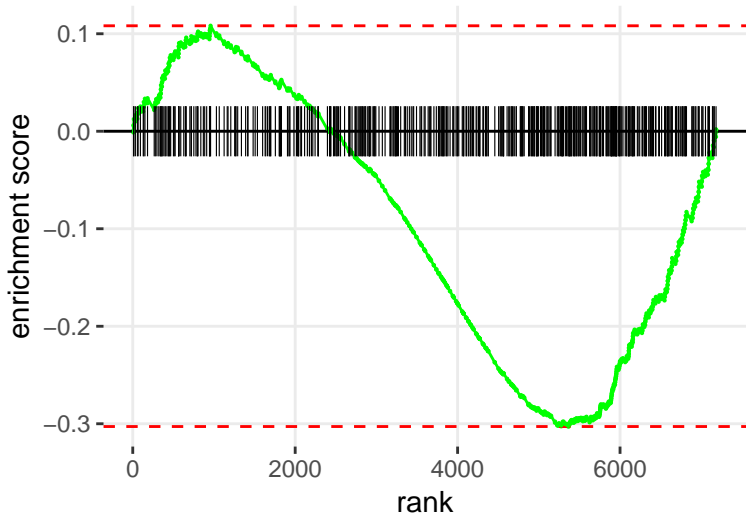


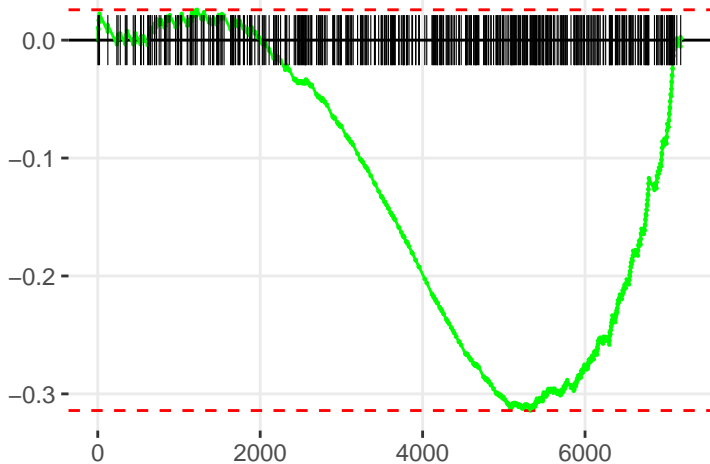
NEGATIVE REGULATION OF SIGNALING%GOBP%GO:0023057



PHOSPHORYLATION%GOBP%GO:0016310

enrichment score

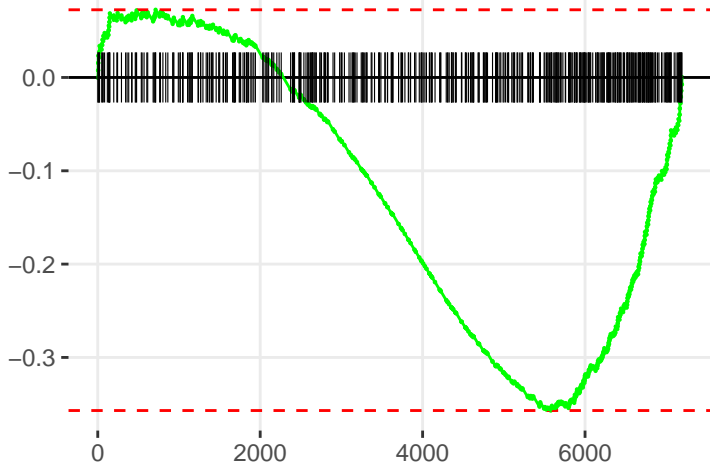
rank



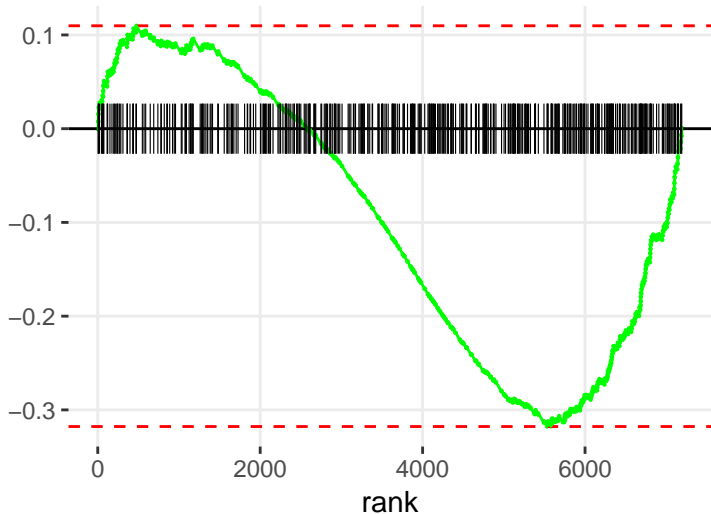
SECRETION%GOBP%GO:0046903

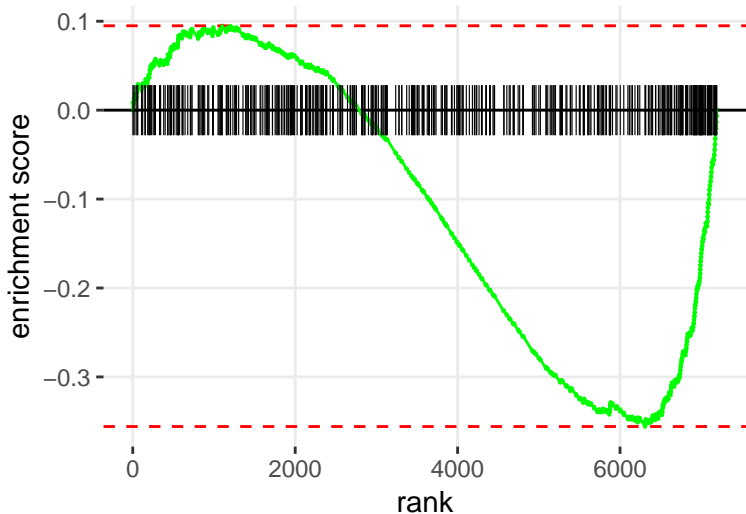
enrichment score

rank

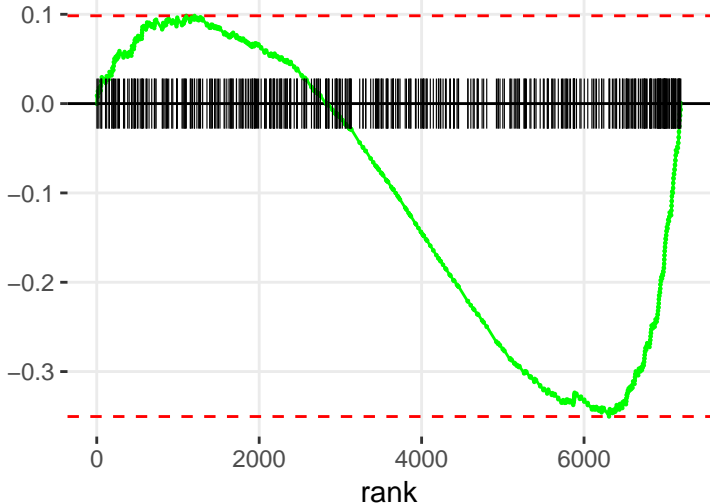


enrichment score



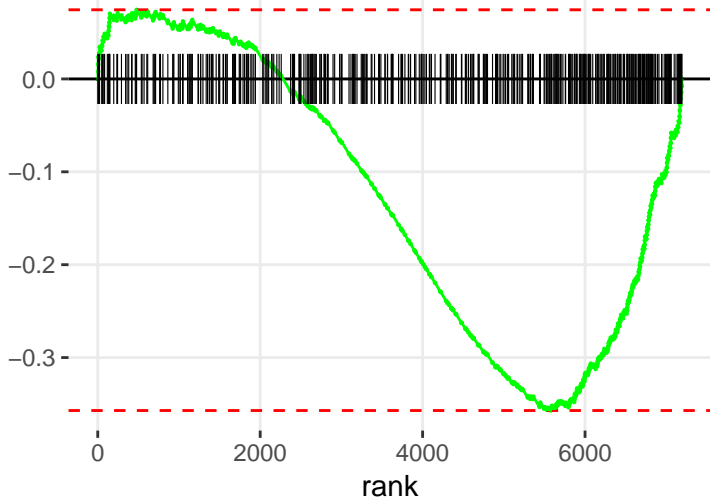


enrichment score

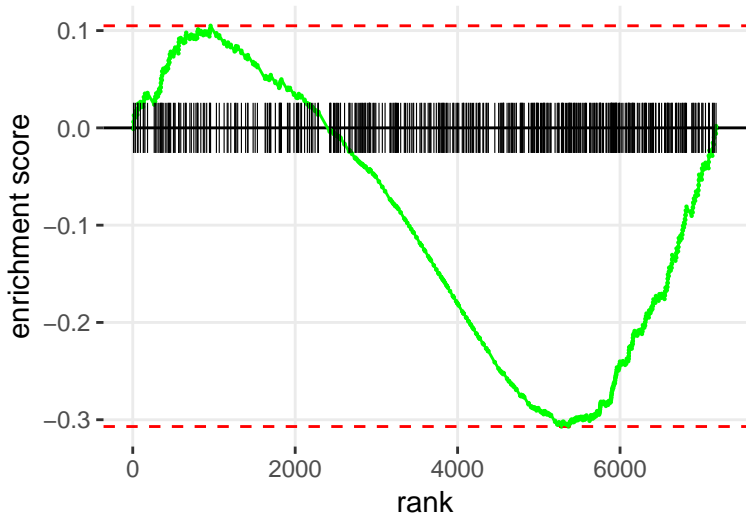


SECRETION BY CELL%GOBP%GO:0032940

enrichment score

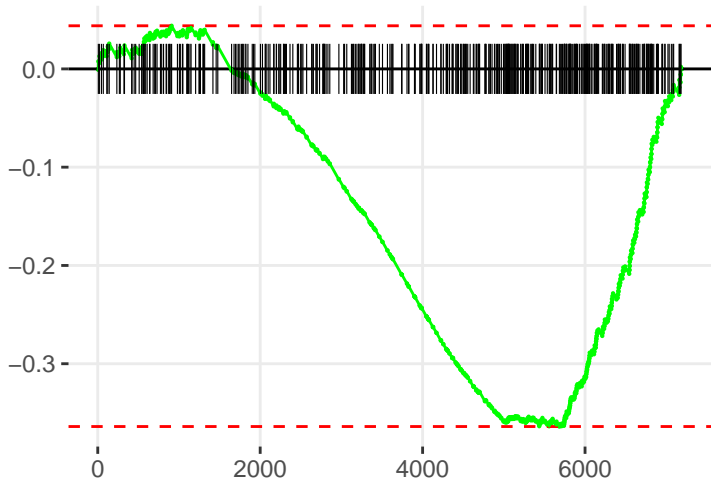


NEGATIVE REGULATION OF SIGNAL TRANSDUCTION%GOBP%GO:0009968

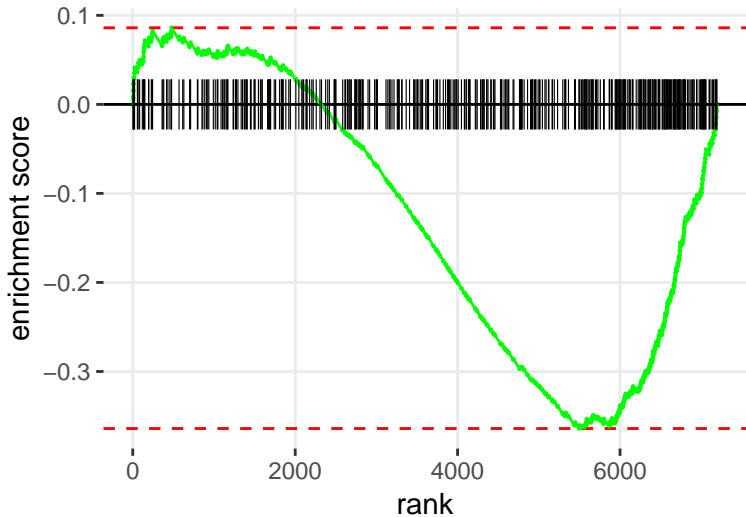


enrichment score

rank



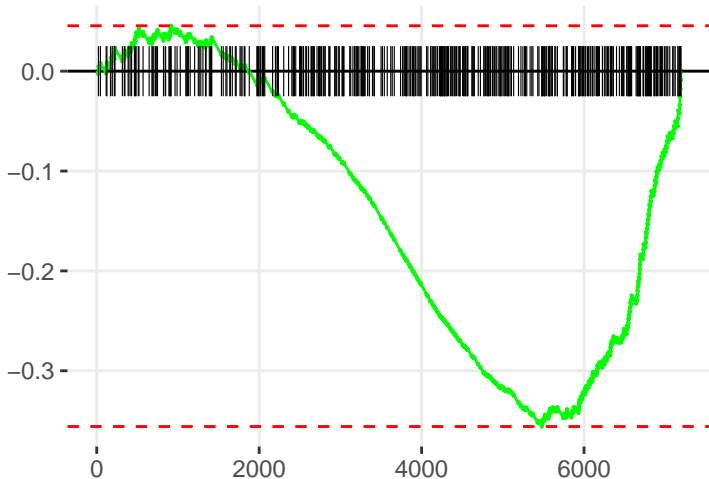
CELL ACTIVATION%GOBP%GO:0001775

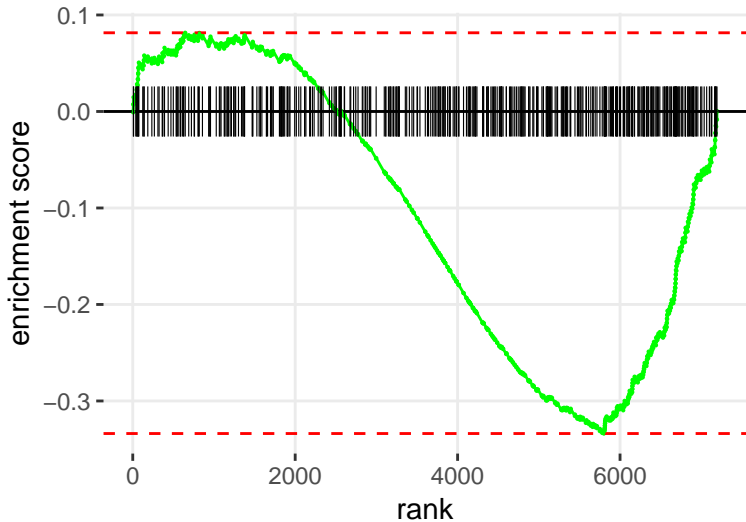


REGULATION OF CELLULAR COMPONENT BIOGENESIS%GOBP%GO:0044087

enrichment score

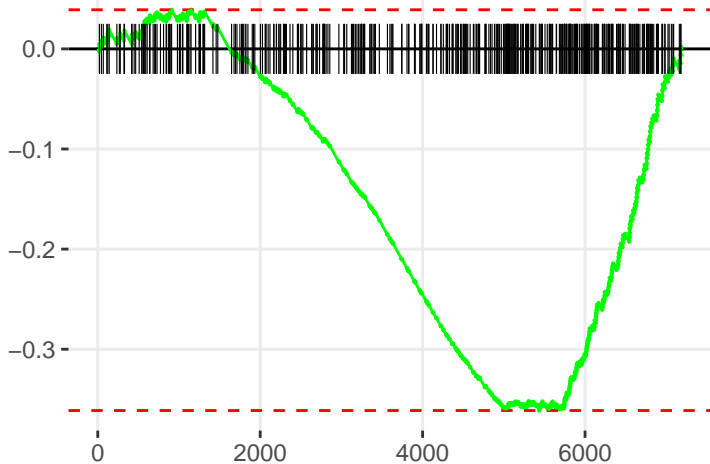
rank

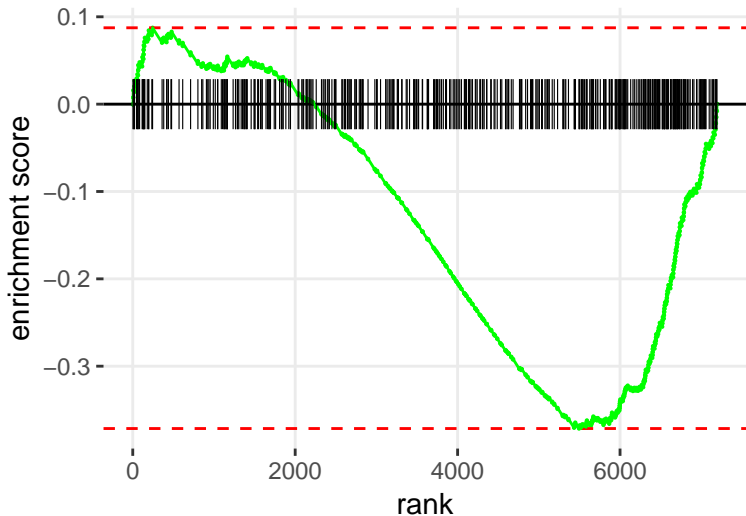


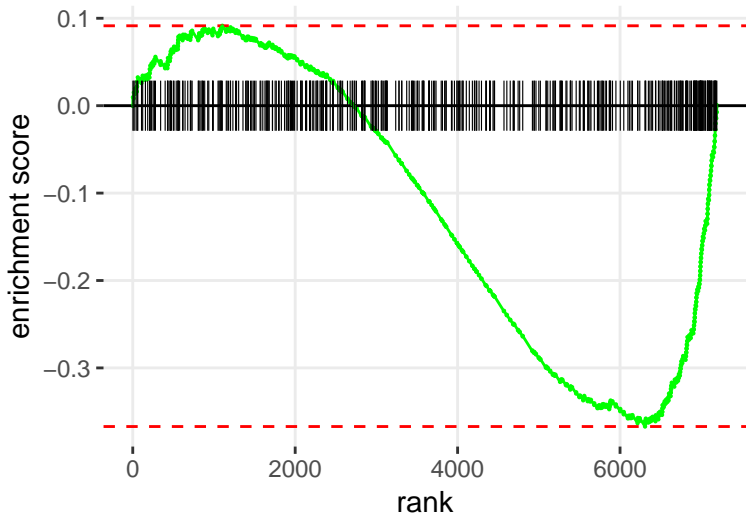


enrichment score

rank

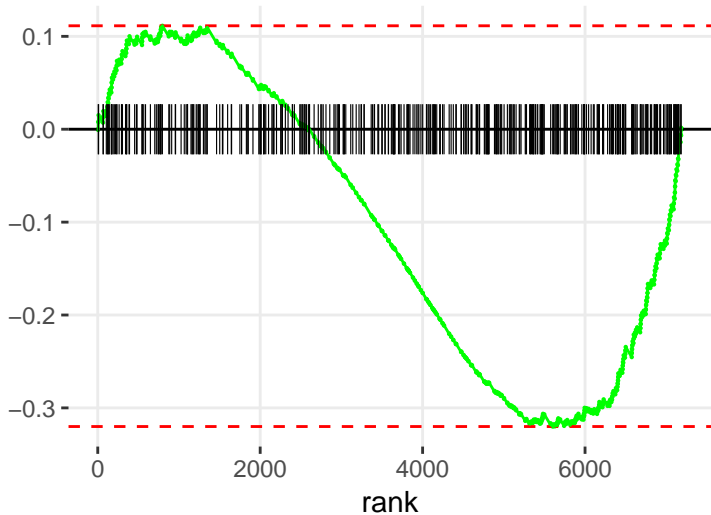






RESPONSE TO ENDOGENOUS STIMULUS%GOBP%GO:0009719

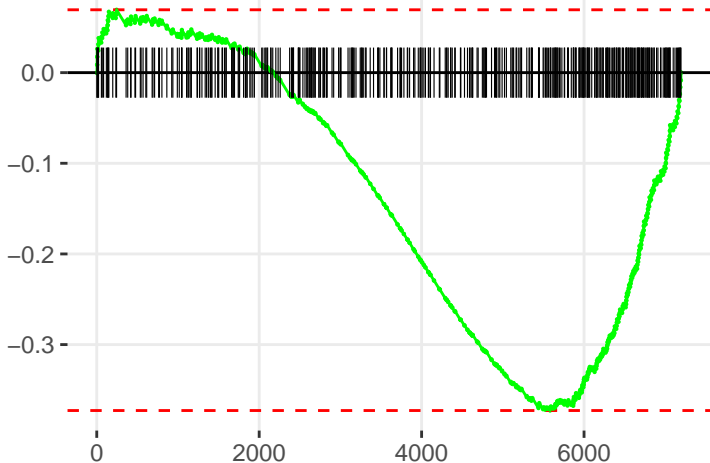
enrichment score



EXOCYTOSIS%GOBP%GO:0006887

enrichment score

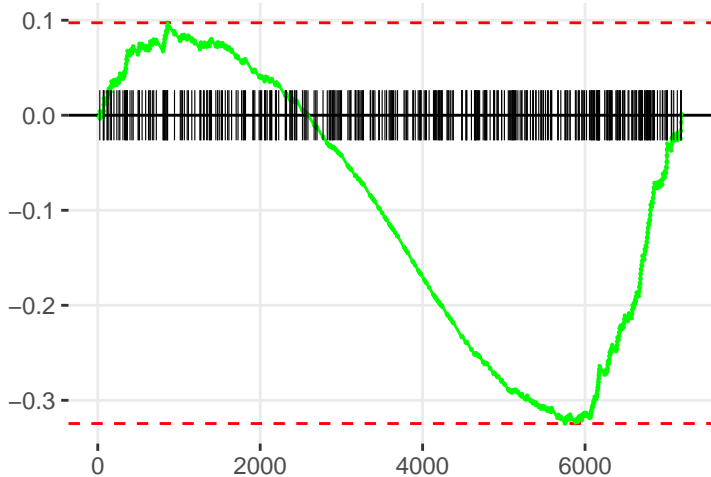
rank



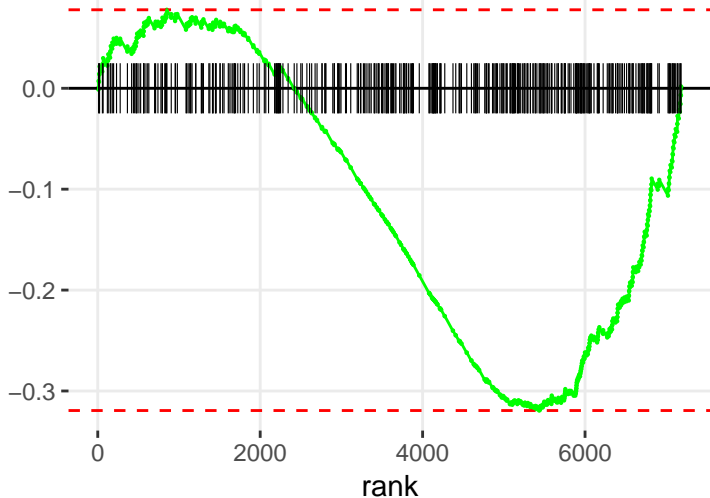
NEUROGENESIS%GOBP%GO:0022008

enrichment score

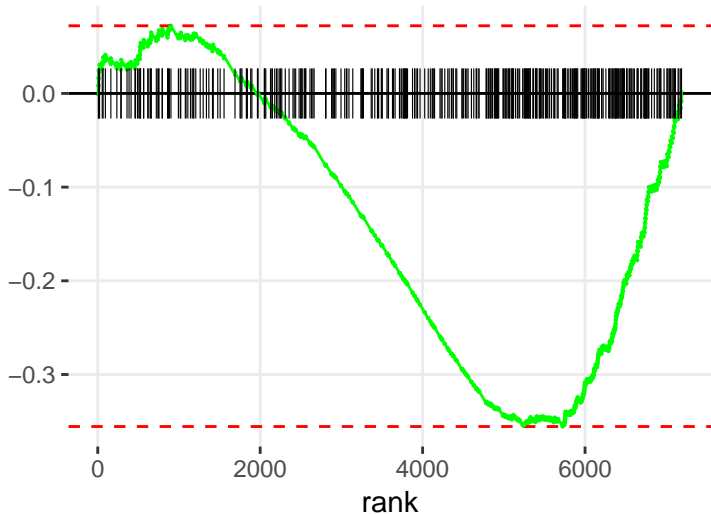
rank



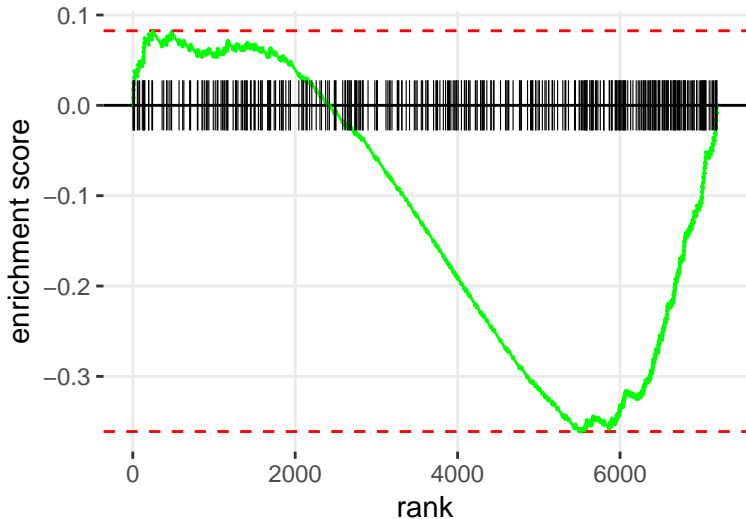
enrichment score



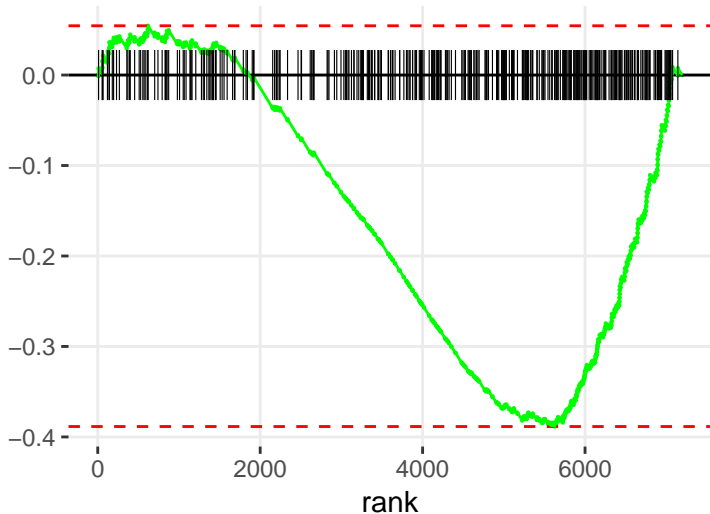
enrichment score



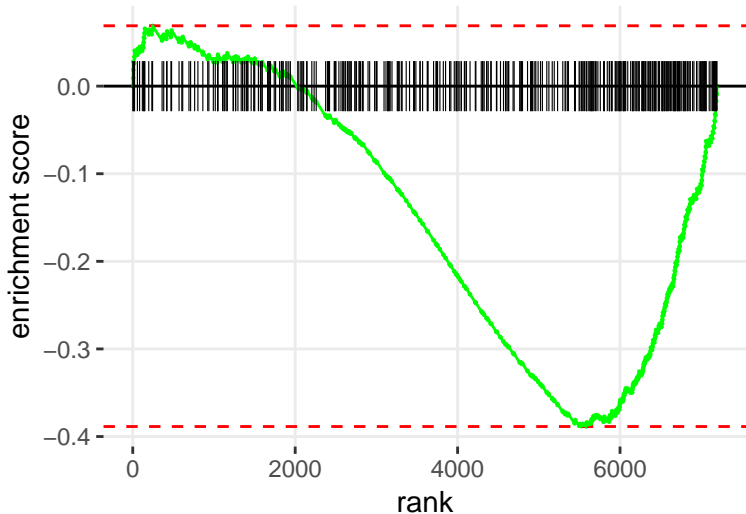
LEUKOCYTE ACTIVATION%GOBP%GO:0045321

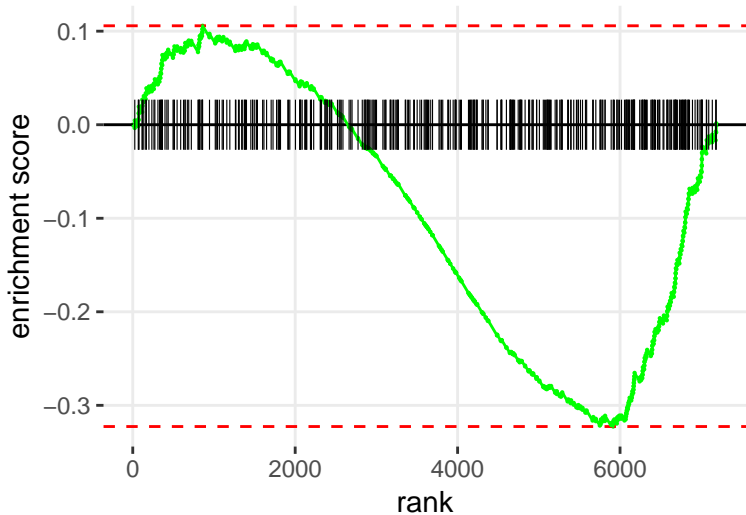


enrichment score

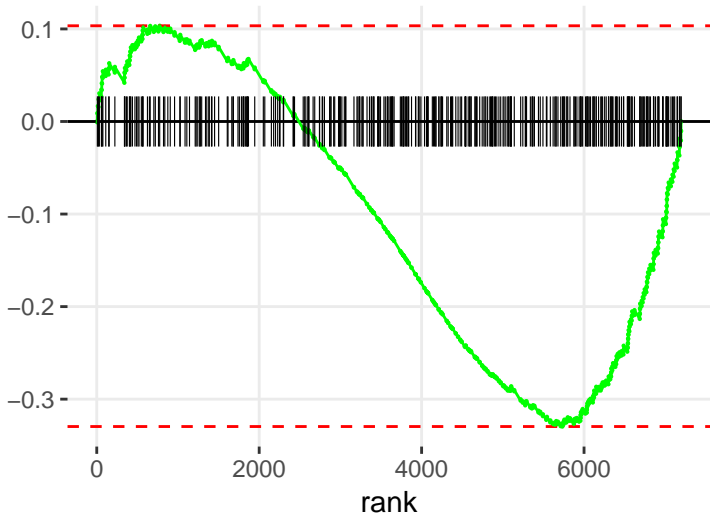


REGULATED EXOCYTOSIS%GOBP%GO:0045055

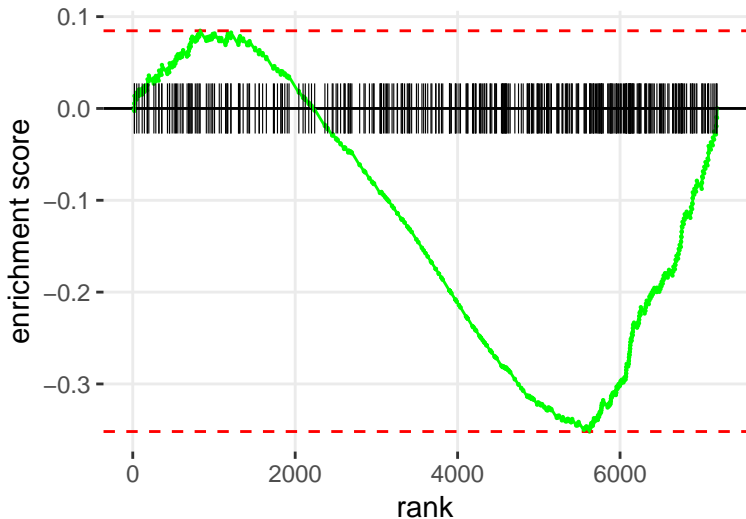




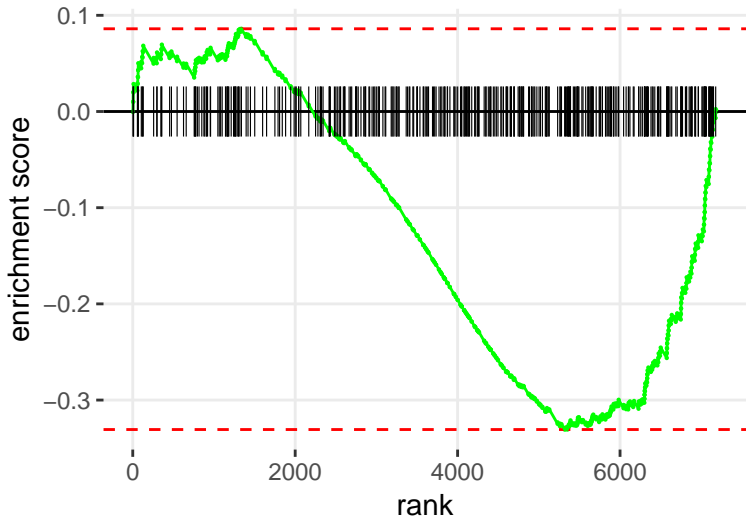
enrichment score



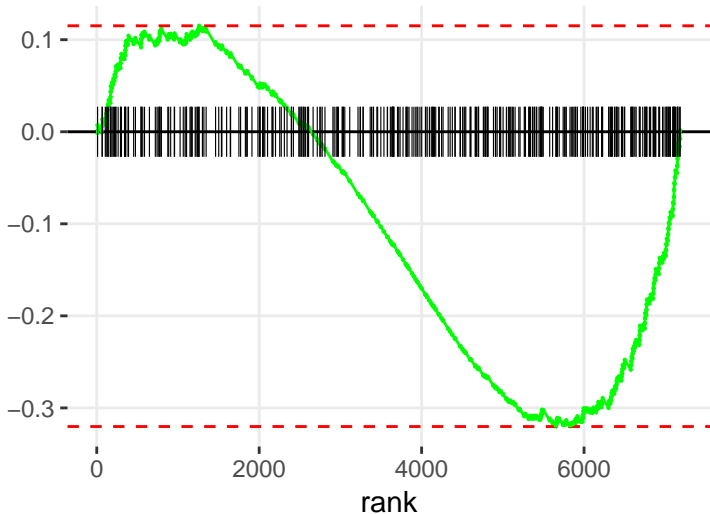
PROTEIN CATABOLIC PROCESS%GOBP%GO:0030163



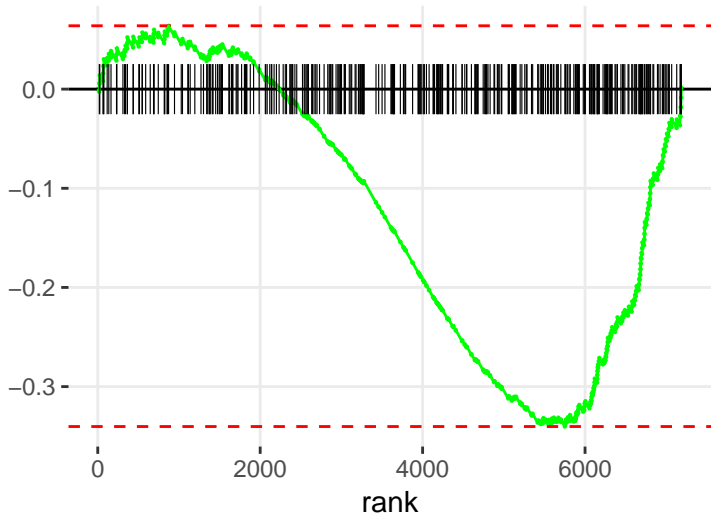
RESPONSE TO OXYGEN-CONTAINING COMPOUND%GOBP%GO:1901700



enrichment score



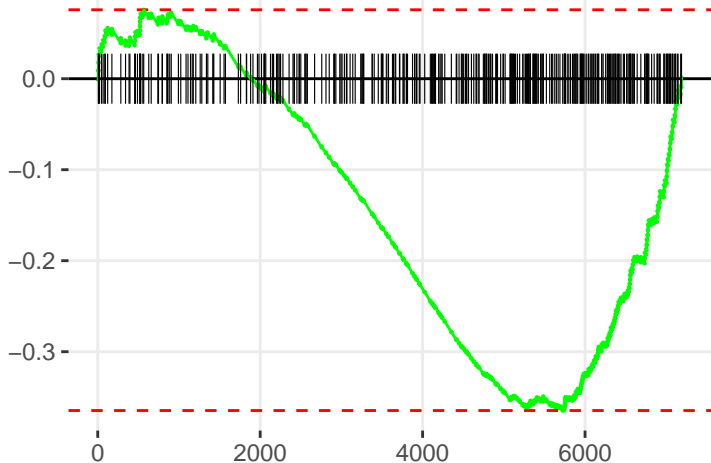
enrichment score



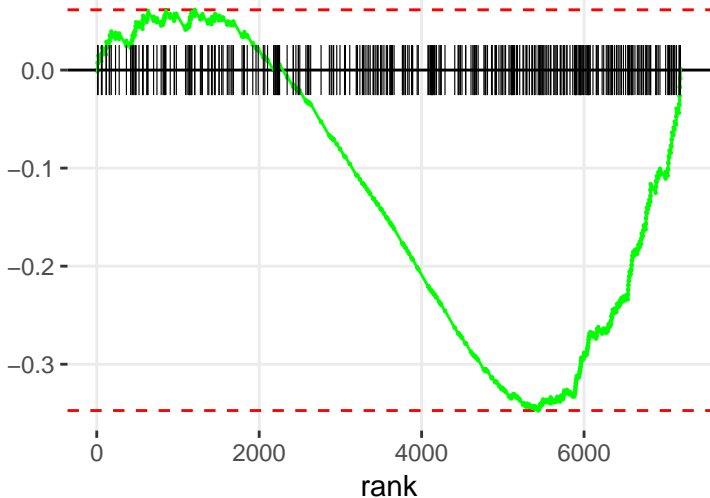
RESPONSE TO CYTOKINE%GOBP%GO:0034097

enrichment score

rank

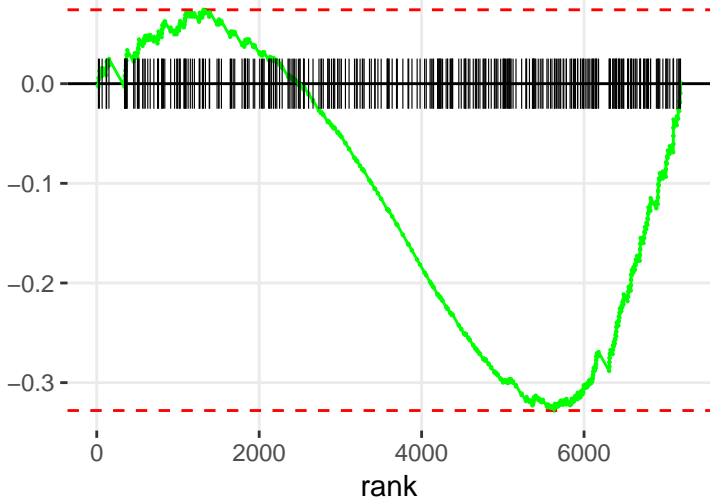


enrichment score



NEGATIVE REGULATION OF CELL DEATH%GOBP%GO:0060548

enrichment score



EGFR1%IOB%EGFR1

enrichment score

rank

0.0

-0.1

-0.2

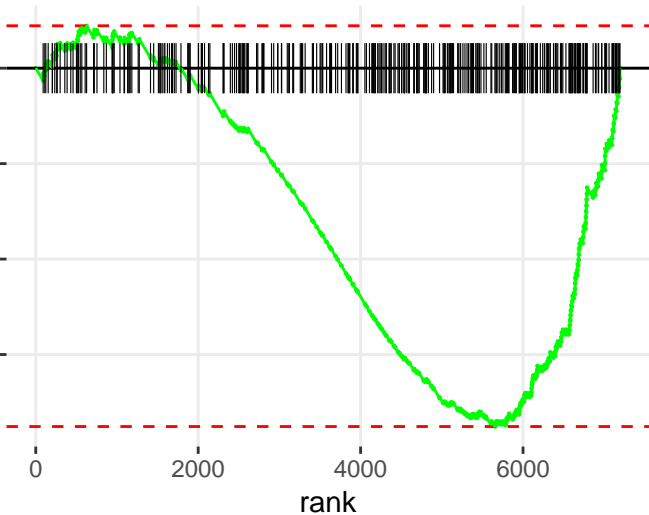
-0.3

0

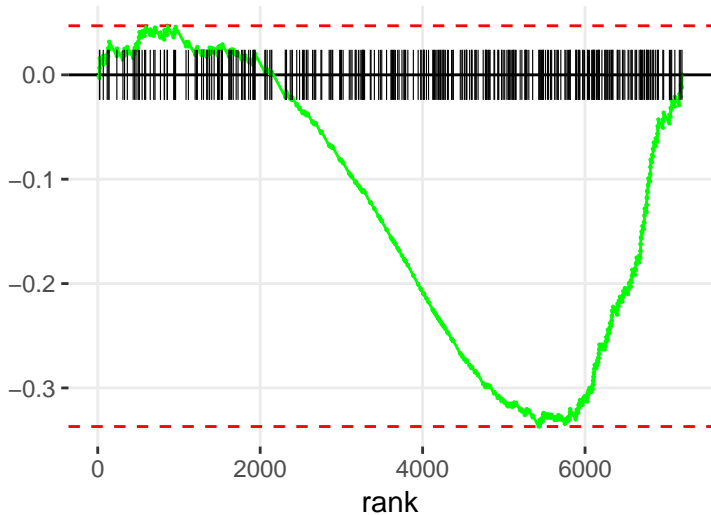
2000

4000

6000



enrichment score



EGFR1%NETPATH%EGFR1

enrichment score

rank

0.0

-0.1

-0.2

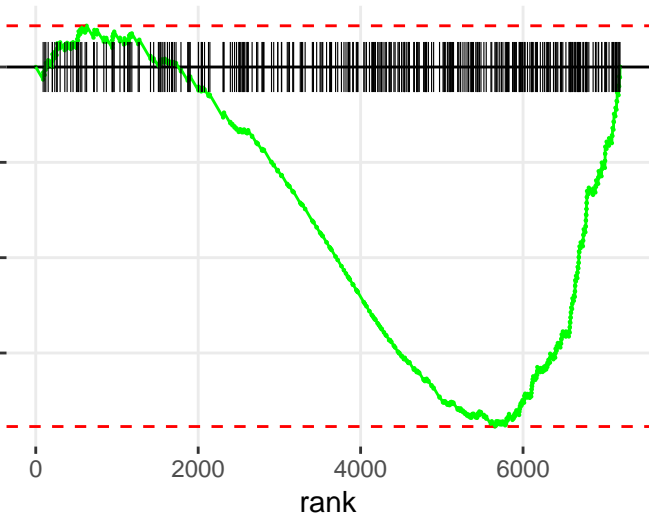
-0.3

0

2000

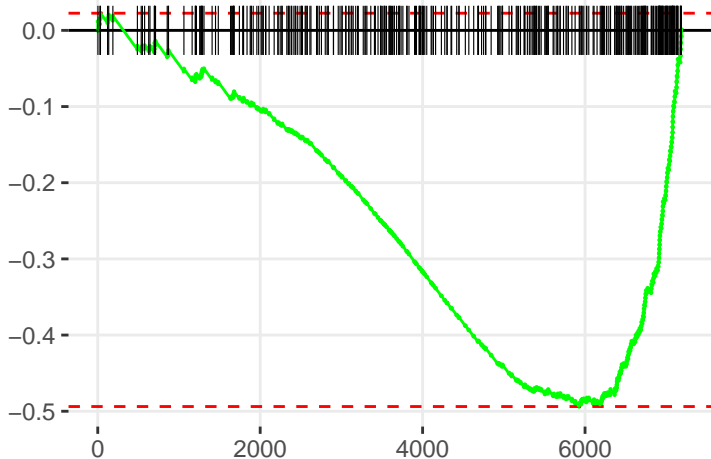
4000

6000

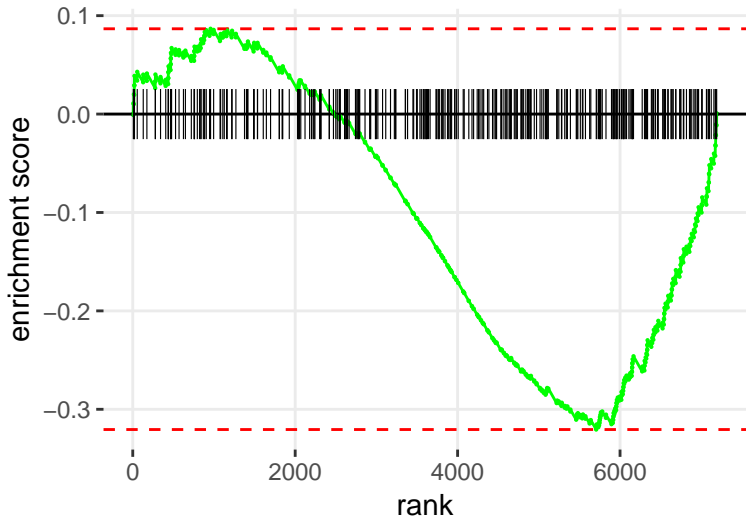


enrichment score

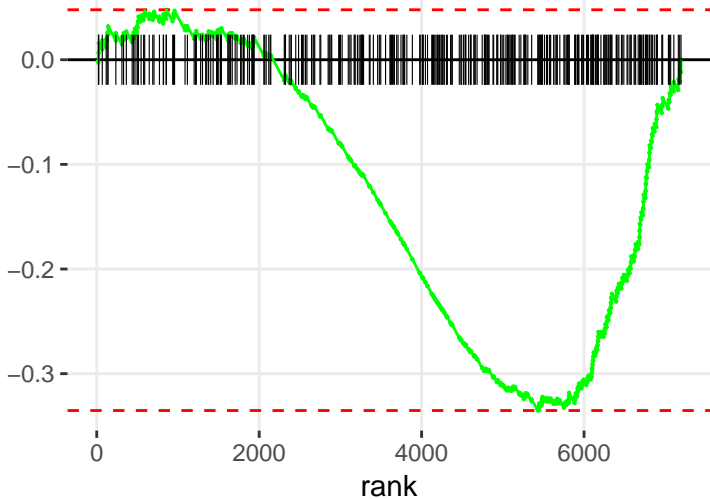
rank

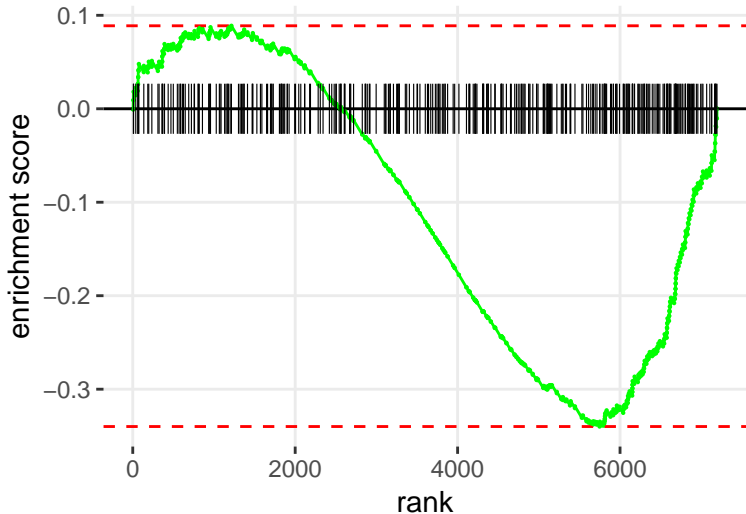


REGULATION OF KINASE ACTIVITY%GOBP%GO:0043549

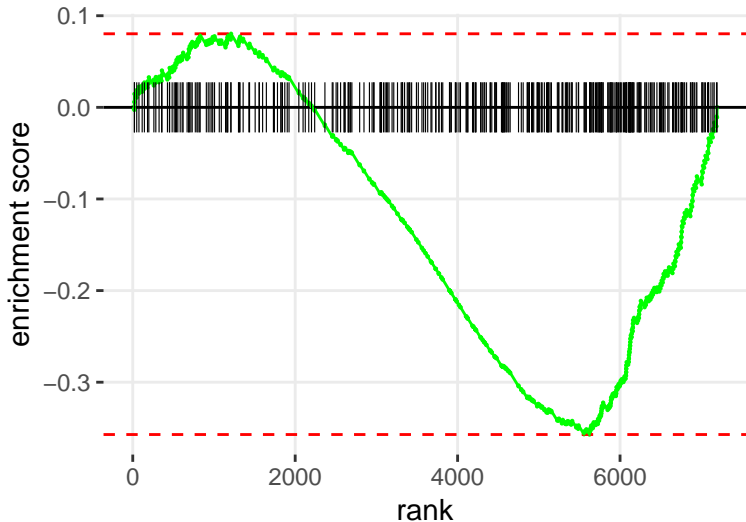


enrichment score





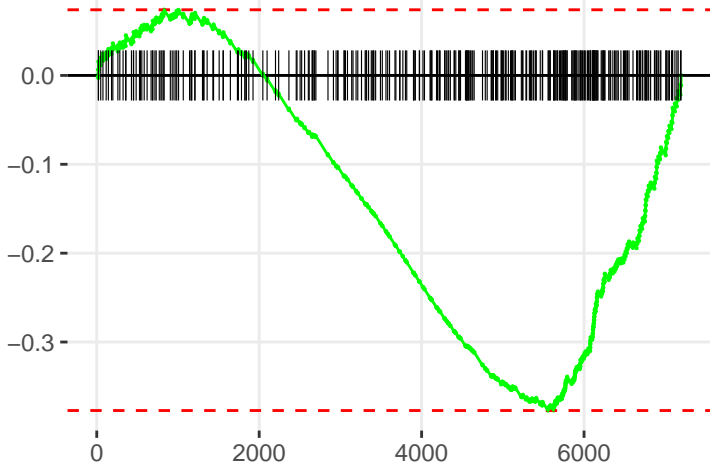
CELLULAR PROTEIN CATABOLIC PROCESS%GOBP%GO:0044257



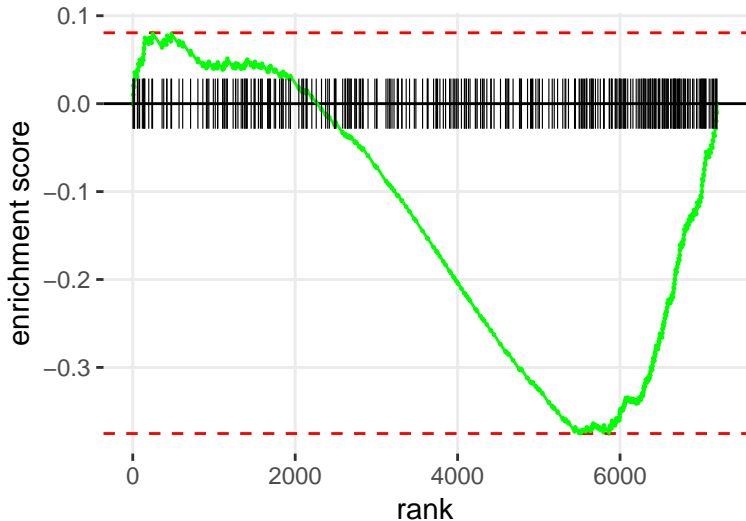
PROTEOLYSIS INVOLVED IN CELLULAR PROTEIN CATABOLIC PROCESS%GOBP%GO:0

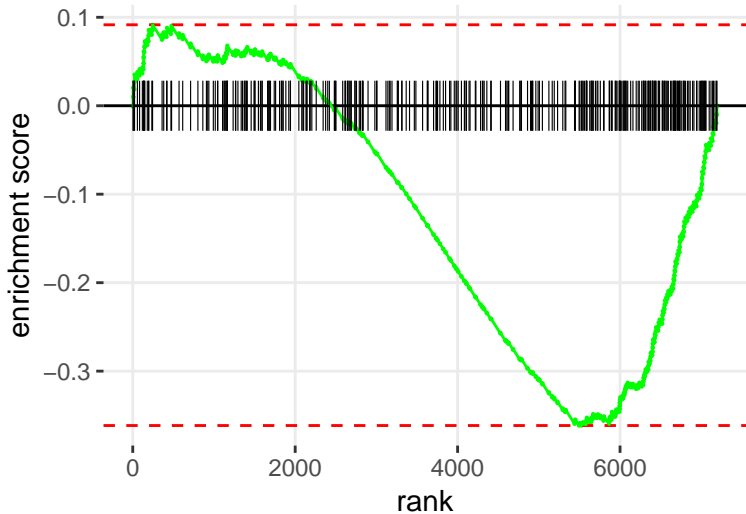
enrichment score

rank



CELL ACTIVATION INVOLVED IN IMMUNE RESPONSE%GOBP%GO:0002263

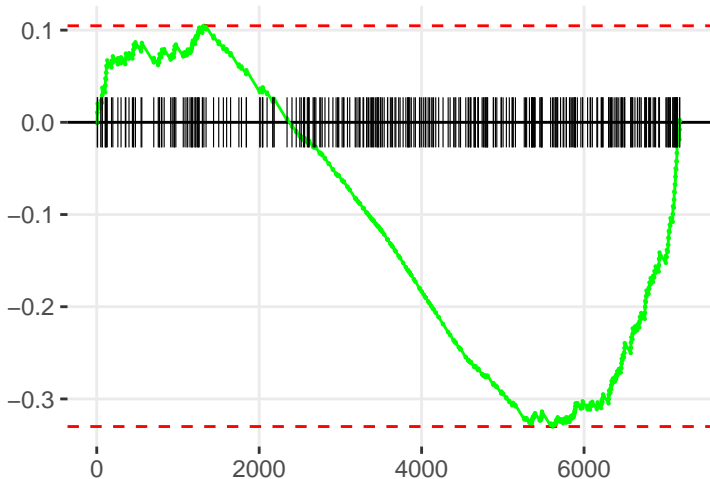




RESPONSE TO NITROGEN COMPOUND%GOBP%GO:1901698

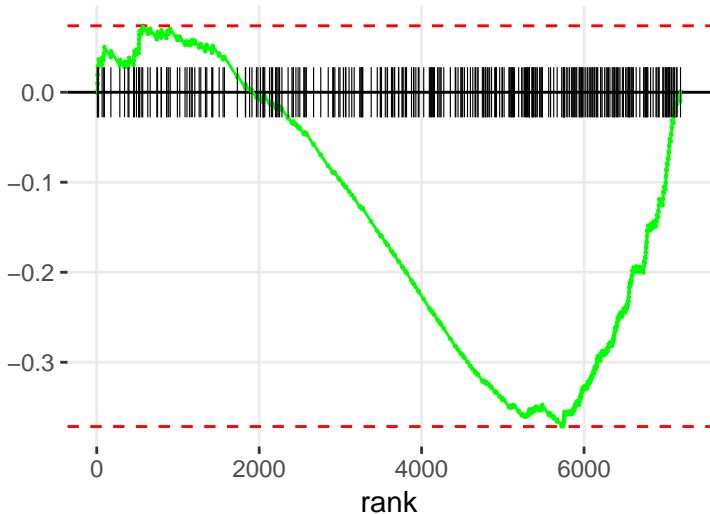
enrichment score

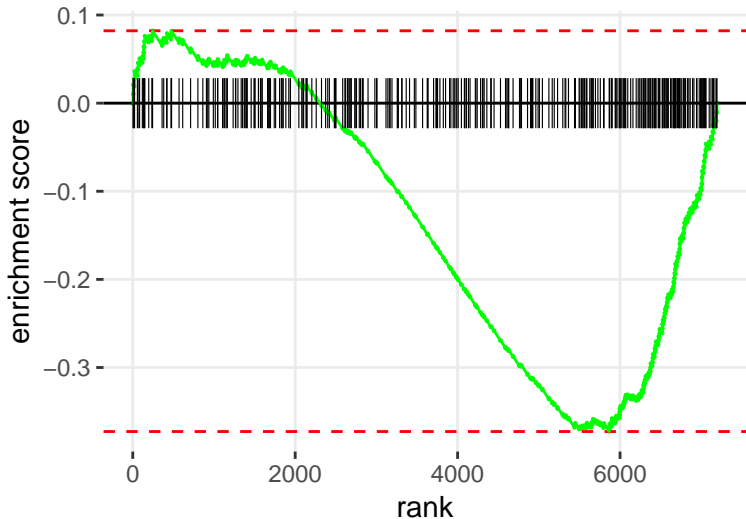
rank



CELLULAR RESPONSE TO CYTOKINE STIMULUS%GOBP%GO:0071345

enrichment score





enrichment score

0.0

-0.1

-0.2

-0.3

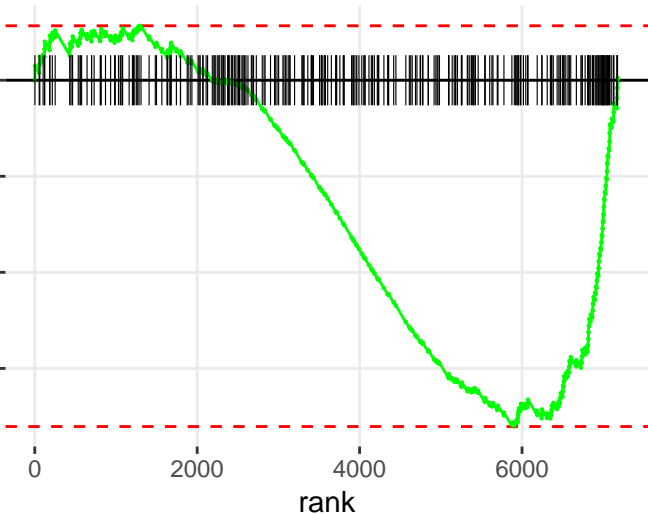
0

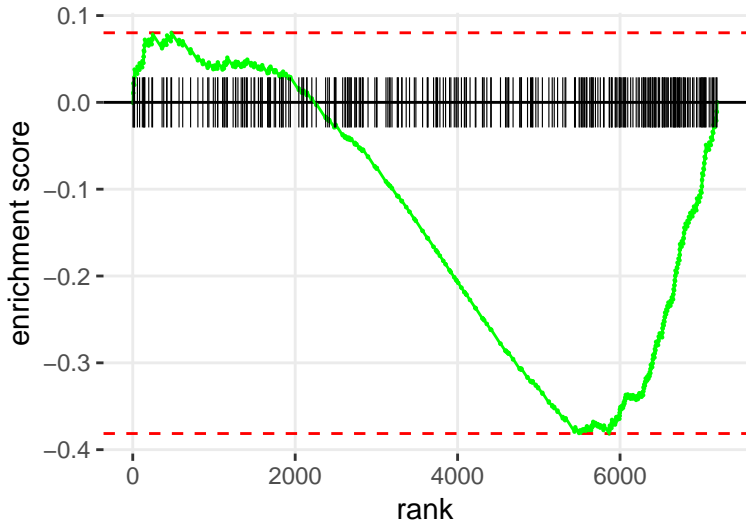
2000

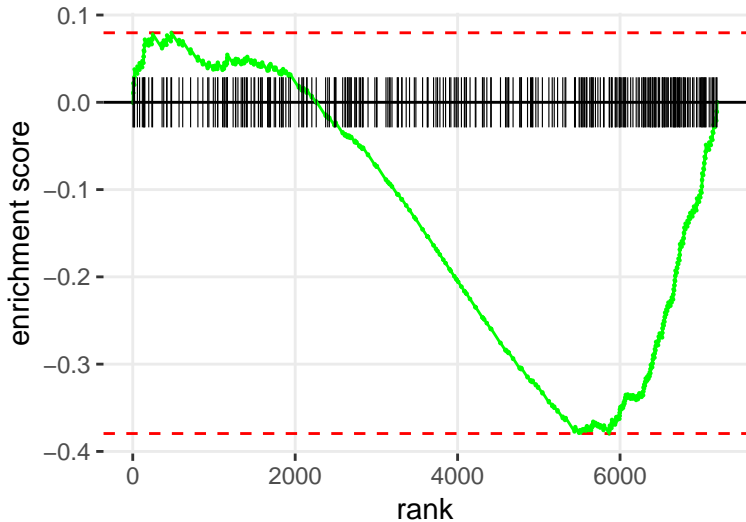
4000

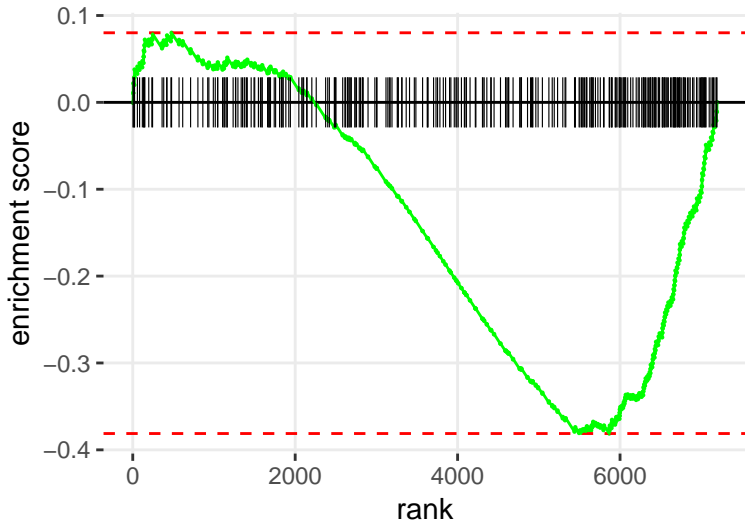
6000

rank

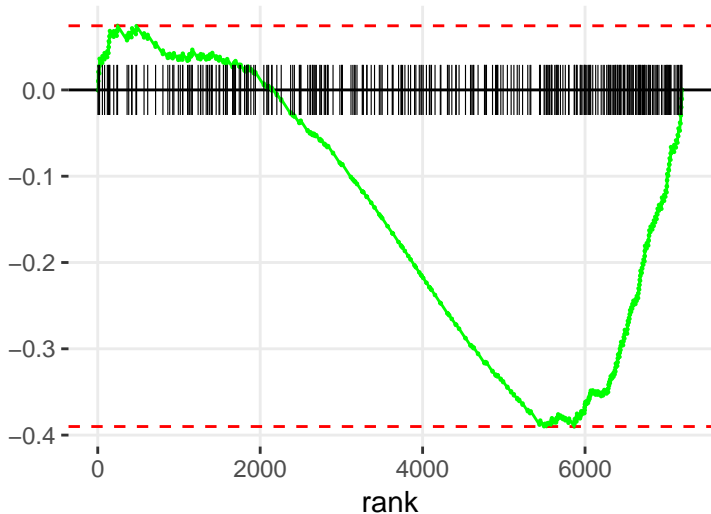


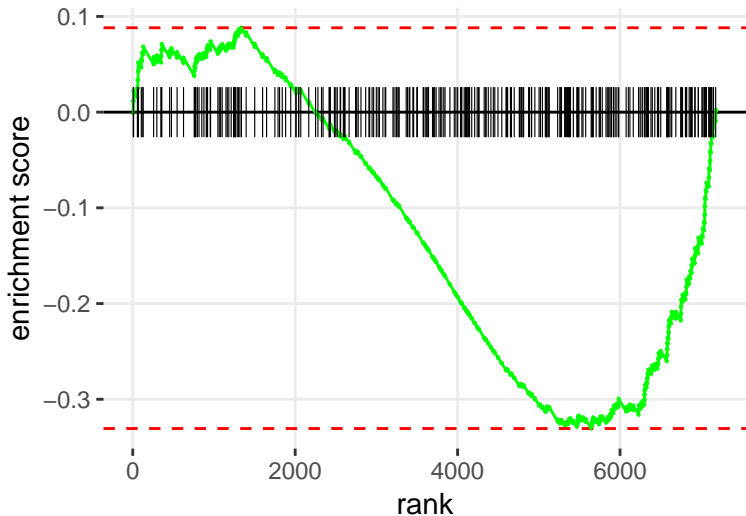


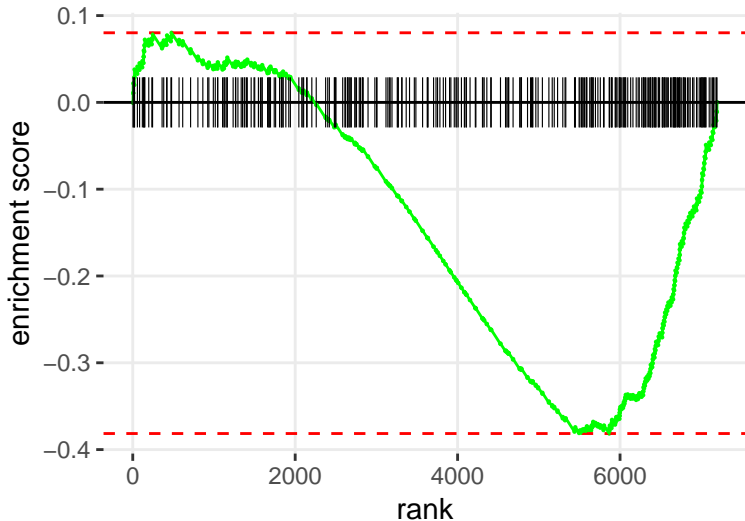




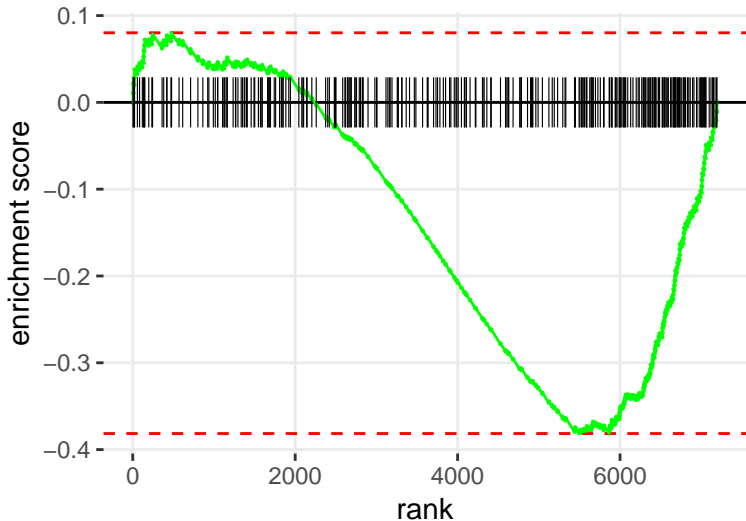
enrichment score



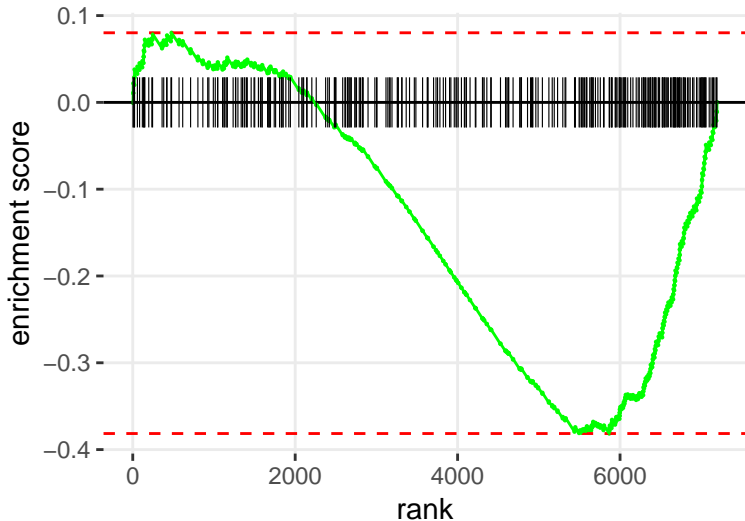




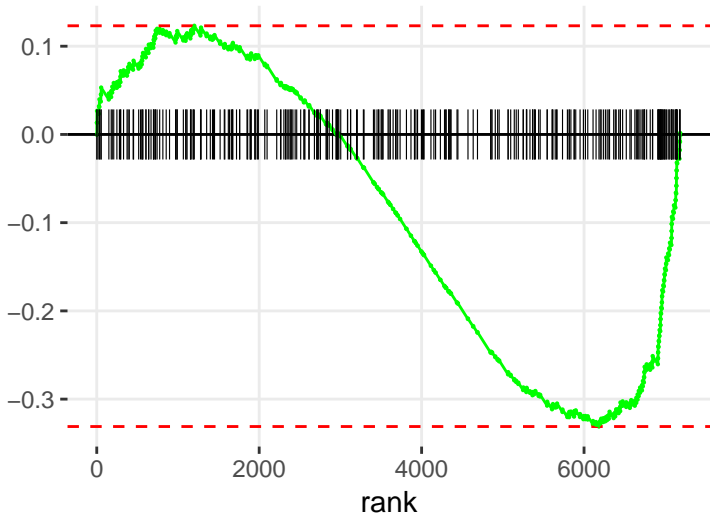
NEUTROPHIL DEGRANULATION%GOBP%GO:0043312

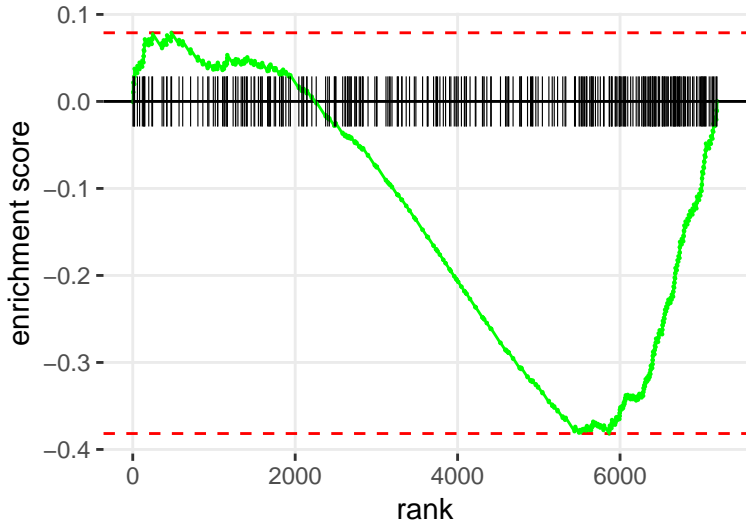


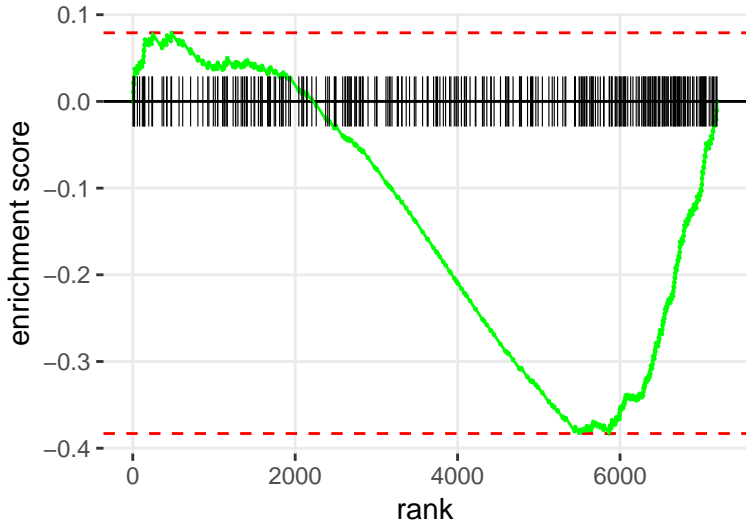
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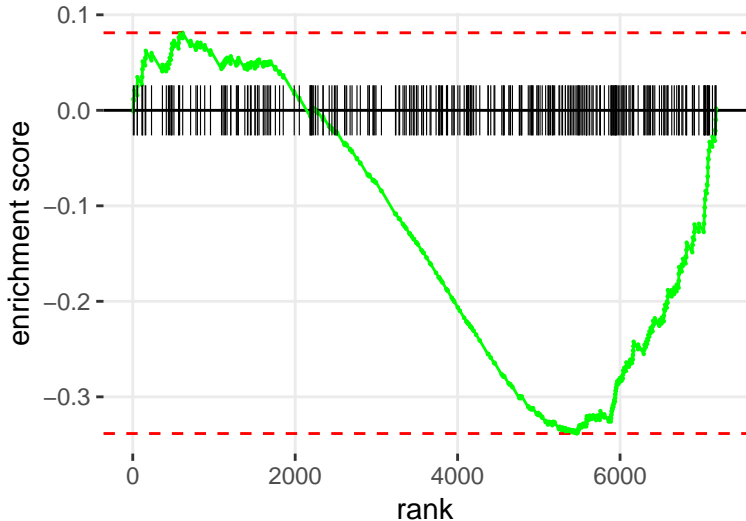


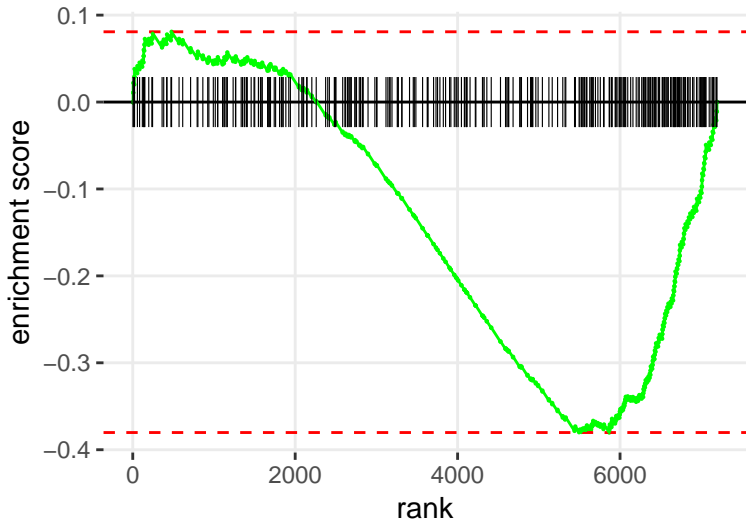
enrichment score



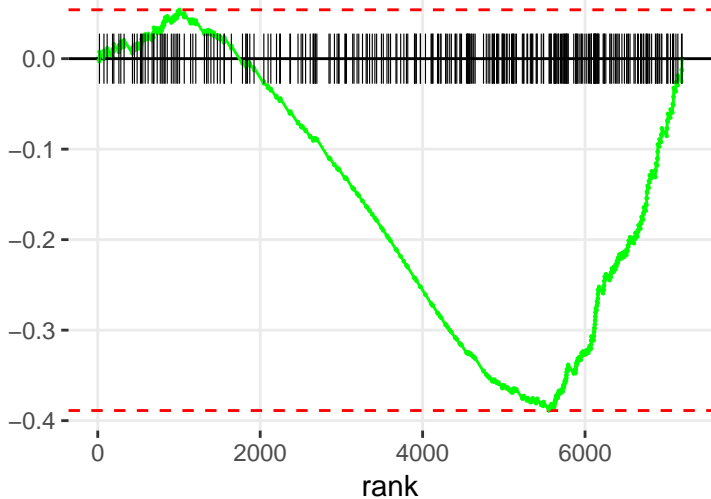








enrichment score



enrichment score

0.0

-0.1

-0.2

-0.3

-0.4

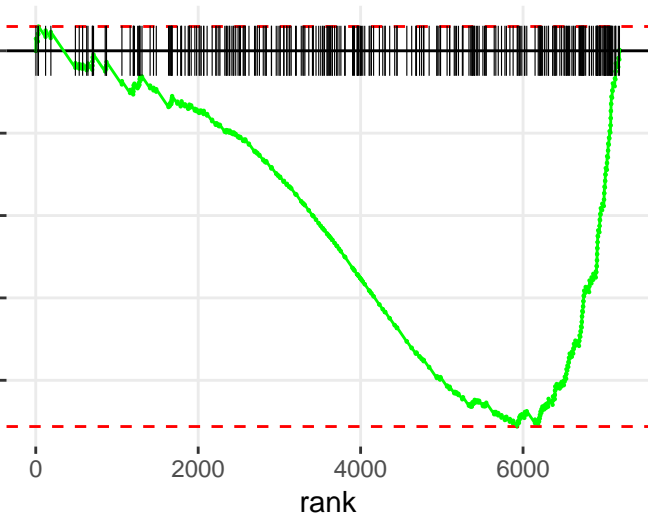
0

2000

4000

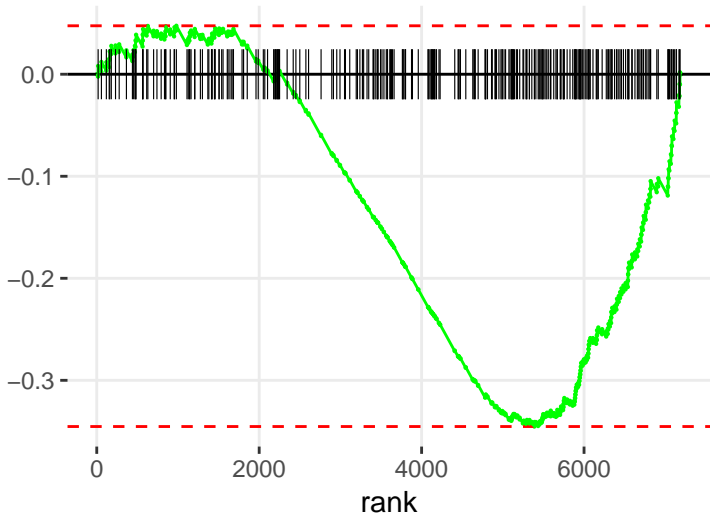
6000

rank



POSITIVE REGULATION OF IMMUNE RESPONSE%GOBP%GO:0050778

enrichment score



enrichment score

0.0

-0.1

-0.2

-0.3

-0.4

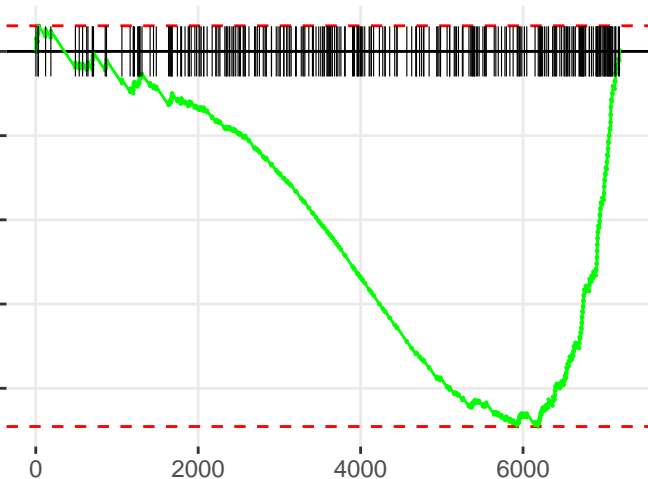
0

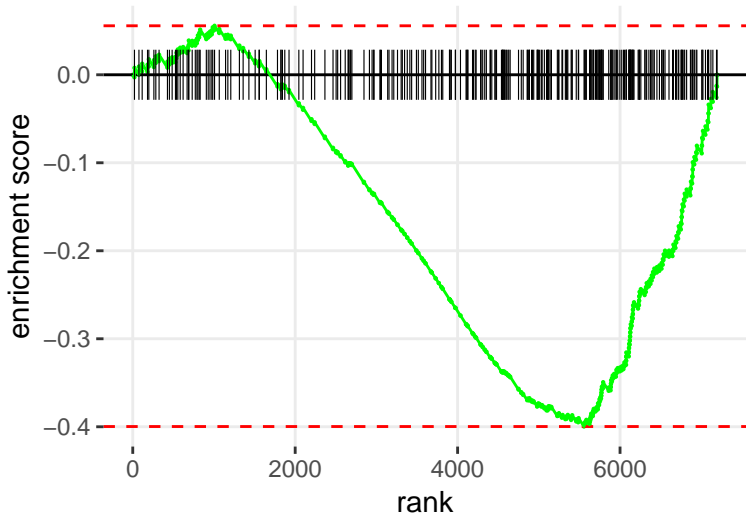
2000

4000

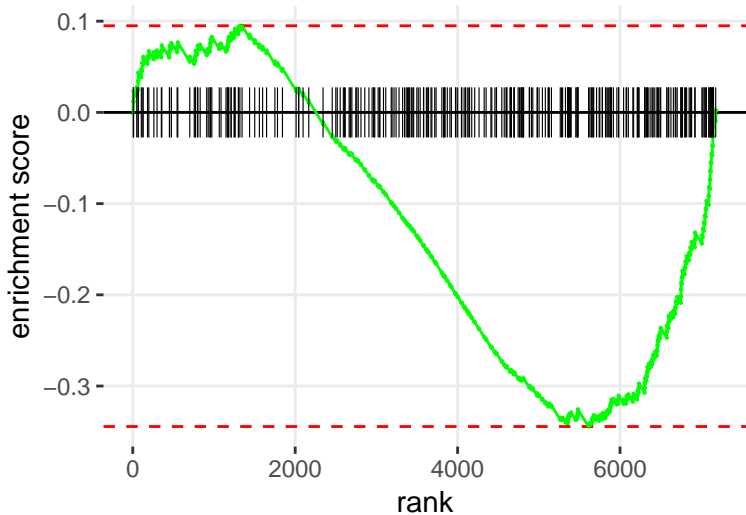
6000

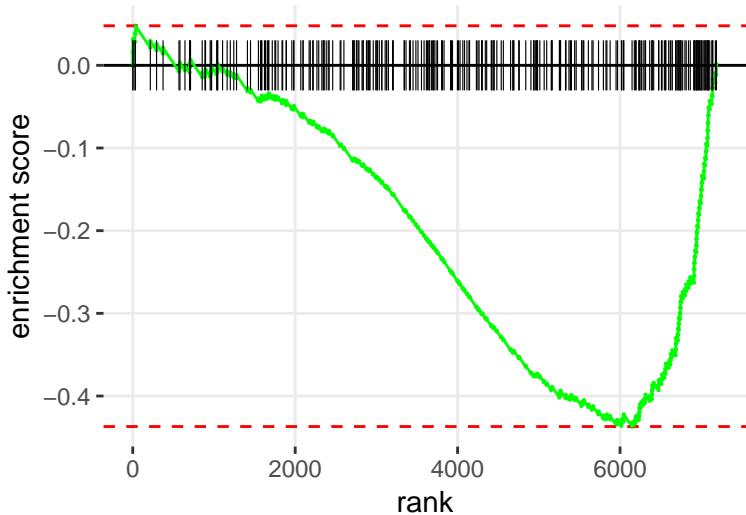
rank



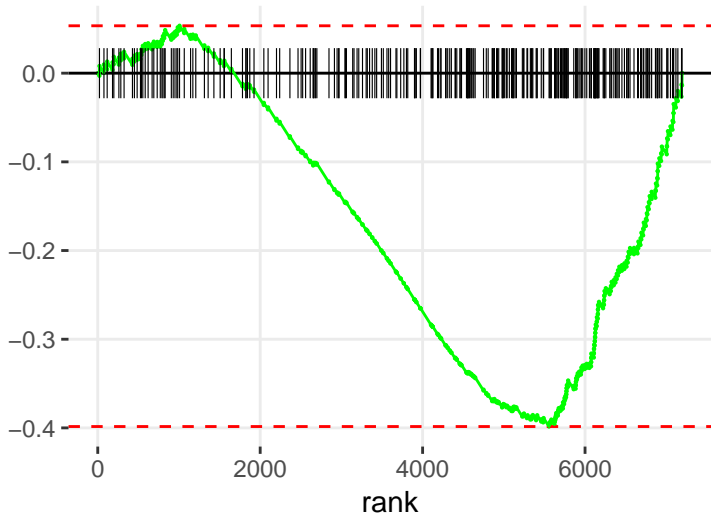


RESPONSE TO ORGANONITROGEN COMPOUND%GOBP%GO:0010243





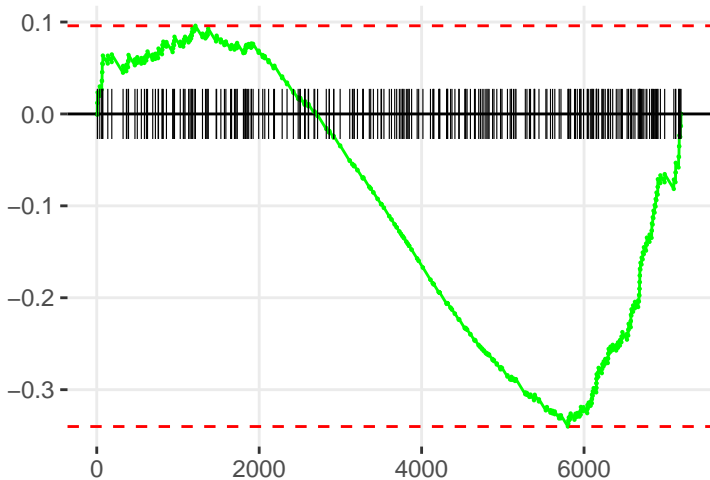
enrichment score



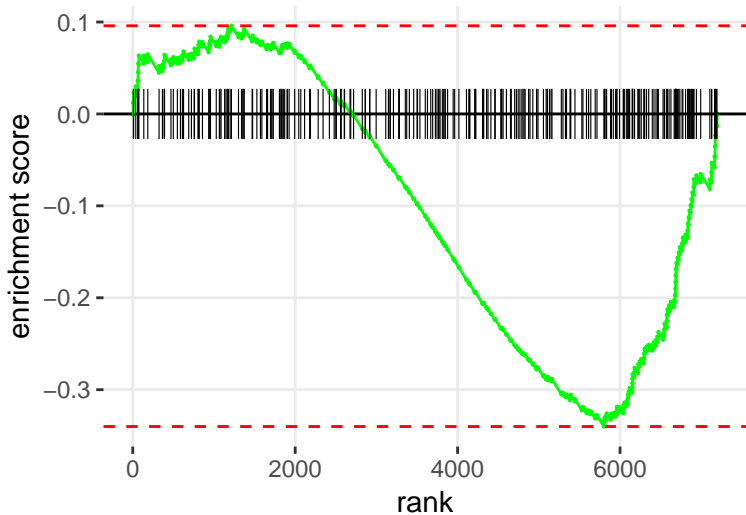
CELL MOTILITY%GOBP%GO:0048870

enrichment score

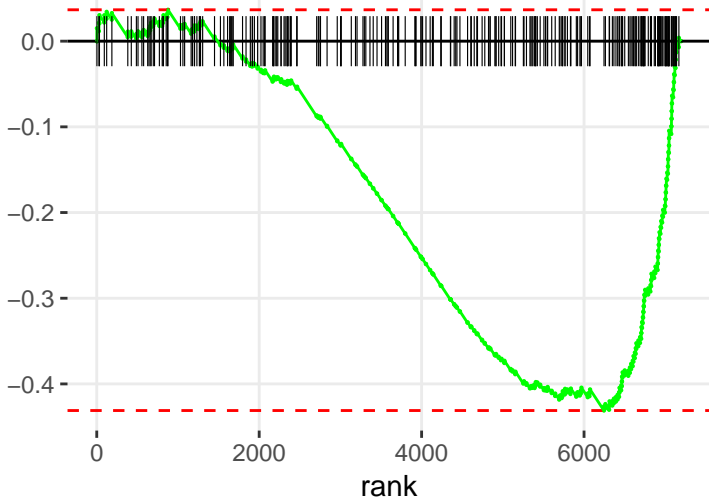
rank



LOCALIZATION OF CELL%GOBP%GO:0051674



enrichment score



enrichment score

0.0

-0.1

-0.2

-0.3

-0.4

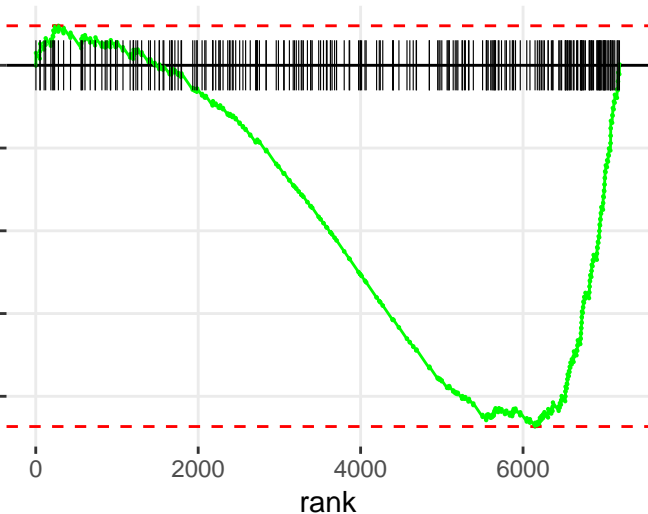
0

2000

4000

6000

rank



enrichment score

0.0

-0.1

-0.2

-0.3

-0.4

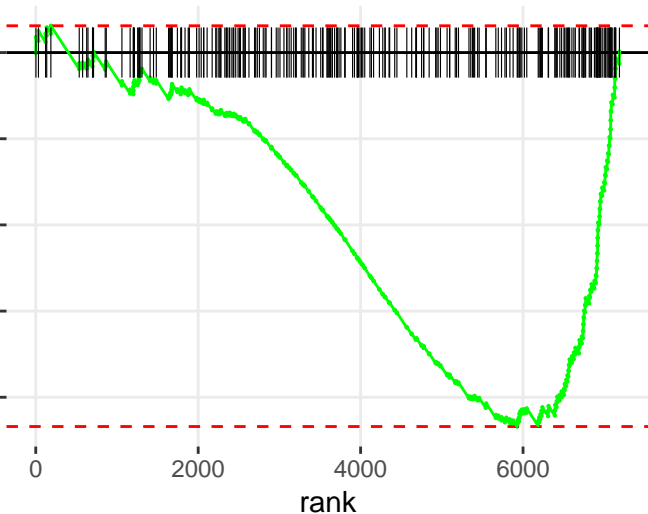
0

2000

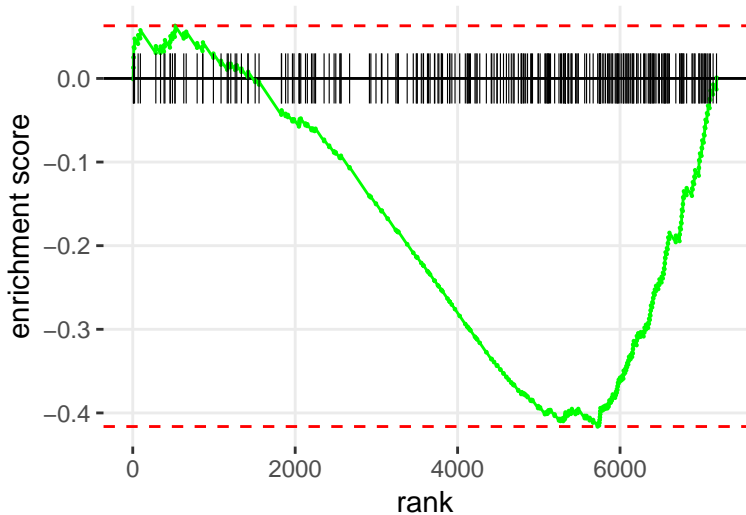
4000

6000

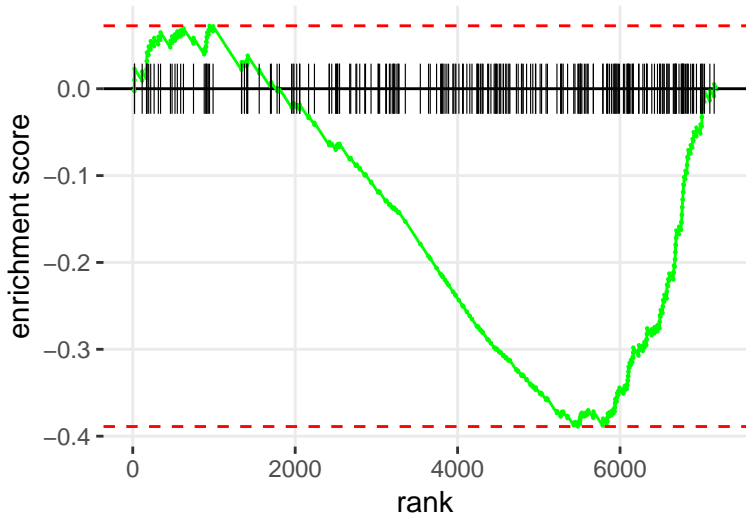
rank



CYTOKINE-MEDIATED SIGNALING PATHWAY%GOBP%GO:0019221



REGULATION OF CYTOSKELETON ORGANIZATION%GOBP%GO:0051493



enrichment score

0.1
0.0
-0.1
-0.2
-0.3

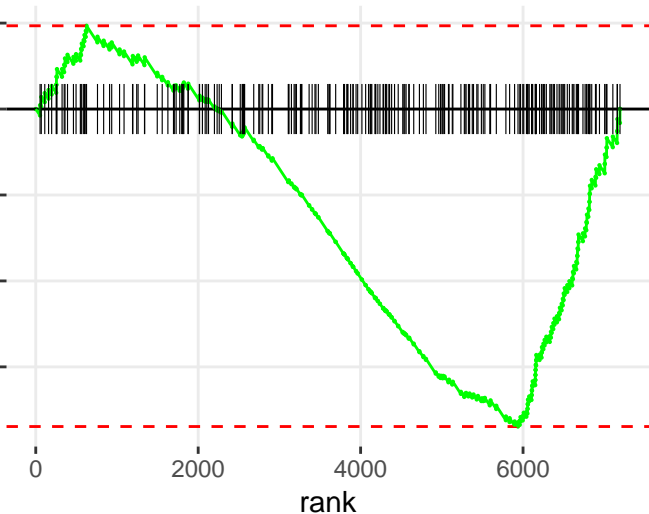
0

2000

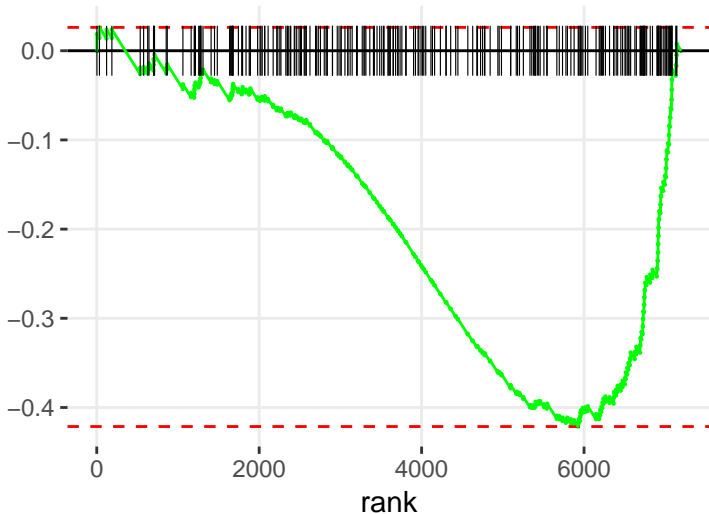
4000

6000

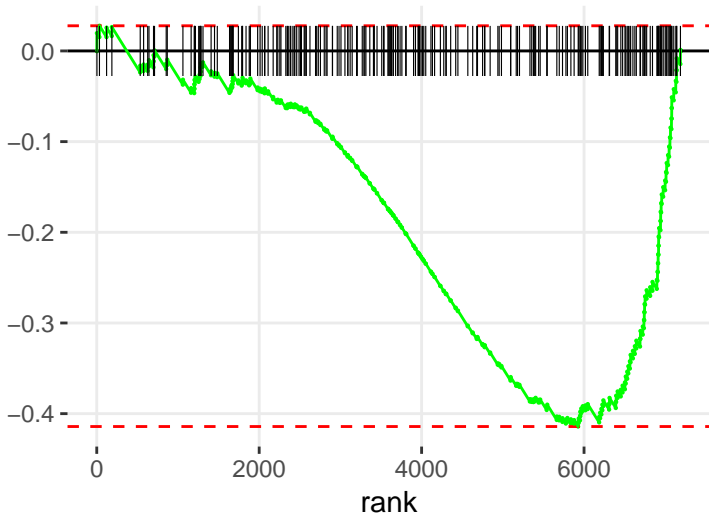
rank

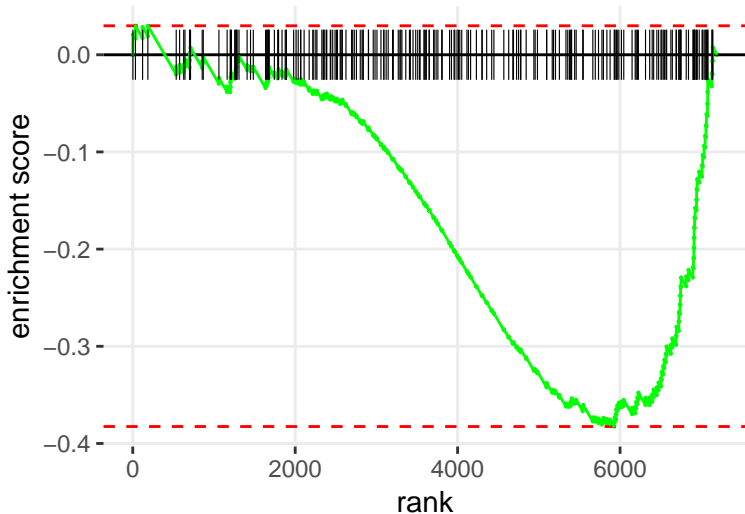


enrichment score

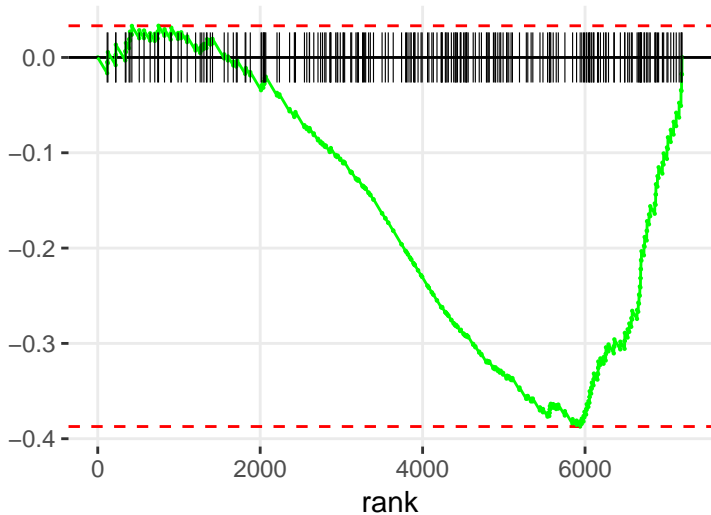


enrichment score

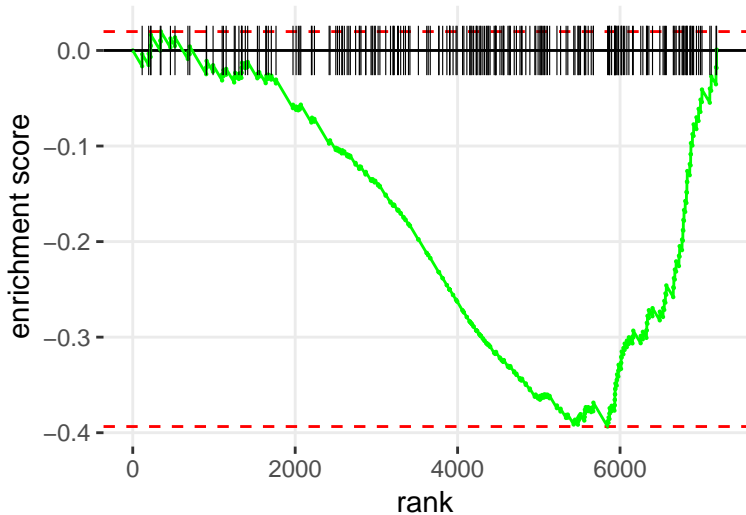


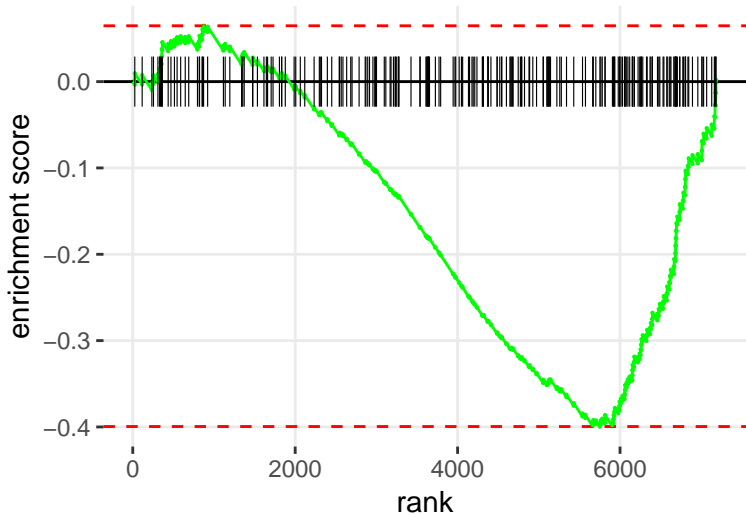


enrichment score

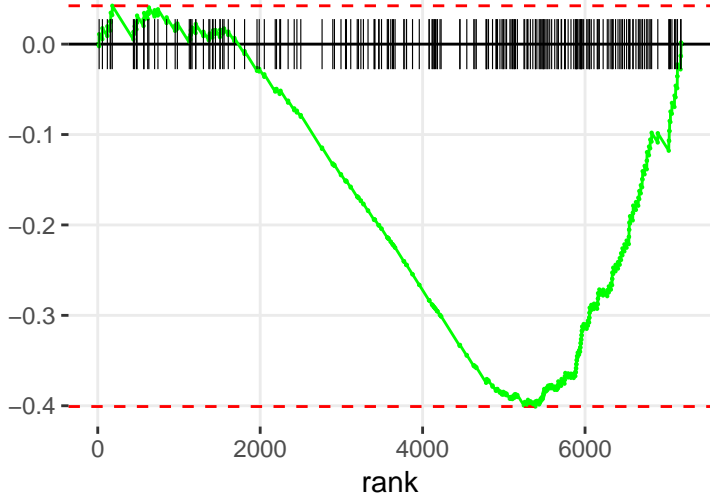


REGULATION OF PROTEIN COMPLEX ASSEMBLY%GOBP%GO:0043254





enrichment score



enrichment score

0.0
-0.1
-0.2
-0.3
-0.4

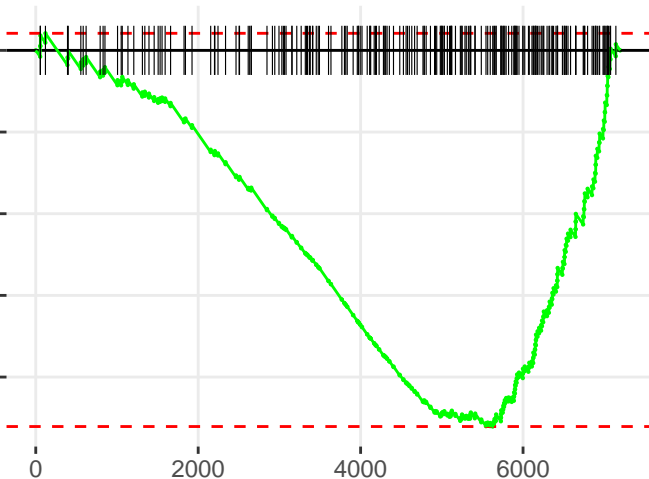
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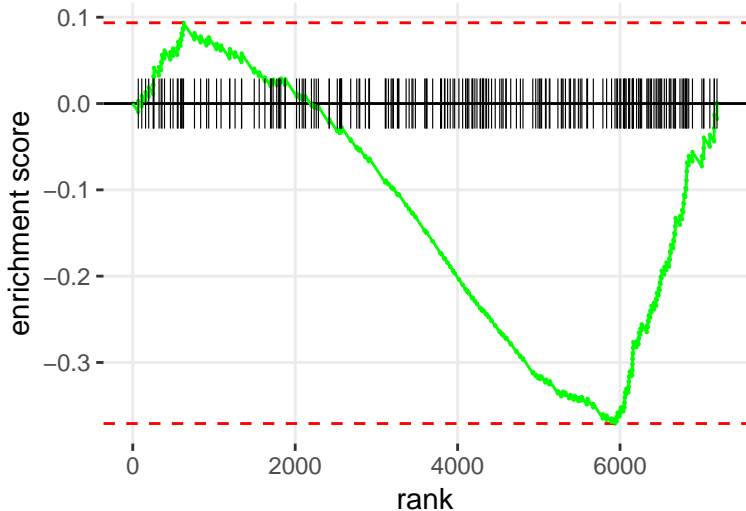
2000

4000

6000

rank





enrichment score

0.0

-0.1

-0.2

-0.3

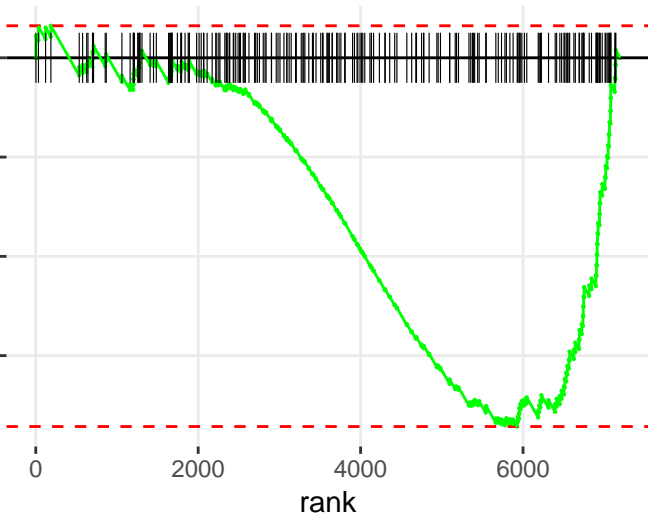
0

2000

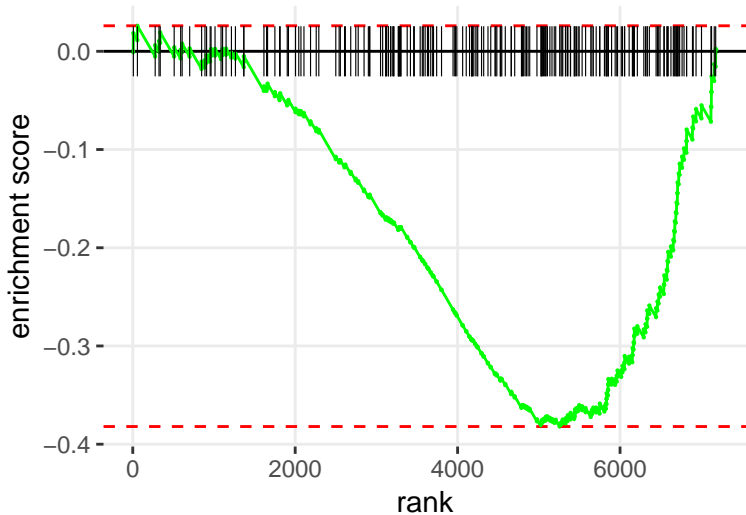
4000

6000

rank



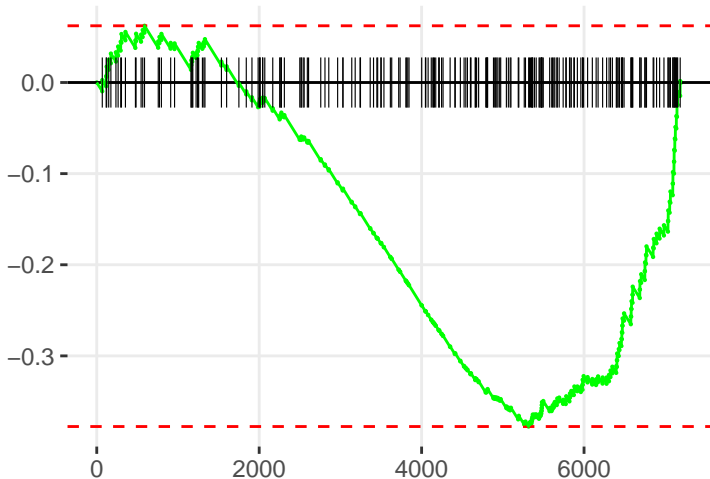
IMPORT INTO CELL%GOBP%GO:0098657

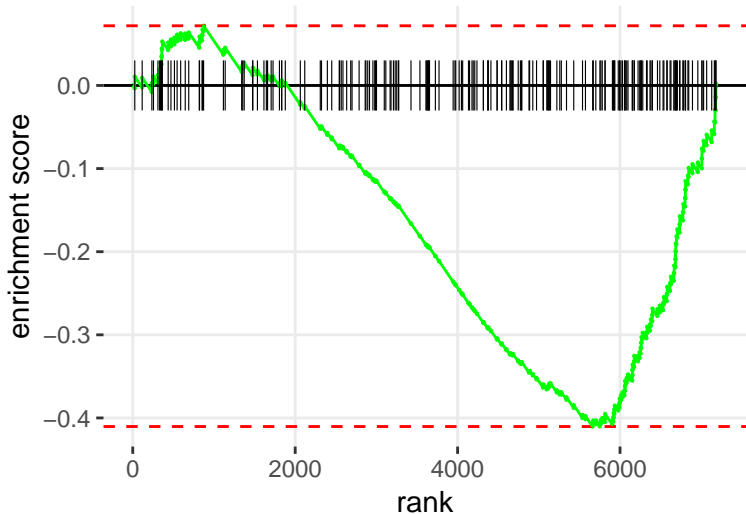


RESPONSE TO HORMONE%GOBP%GO:0009725

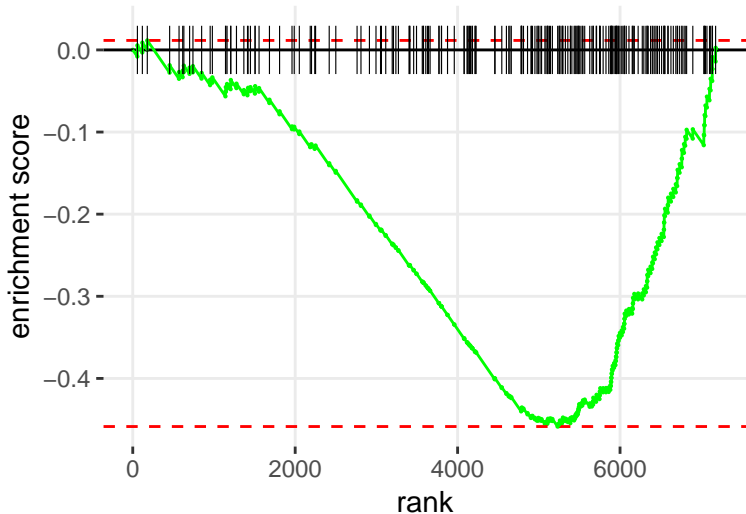
enrichment score

rank





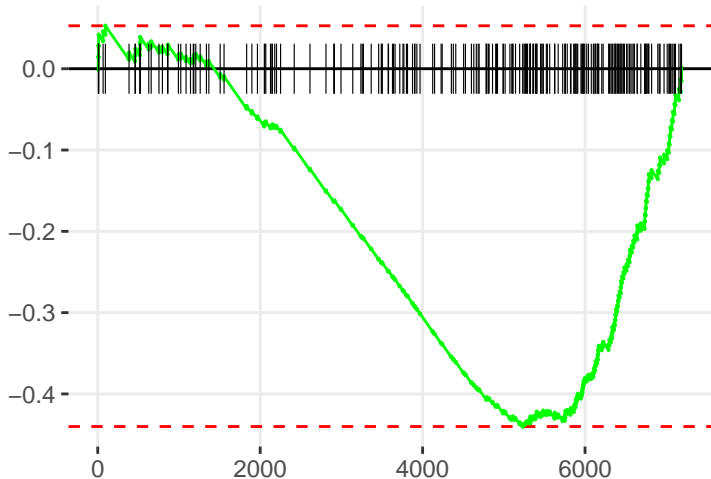
IMMUNE RESPONSE-REGULATING SIGNALING PATHWAY%GOBP%GO:0002764



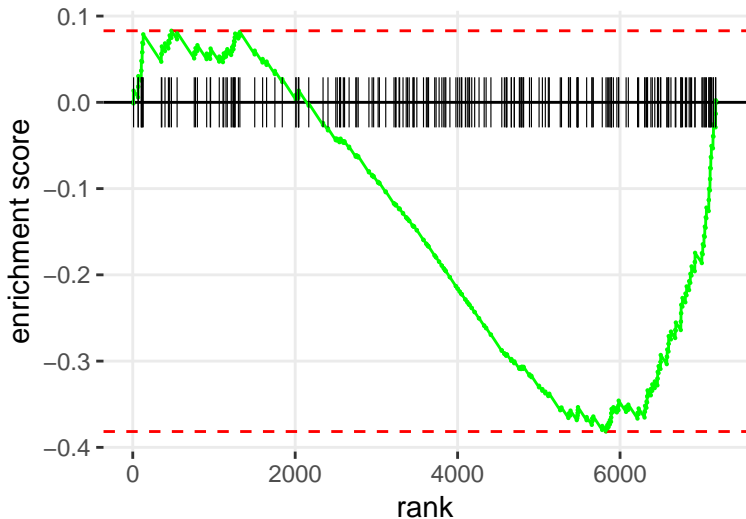
SIGNALING BY INTERLEUKINS%REACTOME%R-HSA-449147.11

enrichment score

rank



CELLULAR RESPONSE TO NITROGEN COMPOUND%GOBP%GO:1901699



enrichment score

0.0
-0.1
-0.2
-0.3
-0.4

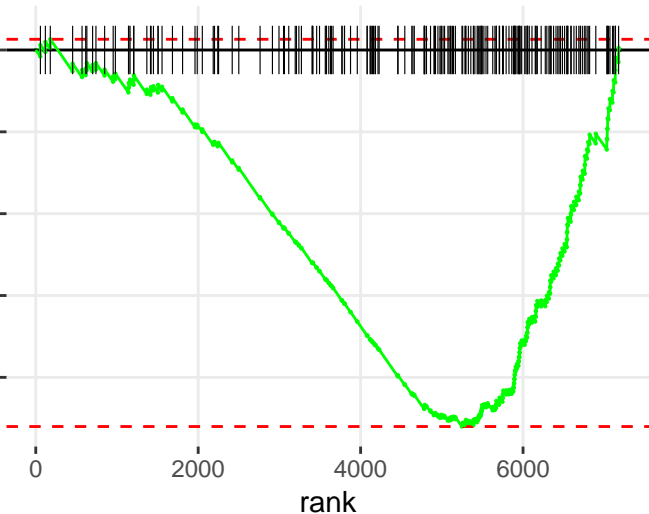
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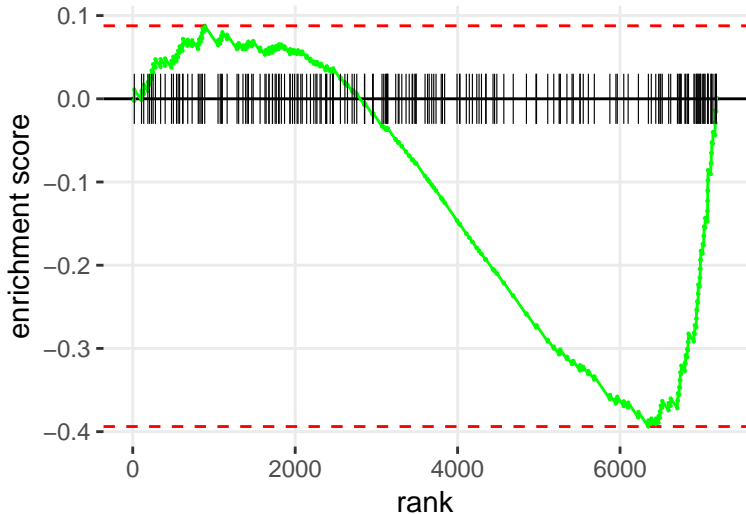
2000

4000

6000

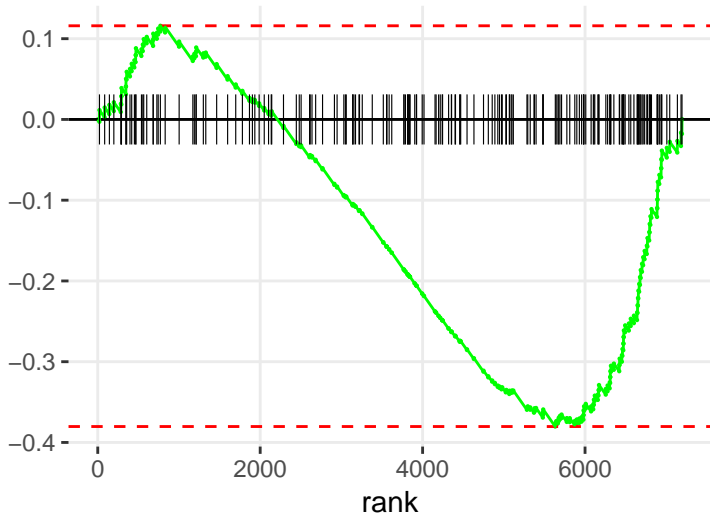
rank

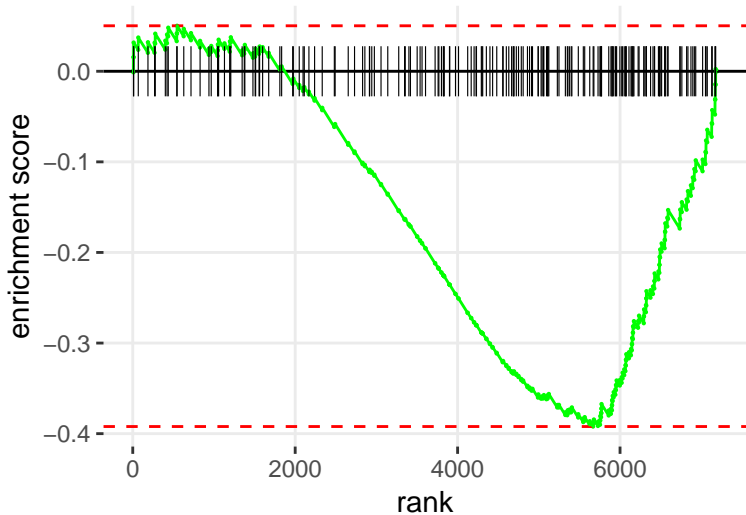




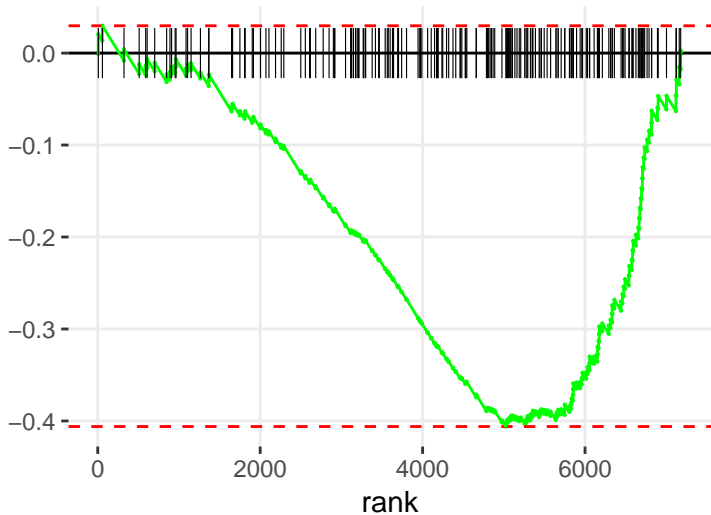
REGULATION OF PROTEIN CATABOLIC PROCESS%GOBP%GO:0042176

enrichment score





enrichment score



REGULATION OF AUTOPHAGY%GOBP%GO:0010506

enrichment score

rank

