

# ATPase activity, coupled to transmembrane movement of ions, rotational mechanism (GO:0044769)

Enrichment Score

0.0

-0.2

-0.4

-0.6

NES: -1.847  
Pval: 0.000e+00  
FDR: 1.535e-01

Ranked metric

5.0

2.5

0.0

-2.5

-5.0

0

2500

5000

Gene Rank

7500

10000

12500

15000

Pos

Zero score at 6714

Neg

