

Running Enrichment Score

0.0

-0.2

-0.4

-0.6

-0.8

Aminoacyl-tRNA biosynthesis

Base excision repair

DNA replication

Homologous recombination

Mismatch repair

Ranked List Metric

3

2

1

0

-1

4000

8000

12000

16000

Rank in Ordered Dataset

