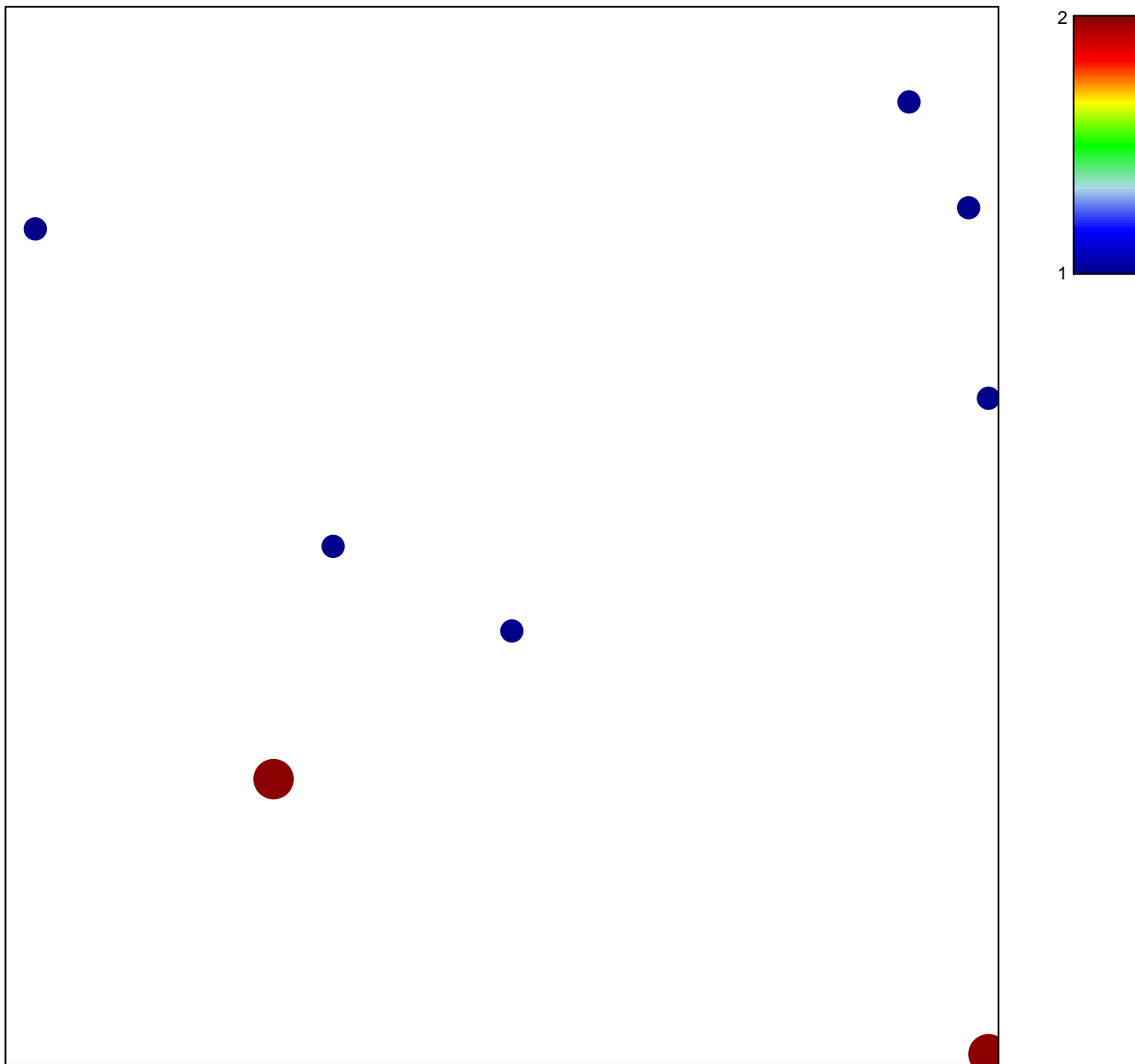


proteoglycan binding



features = 10
chi-square p = 0.81

proteoglycan binding



features = 10 , max = 2