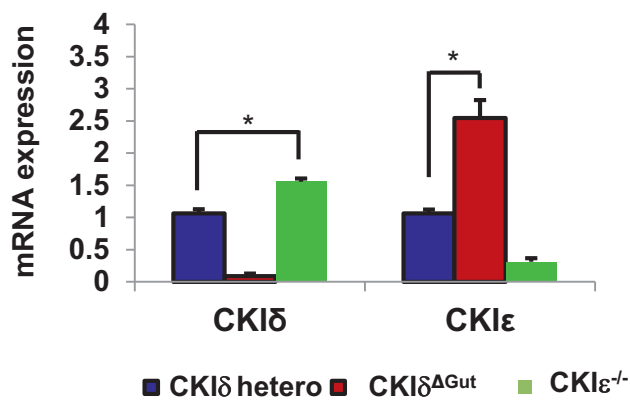


Expanded View Figures

**Figure EV1. Functional redundancy between CKI δ and CKI ϵ in the gut.**

qRT-PCR analysis of CKI δ and CKI ϵ in IECs isolated from CKI $\delta^{+/fl\ ER}$ (CKI δ hetero, $n = 3$), CKI $\delta^{\Delta\text{Gut}}$ ($n = 3$), and CKI $\epsilon^{-/-}$ mice ($n = 3$, mean \pm SEM); t-test was performed, $*P < 0.05$.

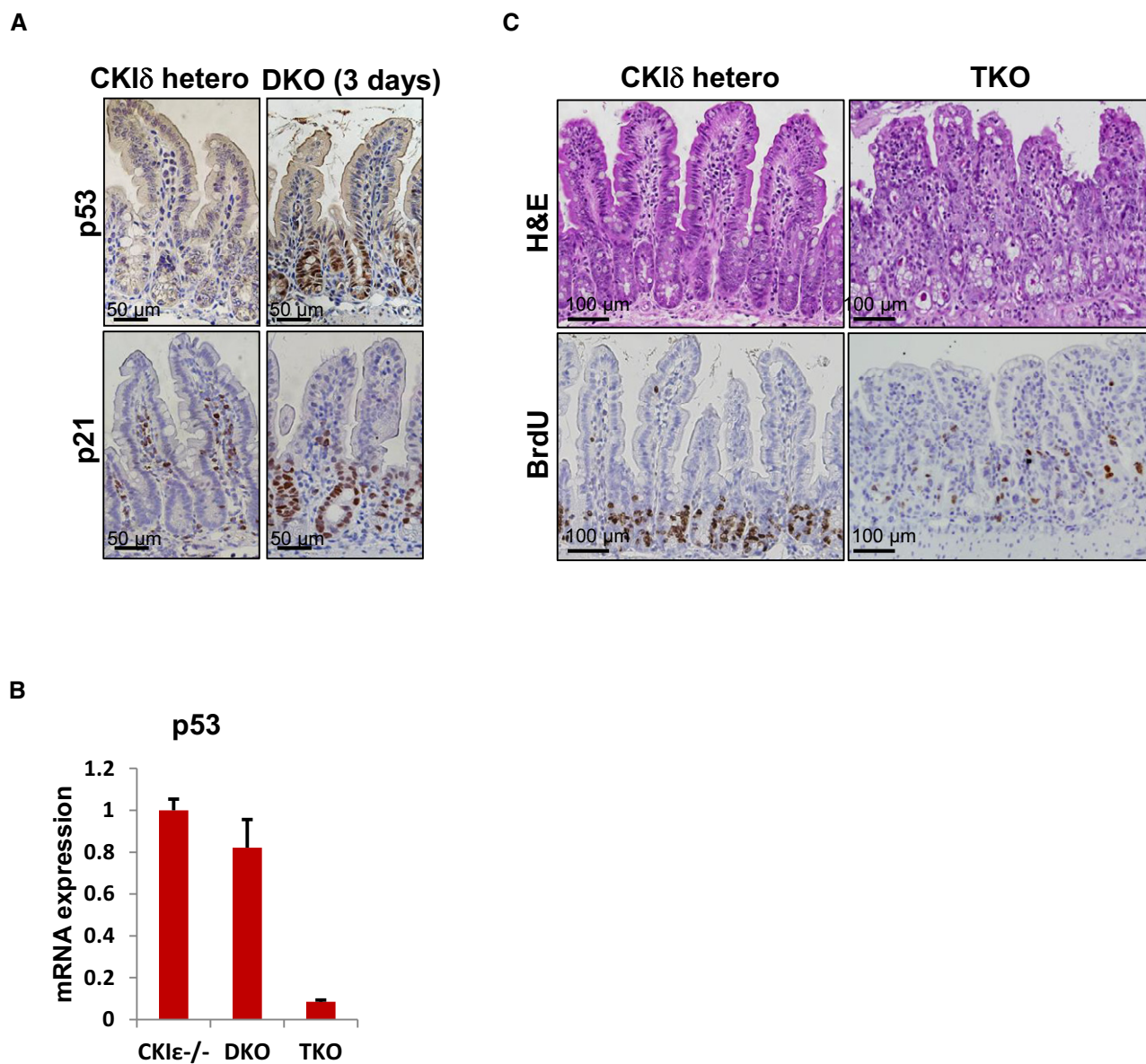


Figure EV2. CKI δ/ϵ ablation triggers a p53-independent cell cycle arrest.

A IHC analysis of p53 and p21 in intestinal sections from CKI δ hetero and DKO mice, 3 days after KO induction. Scale bar, 50 μ m.

B qRT-PCR analysis of p53 levels after KO induction in CKI $\epsilon^{-/-}$, DKO, and CKI δ /CKI ϵ /p53 triple KO mice (TKO) ($n = 3$, mean \pm SEM).

C H&E and IHC of BrdU in intestinal sections from CKI δ hetero and CKI δ /CKI ϵ /p53 triple KO mice (TKO), 5 days after KO induction. Scale bar, 100 μ m.

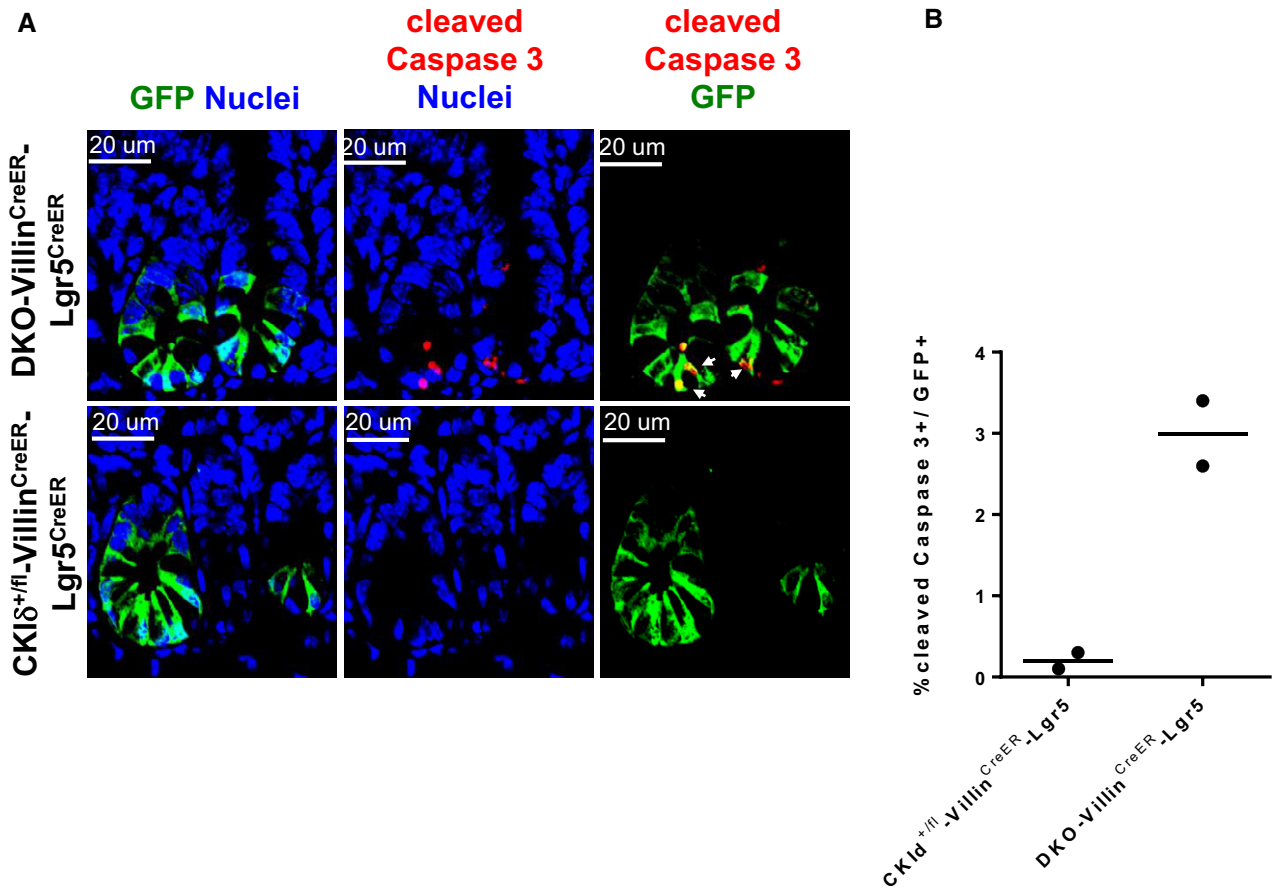


Figure EV3. CKIδ/ε depletion induces ISC apoptosis.

A IF analysis of GFP (green) and cleaved Caspase 3 (red) in intestinal sections of CKIδ^{+/fl}-Villin^{CreER}-Lgr5^{CreER} and DKO-Villin^{CreER}-Lgr5^{CreER} mice 3 days after KO induction; nuclear counterstain Hoechst. Scale bar, 20 μm. Arrows point to cells doubly stained by GFP and cleaved Caspase 3 representing apoptotic Lgr5-GFP cells.

B Quantification of cleaved Caspase 3 IF staining, based on %cleaved Caspase 3⁺ cells out of 300 GFP⁺ cells in CKIδ^{+/fl}-Villin^{CreER}-Lgr5^{CreER} (*n* = 2) and DKO-Villin^{CreER}-Lgr5^{CreER} mice (*n* = 2).

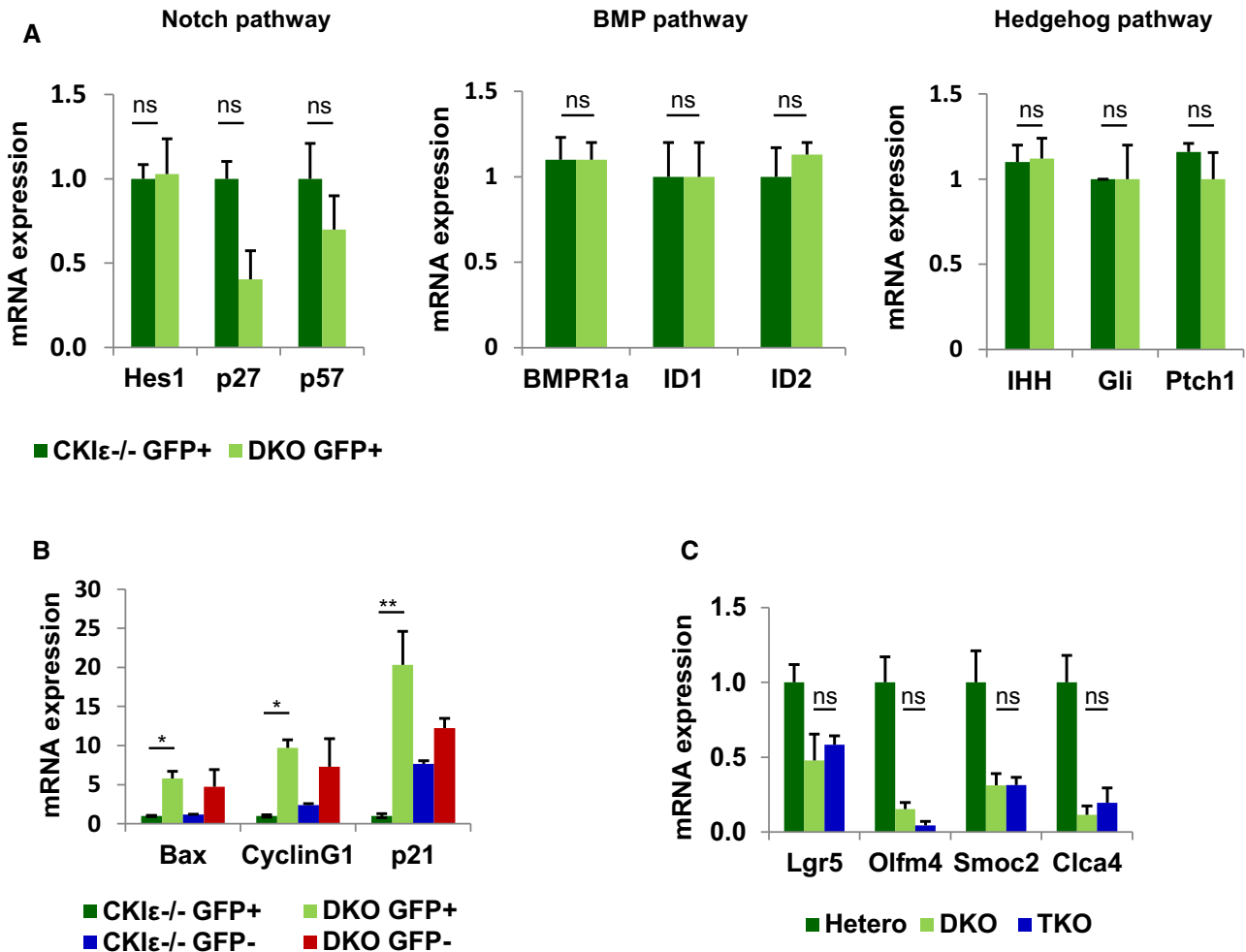


Figure EV4. Extinction of DKO ISCs is not mediated by Notch signaling downregulation or p53 activation.

- A qRT-PCR analysis of Notch, BMP, and Hedgehog target genes in sorted GFP⁺ cells isolated from crypts of CK1 ϵ ^{-/-}-Villin^{CreER}-Lgr5^{CreER} (CK1 ϵ ^{-/-}) and DKO-Villin^{CreER}-Lgr5^{CreER} mice (DKO) 2.5 days after KO induction. Data represent mean of three independent sorting experiments, each with a pool of six mice; t-test was performed, ns = non-significant. Error bars indicate SEM.
- B qRT-PCR analysis of p53 targets in sorted GFP⁺ and GFP⁻ cells isolated from crypts of CK1 ϵ ^{-/-}-Villin^{CreER}-Lgr5^{CreER} (CK1 ϵ ^{-/-}) and DKO-Villin^{CreER}-Lgr5^{CreER} (DKO), 2.5 days after KO induction. Data represent mean of three independent sorting experiments, each with a pool of six mice; t-test was performed, **P* < 0.05 ***P* < 0.01. Error bars indicate SEM.
- C qRT-PCR analysis of ISC markers in IECs isolated from CK1 δ hetero (hetero; *n* = 3), DKO (*n* = 3) and TKO (*n* = 3) mice (mean \pm SEM), 5 days after KO induction; t-test was performed, ns = non-significant.

Figure EV5. Decreased levels of Dvl2 in crypt DKO IECs.

- A Survival analysis was done by scoring 100 organoids of CK1 ϵ ^{-/-} and DKO organoids treated with ENR or ENR+VC. Data represent mean \pm SEM of three independent experiments.
- B Western blot analysis of nuclear (N) and cytoplasmic (C) fractions of CK1 ϵ ^{-/-} and DKO mouse crypt IECs; p-Dvl2 identified by electrophoretic shift in Dvl mobility. WB quantification shown at the bottom.
- C qRT-PCR analysis of Dvl isoforms in sorted GFP⁺ cells isolated from crypts of WT, CK1 ϵ ^{-/-}, CK1 δ KO, and DKO mice 2.5 days after KO induction. Data represent mean of three independent sorting experiments, each with a pool of six mice.
- D qRT-PCR analysis of Fzd isoforms in WT, CK1 ϵ ^{-/-}, DKO, CK1 ϵ ^{-/-} Dvl-NLS, and DKO Dvl-NLS expressing organoids 4 days after tamoxifen treatment and KO induction (*n* = 3, mean \pm SEM).
- E Representative bright field images of CK1 ϵ ^{-/-} and DKO organoids expressing Dvl-NLS and shFzd7 at day 4 or 5 after KO and Dvl induction. Insets represent organoids counted as live. Scale bar, 400 μ m.

Source data are available online for this figure.

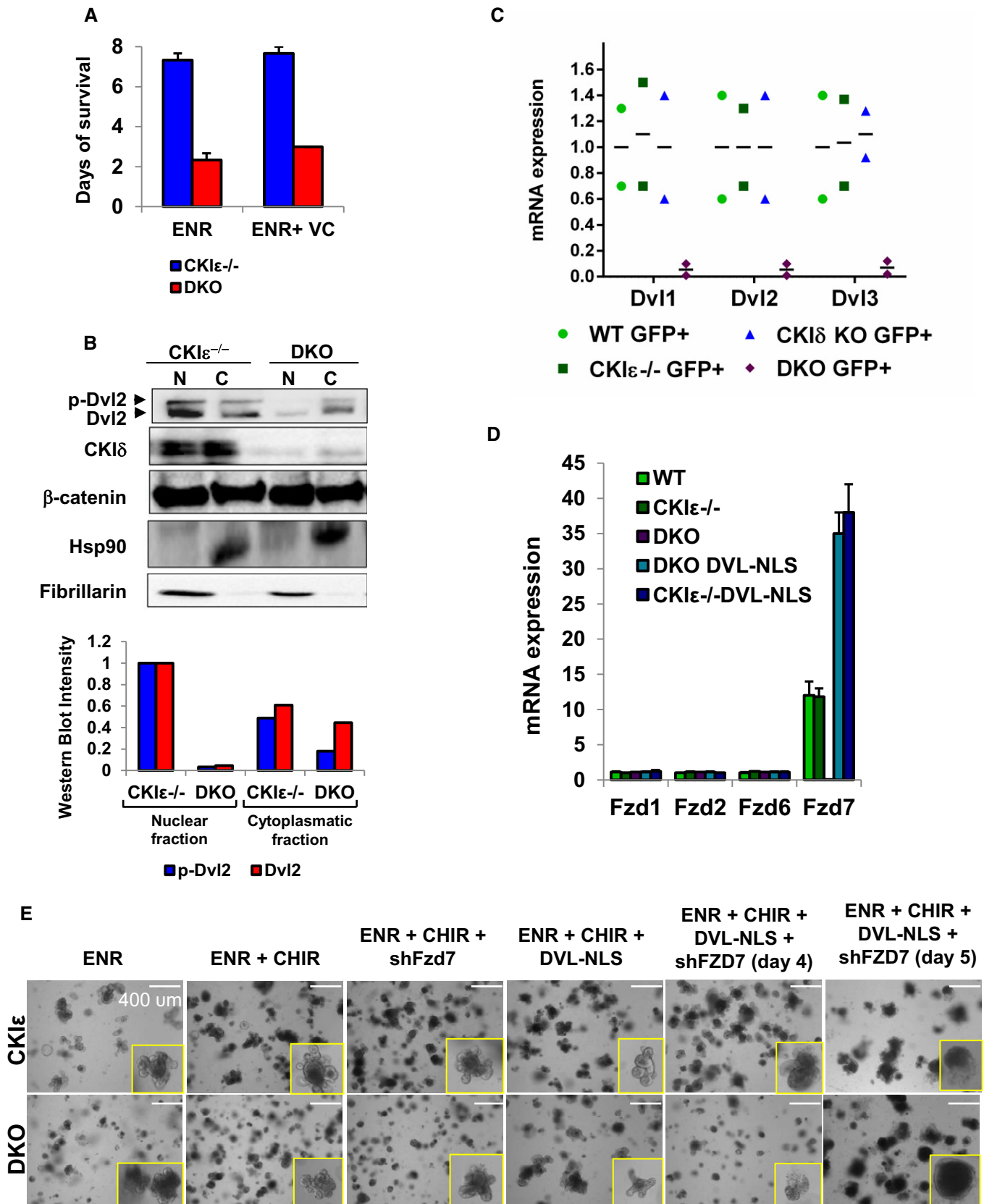


Figure EV5.