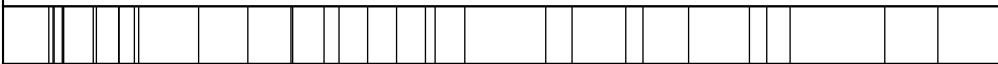
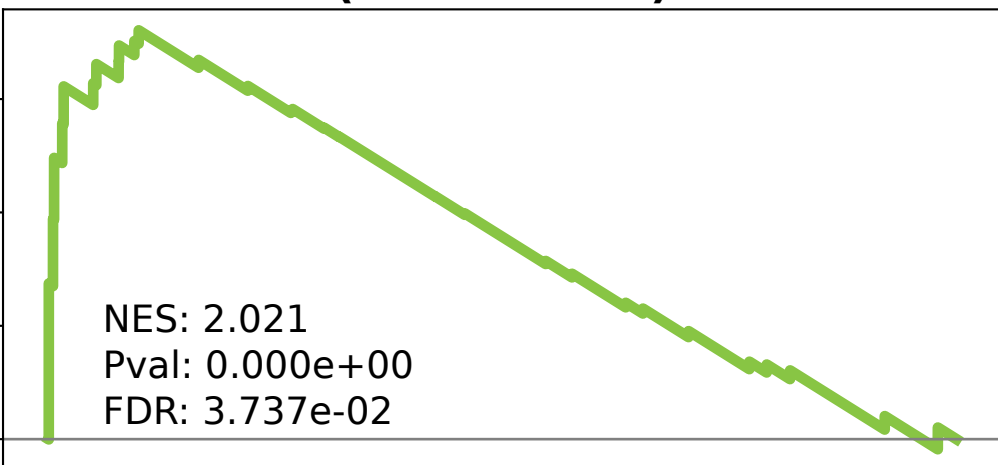


positive regulation of ATP metabolic process (GO:1903580)

Enrichment Score

0.6
0.4
0.2
0.0

NES: 2.021
Pval: 0.000e+00
FDR: 3.737e-02



Pos

Zero score at 5270

Neg

0 2500 5000 7500 10000 12500 15000

Gene Rank