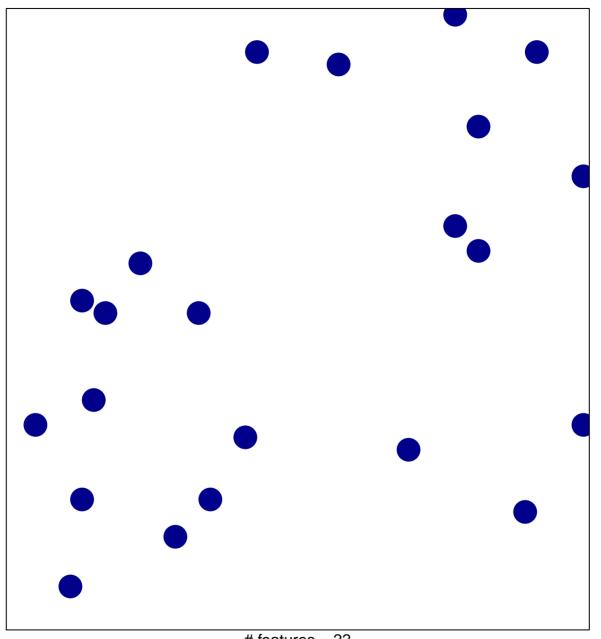
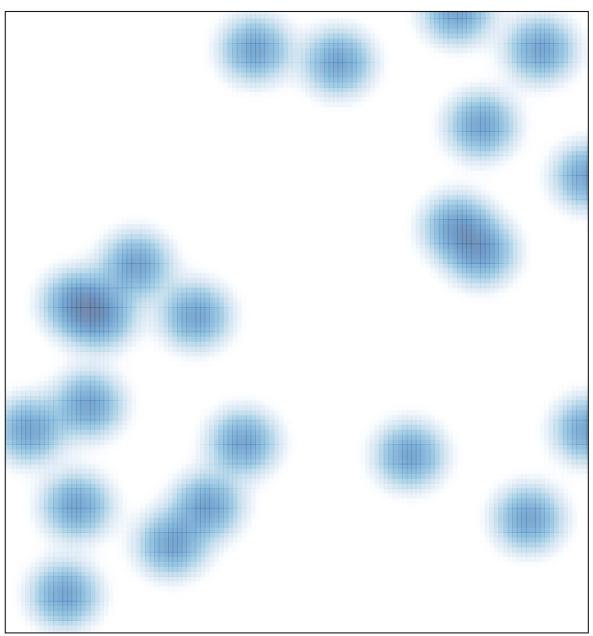
## histone methylation



# features = 22 chi-square p = 0.85

## histone methylation



# features = 22 , max = 1