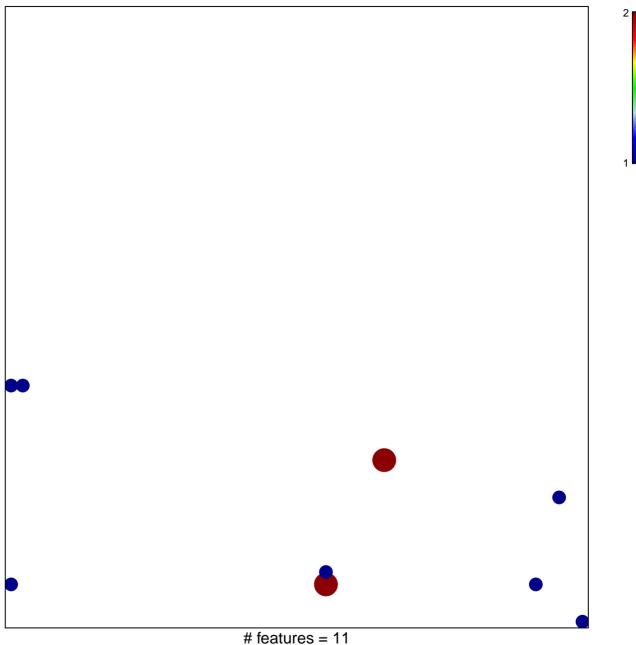
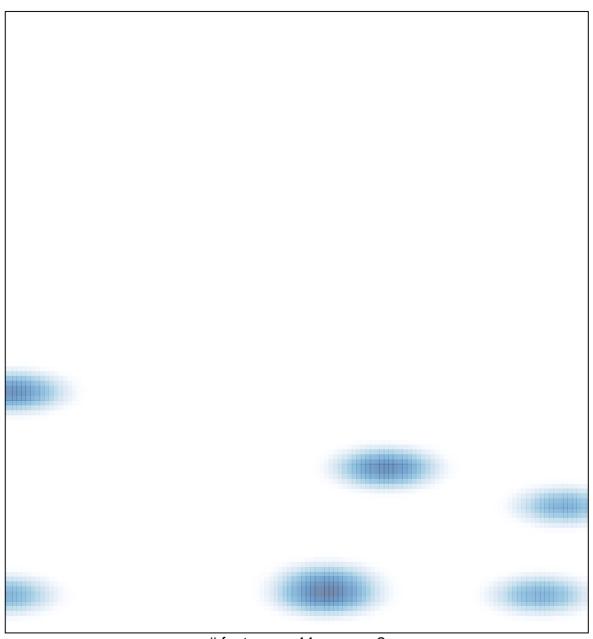
## REACTOME\_HDL\_MEDIATED\_LIPID\_TRANSPORT



# features = 11 chi-square p = 0.78

## REACTOME\_HDL\_MEDIATED\_LIPID\_TRANSPORT



# features = 11, max = 2