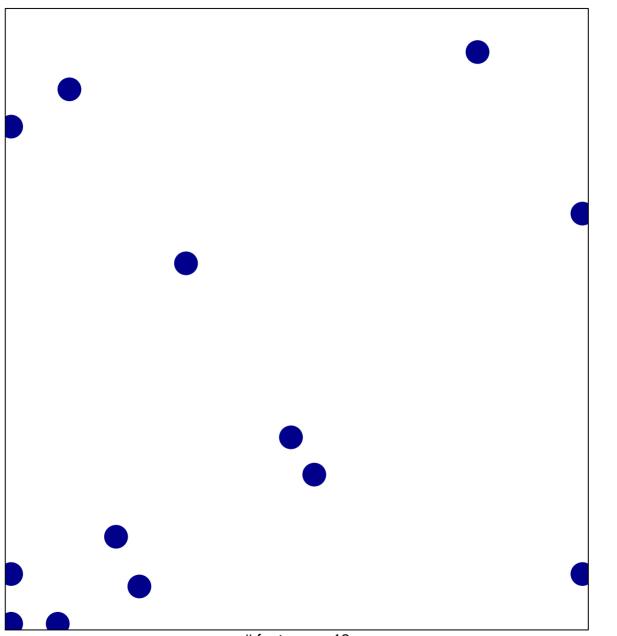
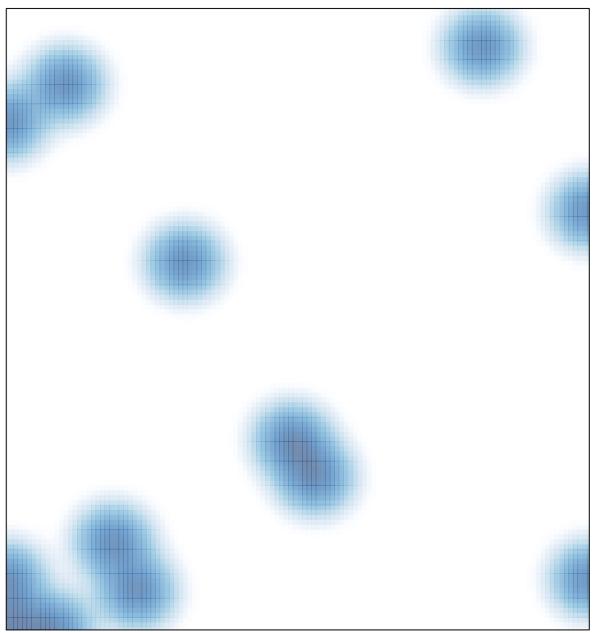
## low-density lipoprotein particle binding



# features = 13 chi-square p = 0.83

## low-density lipoprotein particle binding



# features = 13 , max = 1