

## Supplementary Information

### Selective CDK9 inhibition overcomes TRAIL resistance by concomitant suppression of cFlip and Mcl-1

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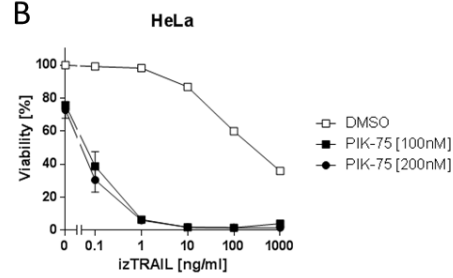
Running Title: CDK9 inhibition overcomes TRAIL resistance

This file contains Figures S1 – S7 with legends and supplementary literature

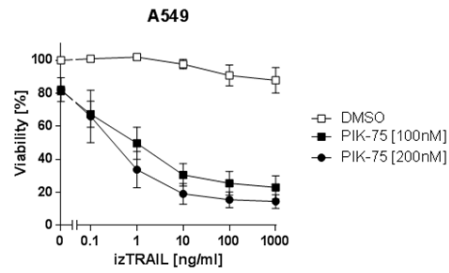
**A**

IC50 [nM]	p110 $\alpha$	p110 $\beta$	p110 $\gamma$	p110 $\delta$	Ref
PIK-75	6	1300	76	510	[1]
TGX-221	5000	5	>10000	100	[2]
AS-252424	940	>10.000	30	>10.000	[3]
IC-87114	>10.000	>10.000	>10.000	130	[1]

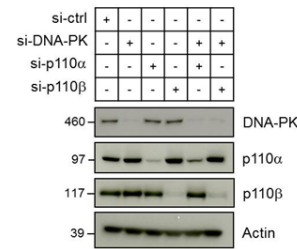
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**C**



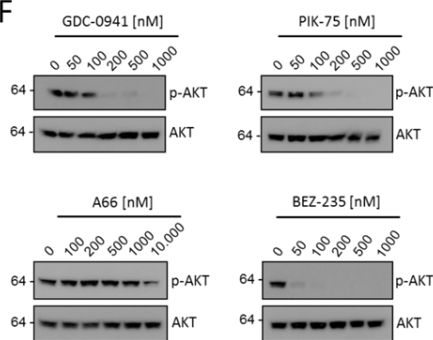
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**E**

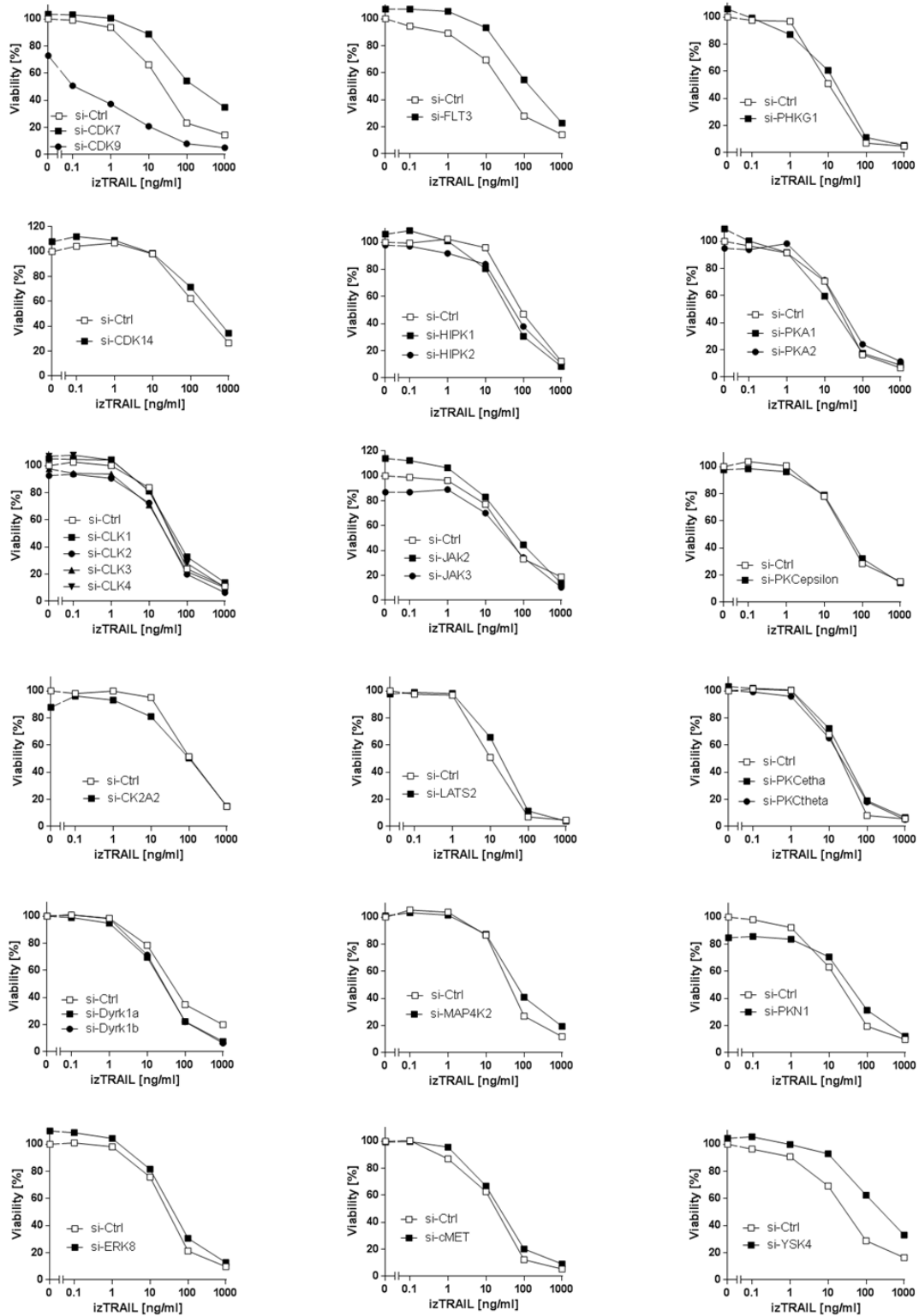
IC50 [nM]	p110 $\alpha$	p110 $\beta$	p110 $\gamma$	p110 $\delta$	Ref.
GDC-0941	3	33	75	3	[4]
A66	32	>10.000	>1250	3480	[5]
BEZ-235	4	75	5	7	[6]

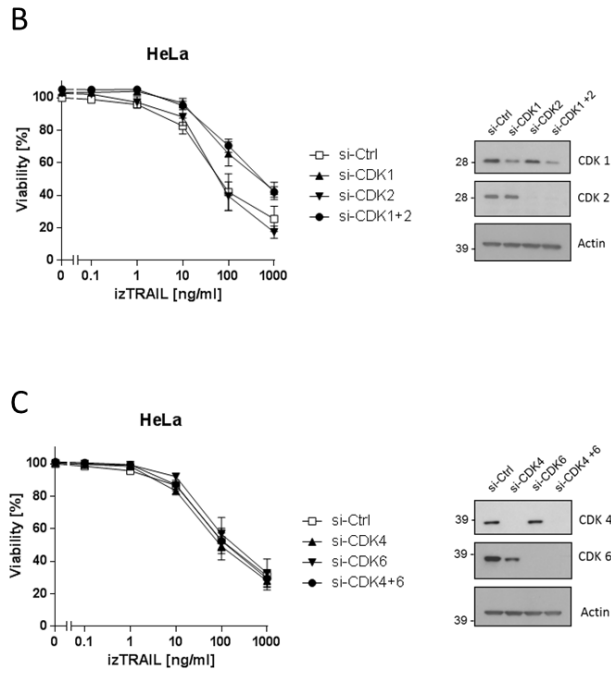
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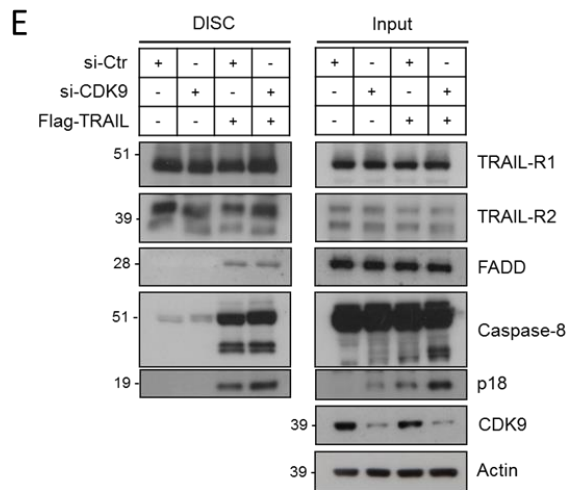
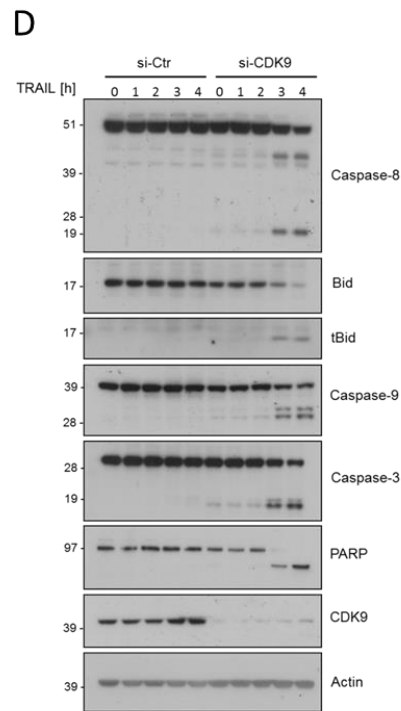
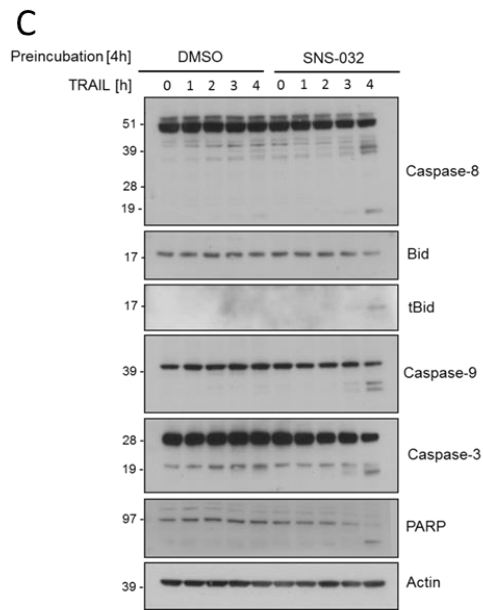
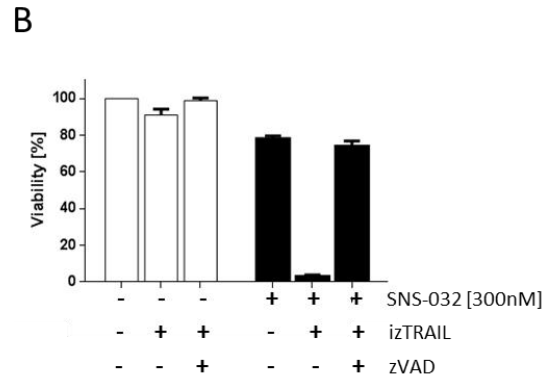
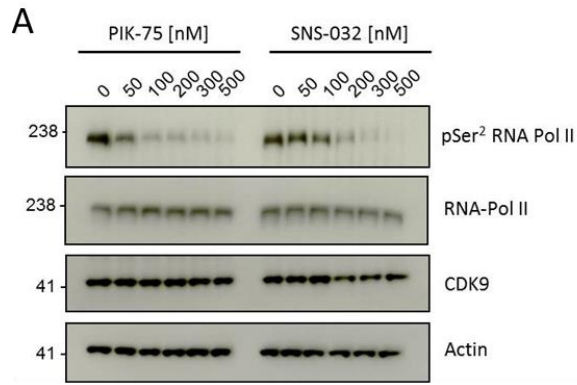
**Figure S1** (a) IC50 values for PI3K isoform-specific inhibitors used in Figure 1a are shown. HeLa (b) or A549 (c) cells were pre-incubated with the indicated concentration of PIK-75 for 1h and subsequently stimulated with the indicated concentration of izTRAIL. Cell viability was quantified after 24h. Values represent means  $\pm$  SEM of three independent experiments. (d) Westernblot demonstrating the knockdown efficiency for Figure 1c is shown. (e) IC50 values for PI3K inhibitors used in Figure 1d are shown. (f) HeLa cells were stimulated with different PI3K inhibitors at the indicated concentrations for 1h. Cells were lysed and subjected to western blotting. One representative of two independent experiments is shown.

A

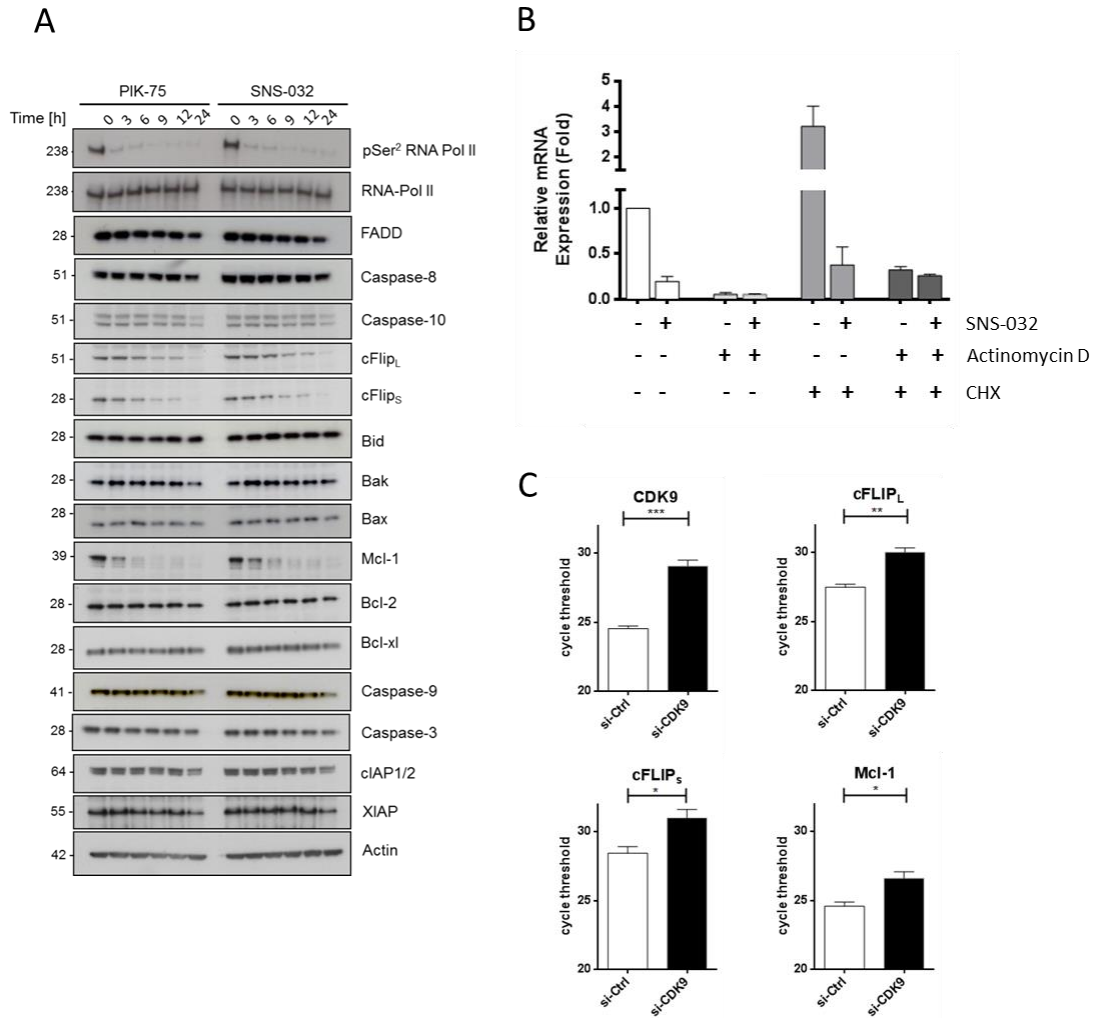




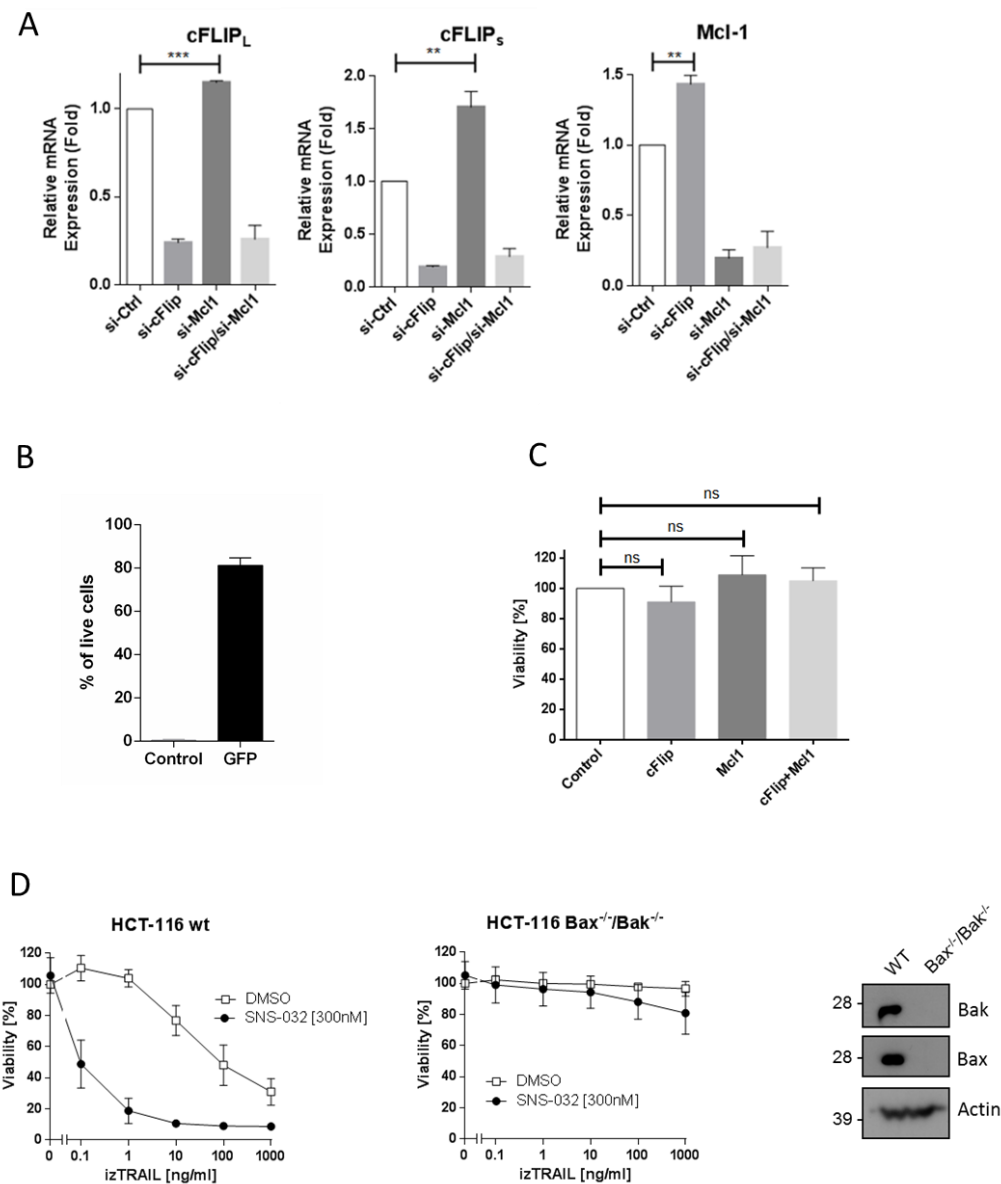
**Figure S2** (a) HeLa cells were subjected to the indicated knockdowns for 48h and subsequently stimulated with the indicated concentrations of izTRAIL. Cell viability was quantified after 24h. (b) HeLa cells were transfected with siRNA targeting CDK1 and/or CDK2 for 48h and subsequently stimulated with the indicated concentrations of izTRAIL. Cell viability was quantified after 24h. (c) HeLa cells were transfected with siRNA targeting CDK4 and/or CDK6 for 48h and subsequently stimulated with the indicated concentrations of izTRAIL. Representative western blots are shown. Values represent means  $\pm$  SEM of three independent experiments (b-c)



**Figure S3** (a) HeLa cells were treated for 3h with PIK-75 or SNS-032 at the indicated concentrations. Cells were subsequently lysed and subjected to western blotting. One representative of two independent experiments is shown. (b) HeLa cells were preincubated with 300nM SNS-032  $\pm$  20  $\mu$ M zVAD for 1h and subsequently stimulated with 10 ng/ml izTRAIL. Cell viability was quantified after 24h. Values represent means  $\pm$  SEM of three independent experiments (c) HeLa cells were preincubated with DMSO or SNS-032 [300 nM] for 4h and subsequently stimulated with izTRAIL [10 ng/ml] for the indicated times. Cells were lysed and subjected to western blotting. One representative of two independent experiments is shown. (d) HeLa cells were transfected with siRNA targeting CDK9 for 72h and subsequently stimulated with izTRAIL [10 ng/ml] for the indicated times. Cells were lysed and subjected to western blotting. One representative of two independent experiments is shown. (e) A549 cells were subjected to indicated knockdowns for 72h, stimulated with Flag-TRAIL (1 $\mu$ g/ml] for 1h and subsequently the TRAIL-DISC was analysed by western blotting. One representative of two independent experiments is shown.



**Figure S4** (a) HeLa cells were treated with PIK-75 (100 nM) or SNS-032 (300 nM) for the indicated times. Cells were lysed and subjected to western blotting. One representative of two independent experiments is shown. (b) A549 cells were preincubated with cycloheximide (CHX; 1  $\mu$ g/ml) and/or actinomycin D (2  $\mu$ g/ml) and/or SNS-032 (300 nM) as indicated. Mcl-1 mRNA expression was quantified by RT-PCR. Values represent means  $\pm$  SEM of at least two independent experiments. (c) HeLa cells were transfected with siRNA targeting CDK9. After 72h mRNA of 500.000 cells per condition was isolated, cDNA was generated and expression of indicated target genes was quantified by RT-PCR. Cycle threshold for the indicated transcripts is shown. Values represent means  $\pm$  SEM of three independent experiments. \*  $P < 0.05$ ; \*\*  $P < 0.01$ ; \*\*\*  $P < 0.001$  Student's t-test



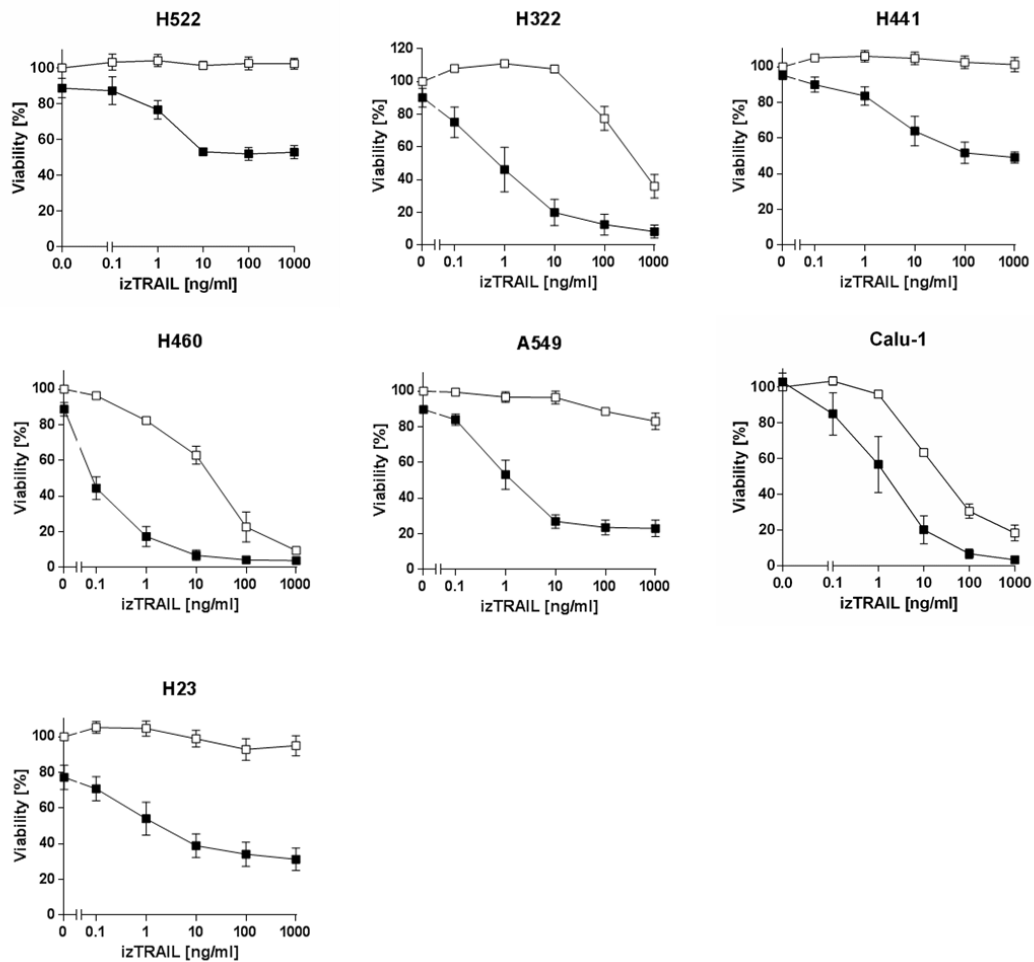
**Figure S5** (a) HeLa cells were subjected to indicated knockdowns and mRNA expression of indicated target genes was quantified by RT-PCR relative to GAPDH. (b) HeLa cells were transfected with a GFP-containing plasmid and GFP positive cells were quantified by flow cytometry. (c) HeLa cells were transfected with expression plasmids for cFlip and/or Mcl-1 or empty vector control for and cell viability was quantified after 48h. (d) HCT116<sup>WT</sup> or HCT116<sup>Bax<sup>-/-</sup>/Bak<sup>-/-</sup></sup> cells were pre-incubated with SNS-032 [300 nM] and subsequently stimulated with izTRAIL at the indicated concentrations. Cell viability was quantified after 24h. Representative western blots to confirm Bax/Bak deficiency are shown. All values represent means  $\pm$  SEM of three independent experiments. ns = not significant; \*\* P < 0.01; \*\*\* P < 0.001 Student's t-test



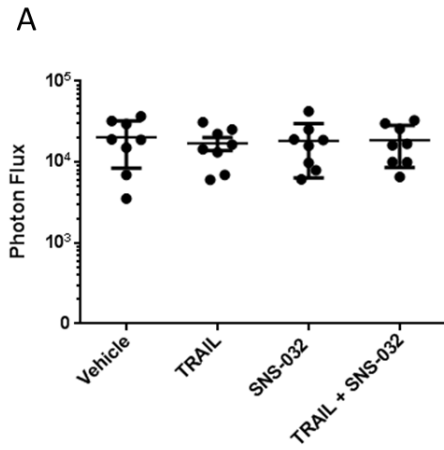
A

NSCLC cell line	KRAS	TP53
A549		
Calu-1		
H460		
H322		
H522		
H23		
H441		

B



**Figure S6** (a) Mutations in KRAS and/or p53 of cell lines included in the NSCLC panel (Figure 6a) are indicated by a black box. (b) Cell lines included in the NSCLC panel (Figure 6a) were preincubated with DMSO (open square) or SNS-032 [300 nM] (filled square) for 1h and subsequently stimulated with izTRAIL at the indicated concentrations. Cell viability was quantified after 24h. Values are means  $\pm$  SEM of three independent experiments.



**Figure S7** Tumor burden was quantified by bioluminescence imaging (Photon Flux) one week after injection of  $2 \times 10^6$  A549-Luc cells. Dots represent individual mice (n=8 per group) +/- SEM before treatment (after randomization).

## Supplementary Literature

1. Knight ZA, Gonzalez B, Feldman ME, Zunder ER, Goldenberg DD, Williams O *et al.* A pharmacological map of the PI3-K family defines a role for p110alpha in insulin signaling. *Cell* 2006; 125(4): 733-47.
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