Christensen_hypomethylated_in_ependymoma
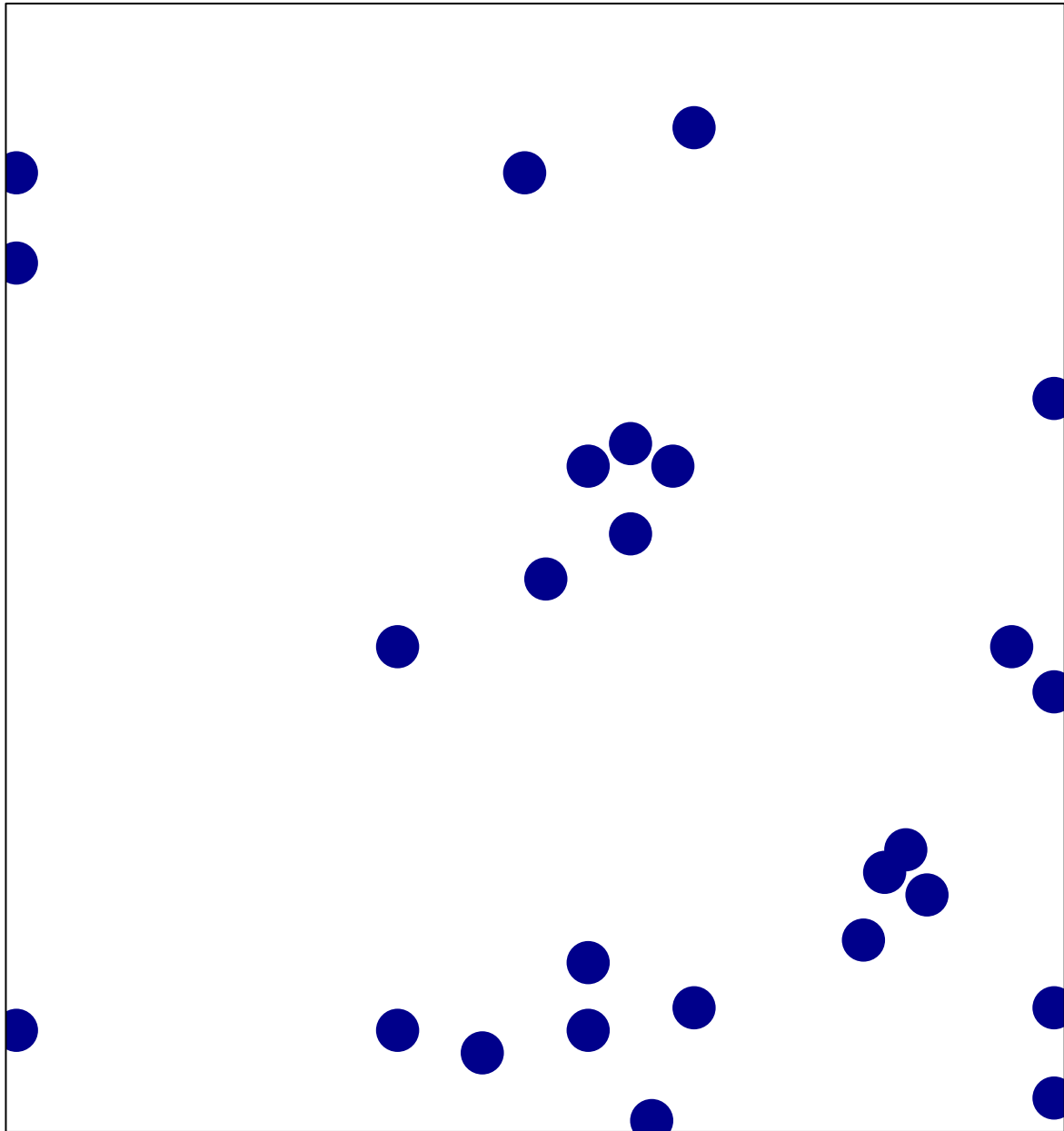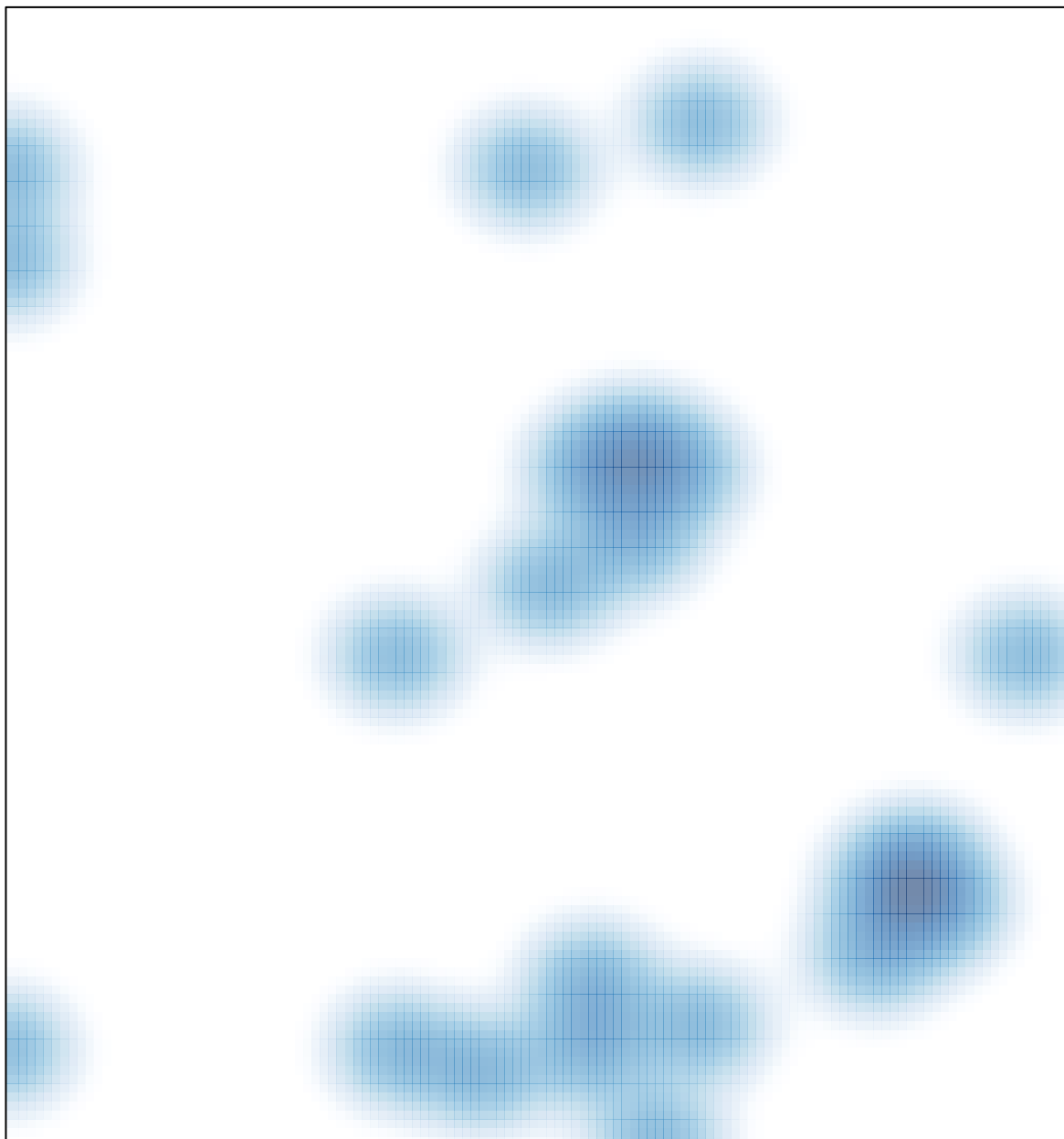

features = 26
chi−square p = 0.82

Christensen_hypomethylated_in_ependymoma

features = 26 , max = 1