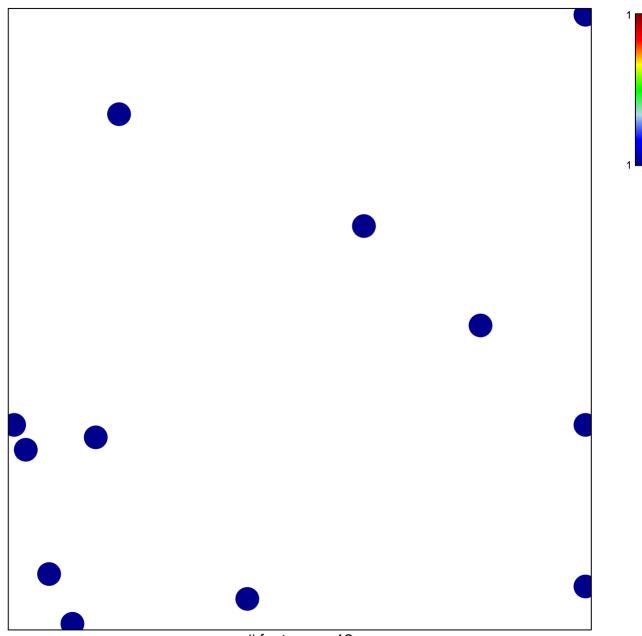
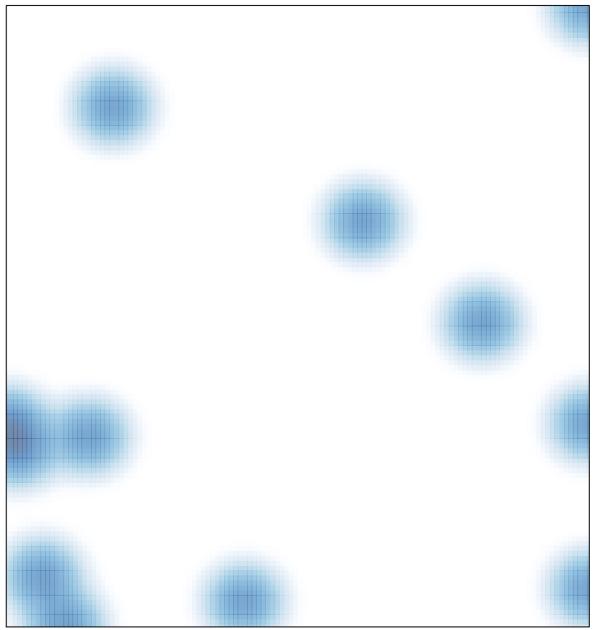
positive regulation of peptidyl-threonine phosphorylation



features = 12 chi-square p = 0.83

sitive regulation of peptidyl-threonine phosphorylat



features = 12 , max = 1