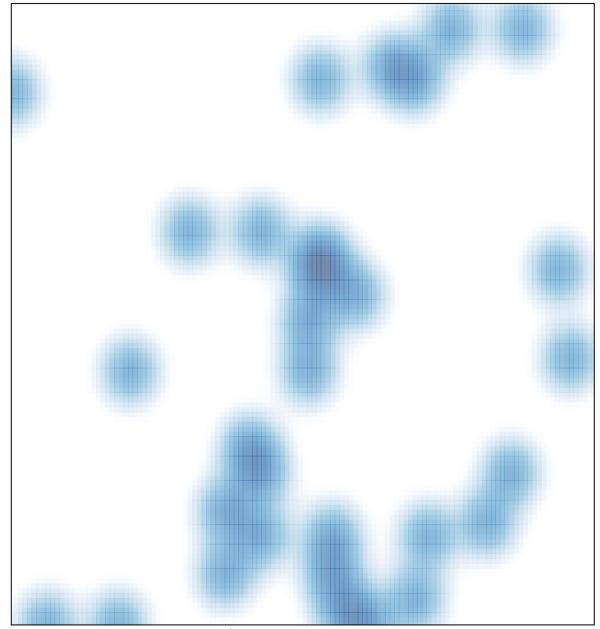


Christensen_hypomethylated_in_secondary_glioblastoma

ristensen_hypomethylated_in_secondary_glioblasto



features = 37, max = 2