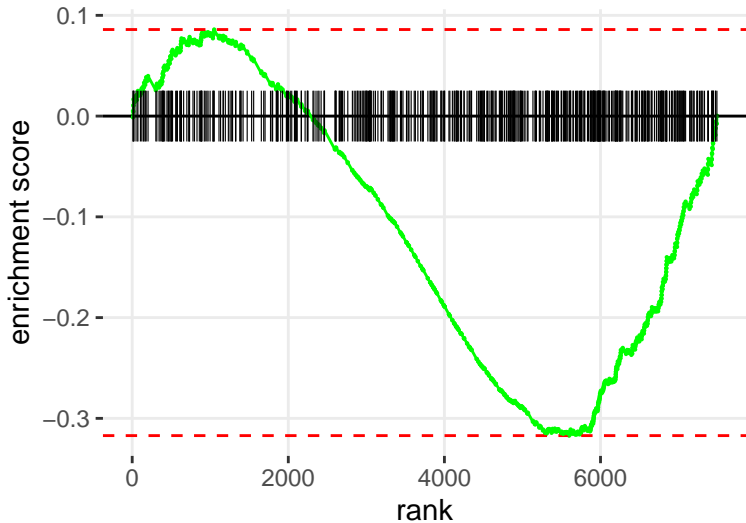


NEGATIVE REGULATION OF SIGNAL TRANSDUCTION%GOBP%GO:0009968



REGULATION OF CELLULAR COMPONENT BIOGENESIS%GOBP%GO:0044087

enrichment score

0.0

-0.1

-0.2

-0.3

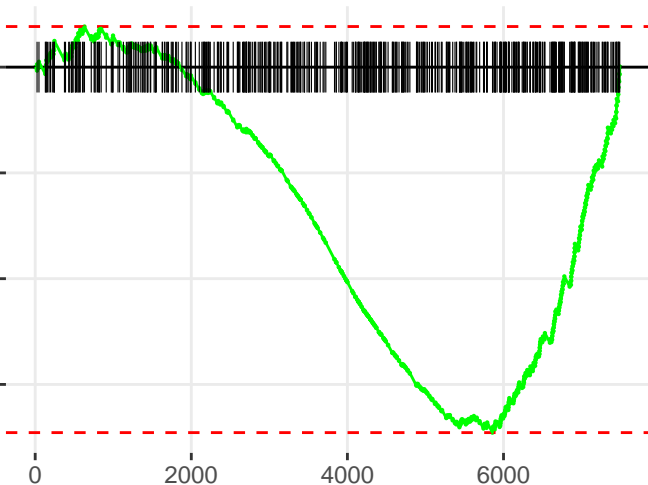
0

2000

4000

6000

rank



enrichment score

0.0

-0.1

-0.2

-0.3

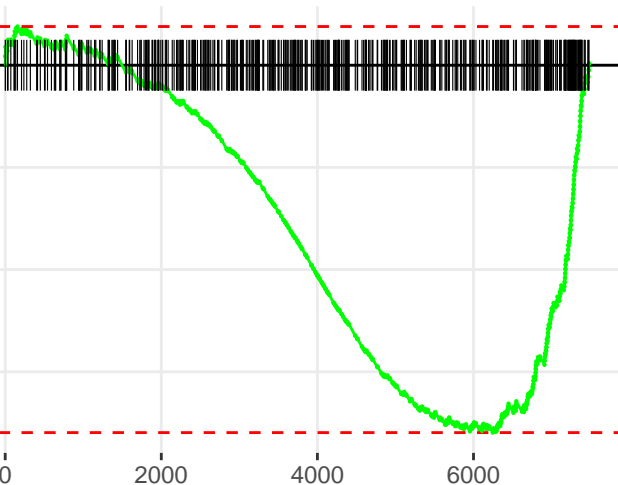
0

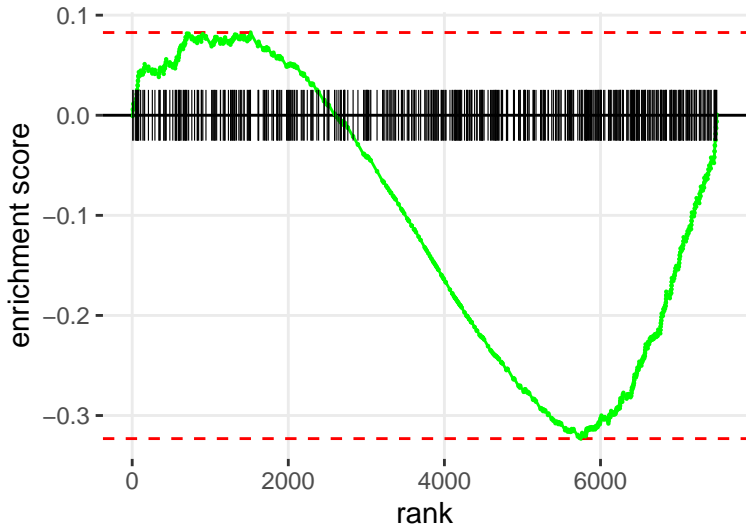
2000

4000

6000

rank





enrichment score

0.0

-0.1

-0.2

-0.3

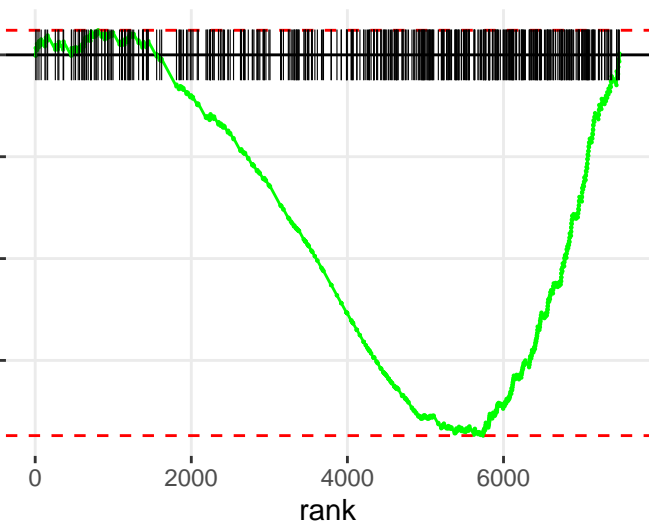
0

2000

4000

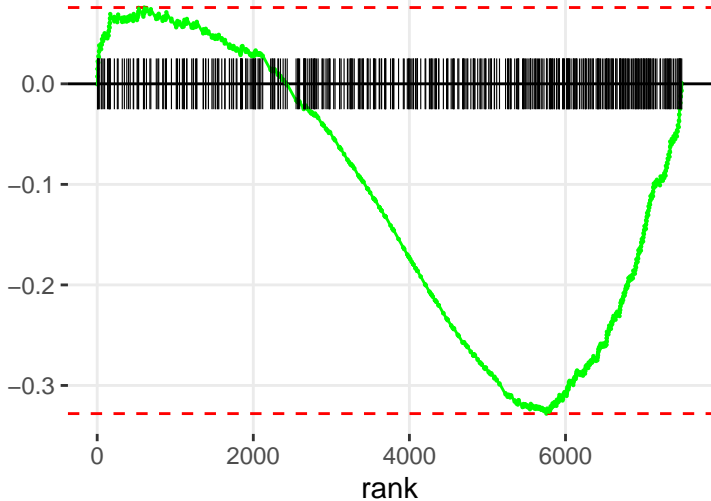
6000

rank

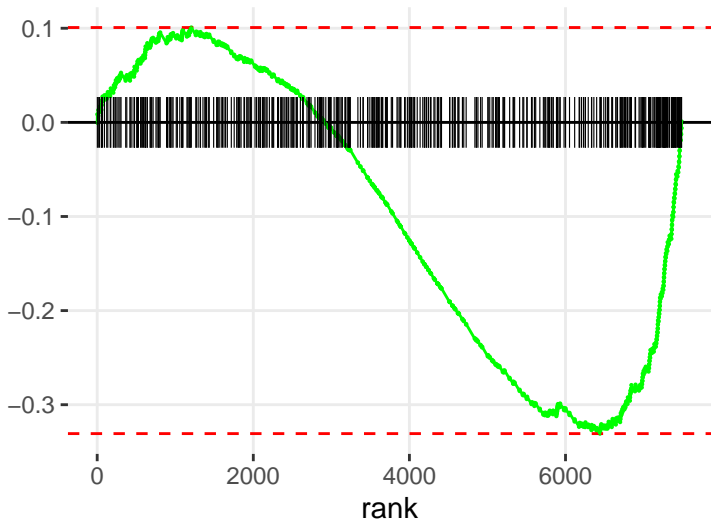


SECRETION BY CELL%GOBP%GO:0032940

enrichment score



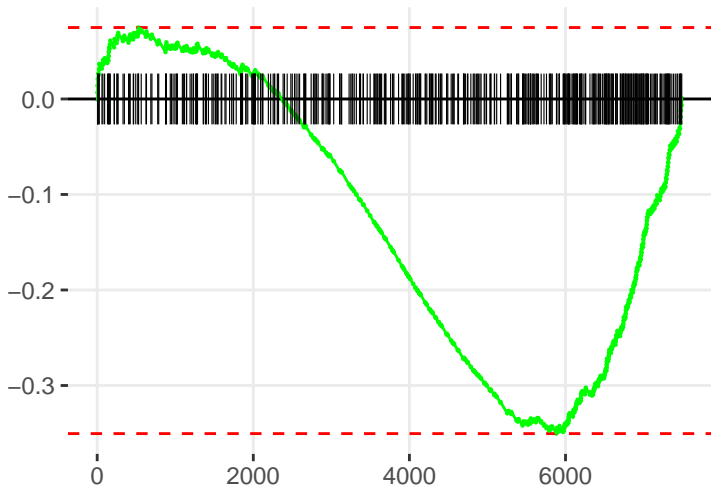
enrichment score



CELL ACTIVATION%GOBP%GO:0001775

enrichment score

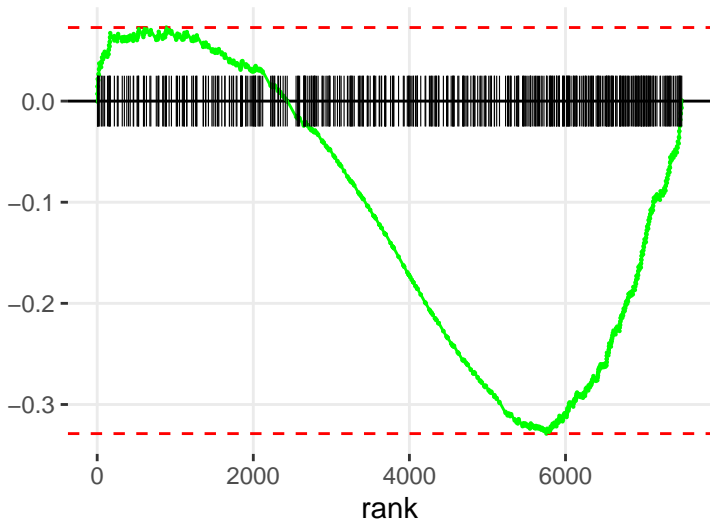
rank



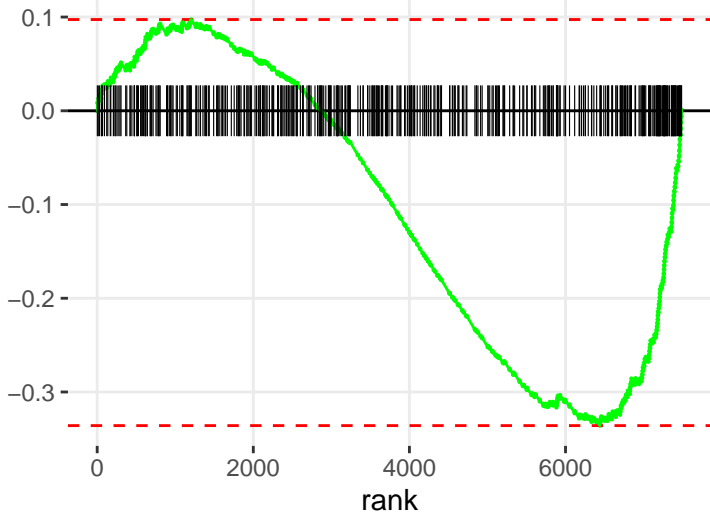


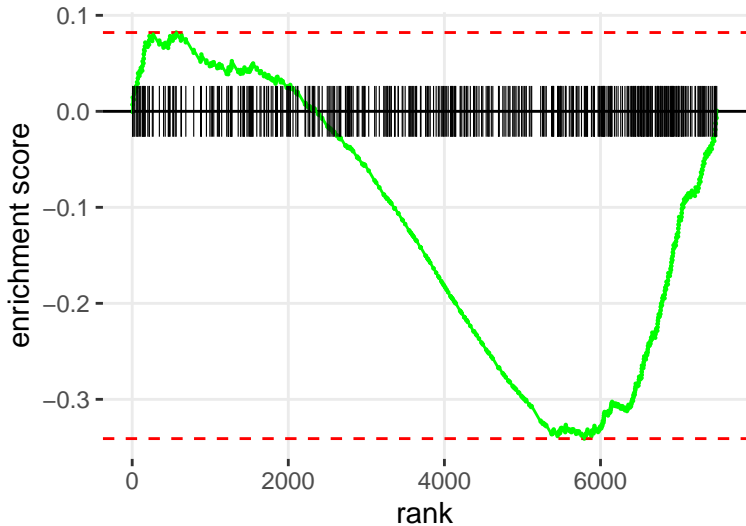
SECRETION%GOBP%GO:0046903

enrichment score

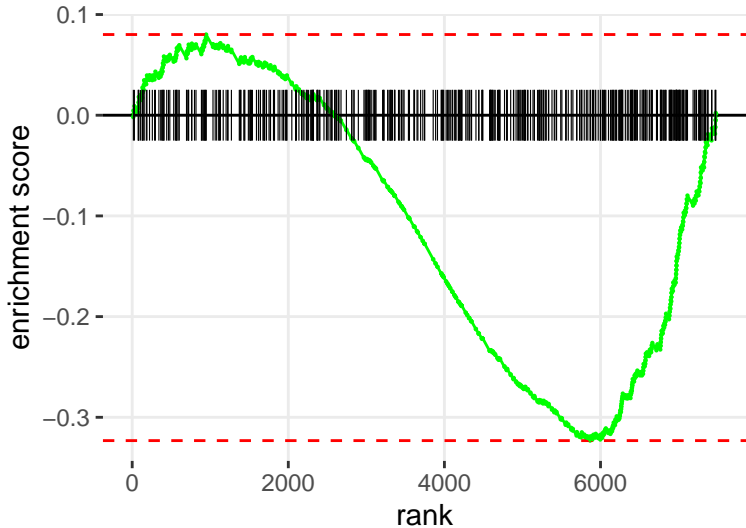


enrichment score





NEUROGENESIS%GOBP%GO:0022008



enrichment score

0.0

-0.1

-0.2

-0.3

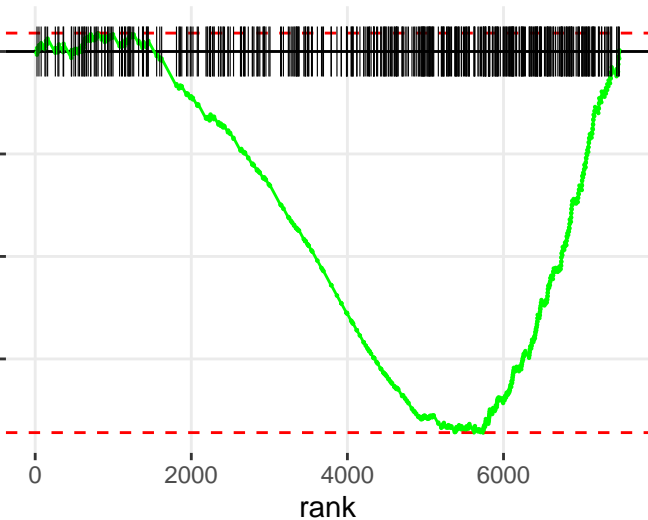
0

2000

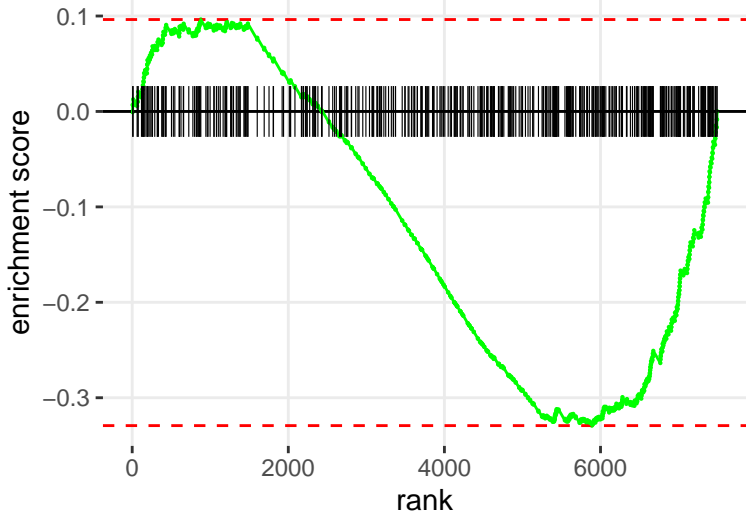
4000

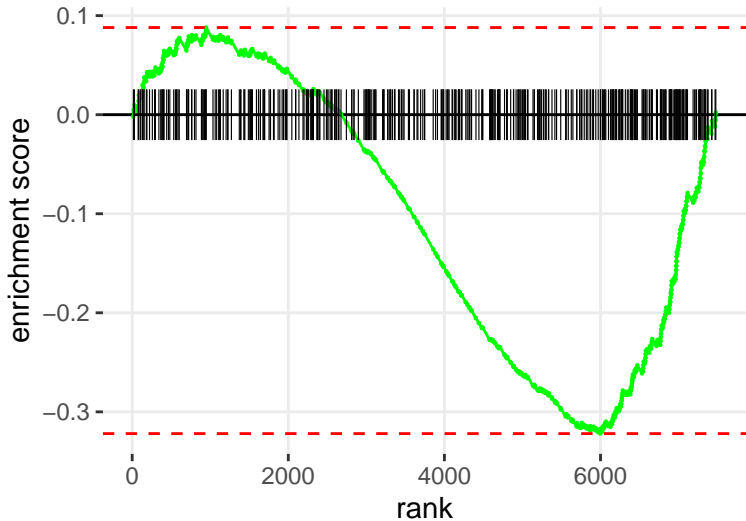
6000

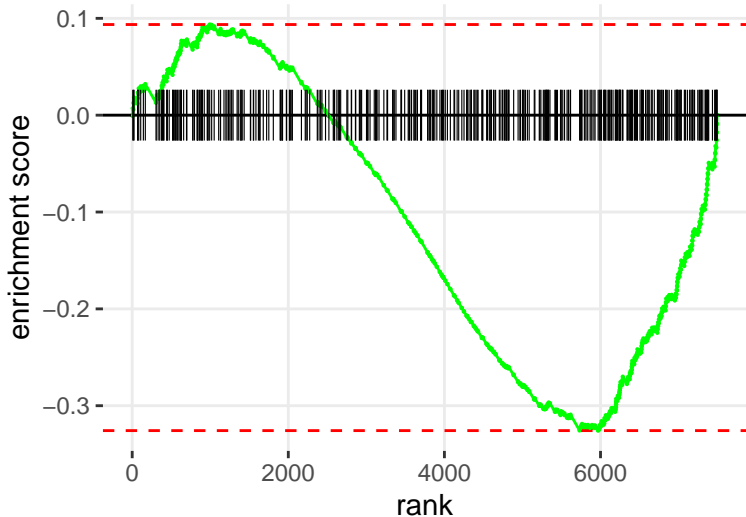
rank



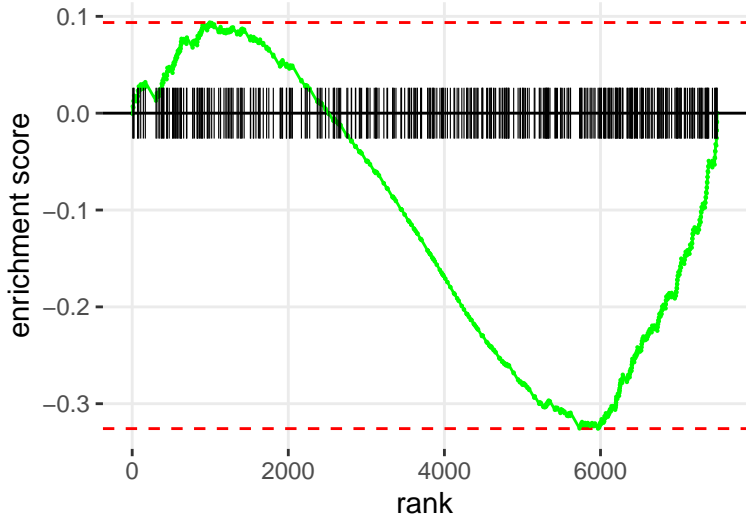
RESPONSE TO ENDOGENOUS STIMULUS%GOBP%GO:0009719



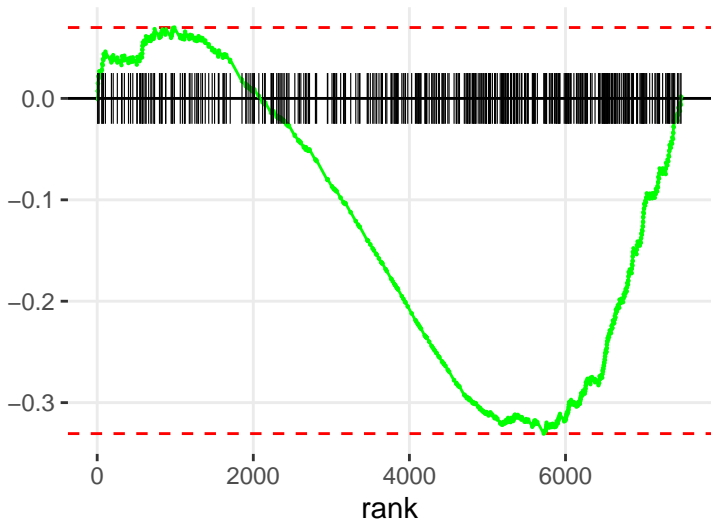


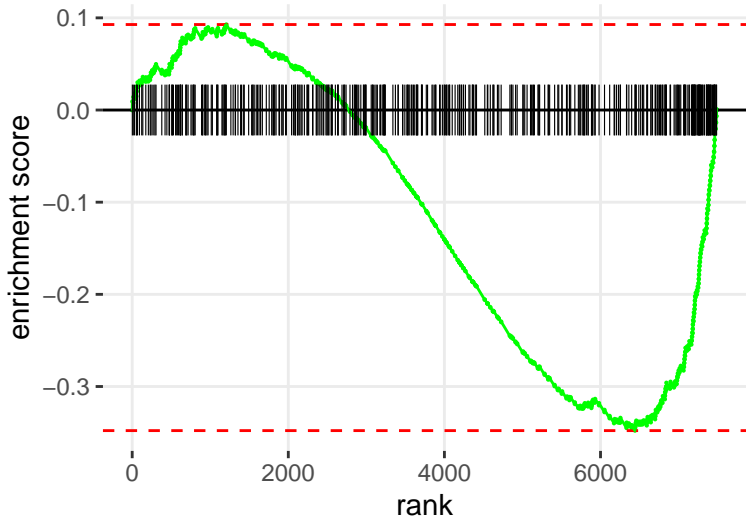






enrichment score





EXOCYTOSIS%GOBP%GO:0006887

enrichment score

0.0

-0.1

-0.2

-0.3

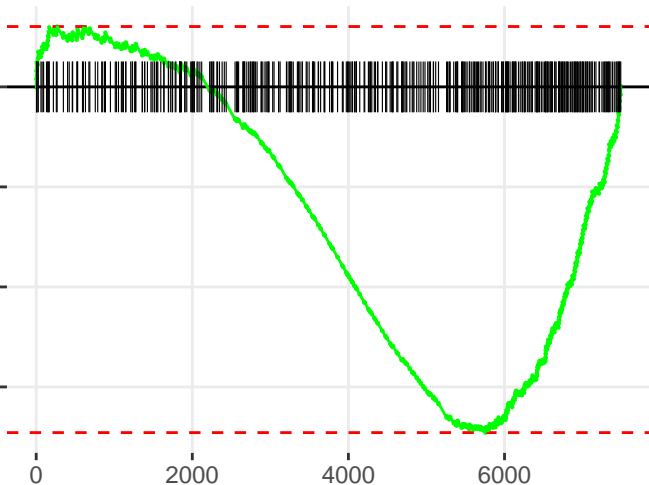
0

2000

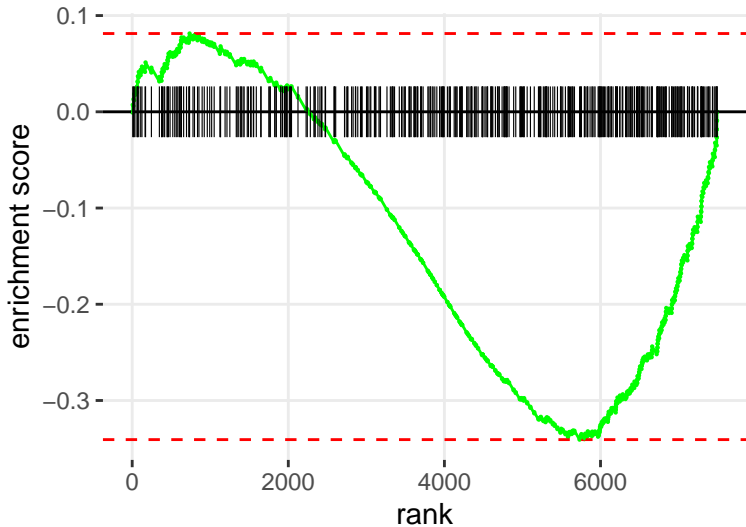
4000

6000

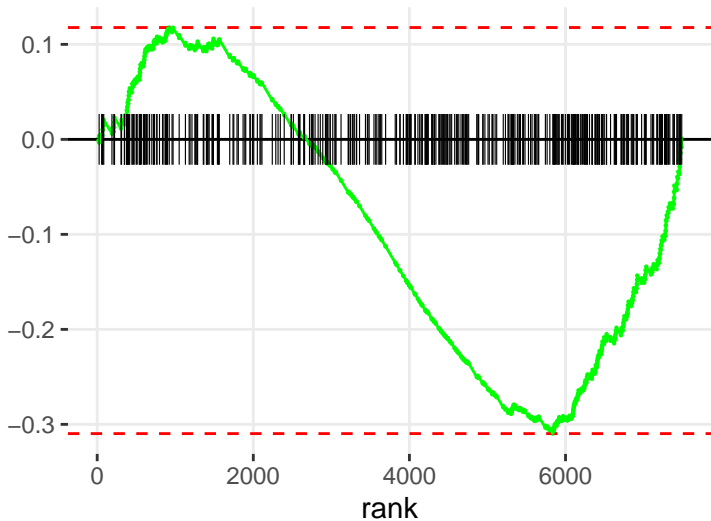
rank



POSITIVE REGULATION OF INTRACELLULAR SIGNAL TRANSDUCTION%GOBP%GO:19



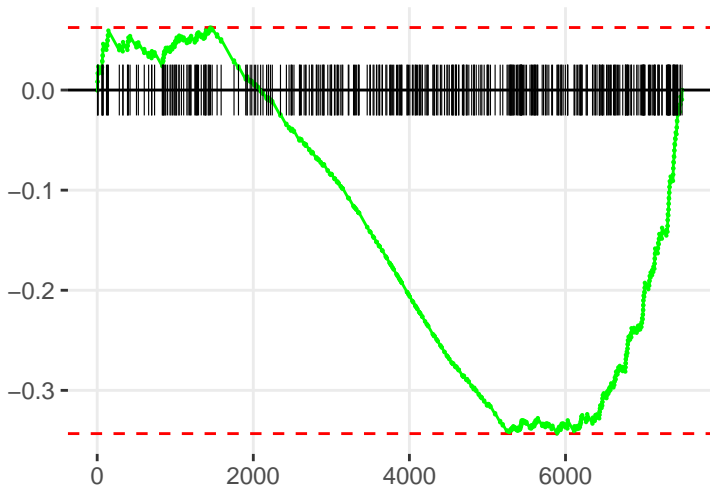
enrichment score



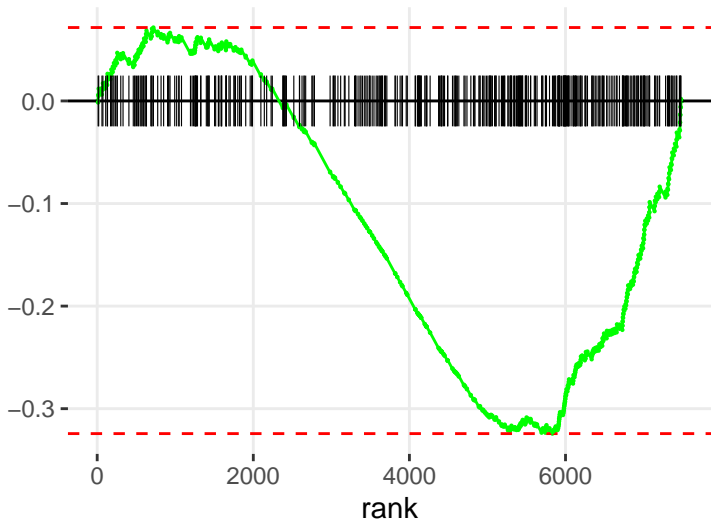
RESPONSE TO OXYGEN-CONTAINING COMPOUND%GOBP%GO:1901700

enrichment score

rank



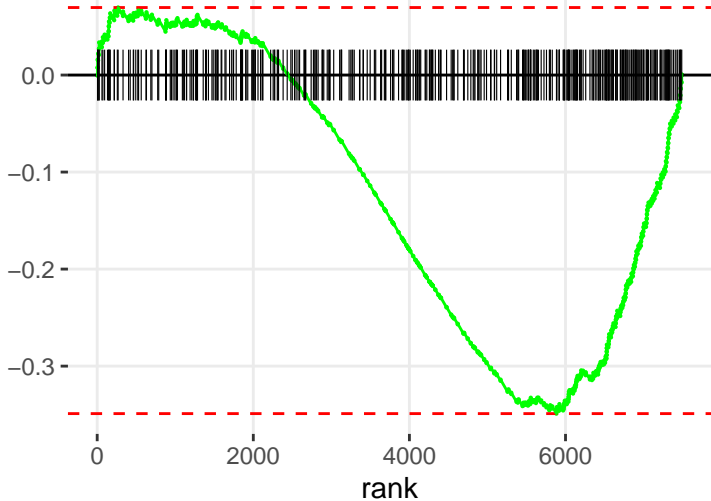
enrichment score





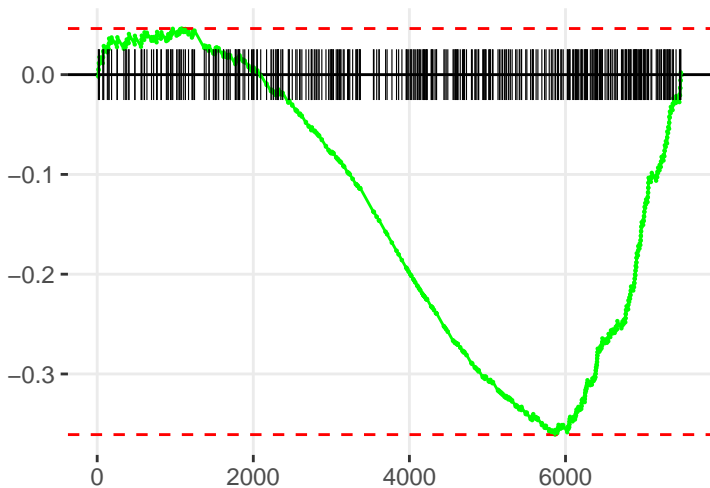
LEUKOCYTE ACTIVATION%GOBP%GO:0045321

enrichment score

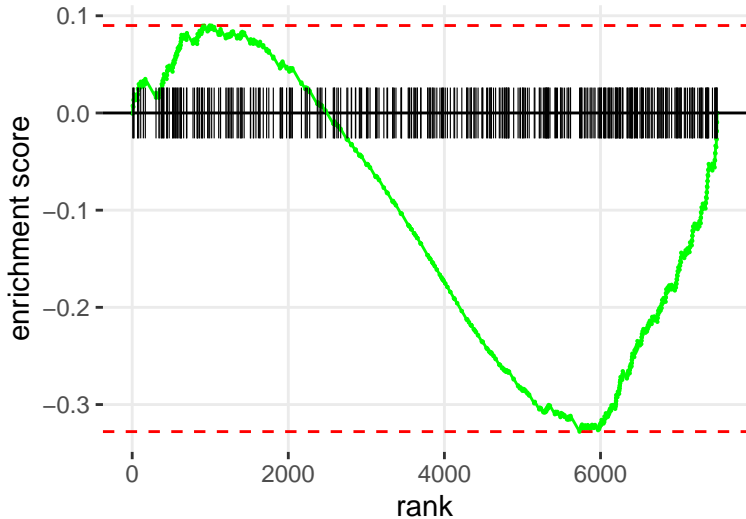


enrichment score

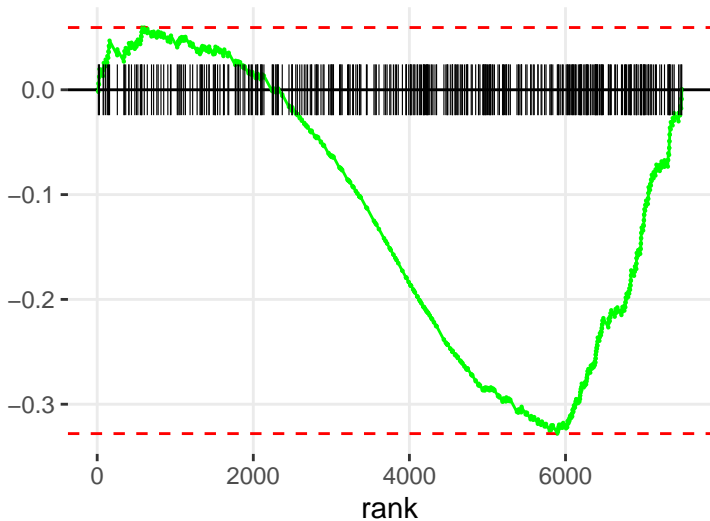
rank

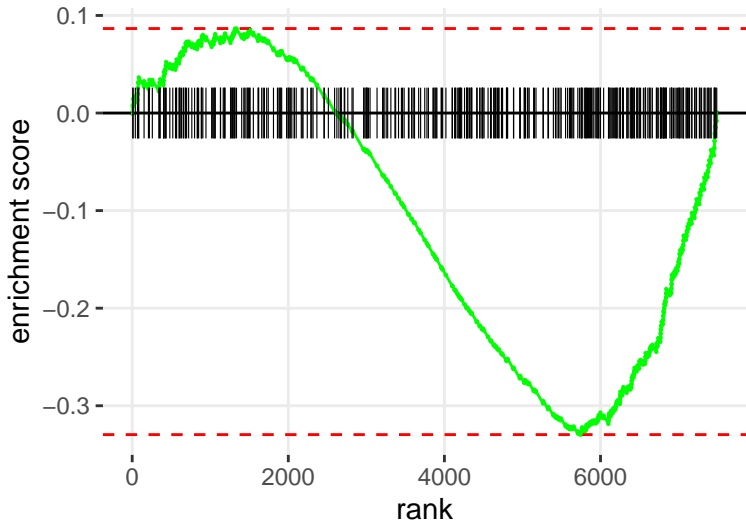


POSITIVE REGULATION OF PHOSPHORYLATION%GOBP%GO:0042327

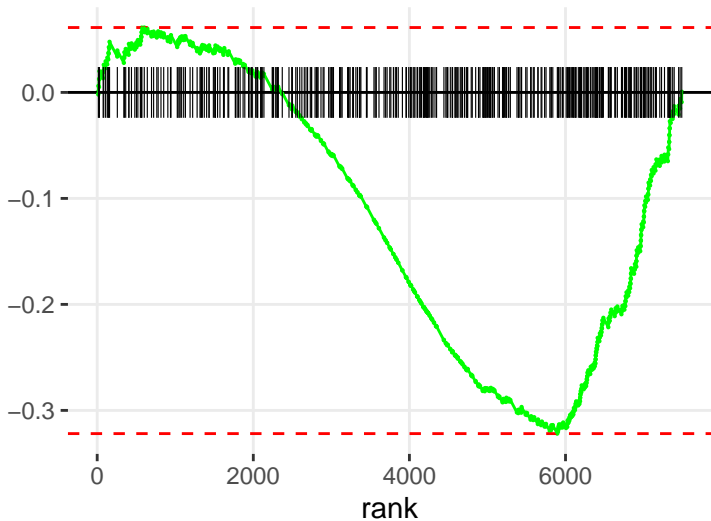


enrichment score





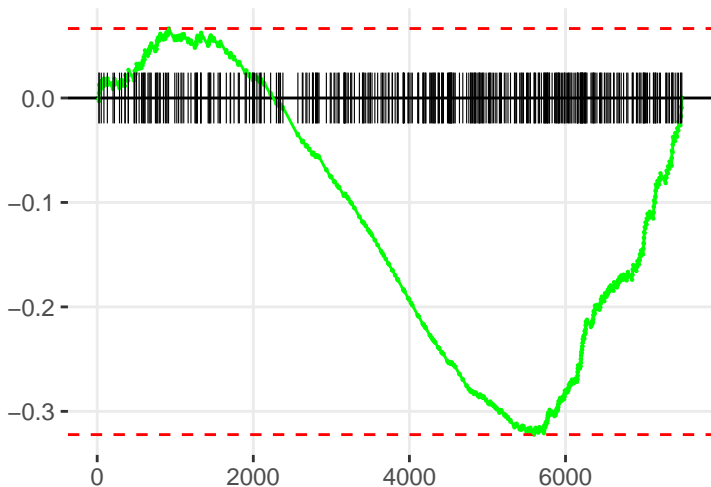
enrichment score



PROTEIN CATABOLIC PROCESS%GOBP%GO:0030163

enrichment score

rank



enrichment score

rank

0.0

-0.1

-0.2

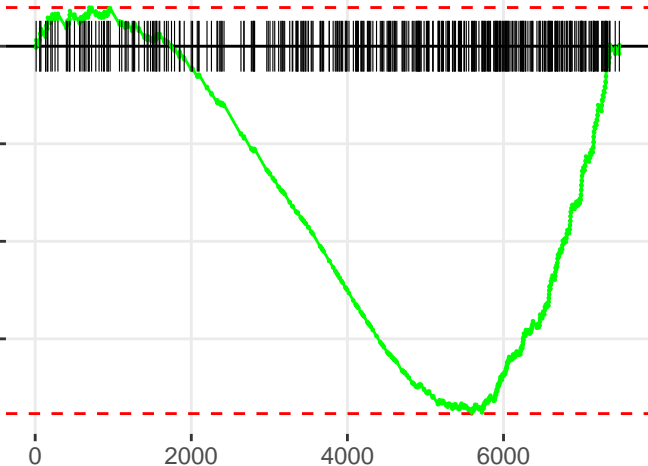
-0.3

0

2000

4000

6000

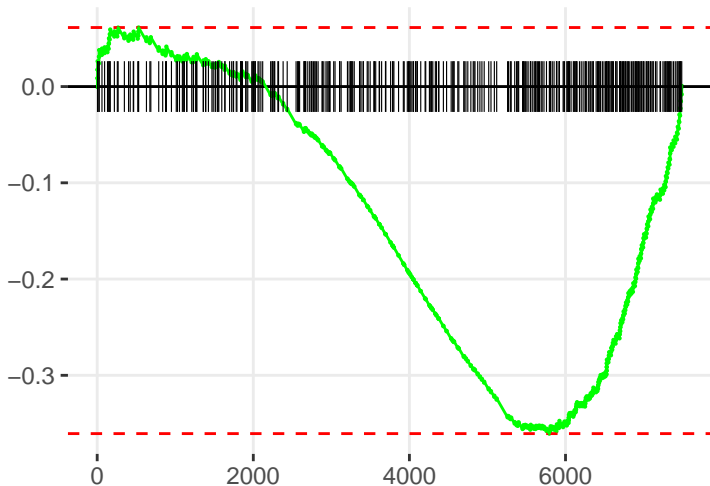




REGULATED EXOCYTOSIS%GOBP%GO:0045055

enrichment score

rank



PROTEIN PHOSPHORYLATION%GOBP%GO:0006468

enrichment score

0.0

-0.1

-0.2

-0.3

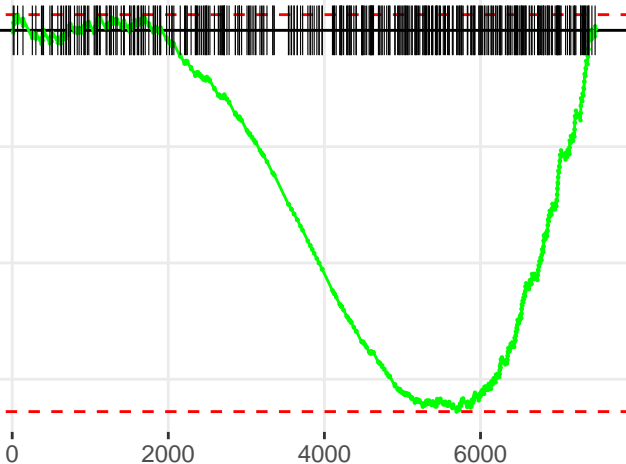
0

2000

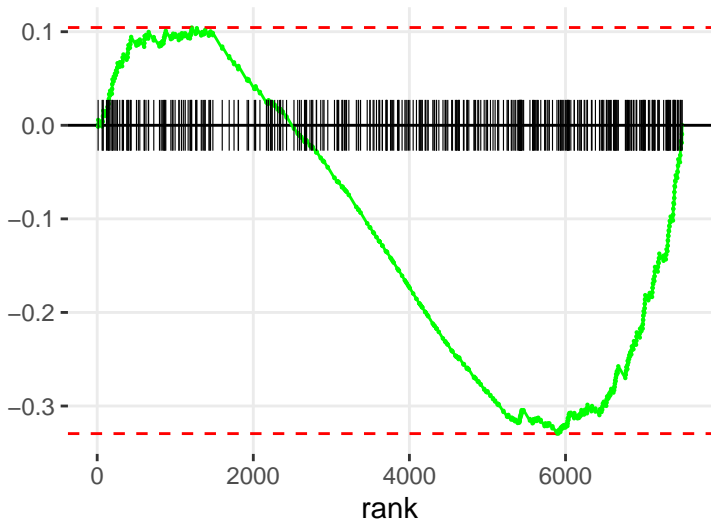
4000

6000

rank



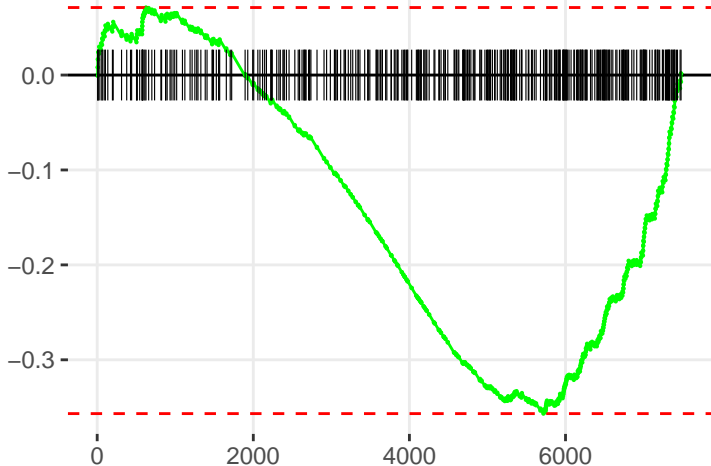
enrichment score



RESPONSE TO CYTOKINE%GOBP%GO:0034097

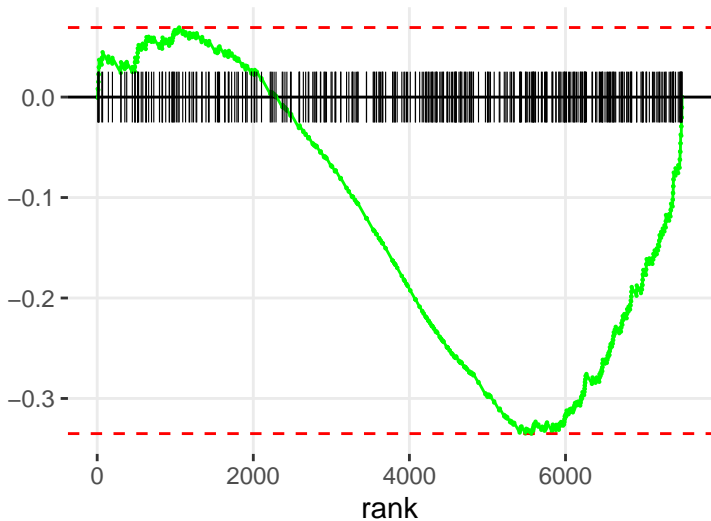
enrichment score

rank



REGULATION OF KINASE ACTIVITY%GOBP%GO:0043549

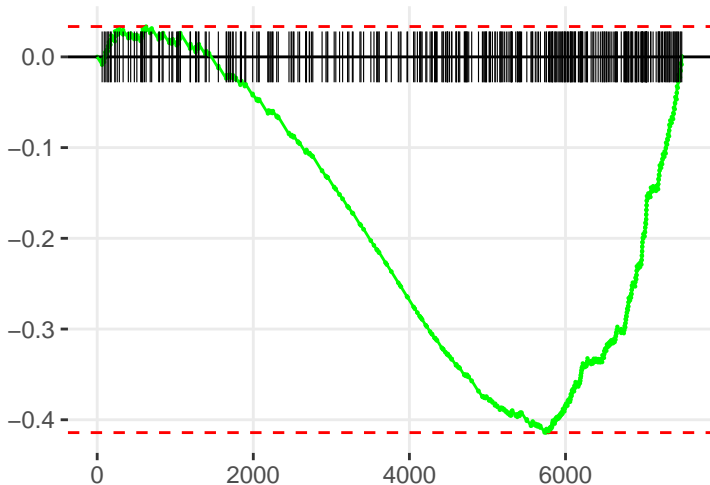
enrichment score



EGFR1%IOB%EGFR1

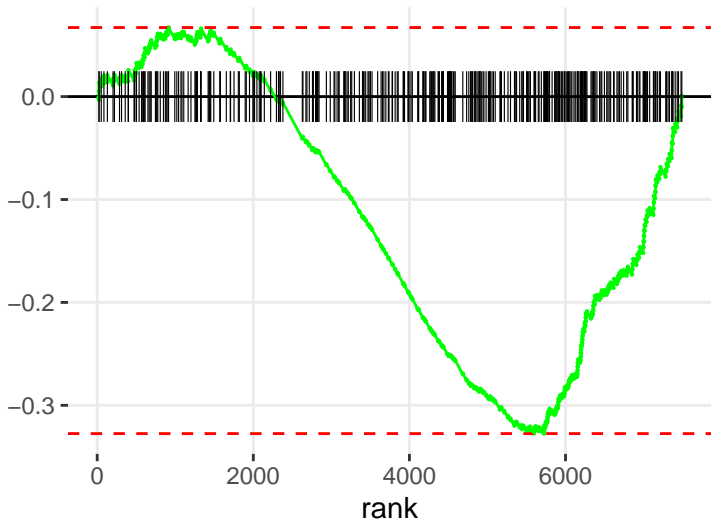
enrichment score

rank



CELLULAR PROTEIN CATABOLIC PROCESS%GOBP%GO:0044257

enrichment score



EGFR1%NETPATH%EGFR1

enrichment score

rank

0.0

-0.1

-0.2

-0.3

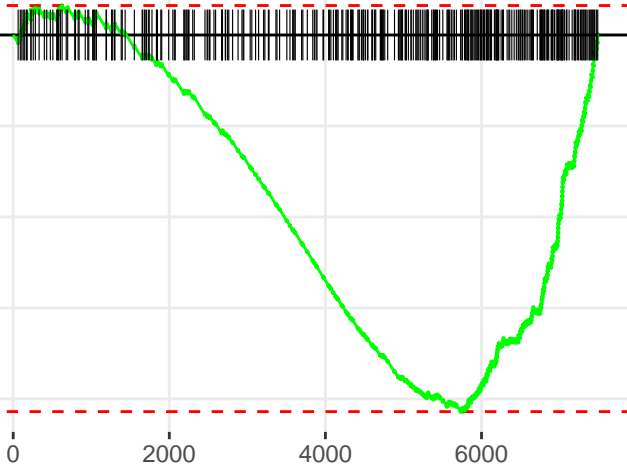
-0.4

0

2000

4000

6000





enrichment score

0.0

-0.1

-0.2

-0.3

-0.4

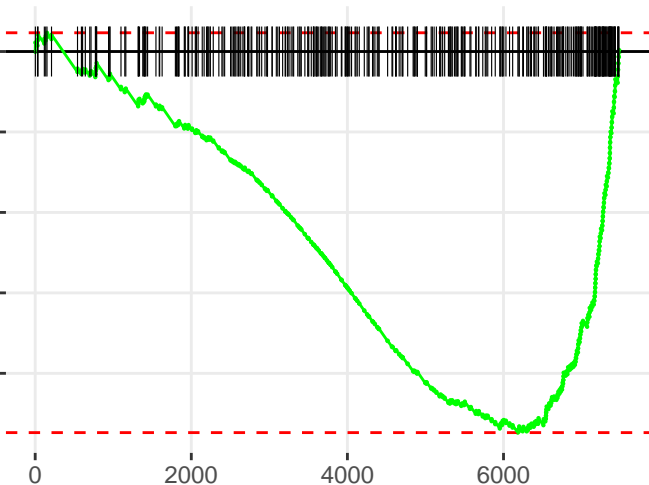
0

2000

4000

6000

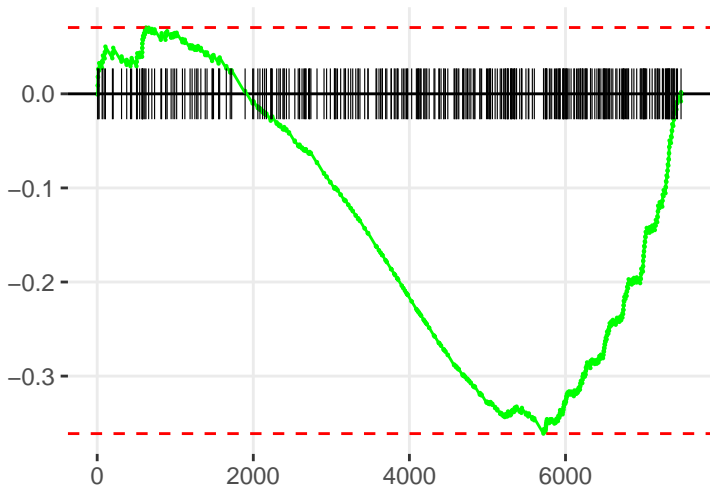
rank



CELLULAR RESPONSE TO CYTOKINE STIMULUS%GOBP%GO:0071345

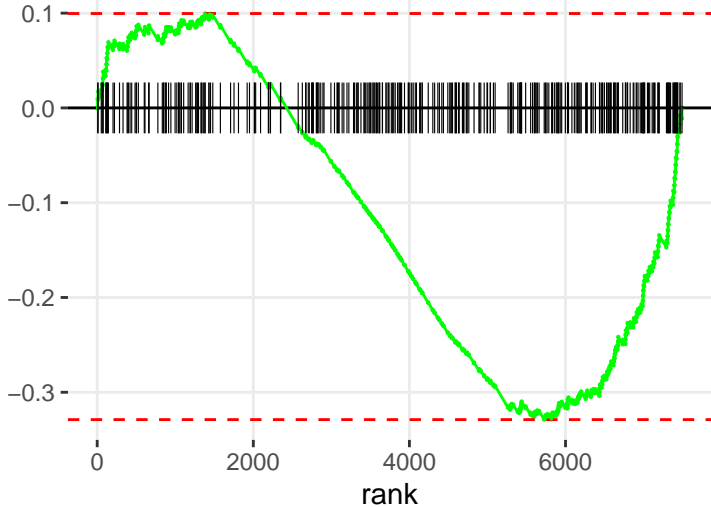
enrichment score

rank



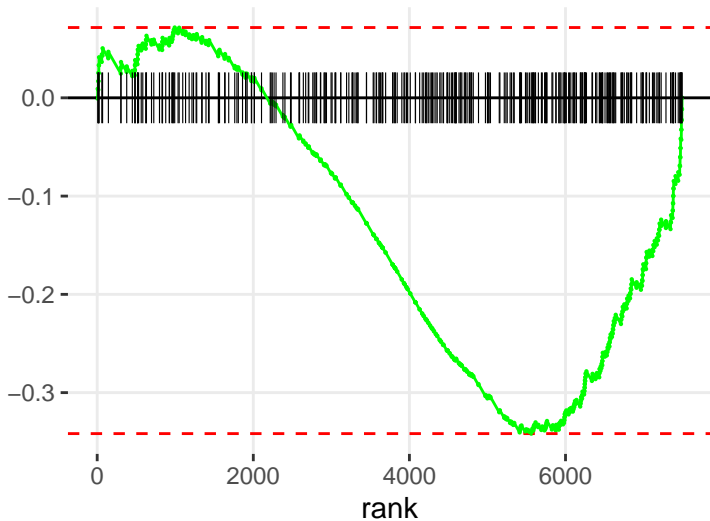
RESPONSE TO NITROGEN COMPOUND%GOBP%GO:1901698

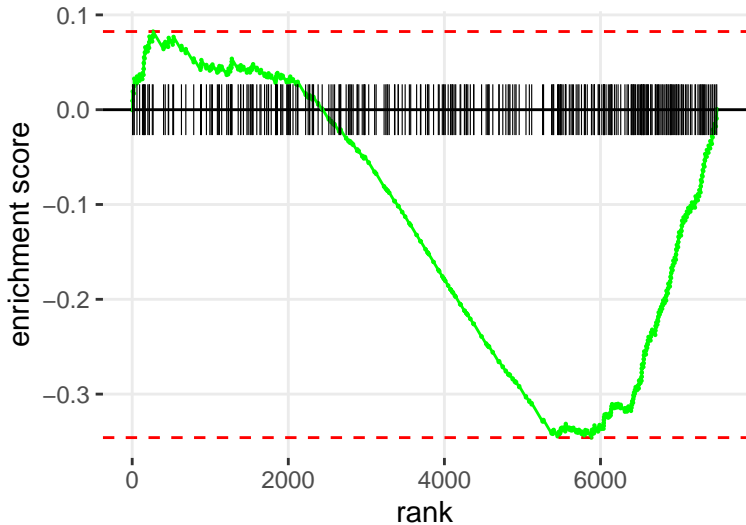
enrichment score



REGULATION OF PROTEIN KINASE ACTIVITY%GOBP%GO:0045859

enrichment score





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

enrichment score

0.0

-0.1

-0.2

-0.3

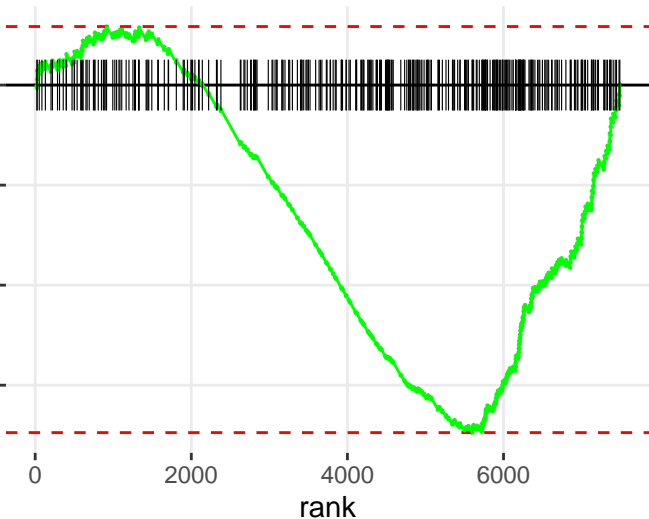
0

2000

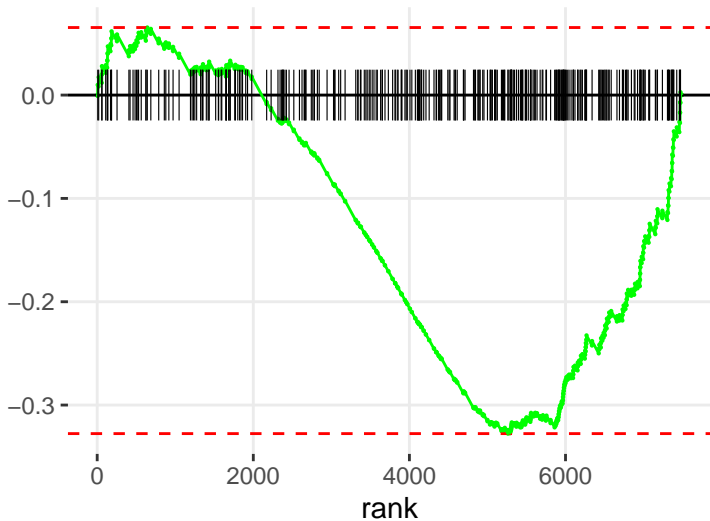
4000

6000

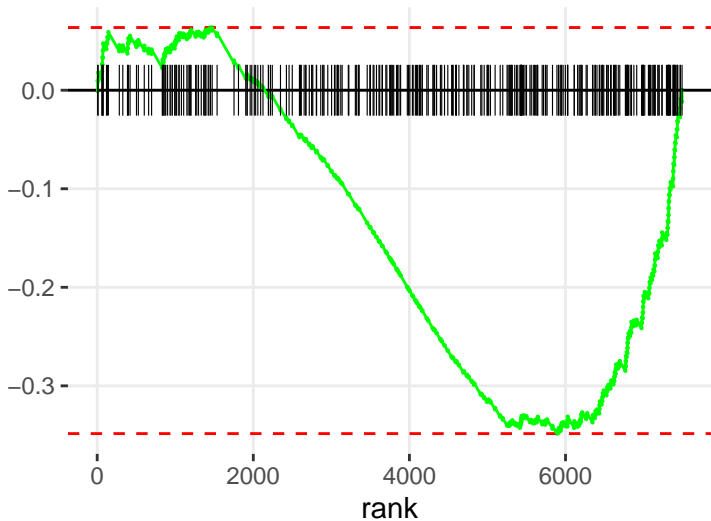
rank



enrichment score

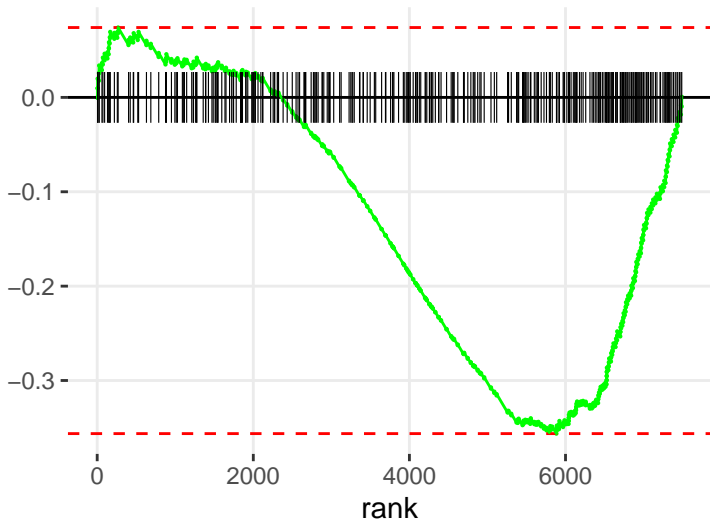


enrichment score





enrichment score



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

enrichment score

