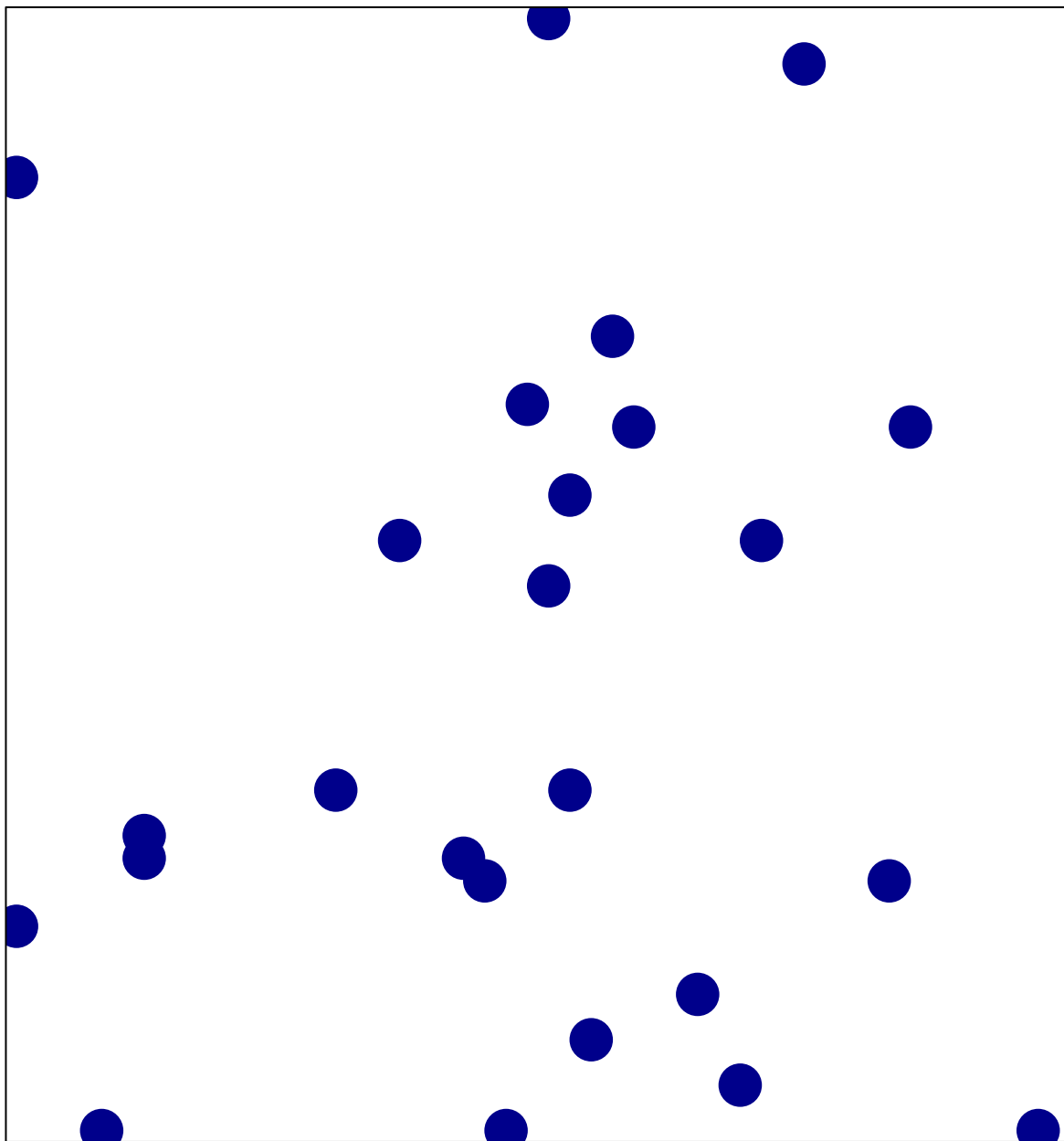
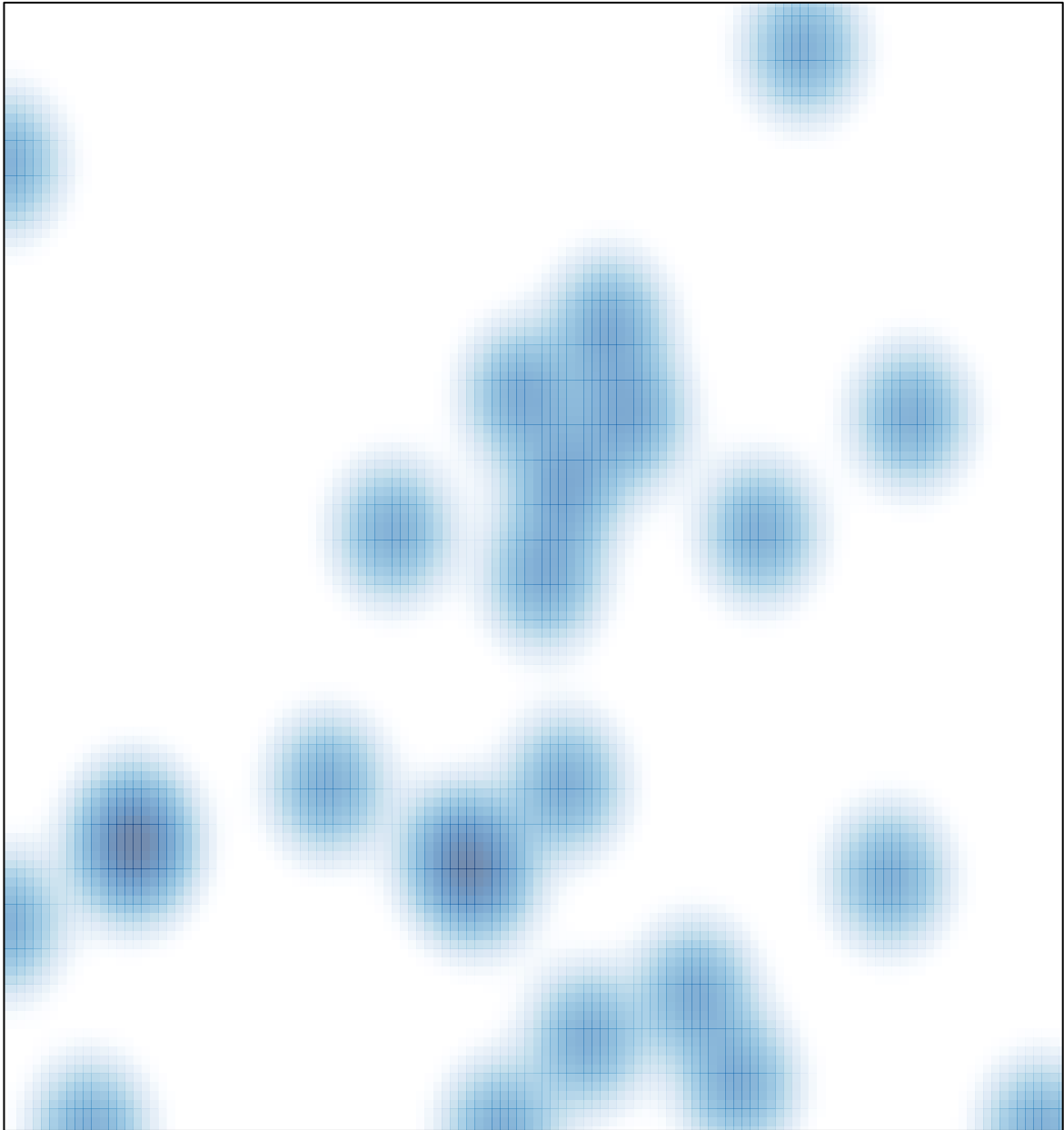


Ang_CRC_Hypomethylated



features = 25
chi-square p = 0.84

Ang_CRC_Hypomethylated



features = 25 , max = 1