## Silibinin overcomes EMT-driven lung cancer resistance to new-generation ALK inhibitors

## **SUPPLEMENTARY INFORMATION**

Table S1. Comparison of ALK-TKIs IC $_{50}$  values (in  $\mu$ mol/L) in ALK-rearranged NSCLC cells cultured in the absence/presence of SBN

	H2228	H2228/CR	H3122	H3122/CR
crizotinib	2.1 ± 0.8 (1.0)	10.6 ± 1.7 (1.0)	$0.29 \pm 0.06 (1.0)$	1.08 ± 0.40 (1.0)
+ 100 μmol/L SBN	$1.3 \pm 0.7 (1.6)$	$3.7 \pm 0.7 \; (2.9)^a$	$0.10 \pm 0.00 (2.9)$	$0.17 \pm 0.03 (6.3)$
brigatinib	0.5 ± 0.08 (1.0)	$3.8 \pm 0.4 (1.0)$	$0.02 \pm 0.00 (1.0)$	0.12 ± 0.03 (1.0)
+ 100 μmol/L SBN	$0.4 \pm 0.07 (1.2)$	$0.6 \pm 0.1 (6.3)$	$0.01 \pm 0.00 (2.0)$	$0.07 \pm 0.01  (1.7)$
lorlatinib	1.0 ± 0.11 (1.0)	83 ± 5.1 (1.0)	$0.004 \pm 0.00 (1.0)$	$0.02 \pm 0.00 (1.0)$
+ 100 μmol/L SBN	$9.0 \pm 2.2  (0.1)$	$19.3 \pm 3.2 (4.3)$	$0.004 \pm 0.00 (1.0)$	$0.003 \pm 0.00 (6.7)$

<sup>&</sup>lt;sup>a</sup> Number in parentheses represent the silibinin (SBN) extent of sensitization expressed as fold-change compared to untreated control cells

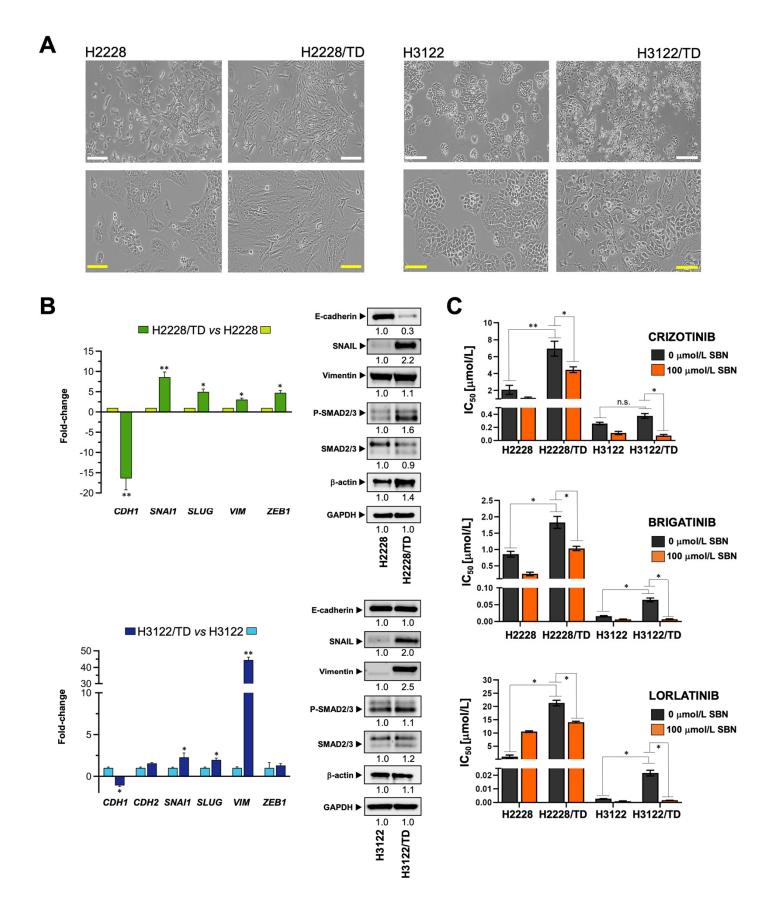


Figure S1. EMT caused by chronic TGF $\beta$  stimulation drives cross-resistance to multiple-generation ALK-TKIs. A. Representative phase contrast microphotographs of H2228/H2228TD and H3122/H3122TD ALK-rearranged NSCLC cell line pairs. TD: transdifferentiated; Scale bar, 100 μm. **B**. *Left*. The transcript abundance of *CDH1*, *CDH2*, *SNAI1*, *SLUG*, *VIM*, and *ZEB1* was calculated using the  $\Delta C_t$  method and presented as fold-change in H2228/H2228TD and H3122/H3122TD cells; \*p<0.05 and \*\*p<0.005, statistically significant differences. *Right*. Expression levels of E-cadherin, SNAIL, vimentin, phospho-SMAD2/3, total SMAD2/3 were detected by immunoblotting in H2228/H2228TD and H3122/H3122TD cells using specific antibodies. The intensity of the bands was measured using the ImageJ software. Fold-change of each protein relative to parental cells was calculated using GAPDH as a loading control. Figure shows representative immunoblots of multiple (*n*=3) independent experiments. **C**. Bar graphs showing the MTT-based IC<sub>50</sub> values of crizotinib, brigatinib, and Iorlatinib for H2228/H2228TD and H3122/H3122TD cells calculated in the absence or presence of 100 μmol/L silibinin. The results are presented as the means (*columns*) ± S.D (*bars*) (n=3, in triplicate). \*p<0.05 and \*\*p<0.005, statistically significant differences; n.s. not significant.

Figure S2. Uncropped western blots

