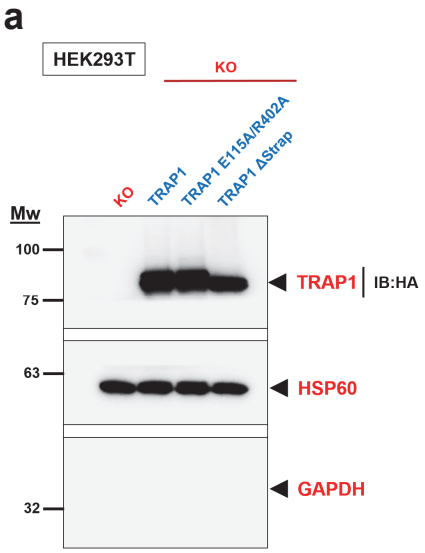
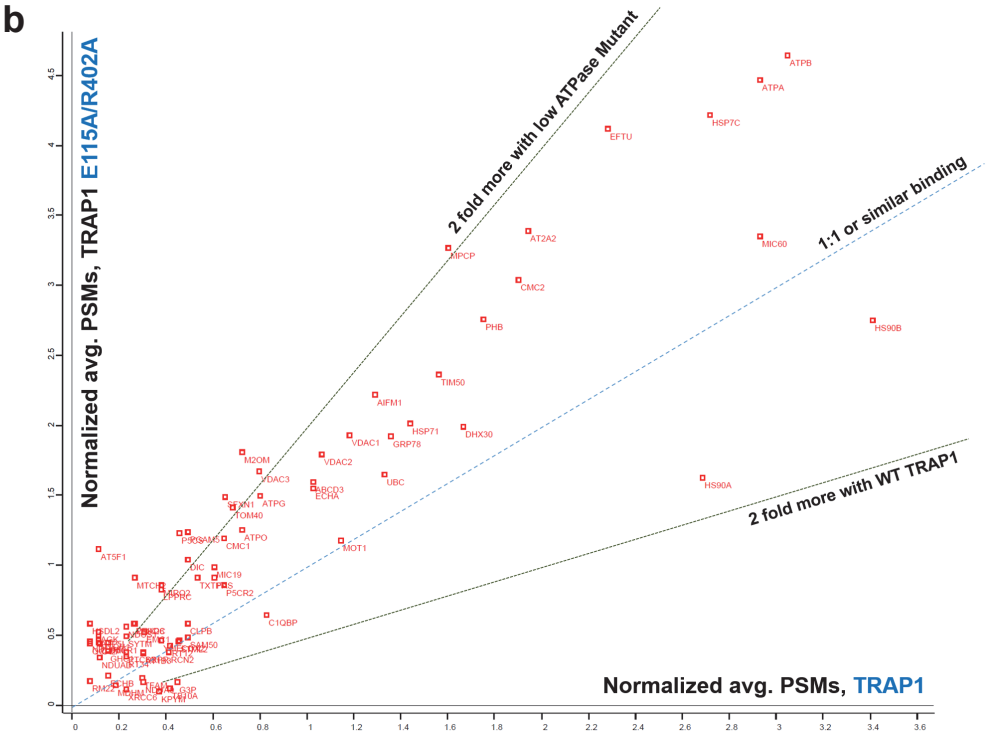
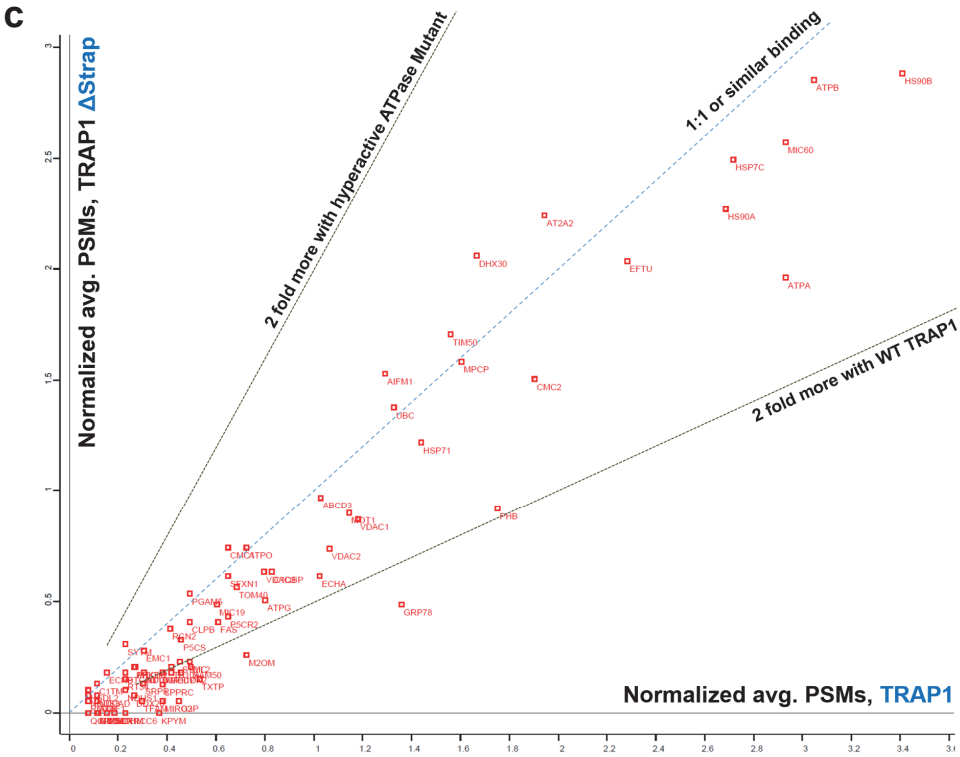
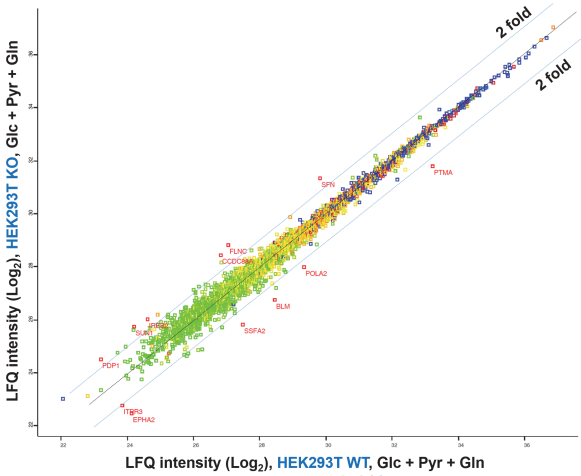
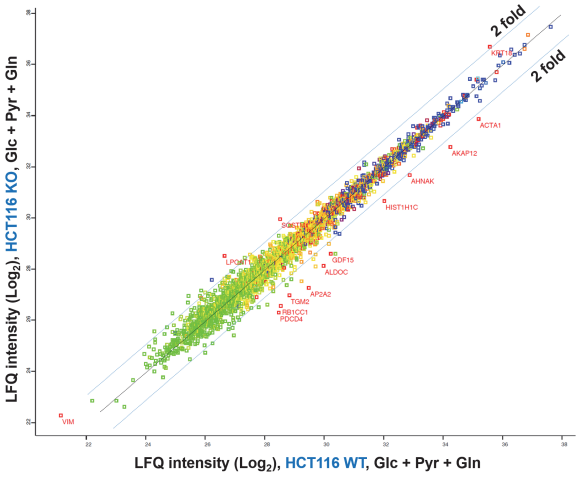
Figure S4

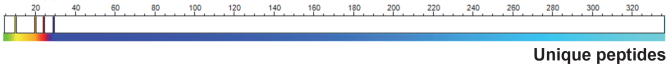

a
HEK293T
KO
KO
TRAP1
TRAP1 E115A/R402A
TRAP1 ΔStrap
Mw
100
75
63
32
TRAP1
IB:HA
HSP60
GAPDH


b
Normalized avg. PSMs, TRAP1 E115A/R402A
Normalized avg. PSMs, TRAP1
2 fold more with low ATPase Mutant
1:1 or similar binding
2 fold more with WT TRAP1


c
Normalized avg. PSMs, TRAP1 ΔStrap
Normalized avg. PSMs, TRAP1
2 fold more with hyperactive ATPase Mutant
1:1 or similar binding
2 fold more with WT TRAP1


d
LFQ intensity (Log2), HEK293T KO, Glc + Pyr + Gln
LFQ intensity (Log2), HEK293T WT, Glc + Pyr + Gln
2 fold
2 fold


e
LFQ intensity (Log2), HCT116 KO, Glc + Pyr + Gln
LFQ intensity (Log2), HCT116 WT, Glc + Pyr + Gln
2 fold
2 fold


Unique peptides