Task:

Prepare a datatset including paired cancer samples and then check TET1/2 SOX1,2,3, YY1 and YY2… Gene expression and then to compare the function difference between these genes. And then find P(A|B,C) network.

Biomarker panel for pathway activation, Idea:

1, build a databset,pair

2, selection differential markers for each pathway

3, build random foreast model to train the model

4, find another dataset to test the model