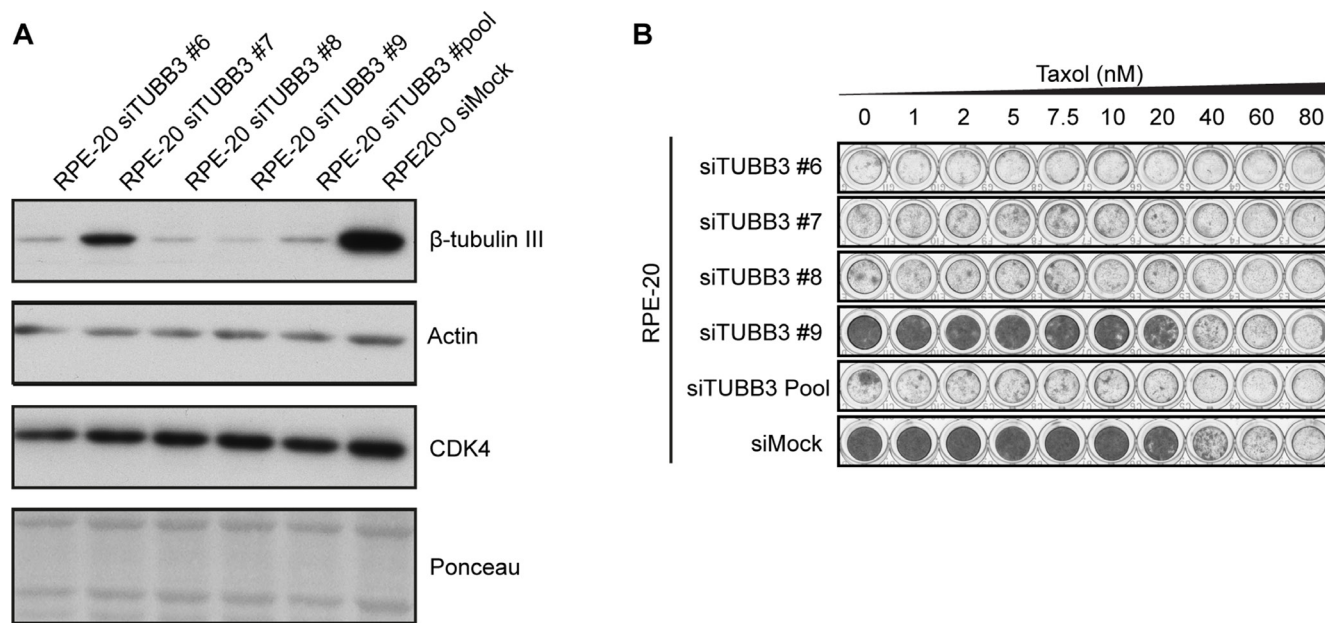
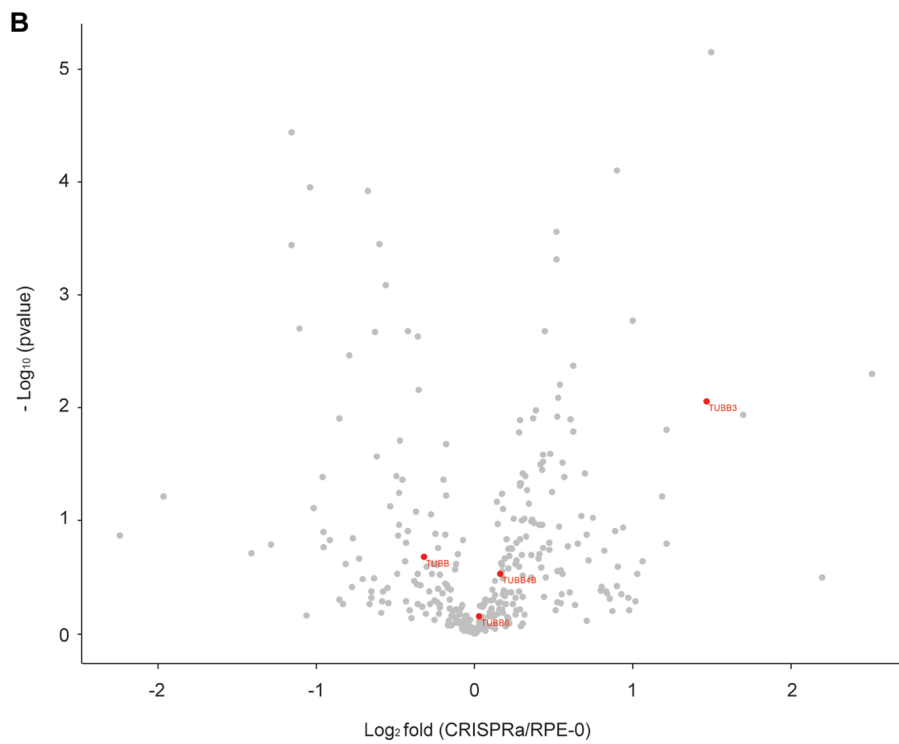
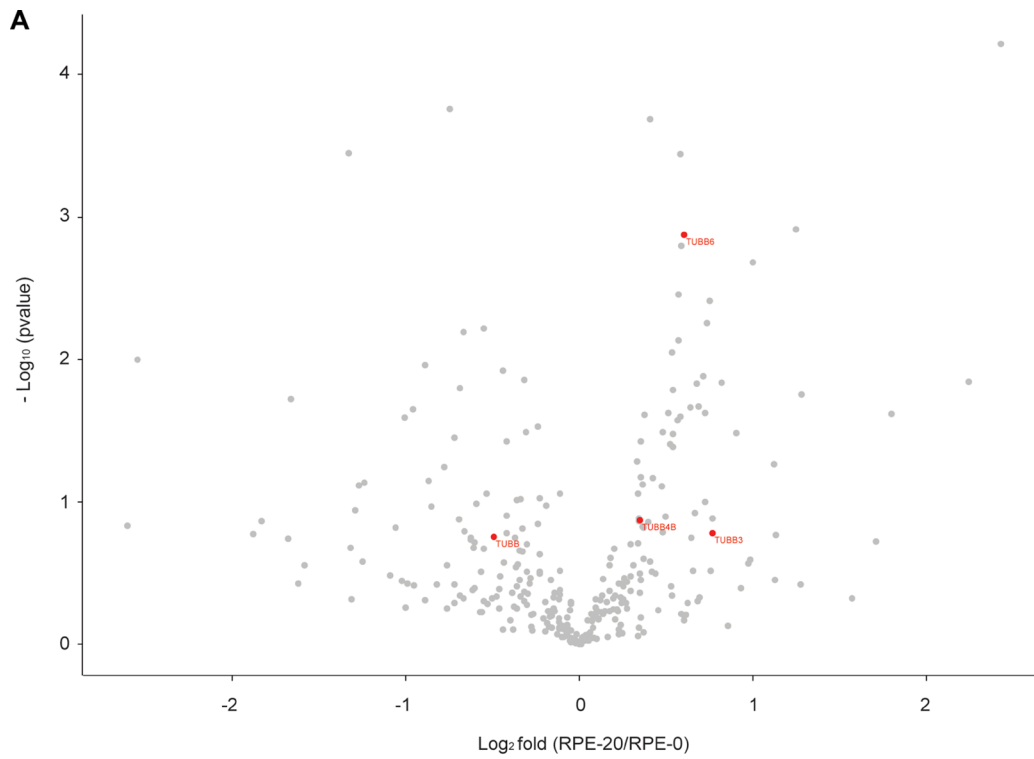


TUBB3 overexpression has a negligible effect on the sensitivity to taxol in cultured cell lines

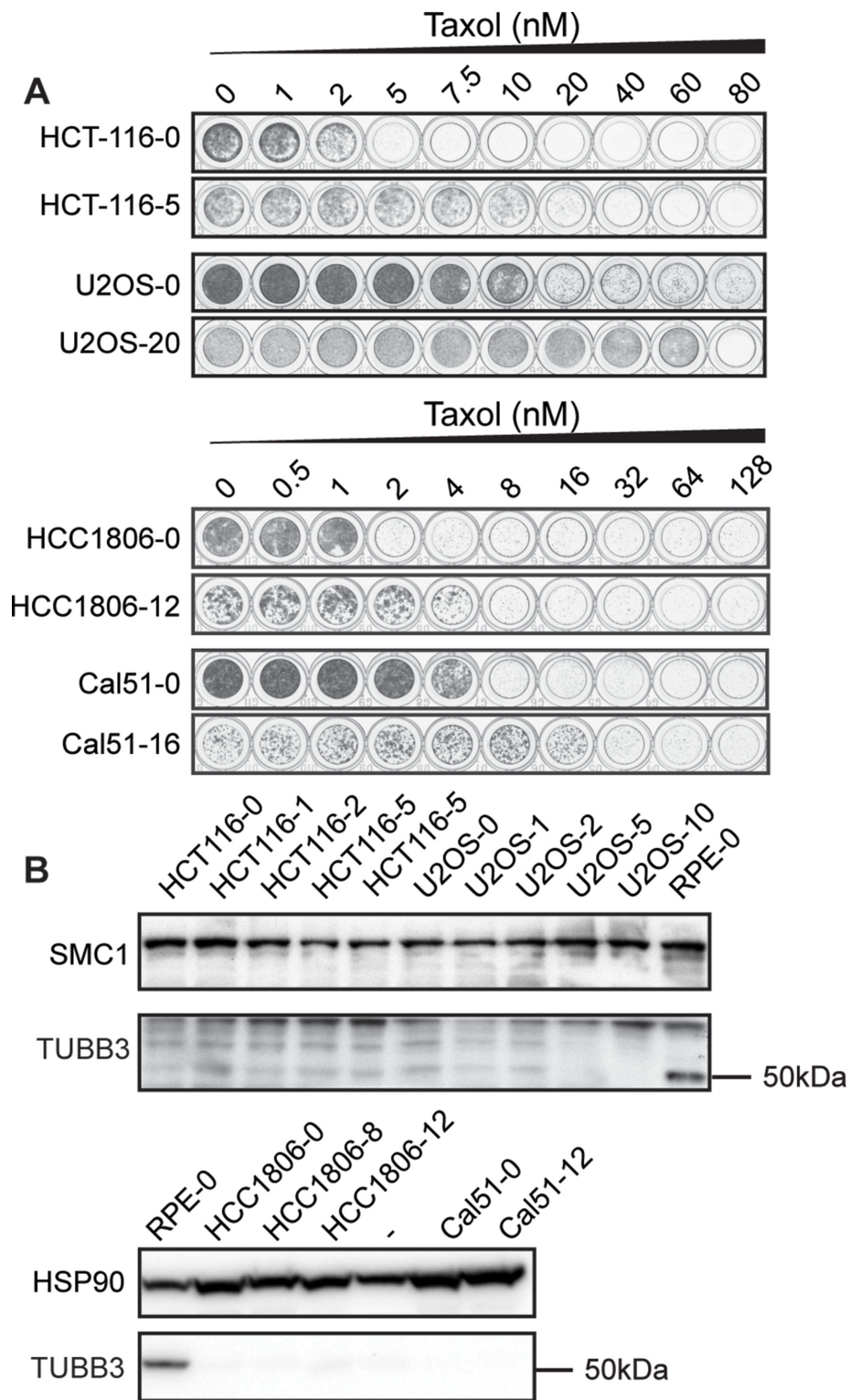
Supplementary Materials



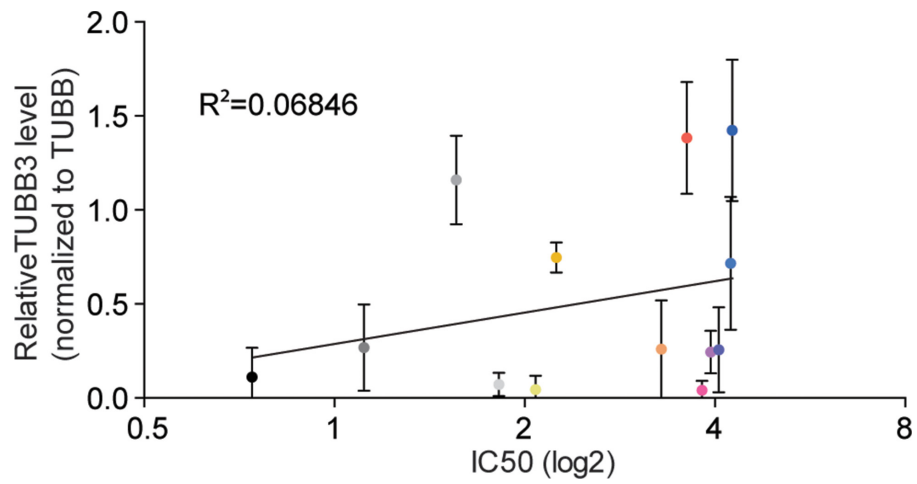
Supplementary Figure 1: βIII-tubulin knockdown does not affect the sensitivity of RPE-20 to taxol. (A) Western blot of RPE-20 cells transfected with single siRNAs targeted against TUBB3 for 48 hours. (B) Relative survival plots of RPE-20 cells treated with different doses of taxol for 7 days after knockdown of TUBB3 as described in A).



Supplementary Figure 2: Protein quantification of b-tubulin isotypes by mass spectrometry. A volcano plot demonstrating magnitude of the protein comparisons between RPE-0 and RPE-20 in (A) and RPE-0 and CRISPRa TUBB3 (exon 3) in (B). The vertical axis indicates $-\log_{10}(p\text{-value})$. The horizontal axis indicates \log_2 fold change. The b-tubulin isotypes found are labeled in red. Data was obtained from two sample replicates.



Supplementary Figure 3: Induction of TUBB3 levels is not observed in multiple taxol-resistant cancer cell lines. (A) Relative survival plots of taxol-resistant cell lines derived from a colorectal carcinoma (HCT116), an osteosarcoma (U2OS), two triple-negative breast cancer cell lines (HCC1806 and Cal51), and their respective taxol-sensitive parental cell lines. The values added behind the cell lines indicate the concentration of taxol at which the resistant cell lines were grown in (e.g. HCT-116-5 indicates an HCT-116 cell line cultured continuously at 5 nM of taxol). (B) Whole cell lysates prepared from the parental and taxol-resistant cancer cell lines showing their TUBB3 levels.



Supplementary Figure 4: Slight correlation between relative TUBB3 levels and IC50. The IC50 values were plotted against the relative TUBB3 levels determined for each breast cancer cell line. TUBB3 levels depicted here are averages from three independent experiments and normalized against whole β -tubulin levels.

Supplementary Table 1: sgRNA target sequences in the putative TSS regions of TUBB3 and ABCB1

TUBB3 exon III	Target sequence	Forward sgRNA oligo	Reverse sgRNA oligo
tubb3-1	ggagctggctgtggggcgccgtt	caccggagctggctgtggggcgcc	aaacggcgccccacagccagctcc
tubb3-2	ggggctgtctcccagaggagcgtt	caccggggctgtctcccagaggagc	aaacgctcctctgggagacagcccc
tubb3-3	gccccgcgattcgaaagggtt	caccgccccgcgattcgaaagg	aaacctttcgatcgcgggggc
tubb3-5	ggtgctgggtgtctctaaacgtt	caccggtgctgggtgtctctaaac	aaacgttagagaccaaccgcacc
tubb3-6	gttgctctaaaccggcgtt	caccgttgctctaaaccggcgt	aaacacgccgttagagaccaac
tubb3-8	gcggaaccgagaggtagcgtt	caccgccaaccgagaggtagc	aaacgctacctctcgttccgc
tubb3-9	ggggatccttgctgaggaggtt	caccgggatccttgctgaggag	aaacctcccgagccaaggatcccc
TUBB3 exon I			
tubb3-a	tgctgaagttgctgggtggagg	caccgtgctgaagttgctgggtggagg	aaacctccaccgcaaaactcaagcac
tubb3-b	gaaaggagctccggaaaacgtgg	caccgaaaggagctccggaaaacgtgg	aaaccacgtttccgggactccttc
tubb3-d	ggagtgaggaggtccaaccagg	caccggagtgaggaggtccaaccagg	aaacctggtgggacctccactcc
tubb3-e	agcctagagggaaggacgagcagg	caccgagcctagagggaaggacgagcagg	aaacctgctcgtccttcttaggctc
tubb3-f	agctaggacagcgtcaagagtg	caccgagctaggacagcgtcaagagtg	aaacctcttgactgctgcttagctc
ABCB1			
abcb1-1	gtagctcctctctgtactgtt	caccgtagctcctctctgtact	aaacagtaccagaggaggagctac
abcb1-2	gtgttatcccagtagcagggtt	caccgtgttatcccagtagcagg	aaacctctgtagctgggataaacac
abcb1-3	gaggagctacatgaactaaggcgtt	caccgaggagctacatgaactaaggc	aaacgccttagtcatgtagctctc
abcb1-4	gctacatgaactaaggcaggcgtt	caccgctacatgaactaaggcaggc	aaacgcctgcttagtcatgtagc
abcb1-5	gcactaatcagtgaaaaccaggtt	caccgcaataatcagtgaaaaccag	aaactgggtttcactgattagtc
abcb1-6	gaaaaccaaggataagttgtt	caccgaaaaccaaggataagtt	aaacaaactatccttgggtttc
abcb1-7	gataagttgggtggaggaagggtt	caccgataagttgggtggaggaagg	aaacctctcctccaccaaaactatc
abcb1-8	gtgatcttttgtaaggtgtt	caccgtgatcttttgtaaggtgt	aaacacaccttagcaaaaagatcac
abcb1-9	gagttacatgcttaggatgtt	caccgagttacatgcttaggat	aaacatccctaaacctgtaactc
abcb1-10	gttgagaagtttagcagaatgtt	caccgttgagaagtttagcagaat	aaacattctgctaaacttcaac

Supplementary Table 2: β -tubulin isoforms and ABCB1 RT-qPCR oligos

	Forward oligo	Reverse oligo
TUBB/tubulin beta class I	GCCTTCCTCCACTGGTACAC	GCCTCCTCACCGAAATCCTC
TUBB1/ tubulin beta 1 class VI	GCTTCCAGATCGTCCACTCC	GTCCGACACCTTGGGAGAAG
TUBB2A/tubulin beta 2A class Iia	CTGAAGATGTCGGCCACCTT	CGGTGAACTCCATCTCGTCC
TUBB2B/tubulin beta 2B class Iib	CTGAAGATGTCGGCCACCTT	CGGTGAACTCCATCTCGTCC
TUBB3/tubulin beta 3 class III	GCCTCAAGATGTCCTCCACC	CGGTGAACTCCATCTCGTCC
TUBB4A/tubulin beta 4A class Iva	GACTCTGGAAACCGCACCTT	TCAGAGGGAGGGGTCAAACA
TUBB6/tubulin beta 6 class V	TCCACCTTCATCGGCAACAG	TTTCATCCATGCCCTCACCC
TUBB8/tubulin beta 8 class VIII	GCCTTCCTCCACTGGTACAC	CTCCTCCTCGGCATACTCCT
ABCB1	CAGTTGAGTGGTGGGCAGAA	GCCTTATCCAGAGCCACCTG