

## Node

Shape



Pathway



Signature set

Size



200



5

# genes in pathway

Color



-4 0 4  
Phenotype A Phenotype B

enrichment score

## Edge

Size



↑ overlap size  
↓

Color



Gene overlap



Signature set

## Options

default gsea n datasets

Color



default generic 1 dataset

Color



1 0  
q-value

default generic n datasets

Color

