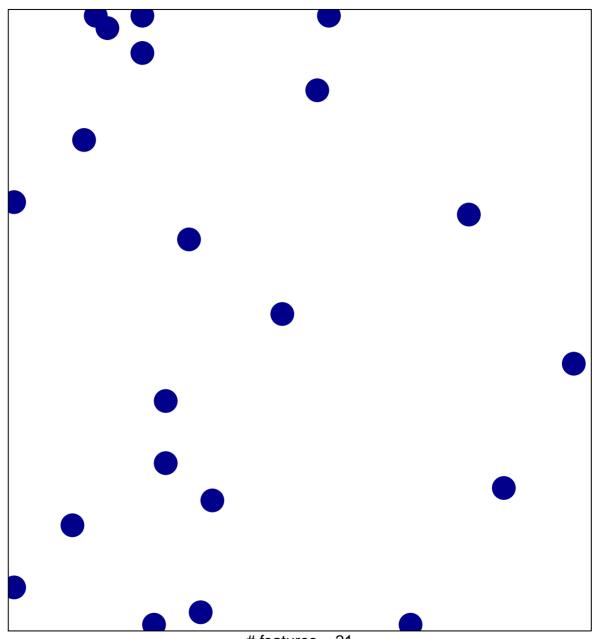
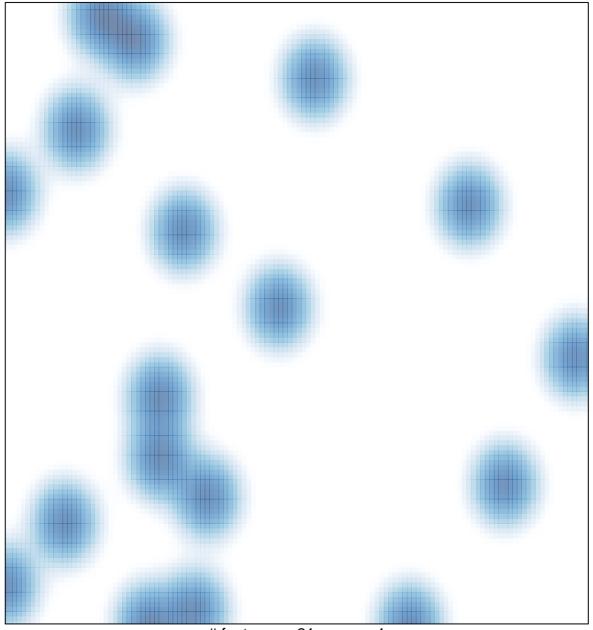
## positive regulation of peptidyl-threonine phosphorylation



# features = 21 chi-square p = 0.85

## sitive regulation of peptidyl-threonine phosphorylat



# features = 21 , max = 1