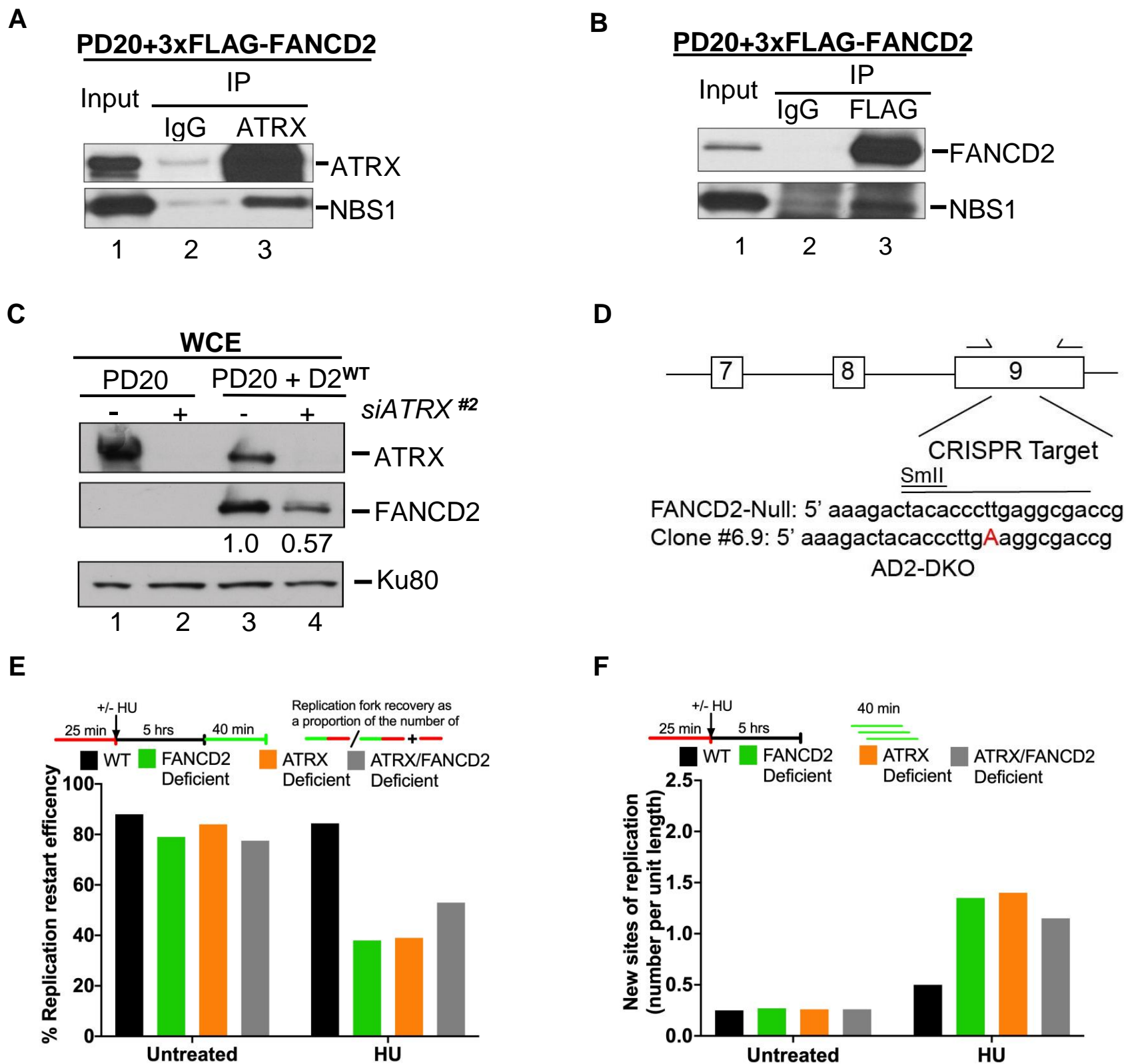
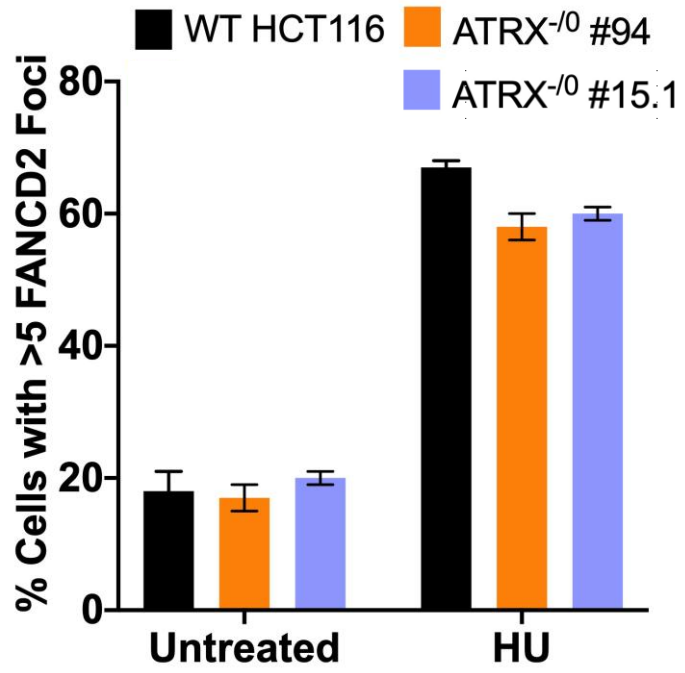


Supplementary Figure 1

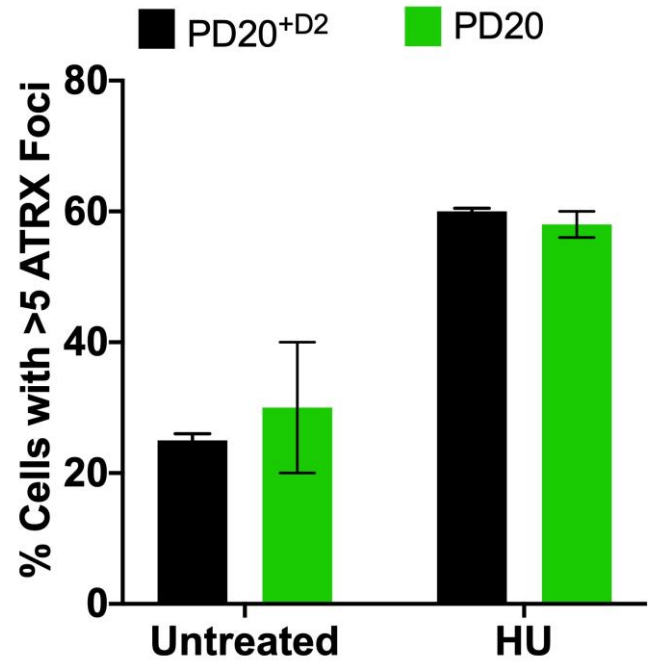


Supplementary Figure 2

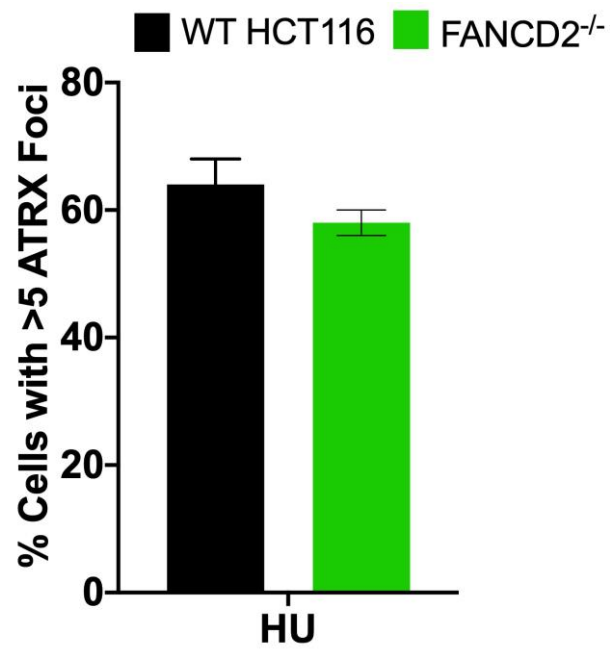
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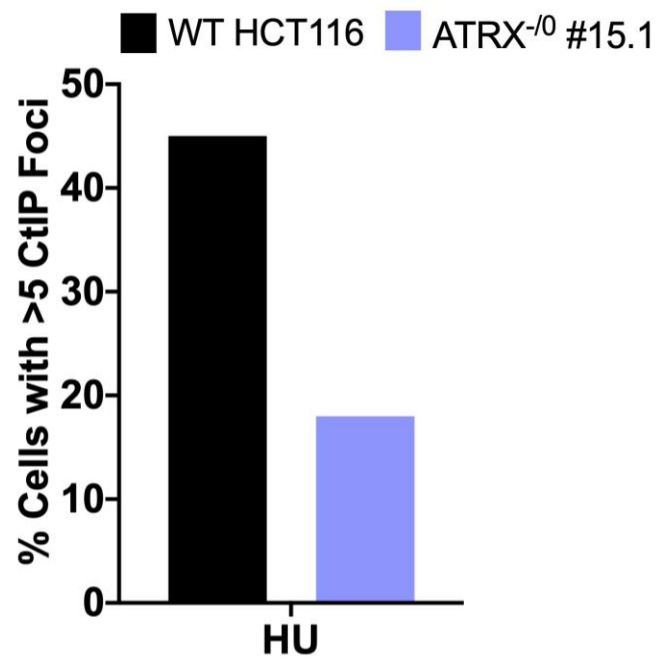
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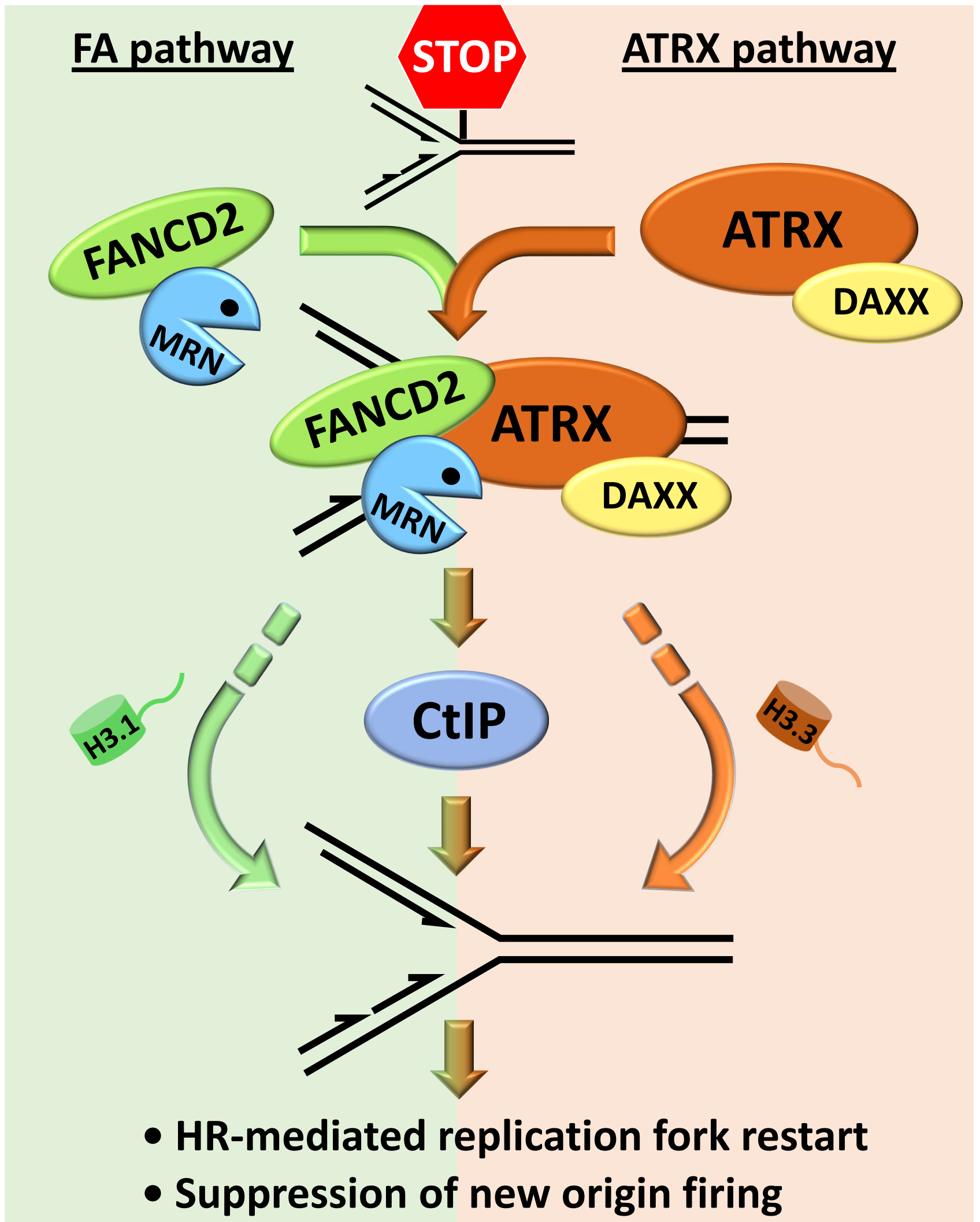
C



D



Supplementary Figure 3



SUPPLEMENTARY FIGURE LEGENDS

Supplementary Figure 1

IP experiments were performed from nuclear extracts (NE) of PD20+3xFLAG-FANCD2 (wild type) cells. **(A)** *NBS1 co-IPs with ATRX*. NEs (lane 1) were subjected to IP with mouse IgG (lane 2) or an anti-ATR_X antibody (lane 3). NE and IP samples were analyzed for the presence of ATR_X and NBS1 by western blot. **(B)** *NBS1 co-IPs with FANCD2*. NEs (lane 1) were subjected to IP with mouse IgG (lane 2) or an anti-FLAG antibody (lane 3). NE and IP samples were analyzed for the presence of FANCD2 and NBS1 by western blot. **(C)** Whole cell extracts (WCE) were prepared from PD20+D2 or PD20 cells that had been treated with control siRNA (lanes 1 and 3) or ATR_X-#2 siRNA (lanes 2 and 4) and analyzed for the presence of ATR_X and FANCD2. Ku-80, loading control. Immunoblot signals for FANCD2 in lanes 3 and 4 were analyzed by densitometry and normalized against Ku80 signals using ImageJ. The relative FANCD2 protein level values are provided underneath each corresponding lane. **(D)** *Schematic of the CRISPR/Cas9 targeting of FANCD2 exon 9 in HCT116 cells*. Sequencing data show a single base pair insertion (in red) in AD2-DKO clone #6.9. **(E-F)** FANCD2 and ATR_X cooperate to promote replication fork restart and suppress new origin firing in a human fibroblast cell line after HU treatment. **(E)** *ATR_X and FANCD2 cooperate to mediate replication fork restart*. Replication fork restart efficiencies were compared between WT (PD20+D2), FANCD2-deficient (PD20), ATR_X-deficient (PD20+D2, siATR_X) and FANCD2/ATR_X double-deficient (PD20, siATR_X) cells. **(F)** *ATR_X and FANCD2 act in concert to suppress new origin firing during replication blockade*. The number of new replication sites originating during BioU labeling after HU treatment was compared between WT (PD20+D2), FANCD2-deficient (PD20), ATR_X-deficient (PD20+D2, siATR_X) and FANCD2/ATR_X double-deficient (PD20, siATR_X) cells.

Supplementary Figure 2

(A) *FANCD2* foci formation occurs independently of *ATR*X during normal DNA replication and following replication stress. HCT116 WT cells and two different *ATR*X⁻⁰ clones (#94 and #15.1) were untreated or treated with 2 mM HU for 20 hr and analyzed for the presence of *FANCD2* foci. Nuclei with > 5 foci were considered positive for *FANCD2* foci formation. (B and C) *ATR*X foci formation occurs independently of *FANCD2* during normal DNA replication and following replication stress. (B) PD20+D2 (*FANCD2*-proficient) and PD20 (*FANCD2*-deficient) fibroblast cells were untreated or treated with 2mM HU for 20 hr and analyzed for the presence of *ATR*X foci. Nuclei with > 5 foci were considered positive for *ATR*X foci formation. (C) Same experiment as in B, but performed in HCT116 WT and *FANCD2*^{-/-} cells. (D, accompanies Main Fig 4B) *ATR*X is required for the HU-inducible *CtIP* foci formation. HCT116 WT cells and a second *ATR*X⁻⁰ clone (#15.1) were treated with 2 mM HU for 20 hours and analyzed for the presence of *CtIP* foci. Nuclei with > 5 foci were considered positive for *CtIP* foci formation.

Supplementary Figure 3

Model of the physical and functional FA-*ATR*X pathway interactions during the restart of HU-stalled replication forks.

Supplementary Table 1

Summary of all P-values for results shown in Main Figures 2 to 6.

Supplementary Table 1

Figure	Samples	p-value (two tales)
2B	WT vs D2 ^{-/-} Day 5	0.001
	WT vs ATRX ^{-/0} Day 5	0.011
	WT vs AD2 ^{DKO} Day 5	0.0001
	ATRX ^{-/0} vs AD2 ^{DKO} Day 5	0.001
	D2 ^{-/-} vs AD2 ^{DKO} Day 5	0.016
2C	WT vs D2 ^{-/-} 150uM HU	0.000149
	WT vs ATRX ^{-/0} 150uM HU	0.022412
	WT vs AD2 ^{DKO} 150uM HU	0.010017
	ATRX ^{-/0} vs D2 ^{-/-} 150uM HU	ns
	ATRX ^{-/0} vs AD2 ^{DKO} 150uM HU	ns
	D2 ^{-/-} vs AD2 ^{DKO} 150uM HU	ns
	D2 ^{-/-} vs AD2 ^{DKO}	ns
3B	WT vs D2 ^{-/-}	ns
	WT vs ATRX ^{-/0}	ns
	WT vs AD2 ^{DKO}	ns
	ATRX ^{-/0} vs D2 ^{-/-}	ns
	ATRX ^{-/0} vs AD2 ^{DKO}	ns
	D2 ^{-/-} vs AD2 ^{DKO}	ns
	WT vs D2 ^{-/-} +HU	0.00063
	WT vs ATRX ^{-/0} +HU	0.00029
	WT vs AD2 ^{DKO} +HU	0.00227
	ATRX ^{-/0} vs D2 ^{-/-} +HU	ns
	ATRX ^{-/0} vs AD2 ^{DKO} +HU	ns
D2 ^{-/-} vs AD2 ^{DKO} +HU	0.01368	

3C	WT vs D2 ^{-/-}	ns
	WT vs ATRX ^{-/0}	ns
	WT vs AD2 ^{DKO}	ns
	ATRX ^{-/0} vs D2 ^{-/-}	ns
	ATRX ^{-/0} vs AD2 ^{DKO}	ns
	D2 ^{-/-} vs AD2 ^{DKO}	ns
	WT vs D2 ^{-/-} +HU	0.00227
	WT vs ATRX ^{-/0} +HU	0.00288
	WT vs AD2 ^{DKO} +HU	0.02518
	ATRX ^{-/0} vs D2 ^{-/-} +HU	ns
	ATRX ^{-/0} vs AD2 ^{DKO} +HU	ns
	D2 ^{-/-} vs AD2 ^{DKO} +HU	0.00728
3D	WT vs D2 ^{-/-} +HU	0.00205
	WT vs ATRX ^{-/0} +HU	0.00224
	WT vs AD2 ^{DKO} +HU	0.00681
	ATRX ^{-/0} vs D2 ^{-/-} +HU	ns
	ATRX ^{-/0} vs AD2 ^{DKO} +HU	ns
	D2 ^{-/-} vs AD2 ^{DKO} +HU	0.00589
	WT vs D2 ^{-/-} +HU +Mirin	ns
	WT vs ATRX ^{-/0} +HU +Mirin	ns
	WT vs AD2 ^{DKO} +HU +Mirin	0.03137
	ATRX ^{-/0} vs D2 ^{-/-} +HU +Mirin	ns
	ATRX ^{-/0} vs AD2 ^{DKO} +HU +Mirin	ns
	D2 ^{-/-} vs AD2 ^{DKO} +HU +Mirin	0.00303
3E	WT vs D2 ^{-/-} +HU	0.00108
	WT vs ATRX ^{-/0} +HU	0.00204

	WT vs AD2-DKO +HU	0.00049
	ATR ^X ^{-/0} vs D2 ^{-/-} +HU	ns
	ATR ^X ^{-/0} vs AD2 ^{DKO} +HU	ns
	D2 ^{-/-} vs AD2 ^{DKO} +HU	0.00165
	WT vs D2 ^{-/-} +HU +Mirin	ns
	WT vs ATR ^X ^{-/0} +HU +Mirin	ns
	WT vs AD2 ^{DKO} +HU +Mirin	0.0333
	ATR ^X ^{-/0} vs D2 ^{-/-} +HU +Mirin	ns
	ATR ^X ^{-/0} vs AD2 ^{DKO} +HU +Mirin	ns
	D2 ^{-/-} vs AD2 ^{DKO} +HU +Mirin	0.00150
4B	WT vs D2 ^{-/-} +HU	0.00119
	WT vs ATR ^X ^{-/0} +HU	0.00031
	WT vs AD2-DKO +HU	0.00122
	ATR ^X ^{-/0} vs D2 ^{-/-} +HU	ns
	ATR ^X ^{-/0} vs AD2-DKO +HU	ns
	D2 ^{-/-} vs AD2-DKO +HU	ns
4C	WT vs D2 ^{-/-}	0.0053
	WT vs ATR ^X ^{-/0}	0.02528
	WT vs AD2 ^{DKO}	0.01945
	ATR ^X ^{-/0} vs D2 ^{-/-}	ns
	ATR ^X ^{-/0} vs AD2 ^{DKO}	ns
	D2 ^{-/-} vs AD2 ^{DKO}	ns
4E	siControl vs siFANCD2	0.00414
	siControl vs siATR ^X	0.00233
	siControl vs siFANCD2+siATR ^X	0.00427

	siControl vs siRAD51	0.00920
	siATRX vs siFANCD2	ns
	siATRX vs siFANCD2+siATRX	ns
	siATRX vs siRAD51	0.00016
	siFANCD2 vs siFANCD2+siATRX	ns
	siFANCD2 vs siRAD51	0.00065
	siFANCD2+siATRX vs siRAD51	0.00173
5B	WT vs D2 Deficient	ns
	WT vs siDAXX	ns
	WT vs D2 Deficient + siDAXX	ns
	D2 Deficient vs siDAXX	ns
	D2 Deficient vs D2 Deficient + siDAXX	ns
	siDAXX vs D2 Deficient + siDAXX	ns
	WT vs D2 Deficient +HU	0.00002
	WT vs siDAXX +HU	0.00157
	WT vs D2 Deficient + siDAXX +HU	0.00002
	D2 Deficient vs siDAXX +HU	ns
	D2 Deficient vs D2 Deficient + siDAXX +HU	0.00326
	siDAXX vs D2 Deficient + siDAXX +HU	ns
5C	WT vs D2 Deficient	ns
	WT vs siDAXX	ns
	WT vs D2 Deficient + siDAXX	ns
	D2 Deficient vs siDAXX	ns
	D2 Deficient vs D2 Deficient + siDAXX	ns

	siDAXX vs D2 Deficient + siDAXX	ns
	WT vs D2 Deficient +HU	0.002269
	WT vs siDAXX +HU	0.002882
	WT vs D2 Deficient + siDAXX +HU	0.007286
	D2 Deficient vs siDAXX +HU	ns
	D2 Deficient vs D2 Deficient + siDAXX +HU	0.025182
	siDAXX vs D2 Deficient + siDAXX +HU	ns
6C	WT vs D2 ^{-/-}	ns
	WT vs +FANCD2 ^{WT}	ns
	WT vs +FANCD2 ^{231R}	ns
	WT vs FANCL ^{-/-}	ns
	D2 ^{-/-} vs +FANCD2 ^{WT}	ns
	D2 ^{-/-} vs +FANCD2 ^{231R}	ns
	D2 ^{-/-} vs FANCL ^{-/-}	ns
	+FANCD2 ^{WT} vs +FANCD2 ^{231R}	ns
	+FANCD2 ^{WT} vs FANCL ^{-/-}	ns
	+FANCD2 ^{231R} vs FANCL ^{-/-}	ns
	WT vs D2 ^{-/-} +HU	0.00273
	WT vs +FANCD2 ^{WT} +HU	ns
	WT vs +FANCD2 ^{231R} +HU	0.00373
	WT vs FANCL ^{-/-} +HU	ns
	D2 ^{-/-} vs +FANCD2 ^{WT} +HU	0.00837
	D2 ^{-/-} vs +FANCD2 ^{231R} +HU	ns
	D2 ^{-/-} vs FANCL ^{-/-} +HU	0.00288
	+FANCD2 ^{WT} vs +FANCD2 ^{231R} +HU	0.00373
	+FANCD2 ^{WT} vs FANCL ^{-/-} +HU	ns

	+FANCD2 ^{231R} vs FANCL ^{-/-} +HU	0.002
6D	WT vs D2 ^{-/-}	ns
	WT vs +FANCD2 ^{WT}	ns
	WT vs +FANCD2 ^{231R}	ns
	WT vs FANCL ^{-/-}	ns
	D2 ^{-/-} vs +FANCD2 ^{WT}	ns
	D2 ^{-/-} vs +FANCD2 ^{231R}	ns
	D2 ^{-/-} vs FANCL ^{-/-}	ns
	+FANCD2 ^{WT} vs +FANCD2 ^{231R}	ns
	+FANCD2 ^{WT} vs FANCL ^{-/-}	ns
	+FANCD2 ^{231R} vs FANCL ^{-/-}	ns
	WT vs D2 ^{-/-} +HU	0.01613
	WT vs +FANCD2 ^{WT} +HU	ns
	WT vs +FANCD2 ^{231R} +HU	0.02381
	WT vs FANCL ^{-/-} +HU	ns
	D2 ^{-/-} vs +FANCD2 ^{WT} +HU	0.01507
	D2 ^{-/-} vs +FANCD2 ^{231R} +HU	ns
	D2 ^{-/-} vs FANCL ^{-/-} +HU	ns
	+FANCD2 ^{WT} vs +FANCD2 ^{231R} +HU	0.002