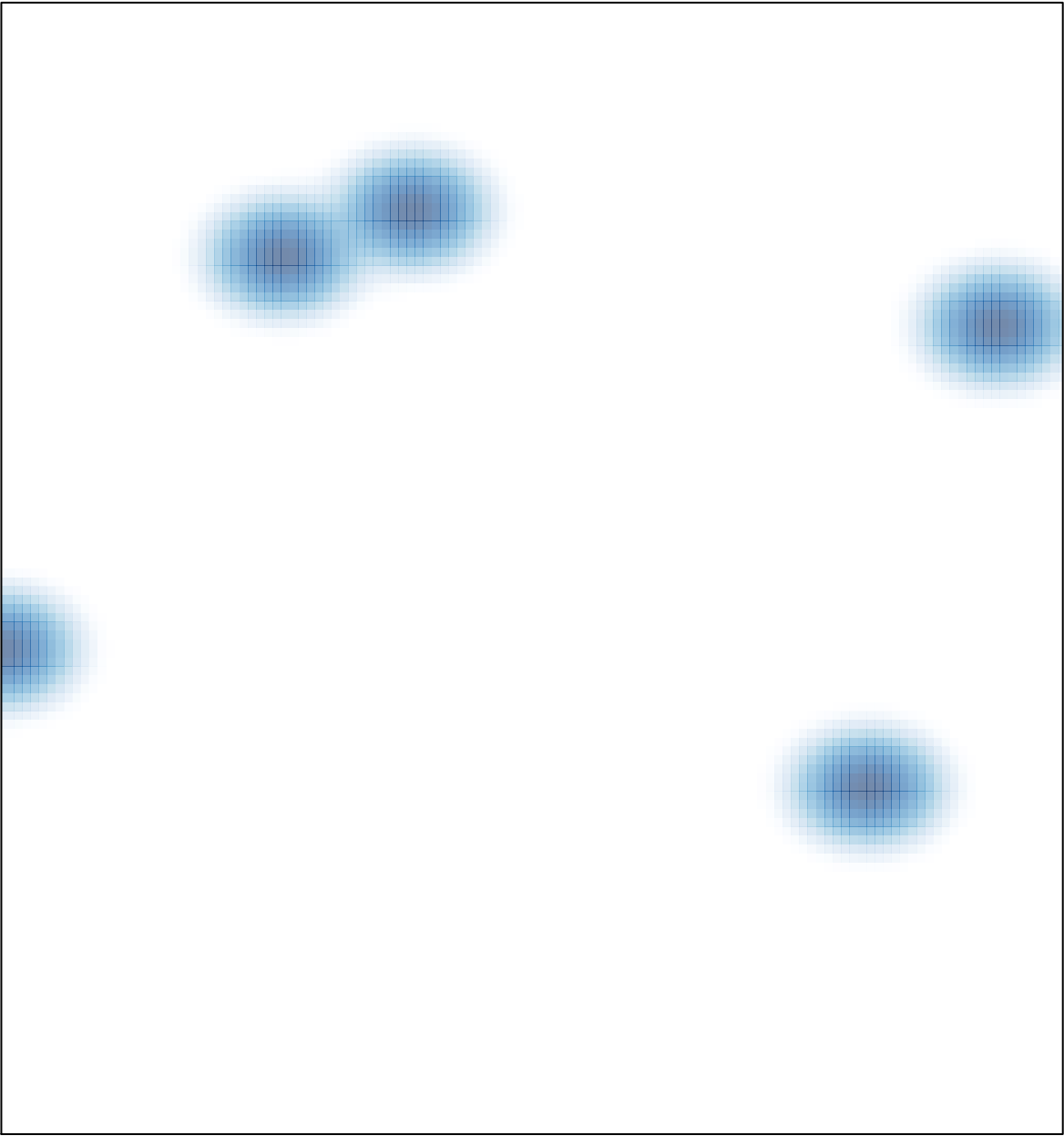


tRNA splicing, via endonucleolytic cleavage and ligation



features = 10
chi-square p = 0.82

RNA splicing, via endonucleolytic cleavage and ligati



features = 10 , max = 1