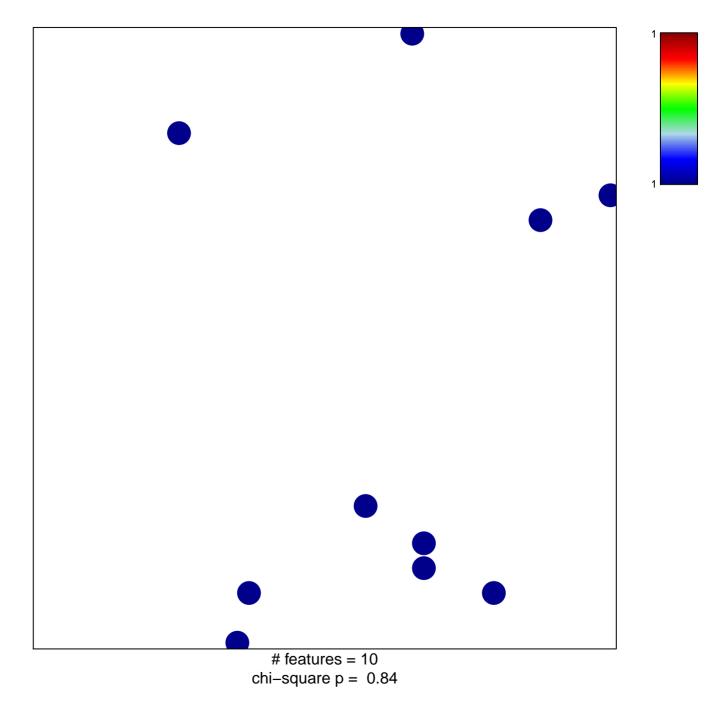
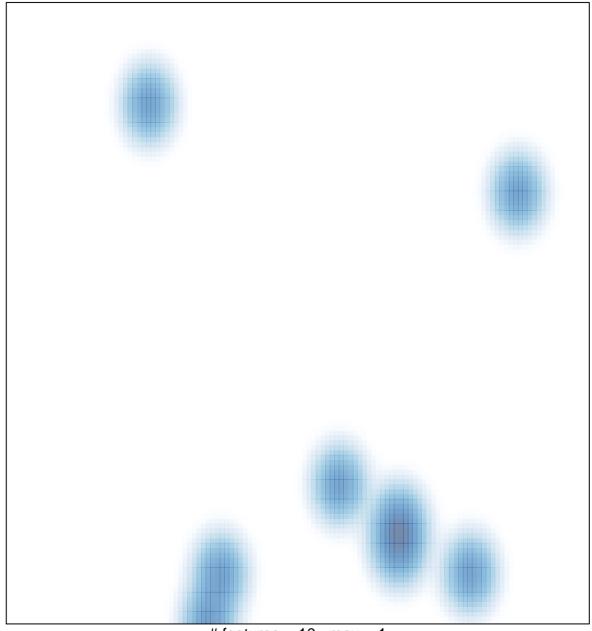
histone H3–K4 trimethylation



## histone H3–K4 trimethylation



# features = 10, max = 1