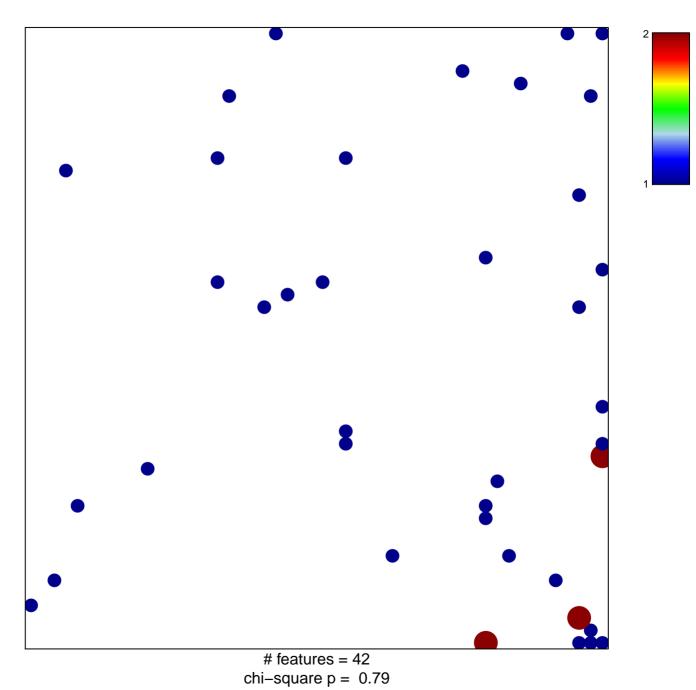
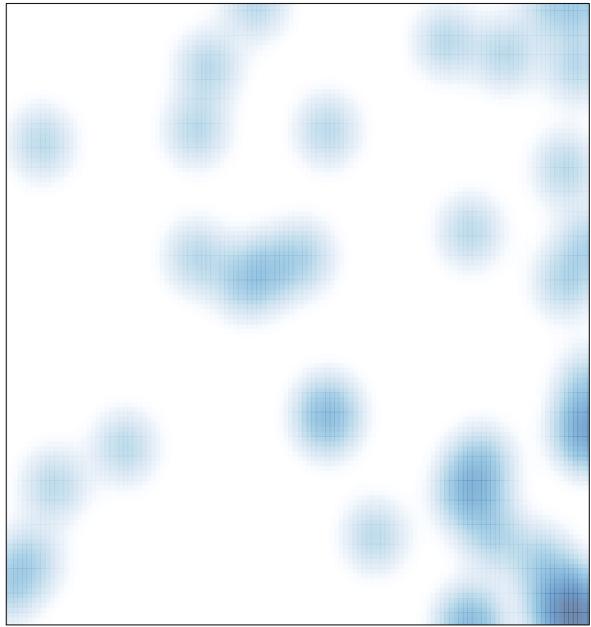
**B** cell differentiation



## **B** cell differentiation



# features = 42, max = 2