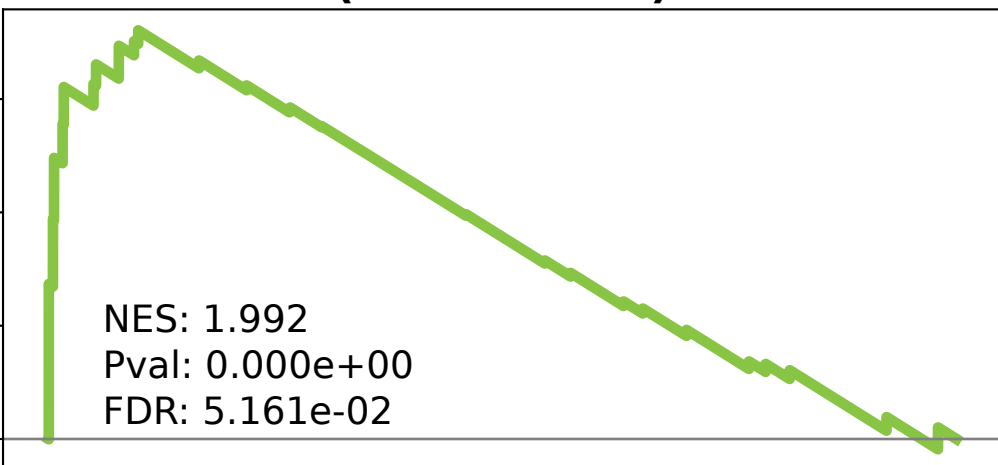


positive regulation of ATP metabolic process (GO:1903580)

Enrichment Score

0.6
0.4
0.2
0.0

NES: 1.992
Pval: 0.000e+00
FDR: 5.161e-02



Pos

Zero score at 5275

Neg

0 2500 5000 7500 10000 12500 15000

Gene Rank