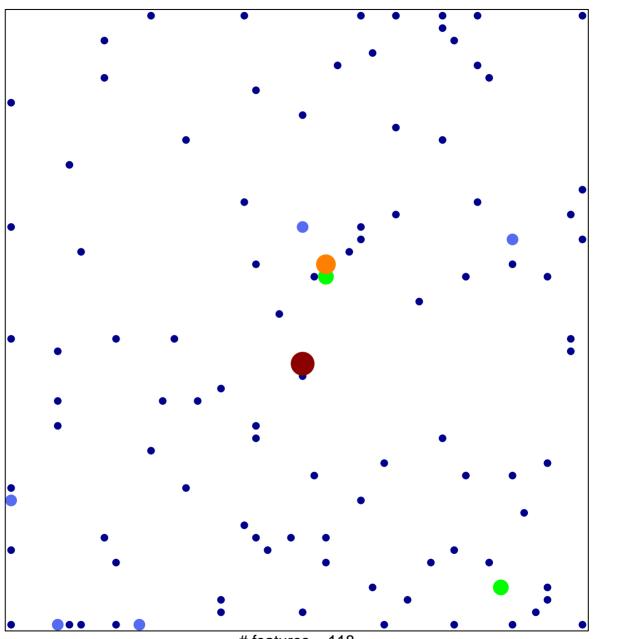
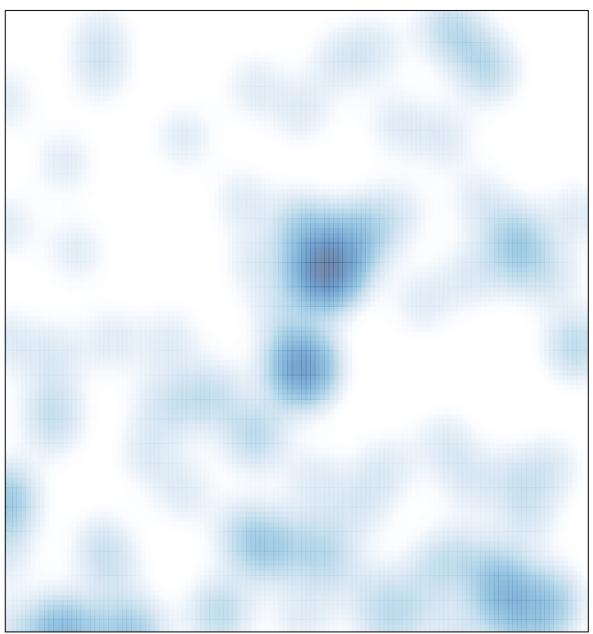
## carbohydrate binding



# features = 118 chi-square p = 0.78

## carbohydrate binding



# features = 118 , max = 5