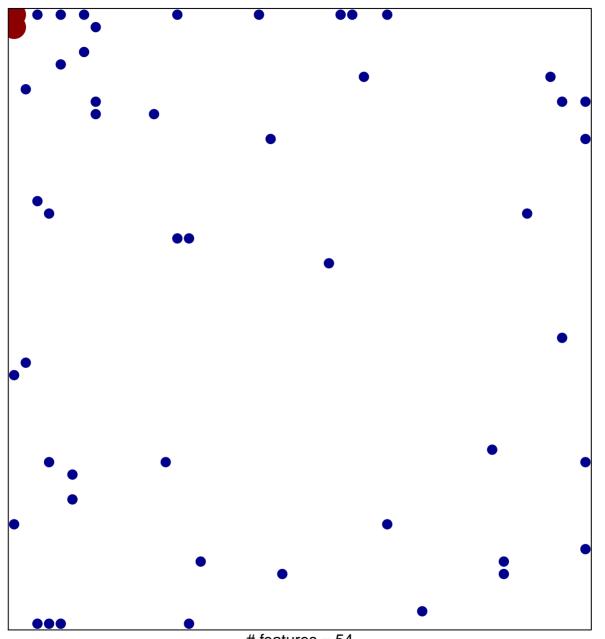
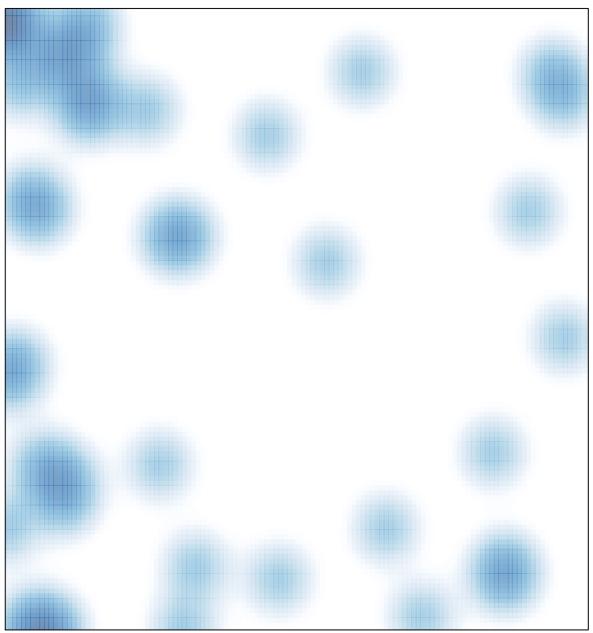
damaged DNA binding



features = 54 chi-square p = 0.76

damaged DNA binding



features = 54 , max = 3