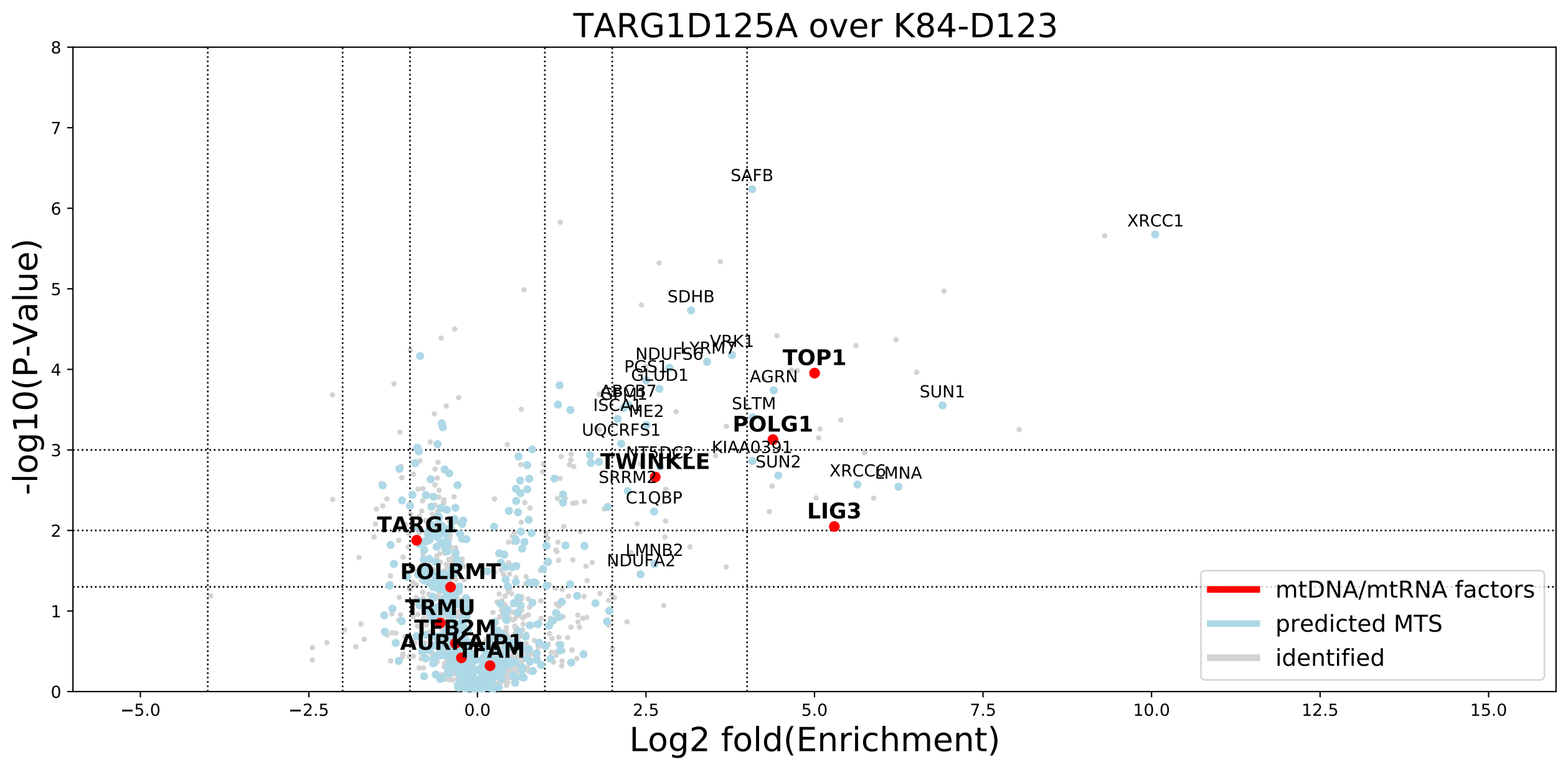
TARG1D125A over K84-D123
-log10(P-Value)
Log2 fold(Enrichment)
mtDNA/mtRNA factors
predicted MTS
identified
TARG1
POLRMT
TRMU
TFB2M
AURKAIP1
TFAM
TWINKLE
POLG1
TOP1
LIG3
SAFB
XRCC1
SDHB
VRK1
LYRM7
NDUFS6
PGS1
GLUD1
AGRN
SUN1
ISCA1
ME2
SLTM
UQCRFS1
KIAA0391
SUN2
XRCC6
LMNA
SRRM2
C1QBP
LMNB2
NDUFA2