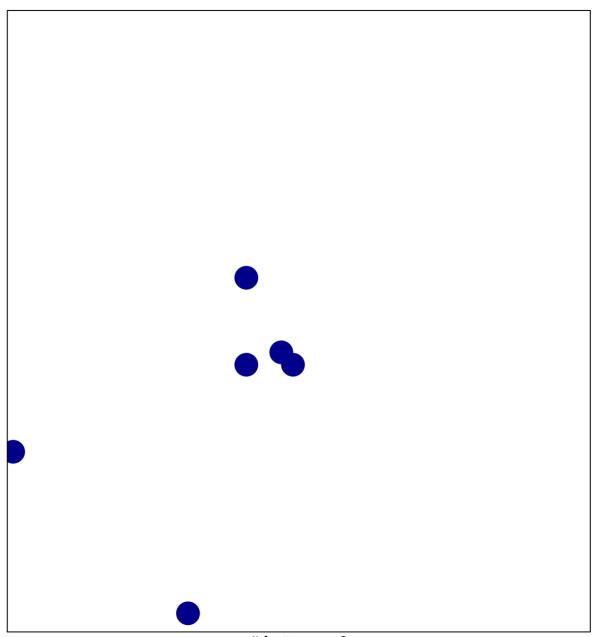
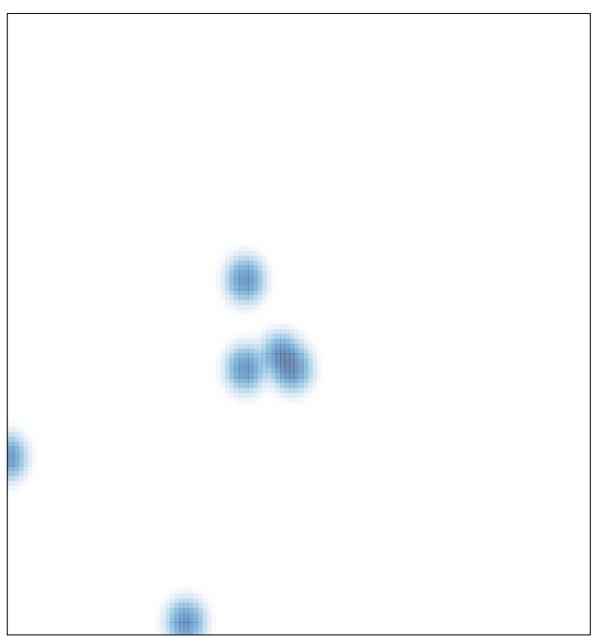
## REACTOME\_GPCR\_LIGAND\_BINDING



# features = 6 chi-square p = 0.83

## REACTOME\_GPCR\_LIGAND\_BINDING



# features = 6, max = 1