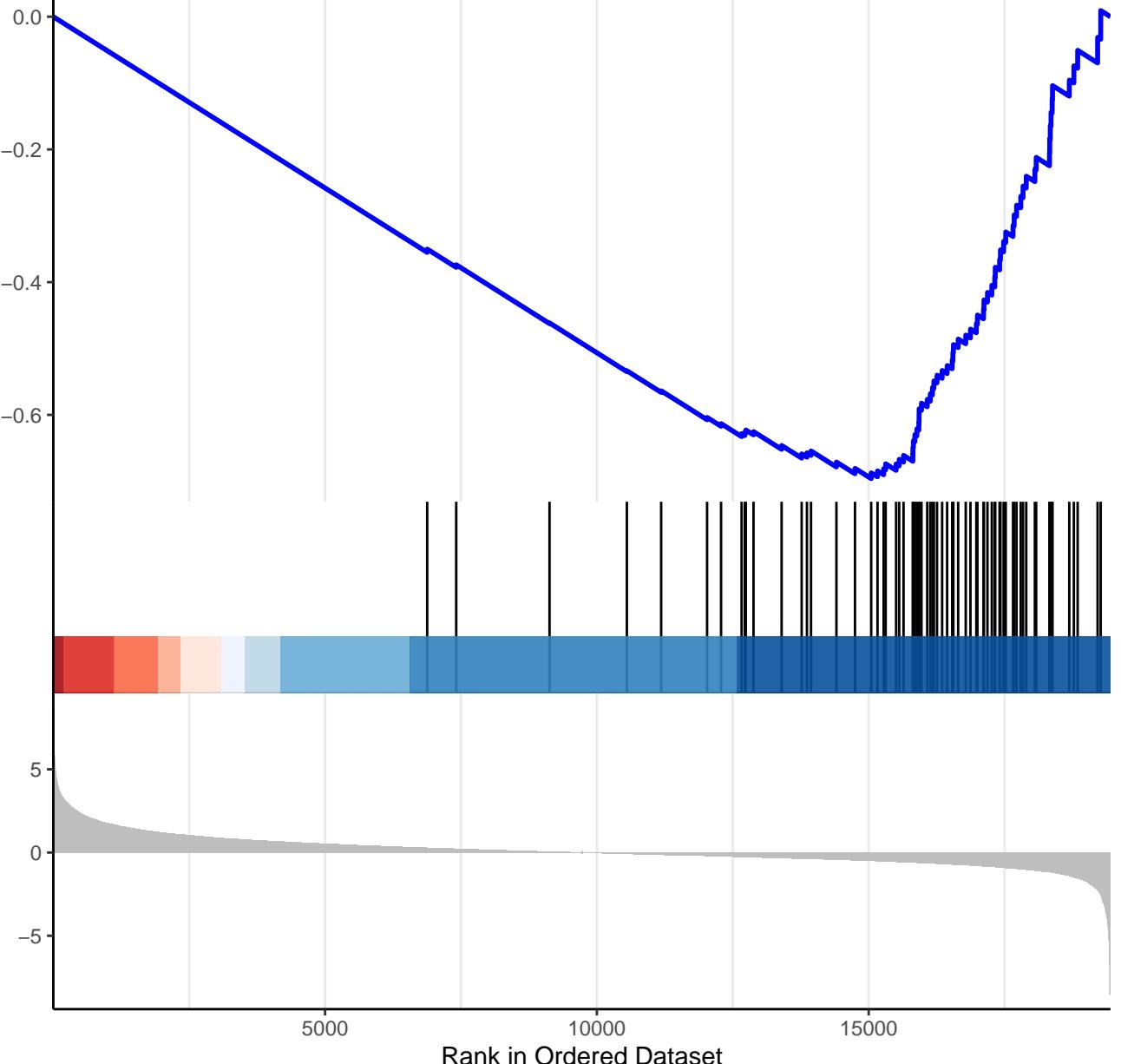


Ribosome biogenesis in eukaryotes

Running Enrichment Score



Ribosome

Running Enrichment Score

0.0
-0.2

-0.4

5

0

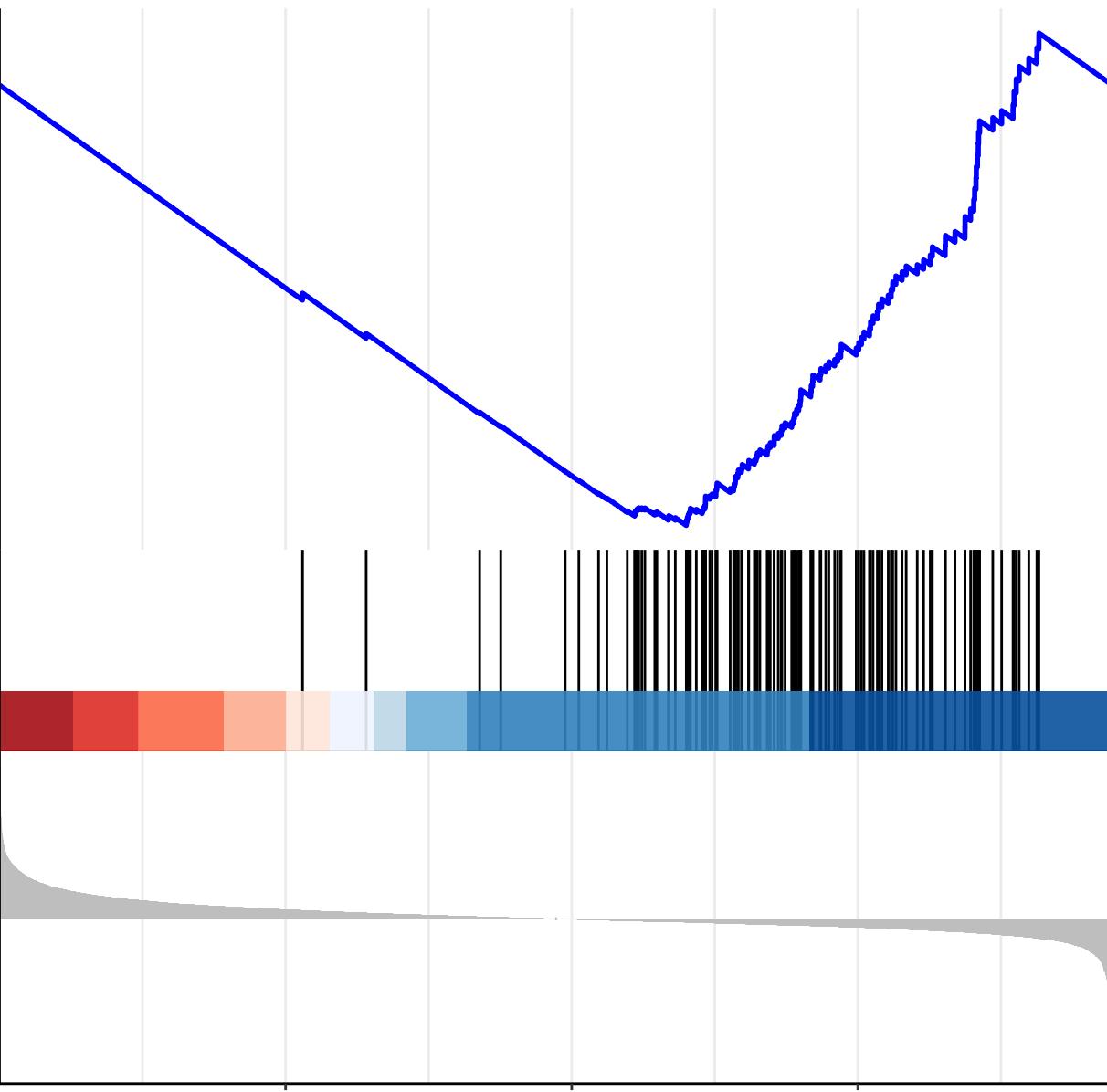
-5

Ranked List Metric

5000 10000 15000

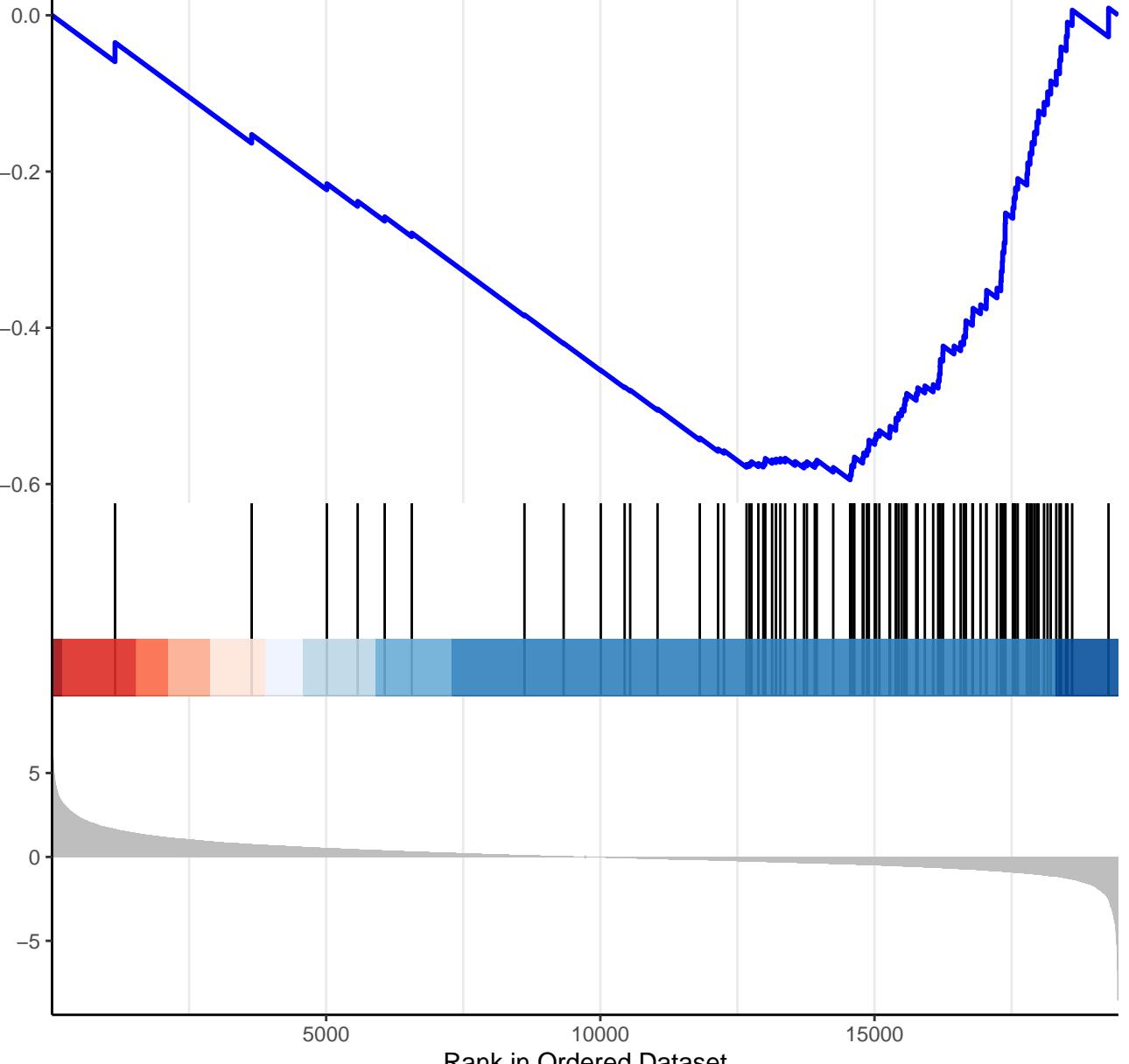
Rank in Ordered Dataset

Pvalue: 0.000 P.adjust: 0.000



Nucleocytoplasmic transport

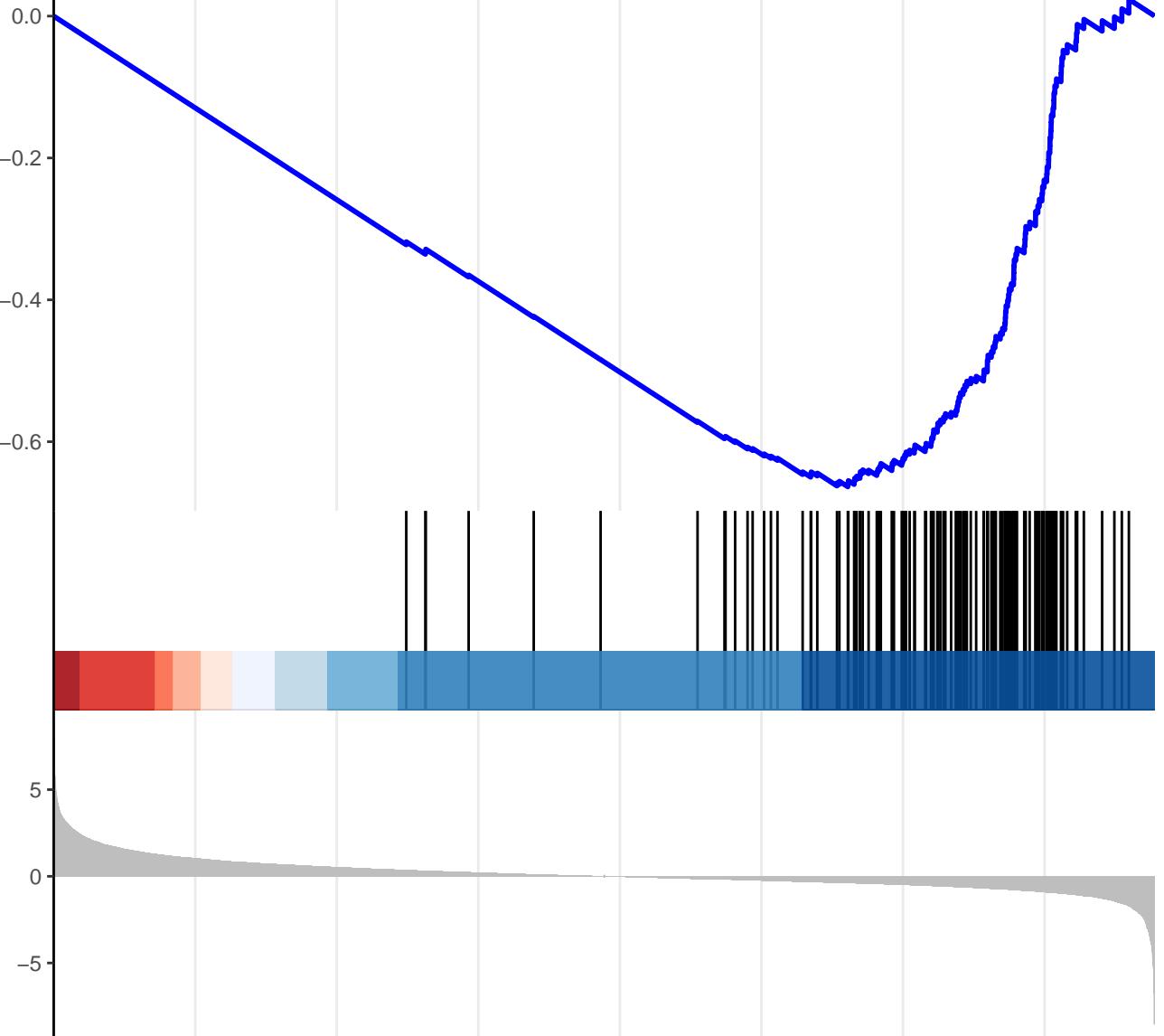
Running Enrichment Score



Pvalue: 0.000 _ P.adjust: 0.000

Spliceosome

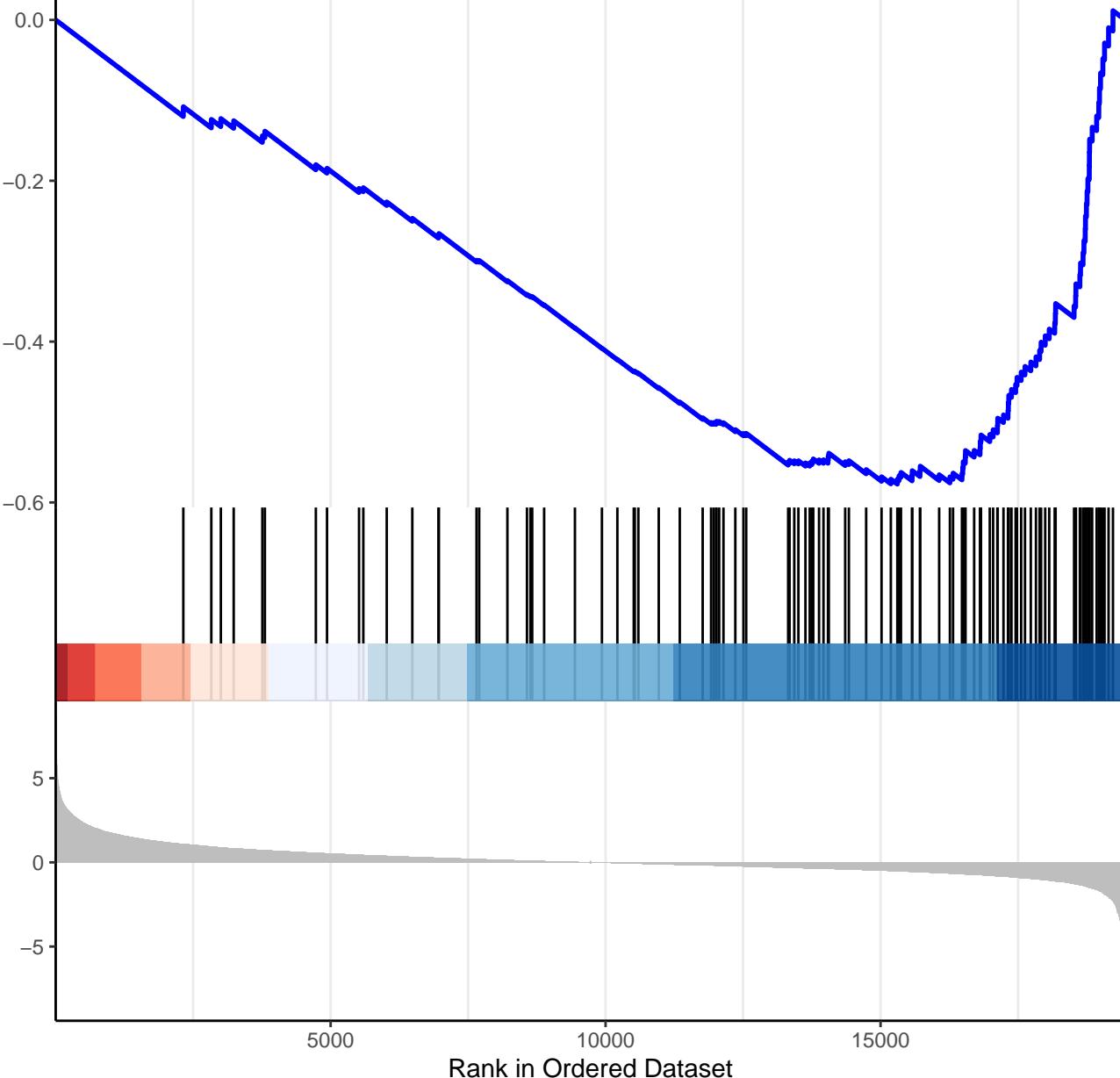
Running Enrichment Score



Pvalue: 0.000 _ P.adjust: 0.000

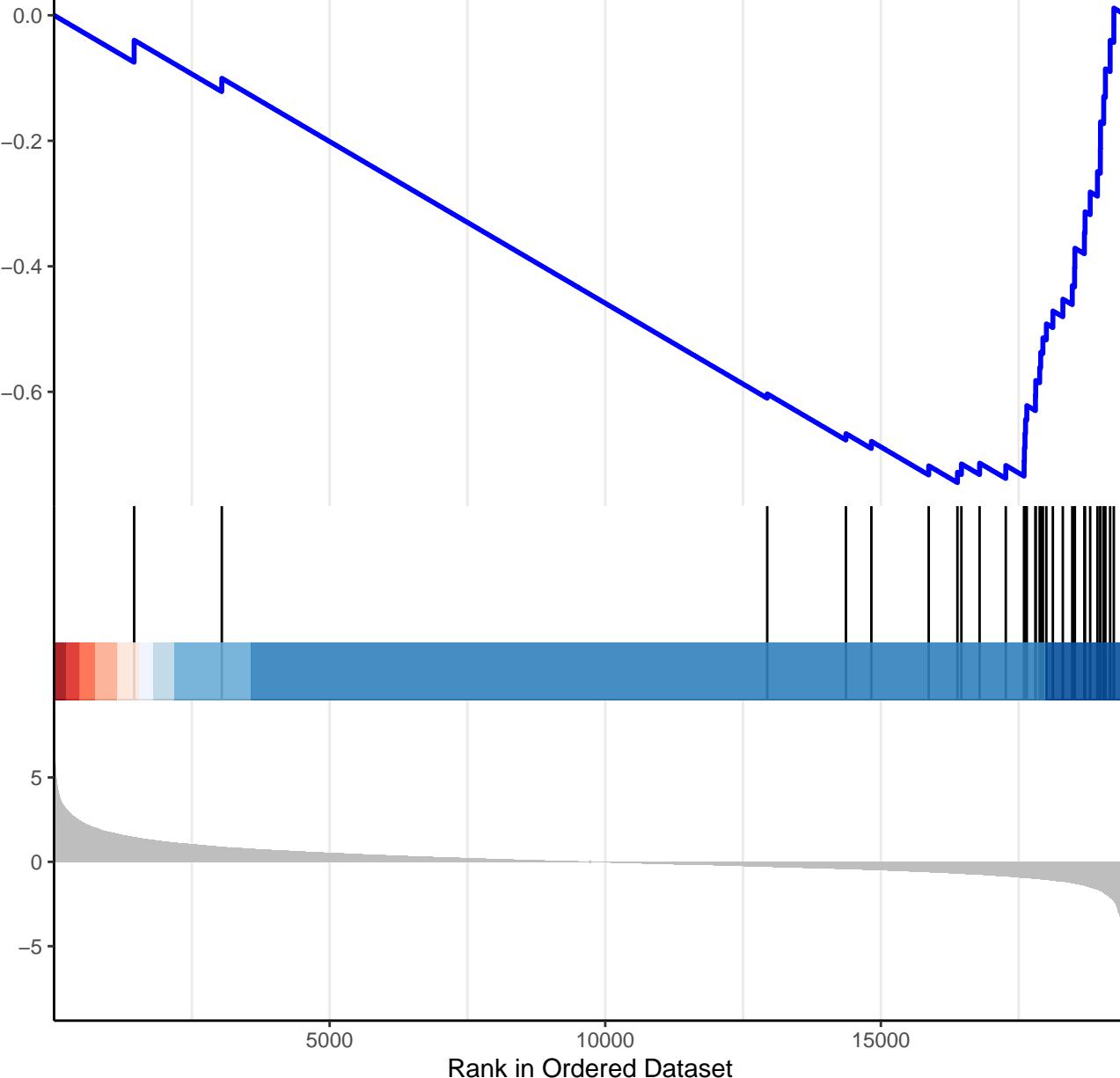
Cell cycle

Running Enrichment Score



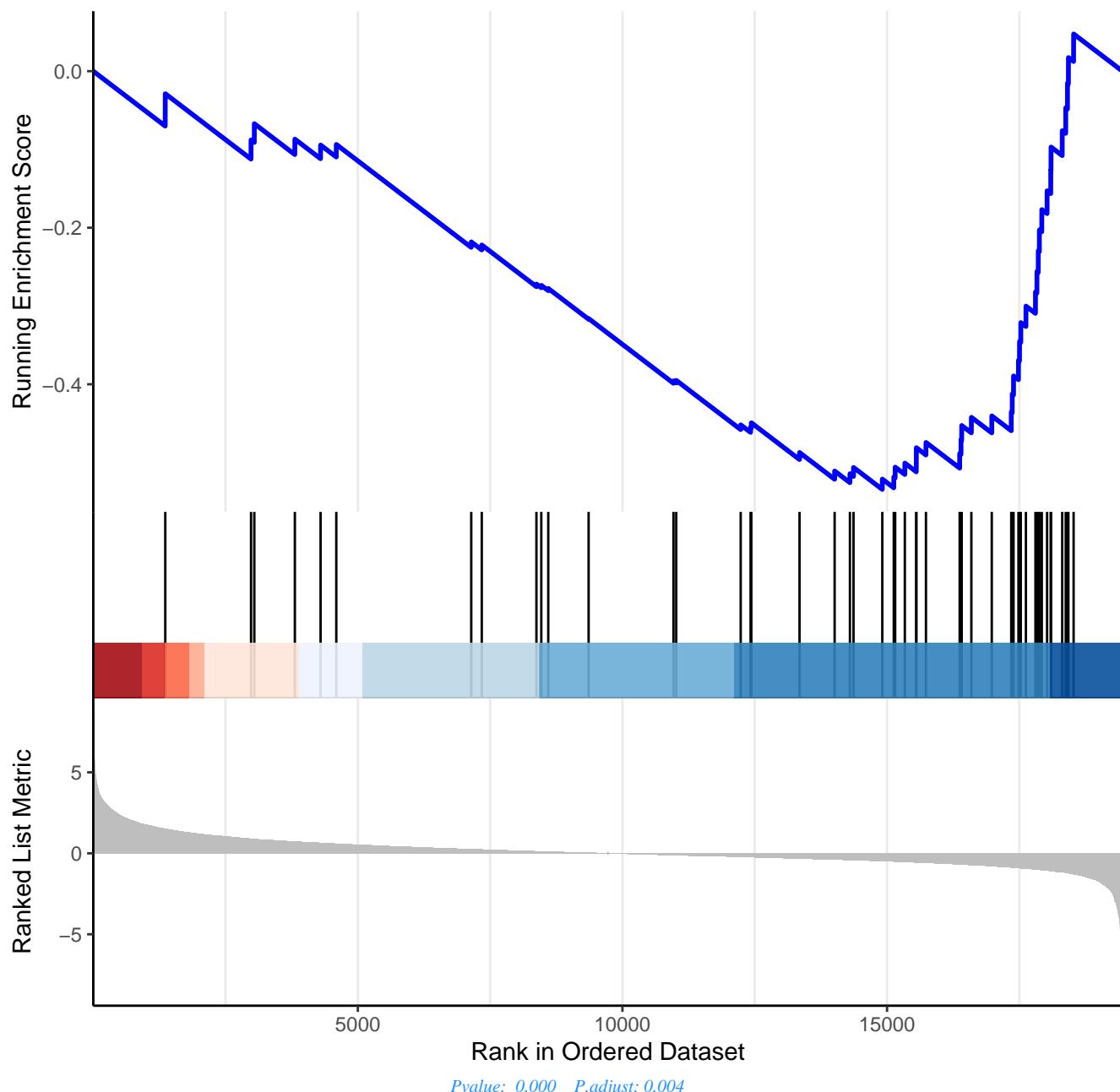
DNA replication

Running Enrichment Score



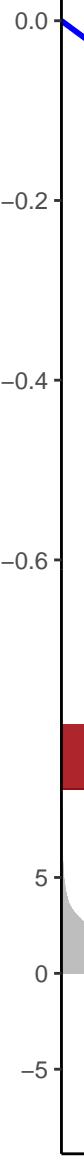
Pvalue: 0.000 _ P.adjust: 0.000

Fanconi anemia pathway



Base excision repair

Running Enrichment Score



Ranked List Metric



Rank in Ordered Dataset

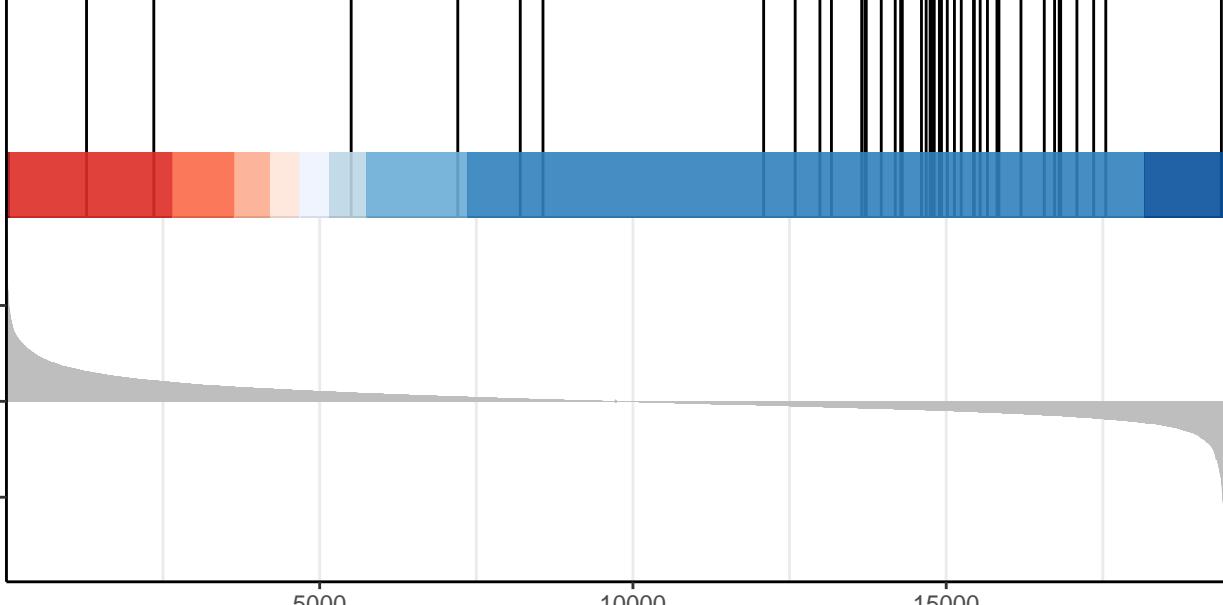
Pvalue: 0.000 _ P.adjust: 0.007

Proteasome

Running Enrichment Score



Ranked List Metric



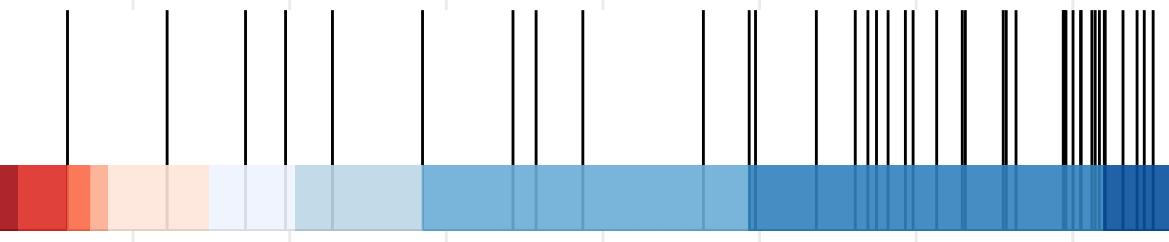
Pvalue: 0.001 _ P.adjust: 0.010

Rank in Ordered Dataset

Homologous recombination

Running Enrichment Score

0.0
-0.2
-0.4



Ranked List Metric

5
0
-5

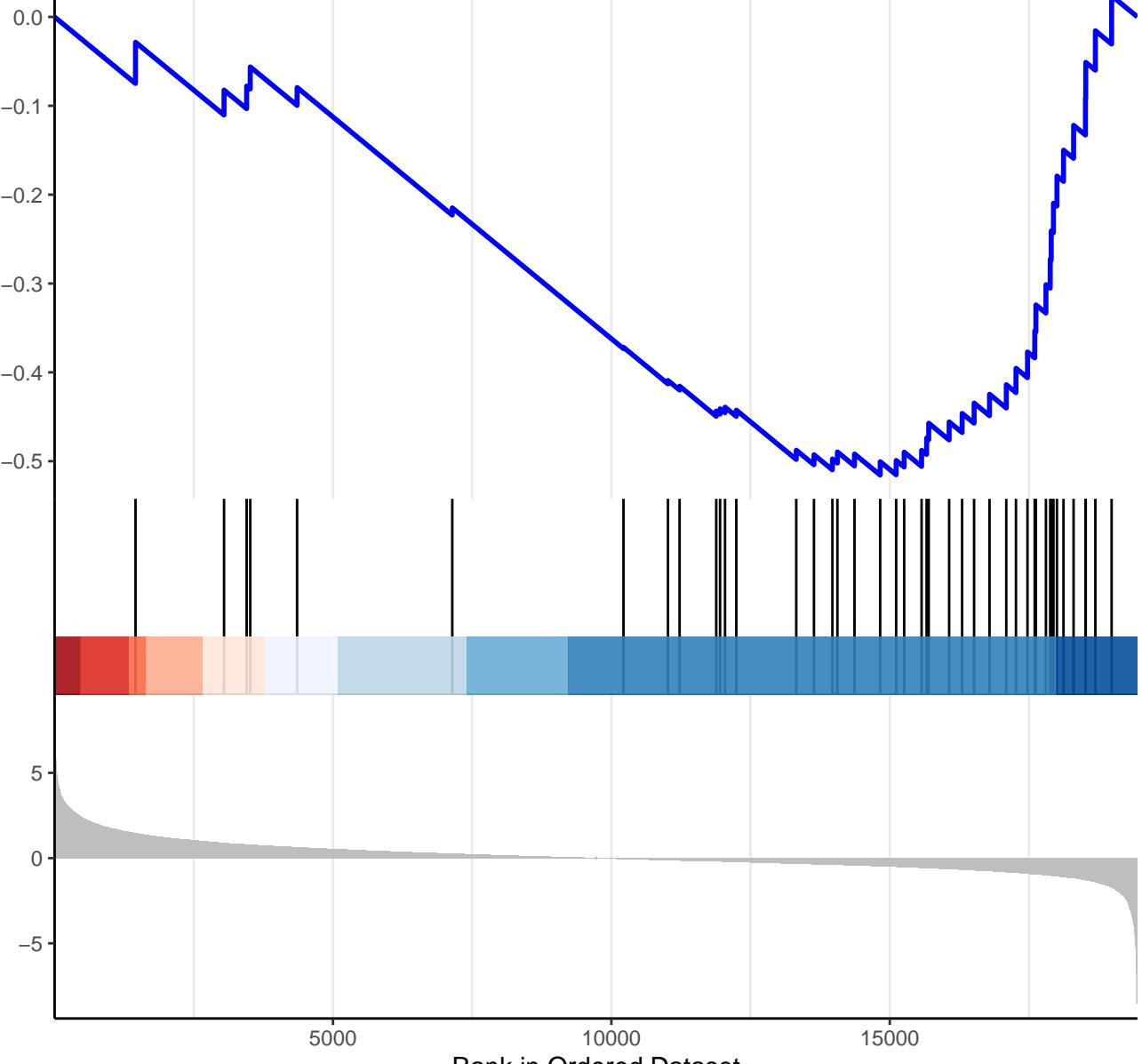
5000 10000 15000

Rank in Ordered Dataset

Pvalue: 0.001 _ P.adjust: 0.011

Nucleotide excision repair

Running Enrichment Score



Pvalue: 0.001 _ P.adjust: 0.012

Aminoacyl-tRNA biosynthesis

Running Enrichment Score

0.0
-0.1
-0.2
-0.3
-0.4
-0.5

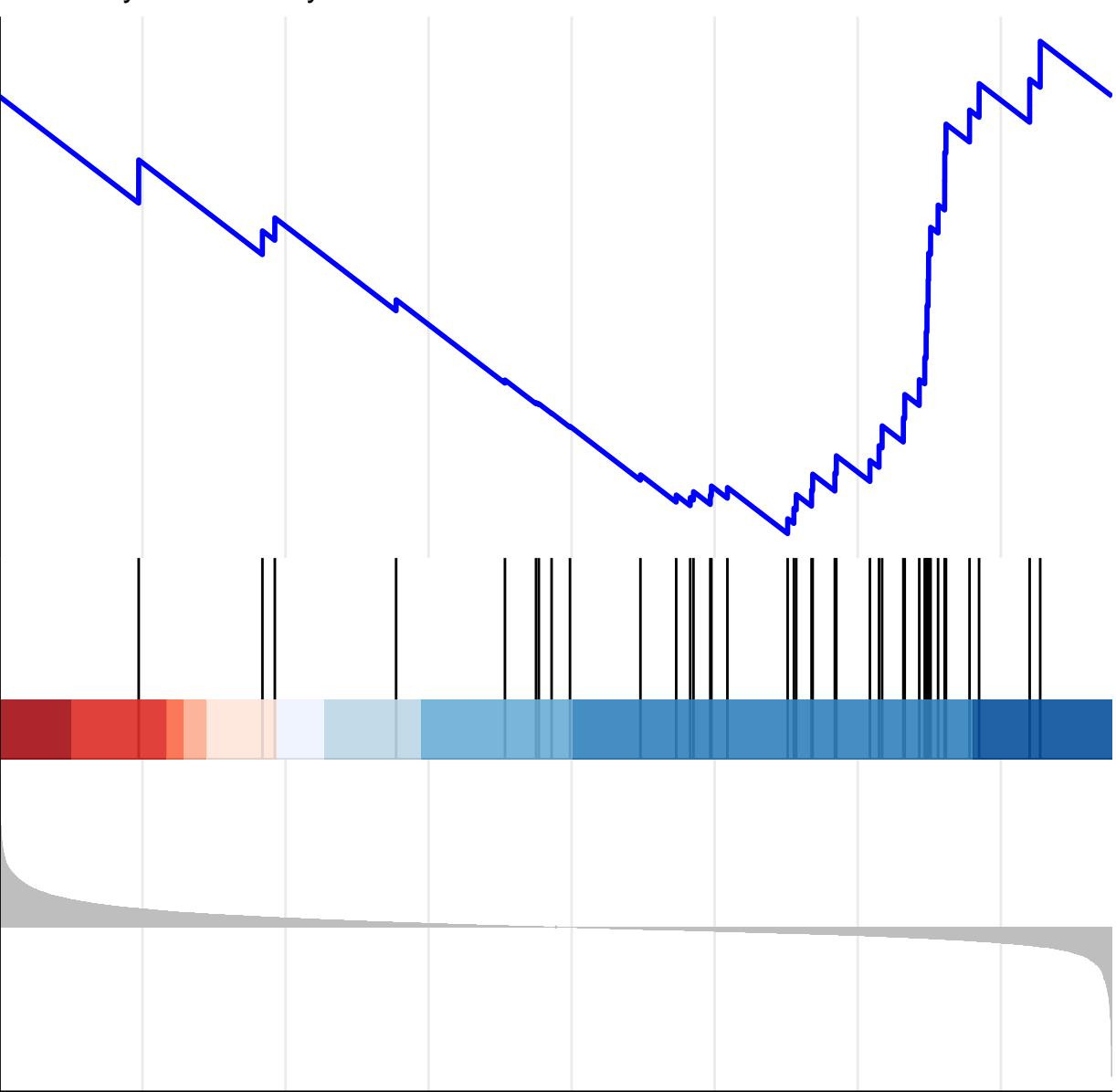
Ranked List Metric

5
0
-5

5000 10000 15000

Rank in Ordered Dataset

Pvalue: 0.001 _ P.adjust: 0.013



RNA polymerase

Running Enrichment Score

0.0

-0.2

-0.4

Ranked List Metric

5

0

-5

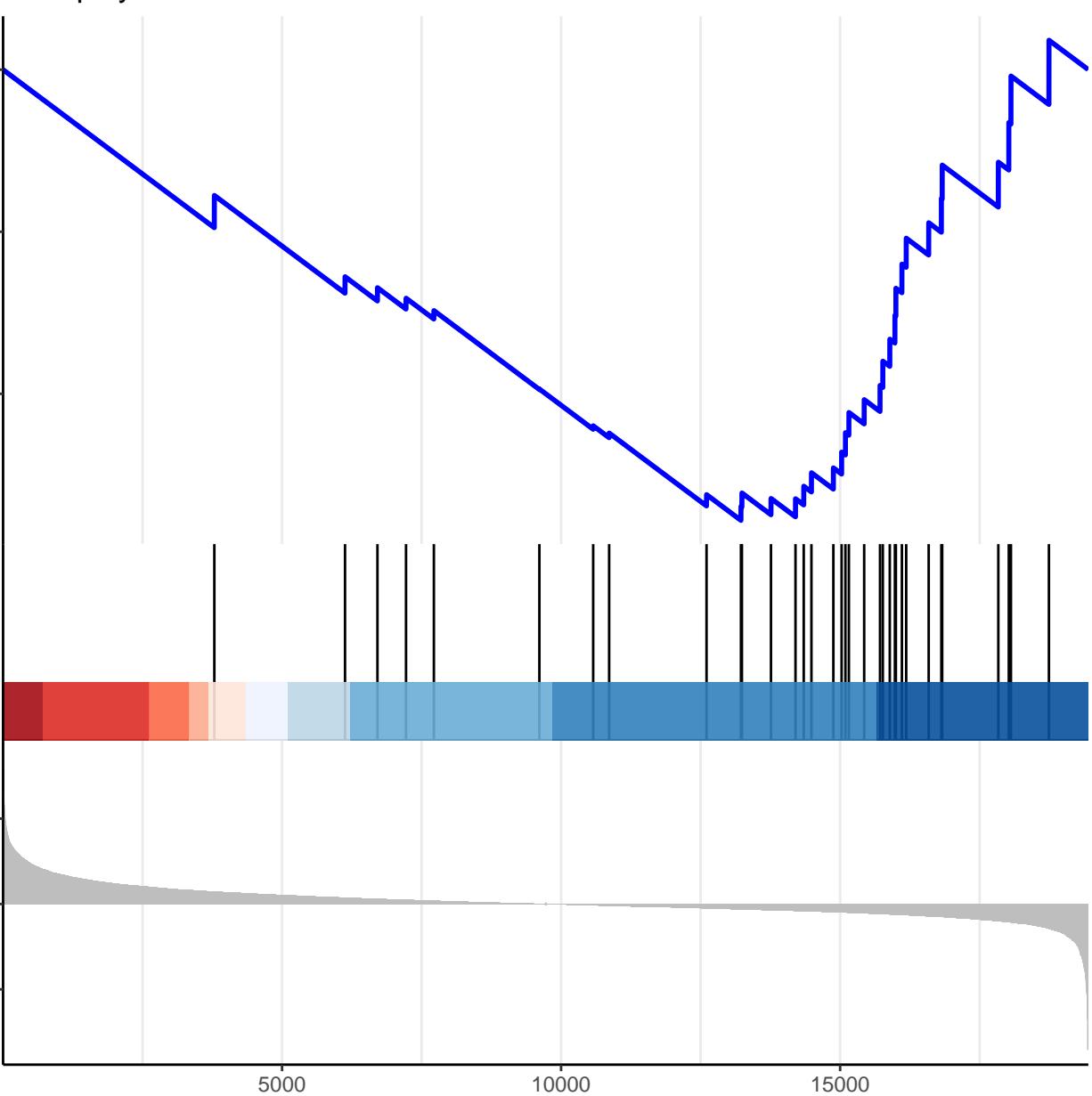
5000

10000

15000

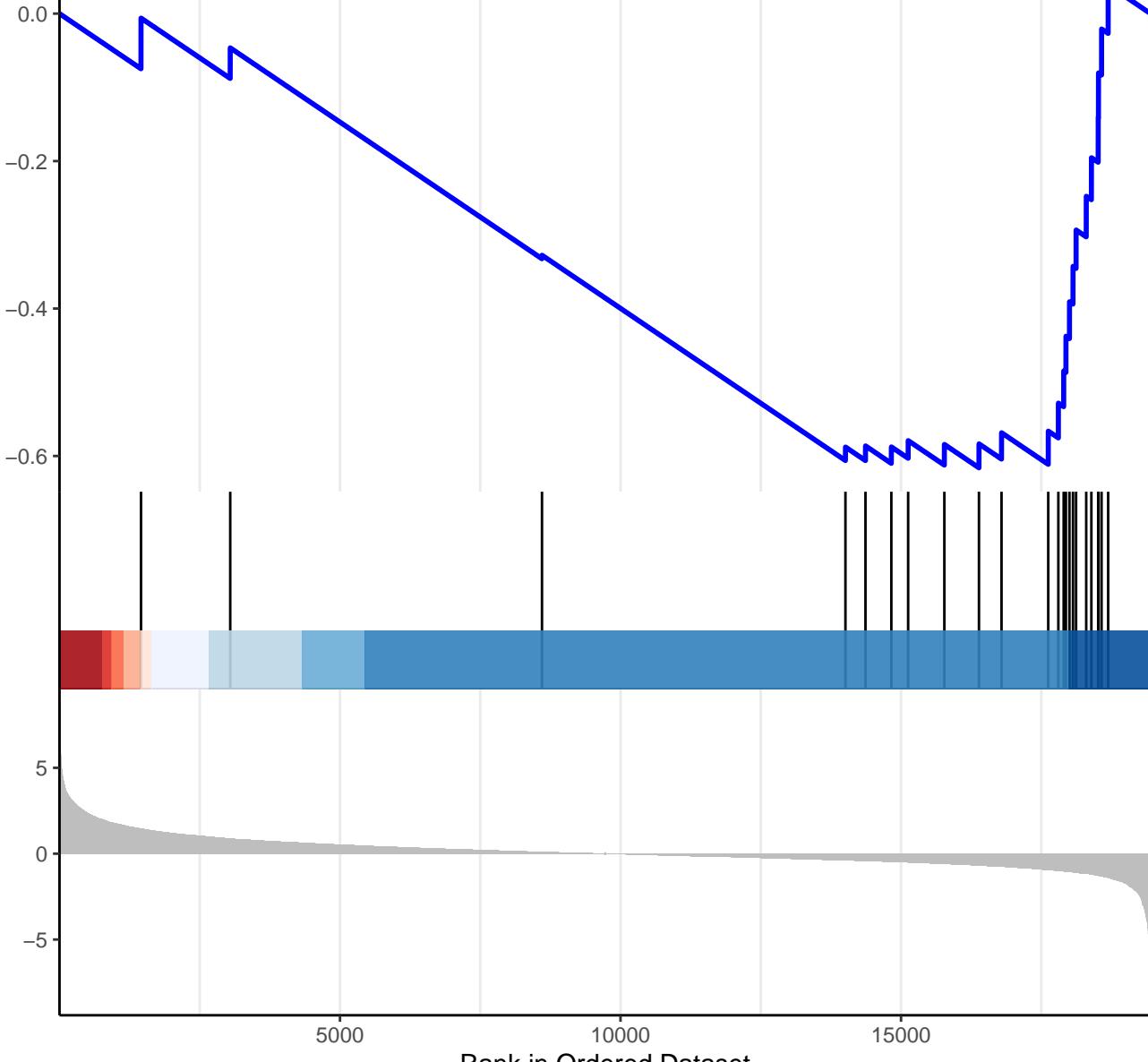
Rank in Ordered Dataset

Pvalue: 0.002 _ P.adjust: 0.019



Mismatch repair

Running Enrichment Score



Pvalue: 0.002 _ P.adjust: 0.021