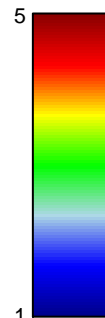
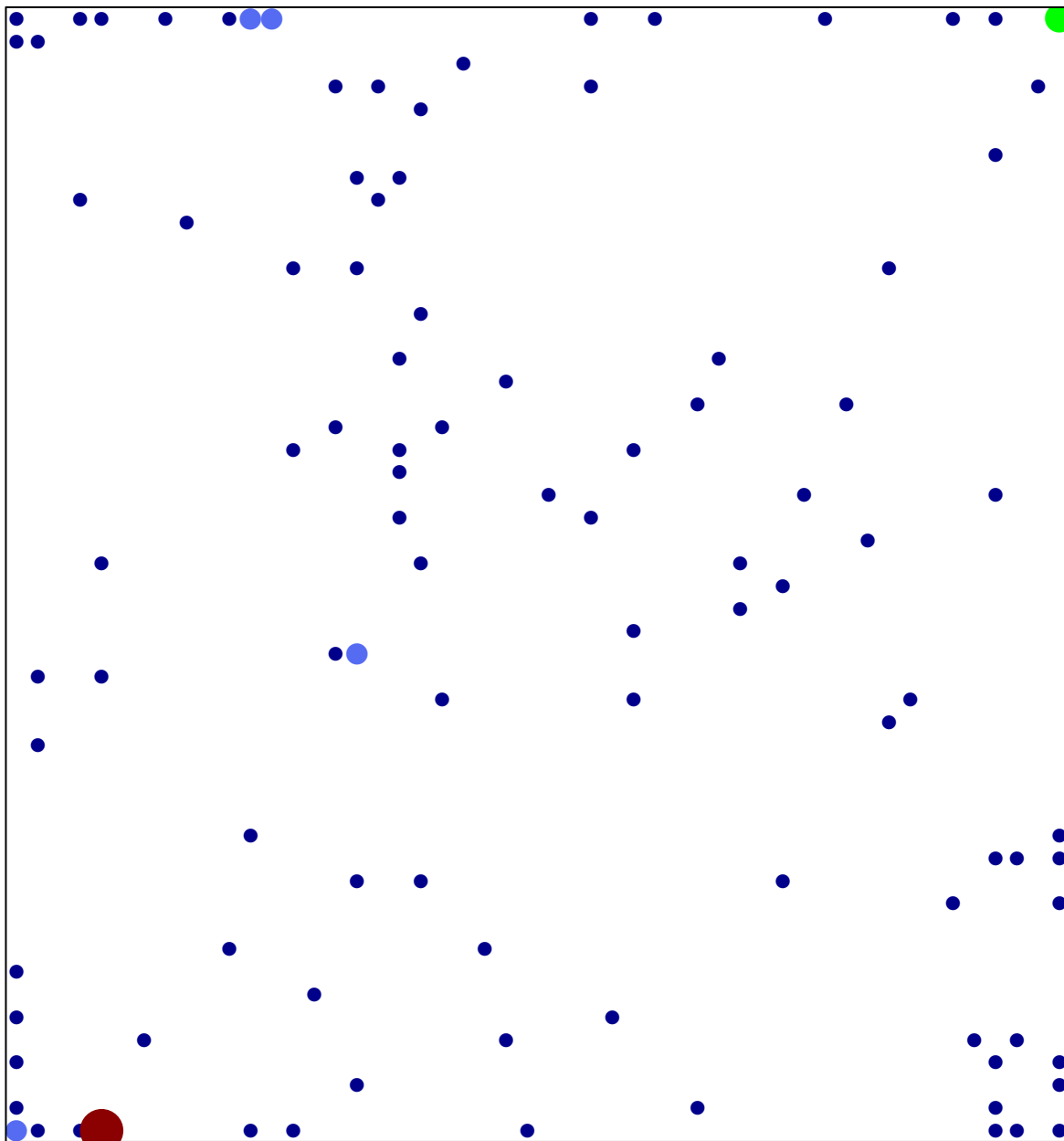
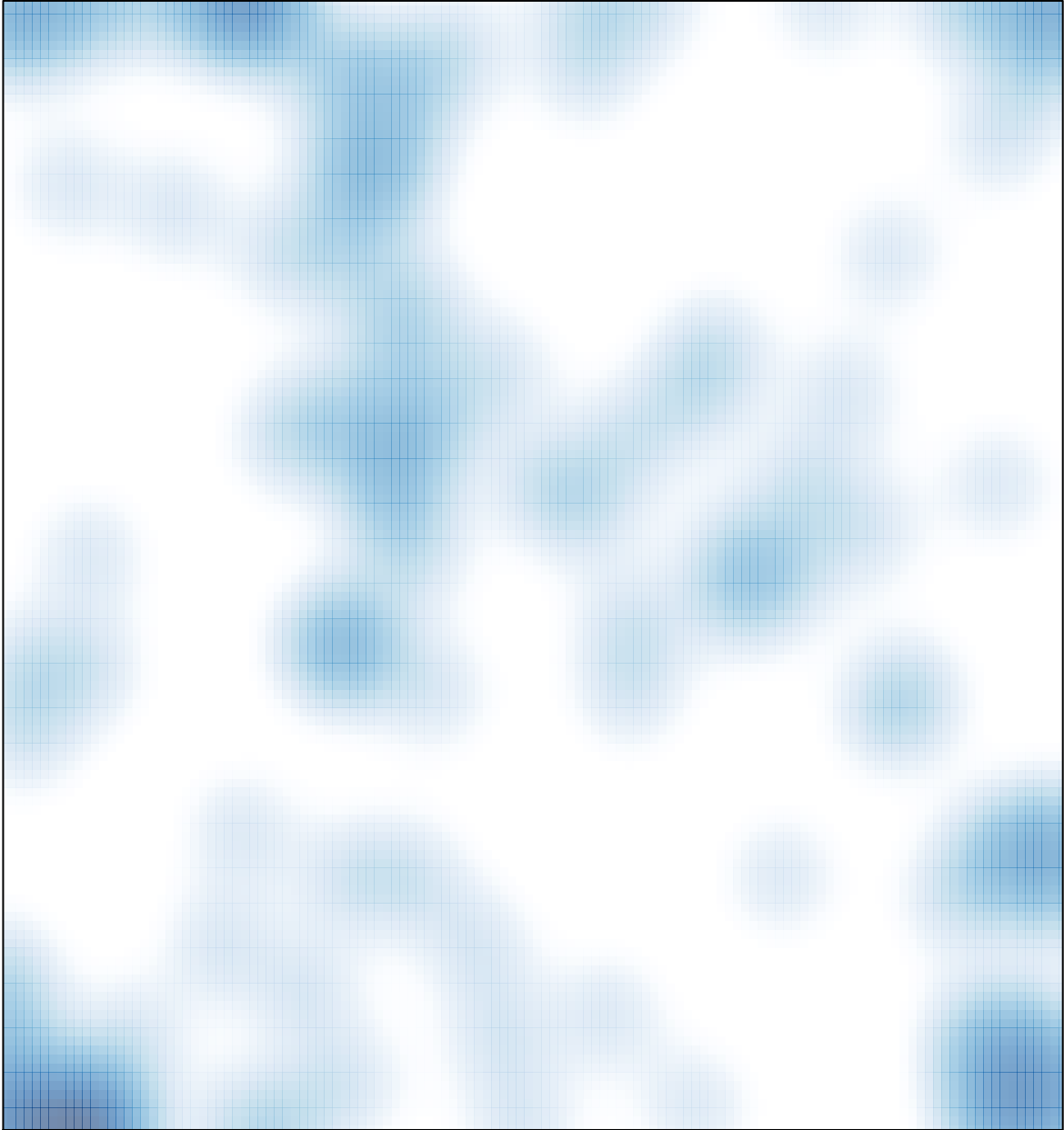


heme binding



features = 111
chi-square p = 0.72

heme binding



features = 111 , max = 5