

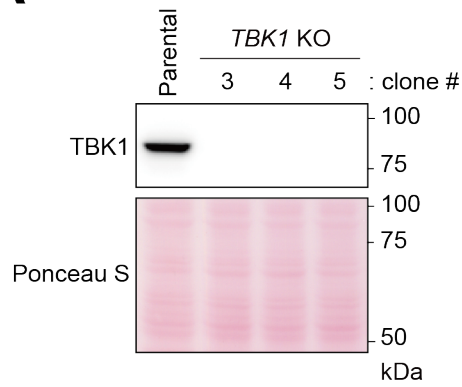
Phosphorylation tunes p62 condensates to drive autophagic degradation of ubiquitinated proteins

Appendix Contexts

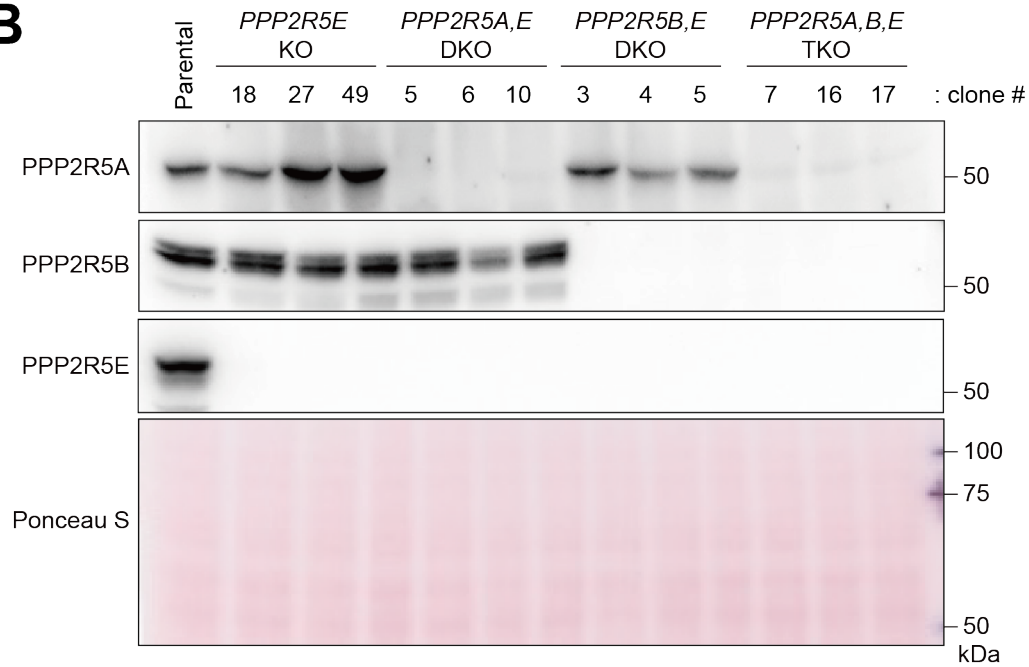
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A



B



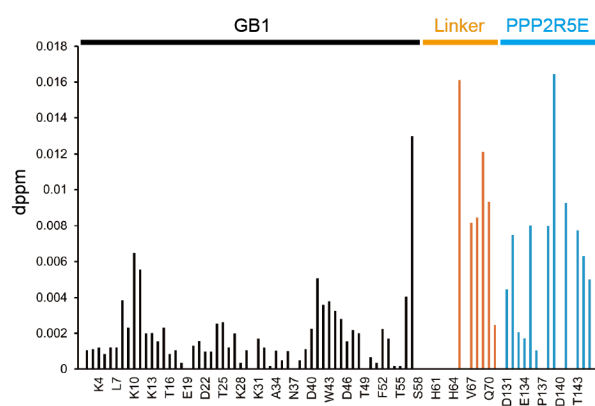
Appendix Figure S1. Generation of knockout cell lines.

(A, B) Immunoblot analysis of *TBK1*-knock-out (A) and *PPP2R5A/B/E* triple-knock-out (B) Huh-1 cells using the indicated antibodies. Results are representative of three independent experiments.



Appendix Figure S2. SDS-PAGE analysis of purified SNAP-KEAP1 and SNAP-PPP2R5E.

20 Purified SNAP-KEAP1 (90.1 kDa) and SNAP-PPP2R5E (74.1 kDa) were analyzed by 12% SDS-PAGE. Lane M, molecular weight marker (Precision Plus Protein™ Unstained Protein Standards (Bio-Rad)); lane 1, SNAP-KEAP1; lane 2, SNAP-PPP2R5E.



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Appendix Figure S3. The chemical shift perturbations.

The chemical shift perturbations upon the addition of KEAP1 in Figure 4E are plotted for each residue.