

Supplementary Information

ZNF506-dependent Positive Feedback Loop Regulates H2AX Signaling after DNA Damage

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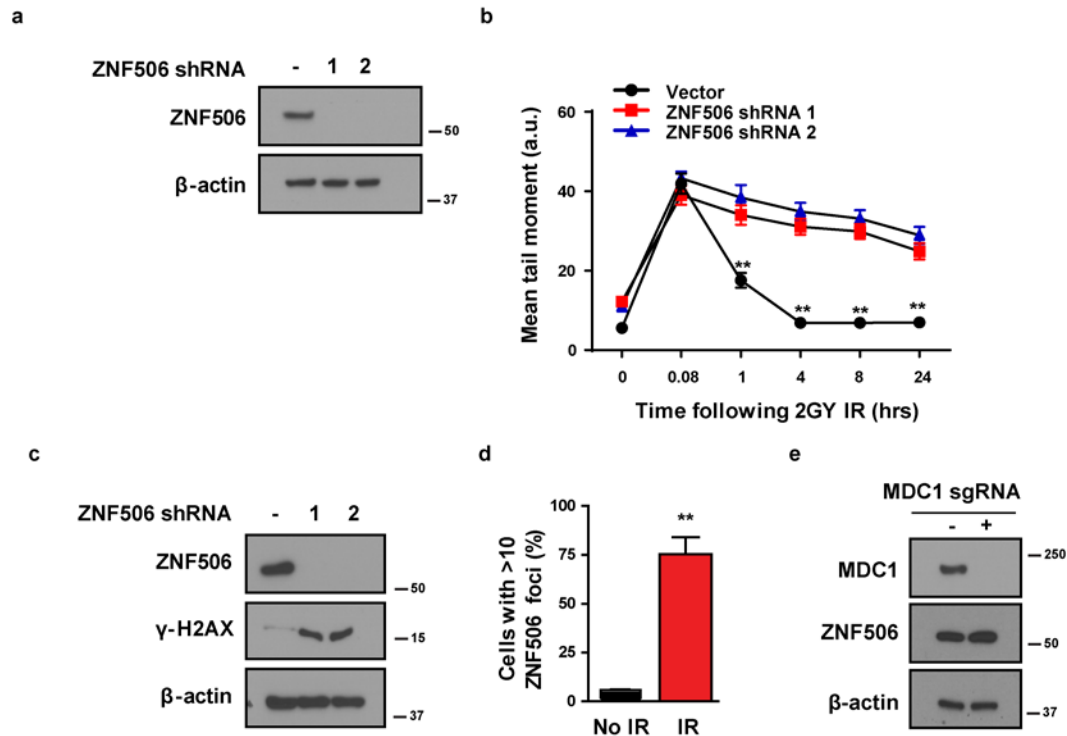
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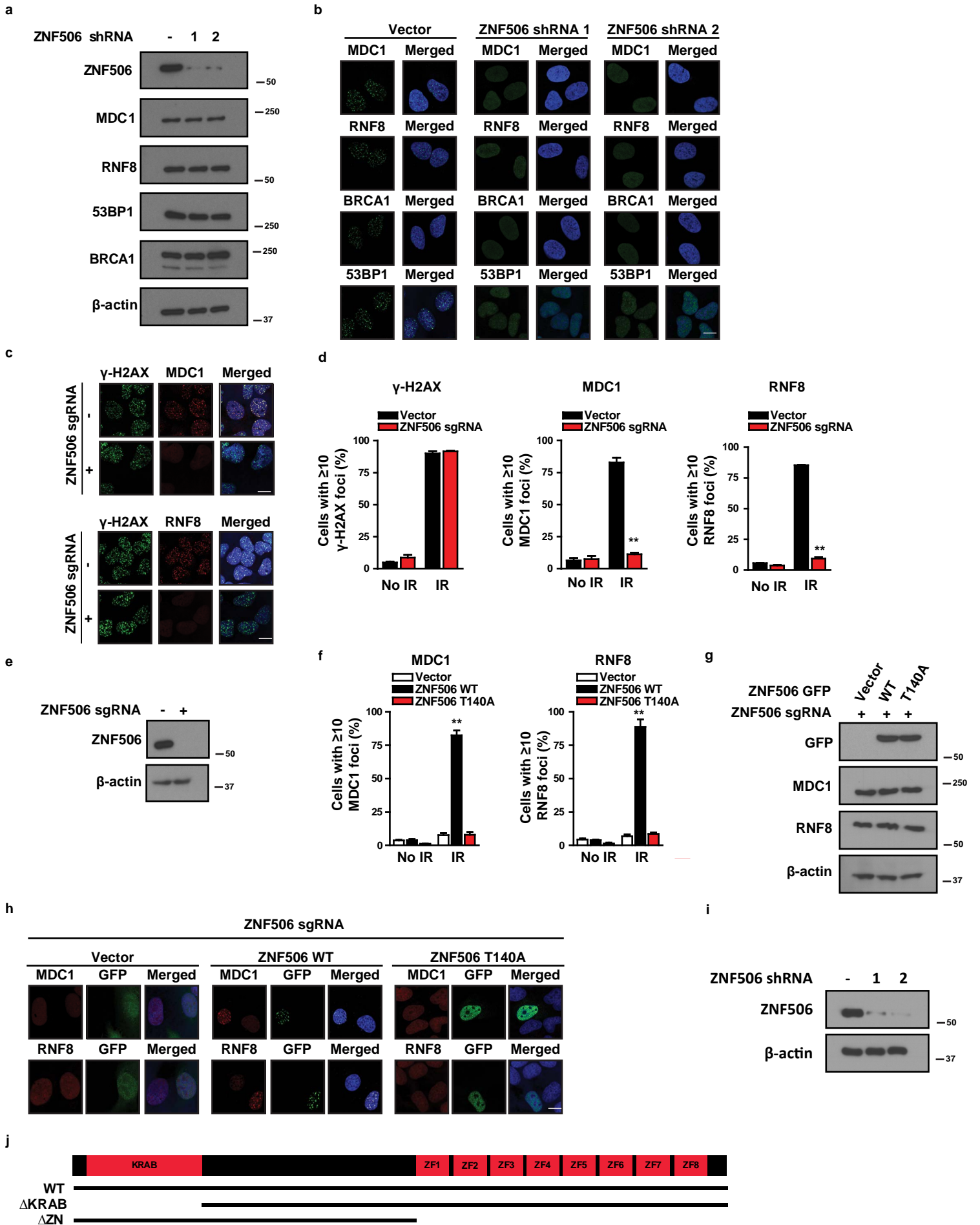


Supplementary Figure 1: ZNF506 regulates DNA damage response. **a-b**, Depletion of ZNF506 leads to DNA repair defect. **a**, Representative western blot showing knockdown efficiency of ZNF506 using shRNAs. **b**, Quantification of mean tail moment of cells subjected to comet assay after the indicated treatment. **c**, Representative western blot showing knockdown efficiency of ZNF506 using shRNAs (**related to Figure 1a-b**). **d**, Quantification of U2OS cells with ZNF506 foci (**related to Figure 1d**). **e**, Representative western blot showing the protein level of ZNF506 in MDC1 knockout U2OS cells (**related to Figure 1e-f**). Shown are the representative data (Mean \pm S.E.M.) from three independent experiments in **b** and **d**. ** $p < 0.01$ by ANOVA for vector vs all other groups at each time point in **b**. ** $p < 0.01$ by t-test for no irradiation (IR) vs IR in **d**. Representative western blots in **a**, **c**, and **e** are provided from 3 biologically independent experiments. Unprocessed blots are provided in **Supplementary Figure 5**.

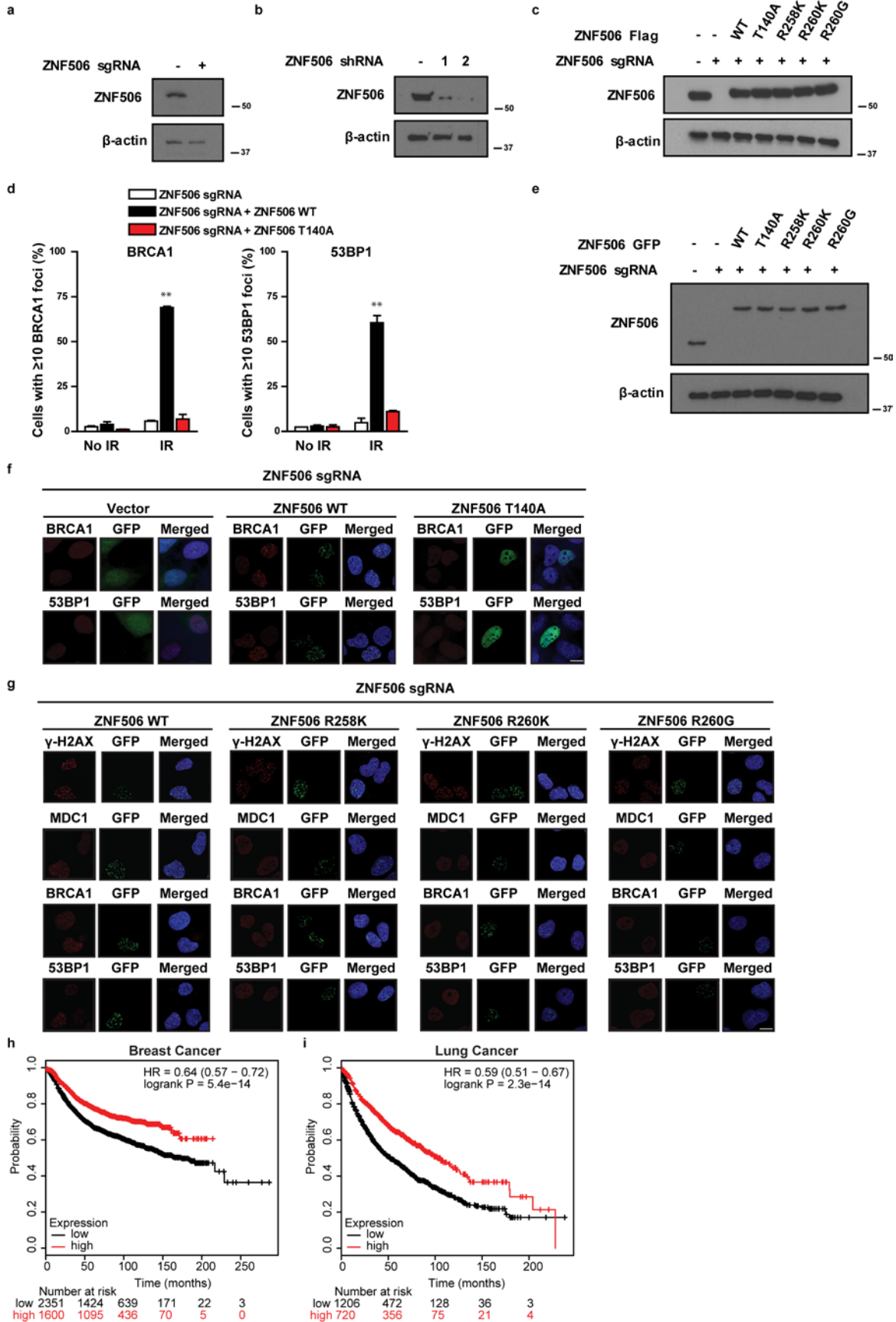
a

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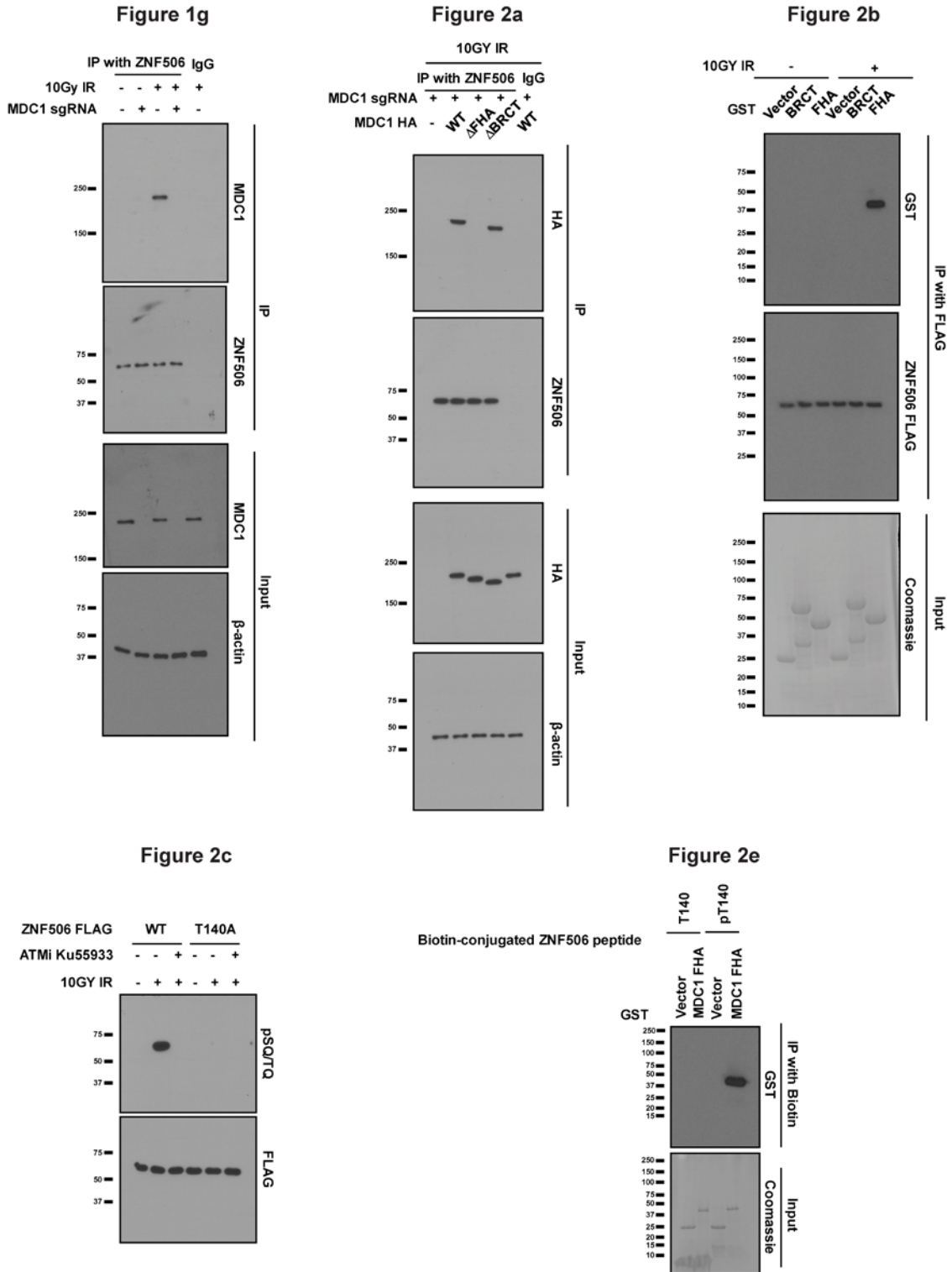
Supplementary Figure 2: a, Protein sequence of ZNF506. The putative ATM phosphorylation site is shown in red and underlined (**related to Figure 2c**).



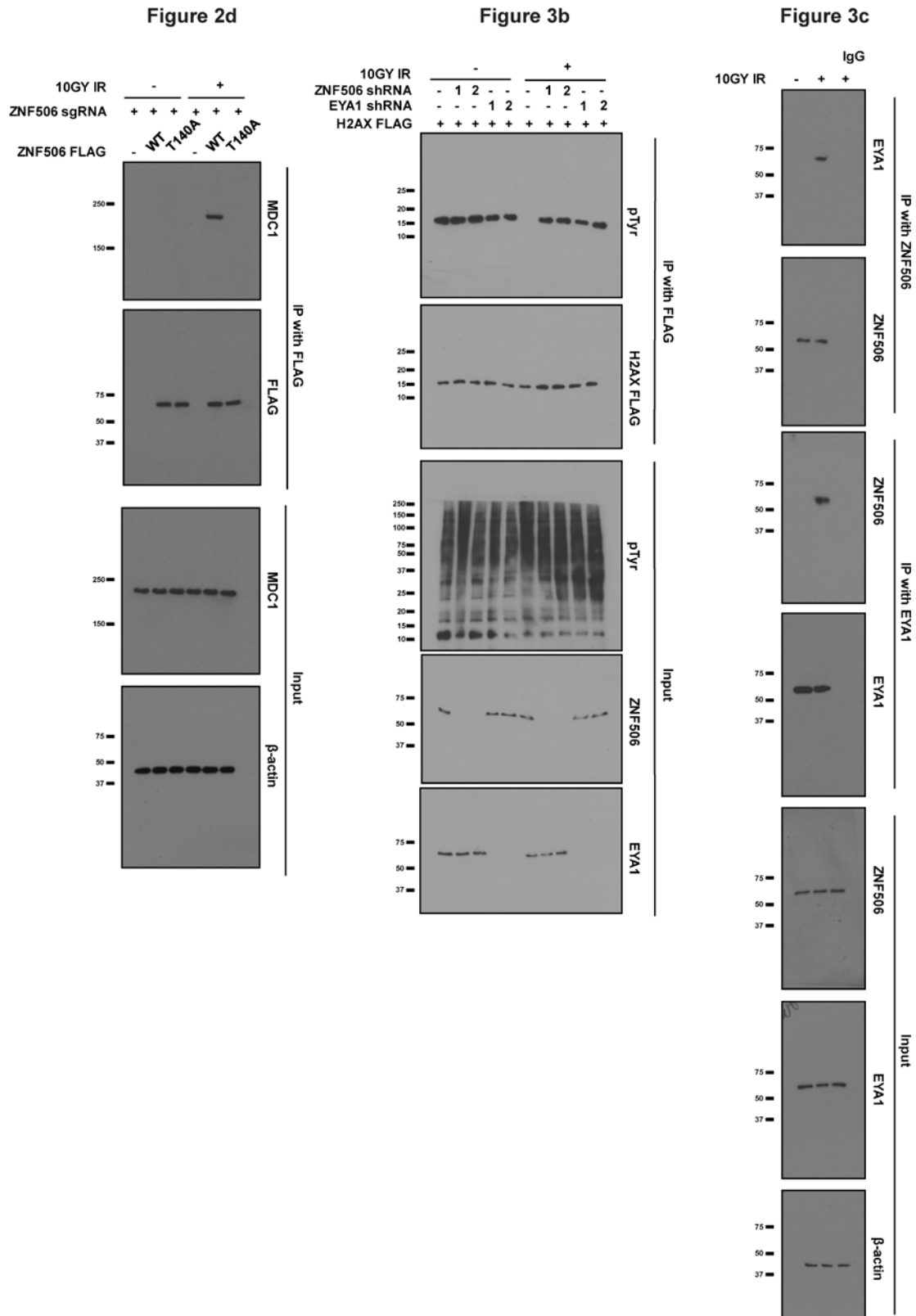
Supplementary Figure 3: ZNF506 modulates DNA damage response. a-b, Knockdown of ZNF506 affects localization of DNA repair proteins to the DNA double strand break site (**related to Figure 3a**). **a,** Representative western blot showing the expression level of various DNA repair proteins is not altered with ZNF506 knockdown. **b,** Representative images of MDC1, RNF8, BRCA1 and 53BP1 foci in U2OS cells after the indicated treatments. Nucleus is stained with DAPI (blue). **c-e,** ZNF506 knockout affects localization of DNA repair proteins to the DNA double strand break site. **c,** Representative images of γ -H2AX, MDC1, and RNF8 foci in U2OS cells after the indicated treatments. Nucleus is stained with DAPI (blue). **d,** Quantification of γ -H2AX, MDC1, and RNF8 foci in U2OS cells after the indicated treatments. **e,** Representative western blot showing the ZNF506 knockout efficiency. **f-h,** Expression of wild-type (WT) but not the phosphorylation mutant (T140A) of ZNF506 (green) restores foci formation of DNA repair proteins. **f,** Quantification of MDC1 and RNF8 foci in U2OS cells after the indicated treatments. **g,** Representative western blot showing the expression level of the indicated plasmids and repair proteins. **h,** Representative images of MDC1 and RNF8 foci in U2OS cells after the indicated treatments. Nucleus is stained with DAPI (blue). **i,** Representative western blot showing the knockdown efficiency of the ZNF506 shRNAs (**related to Figure 3d-e**). **j,** Structure of ZNF506 (**related to Figure 3g-h**). ZNF506 has a KRAB and eight zinc finger (ZN) domains. Shown are the representative data (Mean \pm S.E.M.) from three independent experiments in **d** and **f**. ****p**<0.01 by ANOVA for vector vs knockout with irradiation (IR) in **d**. ****p**<0.01 by ANOVA for ZNF506 WT vs all other groups with IR in **f**. Representative images of three independent experiments are shown in **b**, **c** and **h**. Scale bars, 10 μ m. Representative western blots in **a**, **e**, **g**, and **i** are provided from 3 biologically independent experiments. Unprocessed blots are provided in **Supplementary Figure 5**.



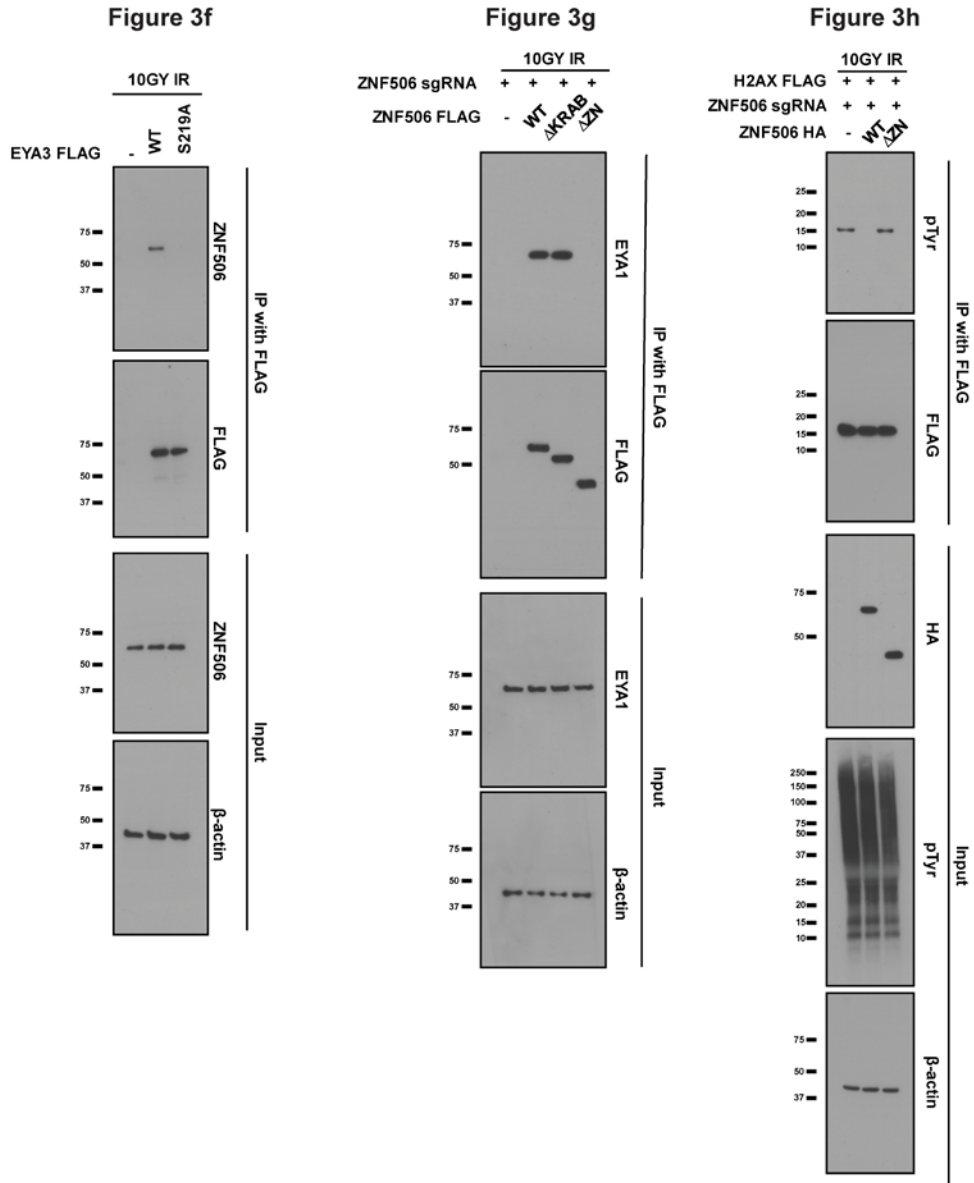
Supplementary Figure 4: ZNF506 is required for the maintenance of genomic integrity. **a**, Representative western blot showing knockout efficiency of ZNF506 using sgRNA in human fibroblasts (**related to Figure 4a**). **b**, Representative western blot showing knockdown efficiency of ZNF506 using shRNA in CH12F3-2a cells (**related to Figure 4b**). **c**, Representative western blot showing the expression level of the indicated plasmids in MDA-MB-231 cells in which ZNF506 was knocked out using sgRNA (**related to Figures 4c and 4g**). **d-f**, Phosphorylation of ZNF506 is required for recruitment of BRCA1 and 53BP1 to DNA damage sites. Cells were stained for the indicated foci (red). Nucleus is stained with DAPI (blue). **d**, Quantification of the indicated foci in U2OS cells after the indicated treatment. **e**, Representative western blot showing the expression level of the indicated plasmids in U2OS cells in which ZNF506 was knocked out using sgRNA (**also related to Figure 4e**). **f**, Representative images of the indicated foci in U2OS cells treated as indicated and fixed an hour after 2GY irradiation (IR). **g**, The mutations in ZNF506 that were identified in patient samples lead to aberrant DNA damage response. Shown are representative images of the indicated foci in U2OS cells after the indicated treatment (**related to Figure 4e**). **h-i**, Kaplan–Meier analysis showing a tight correlation between ZNF506 expression levels and patient survival. Data were from breast and lung cancer patients with low and high ZNF506 expression. Patient number at risk at different times of analyses is indicated at the bottom of the plots. The plots were generated using the KmPlot tool (<http://www.kmplot.com/lung> and <http://www.kmplot.com/breast>). Shown are the representative data (Mean \pm S.E.M.) from three independent experiments in **d**. ** $p < 0.01$ by ANOVA for ZNF506 sgRNA+ZNF506 WT vs all other groups with IR in **d**. Representative images of three independent experiments are shown in **f-g**. Scale bars, 10 μ m. Representative western blots in **a-c**, and **e** are provided from 3 biologically independent experiments. Unprocessed blots are provided in **Supplementary Figure 5**.



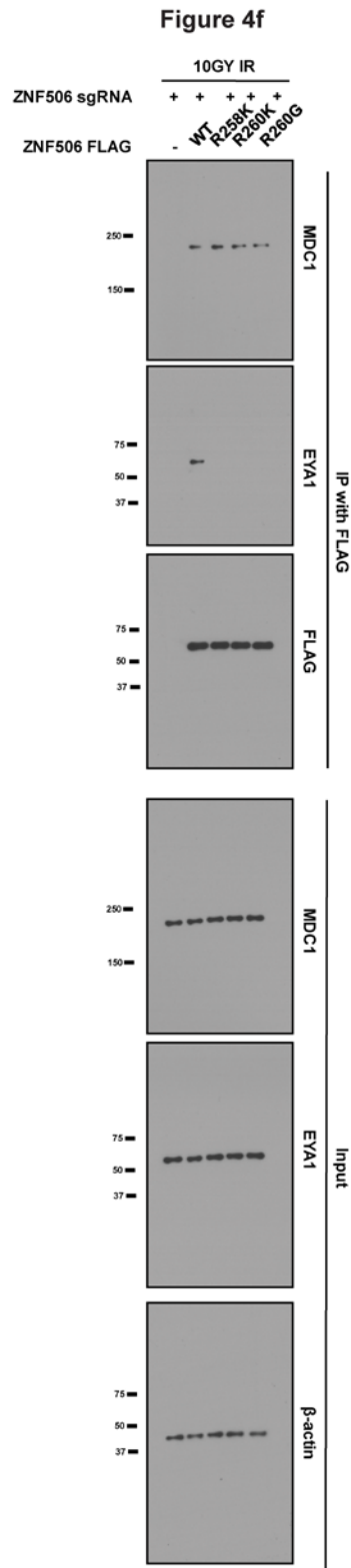
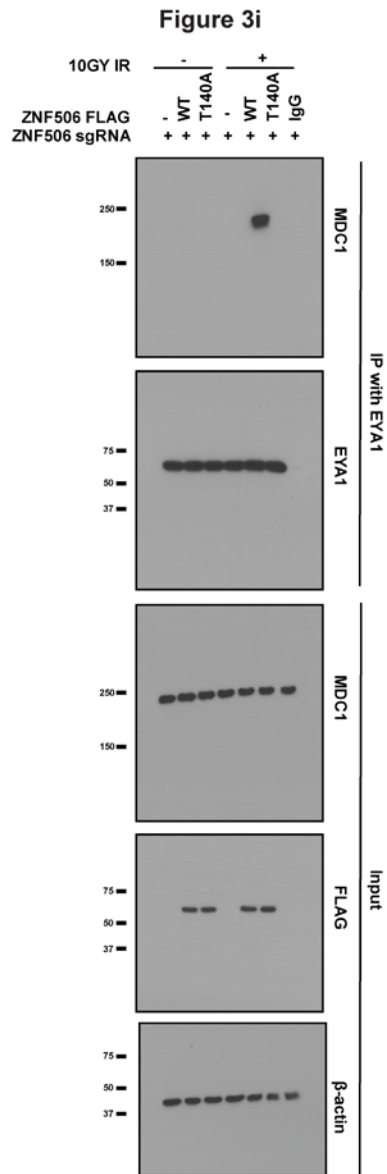
Supplementary Figure 5: Unprocessed images of all blots in this manuscript.



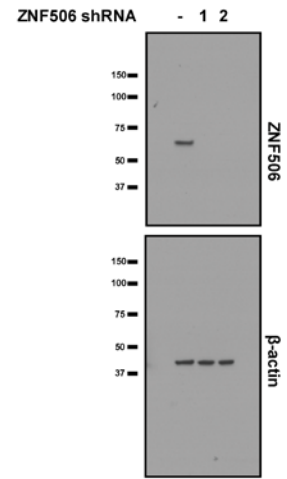
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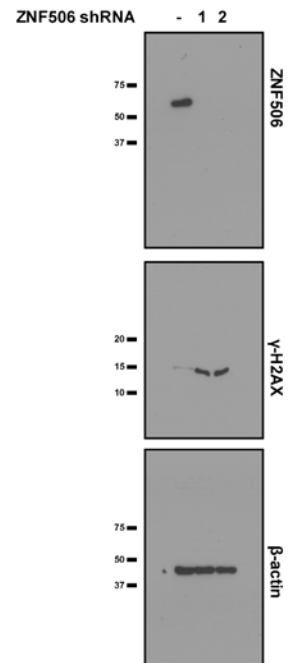
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Supplementary Figure 1a

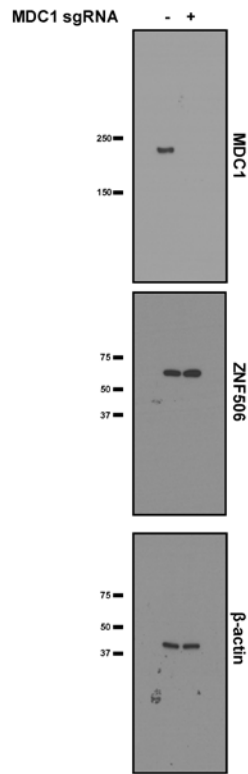


Supplementary Figure 1c

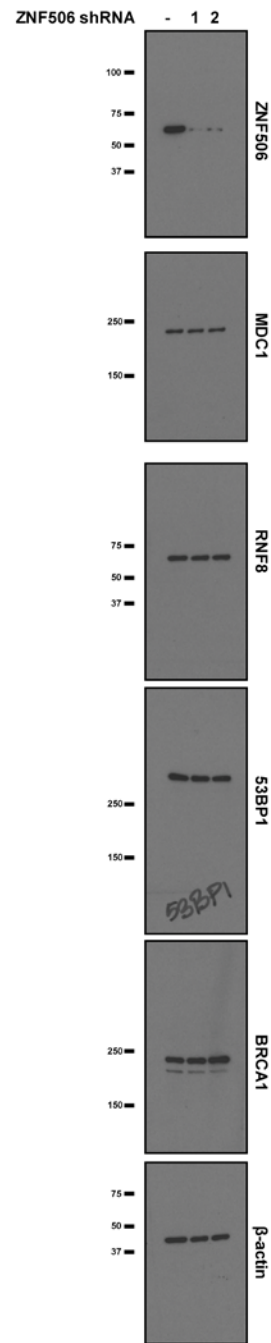


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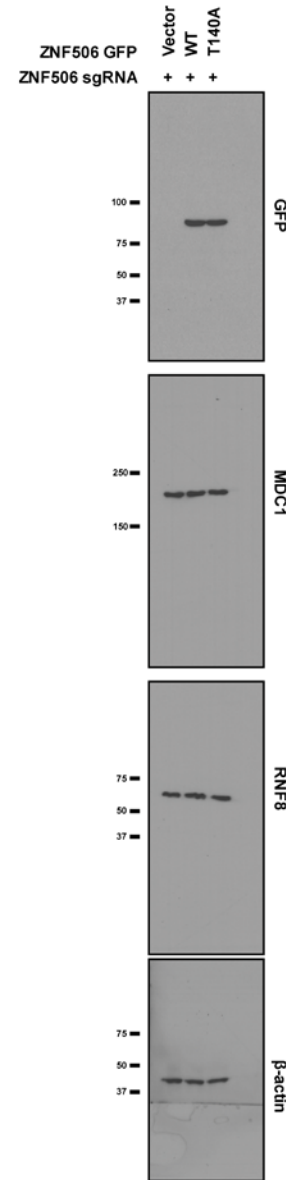
Supplementary Figure 1e



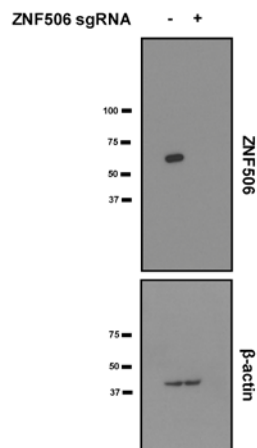
Supplementary Figure 3a



Supplementary Figure 3g

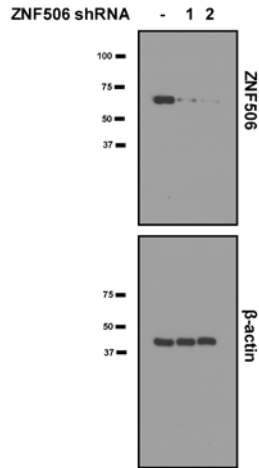


Supplementary Figure 3e

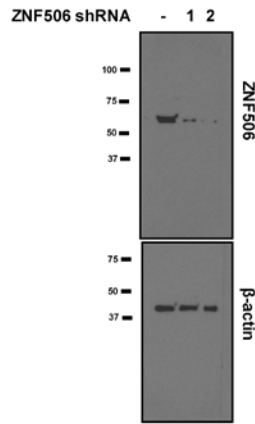


Supplementary Figure 5 continued: Unprocessed images of all blots in this manuscript.

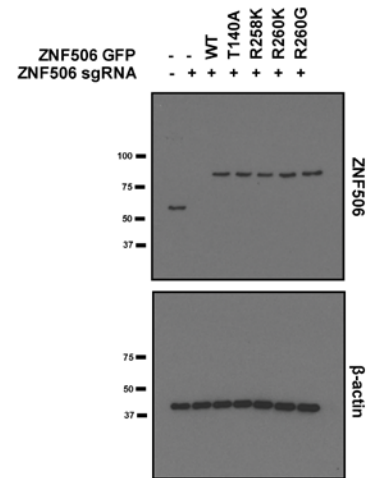
Supplementary Figure 3i



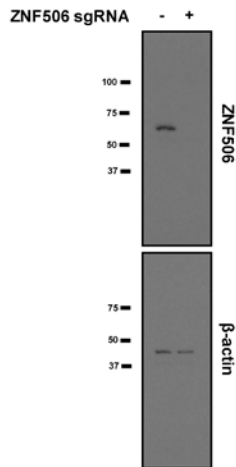
Supplementary Figure 4b



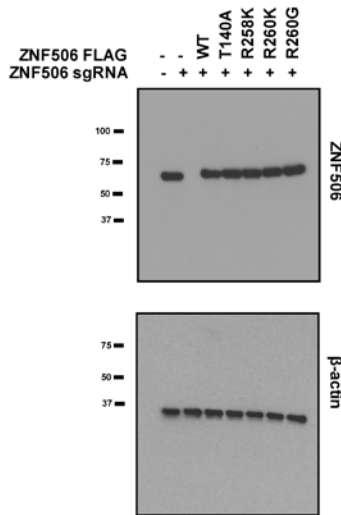
Supplementary Figure 4e



Supplementary Figure 4a



Supplementary Figure 4c



Supplementary Figure 5 continued: Unprocessed images of all blots in this manuscript.