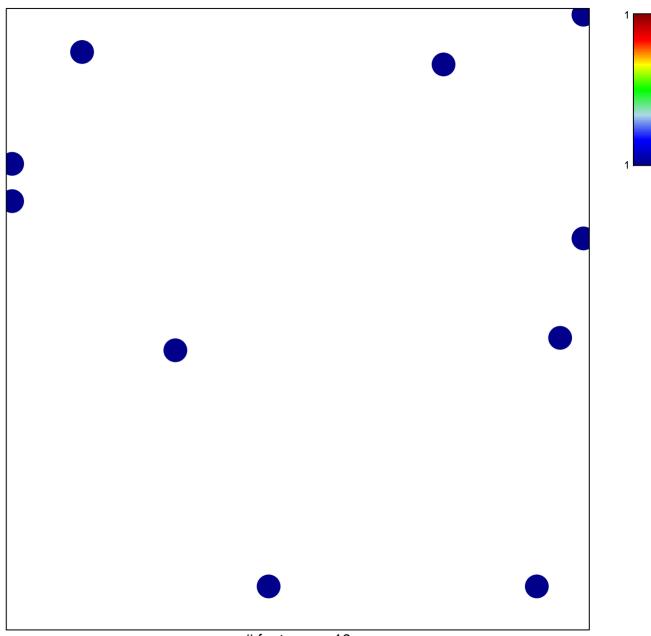
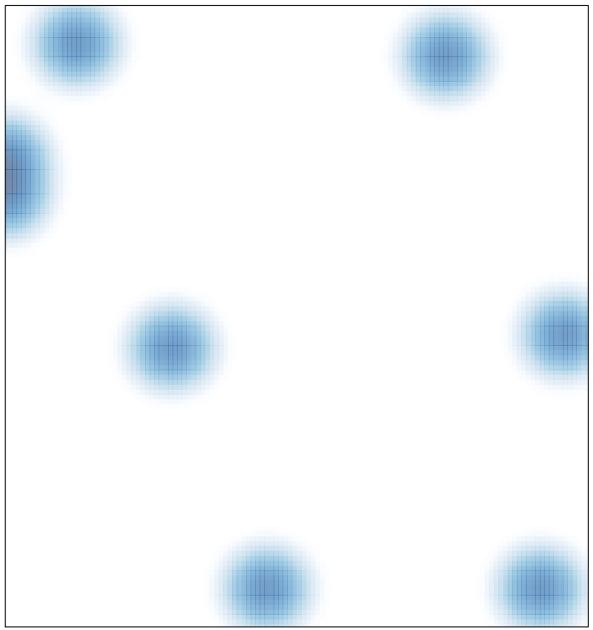
leucine zipper domain binding



features = 10 chi-square p = 0.83

leucine zipper domain binding



features = 10 , max = 1