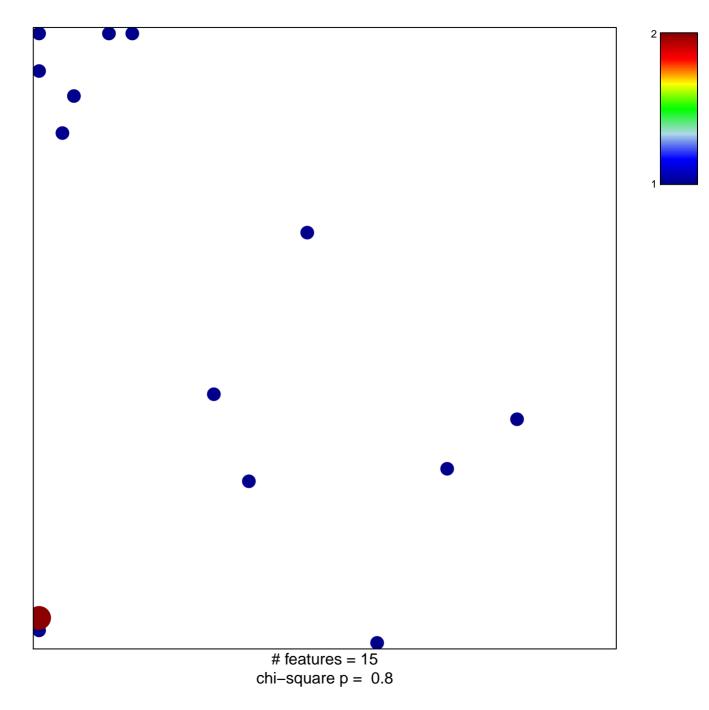
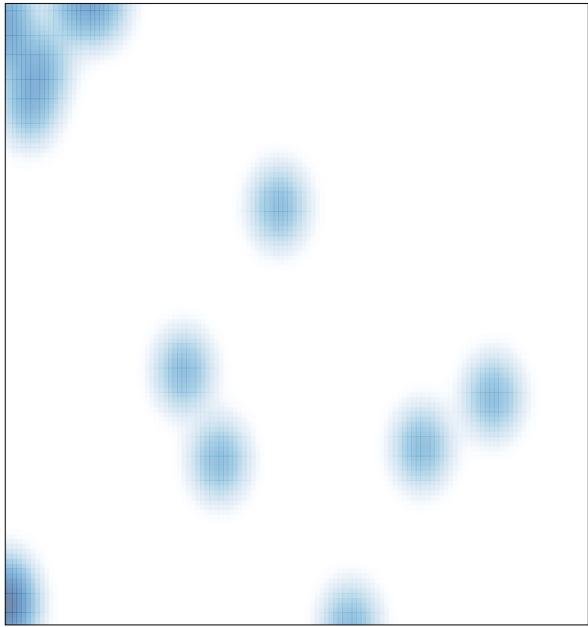
## fibrinolysis



## fibrinolysis



# features = 15, max = 2