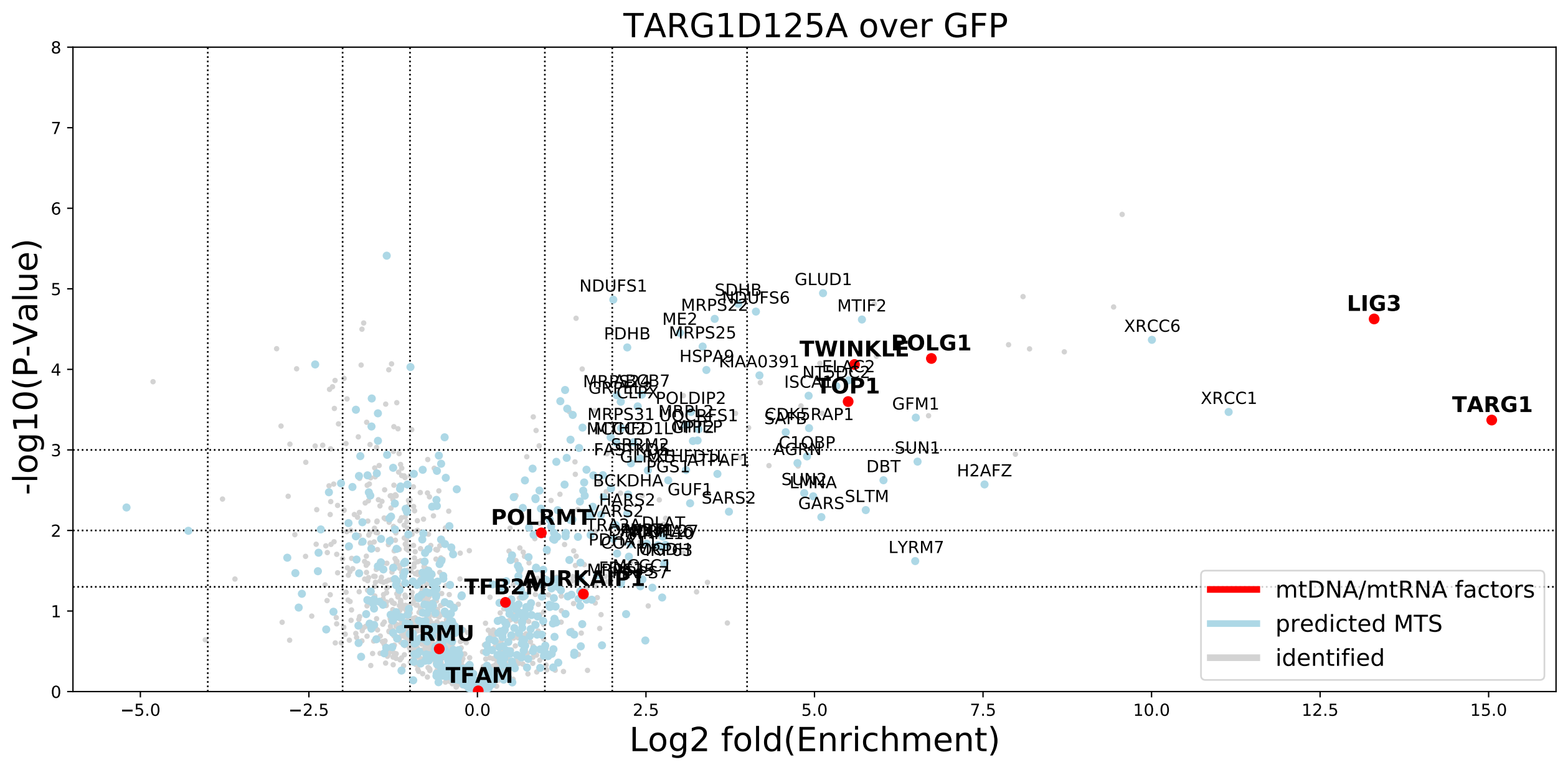
TARG1D125A over GFP
-log10(P-Value)
Log2 fold(Enrichment)
mtDNA/mtRNA factors
predicted MTS
identified
LIG3
TARG1
POLG1
TWINKLE
TOP1
POLRMT
TFB2M
AURKAIP1
TRMU
TFAM
NDUFS1
SDHB
NDUFS6
GLUD1
MTIF2
MRPS22
ME2
PDHB
MRPS25
HSPA9
KIAA0391
XRCC6
XRCC1
GFM1
POLDIP2
MRPS31
SAFB
CDK5RAP1
C1QBP
AGRN
SUN1
DBT
H2AFZ
PGS1
BCKDHA
GUF1
SARS2
HARS2
VARS2
GARS
SLTM
LYRM7
−5.0
−2.5
0.0
2.5
5.0
7.5
10.0
12.5
15.0
0
1
2
3
4
5
6
7
8