

A Novel benzopyrane derivative targeting cancer cell metabolic and survival pathways

Dana M. Zaher^{1,2,#}, Wafaa S. Ramadan^{1,2,#}, Raafat El-Awady^{1,3}, Hany A. Omar^{1,3,4}, Fatema Hersi^{1,2}, Vunnam Srinivasulu¹, Ibrahim Y. Hachim^{1,2}, Farah I. Almarzooq^{1,5}, Cijo G. Vazhappilly^{1,6}, Salim Merali⁷, Carmen Merali⁷, Nelson C. Soares^{1,3}, Paul Schilf⁸, Saleh M. Ibrahim^{1,2,8}, Taleb H. Al-Tel^{1,3,*}

¹Sharjah Institute for Medical Researches, University of Sharjah, Sharjah, United Arab Emirates.

²College of Medicine, University of Sharjah, Sharjah, United Arab Emirates.

³College of Pharmacy, University of Sharjah, Sharjah, United Arab Emirates.

⁴Faculty of Pharmacy, Beni-Suef University, Beni-Suef, 62511 Egypt

⁵United Arab Emirates University, Department of Medical Microbiology and Immunology, College of Medicine and Health Sciences, Al Ain, United Arab Emirates.

⁶School of Arts and Sciences, American University of Ras Al Khaimah, P.O.Box:10021, Ras Al Khaimah, United Arab Emirates.

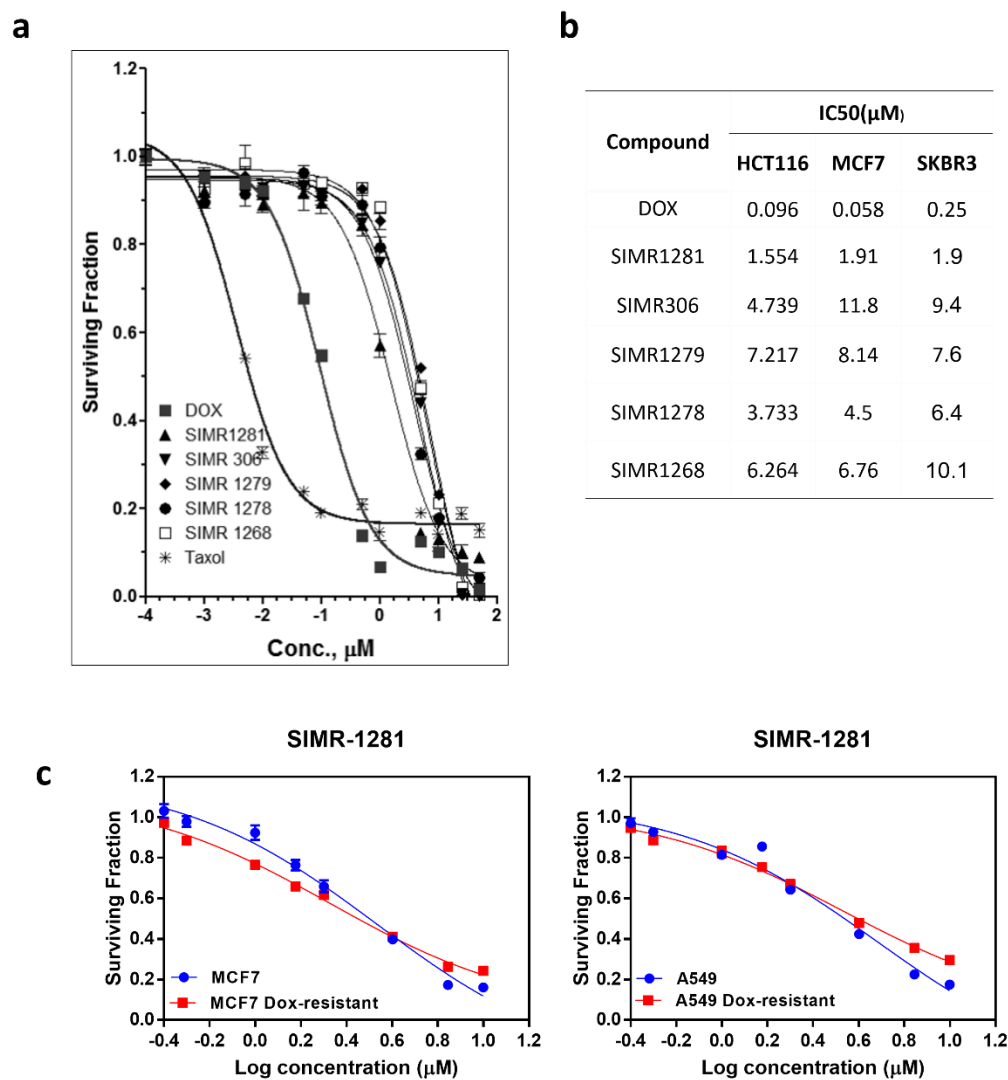
⁷Temple University, School of Pharmacy, 3307 N Broad Street, Room 552, Philadelphia, PA 19140, United States.

⁸Lübeck Institute of Experimental Dermatology, University of Lübeck, Ratzeburger Allee 160 23538 Lübeck, Germany.

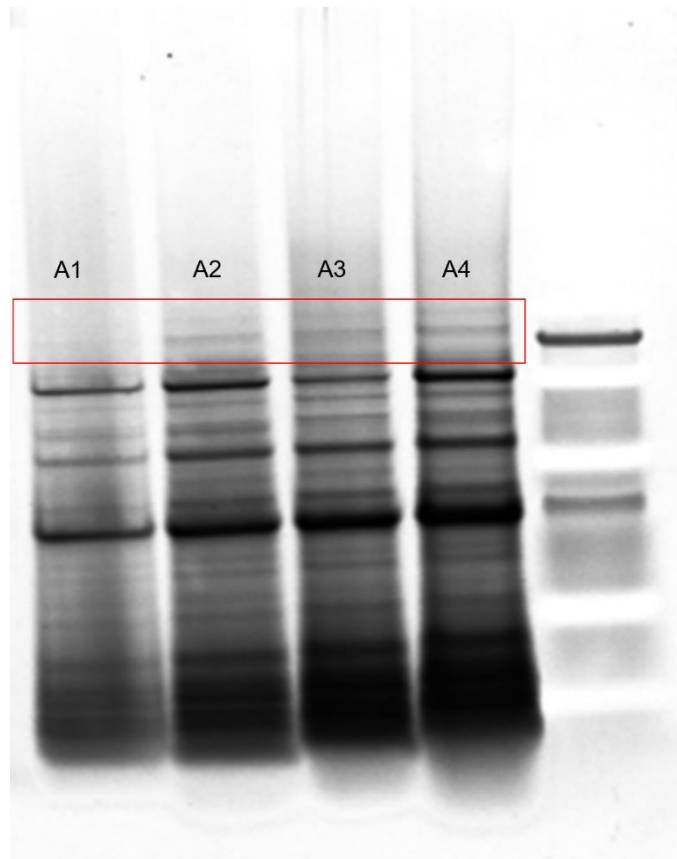
*Correspondance: taltal@sharjah.ac.ae, Tel: +97165057417

These coauthors contributed equally to this work and are both first authors

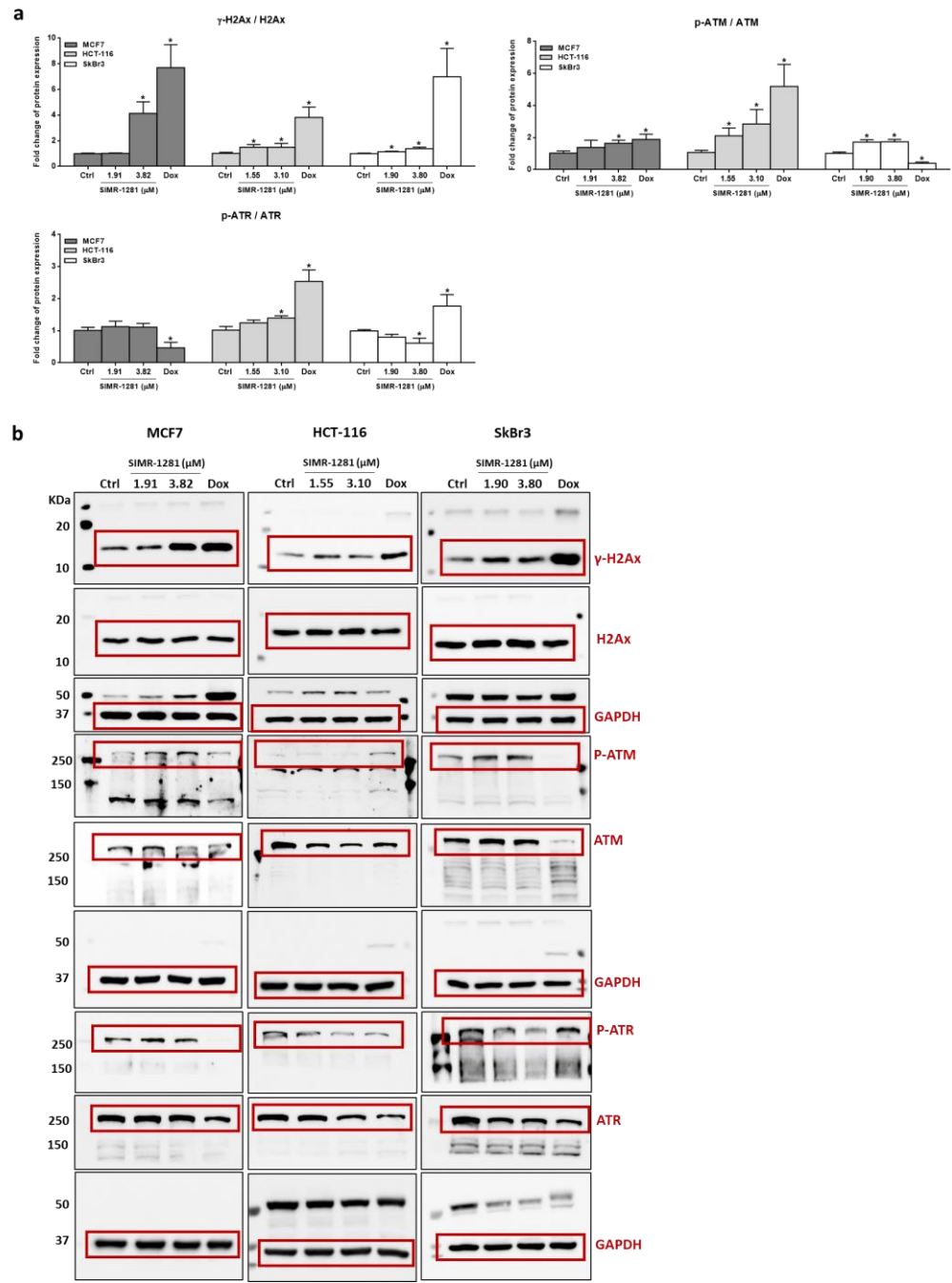
Supplementary Figure S1: Screening of a small library of polysubstituted benzopyranes, for potential anti-cancer activity on cancer cell lines. **(a)** Dose dependent effect of the synthesized SIMR compounds on MCF7, SKBR3 and HCT-116 after 48 h treatment, as analyzed by SRB assay. Points, mean; bars, SEM ($n=6$). **(b)** The IC_{50} of SIMR compounds in cancer cell lines after 48 h treatment. Data are mean \pm SEM ($n=3$). **(c)** Dose dependent effect of the synthesized SIMR - 1281 on MCF7, MCF7 dox resistant, A549 and A549 dox resistant cell lines after 48 h treatment, as analyzed by MTT assay. Points, mean; bars, SEM ($n=6$).



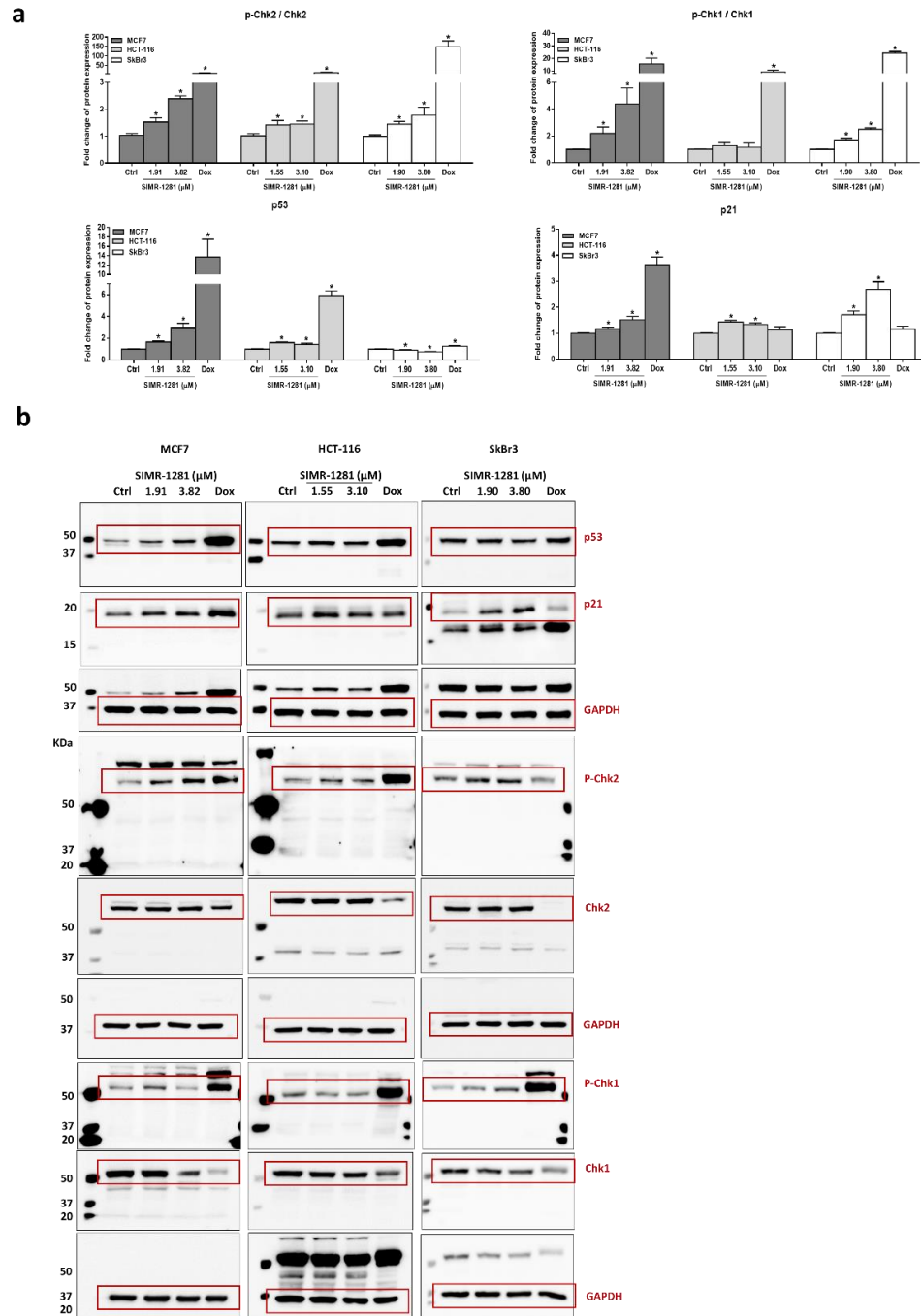
Supplementary Figure S2: DARST Assay showing the binding of SIMR1281 with protein targets



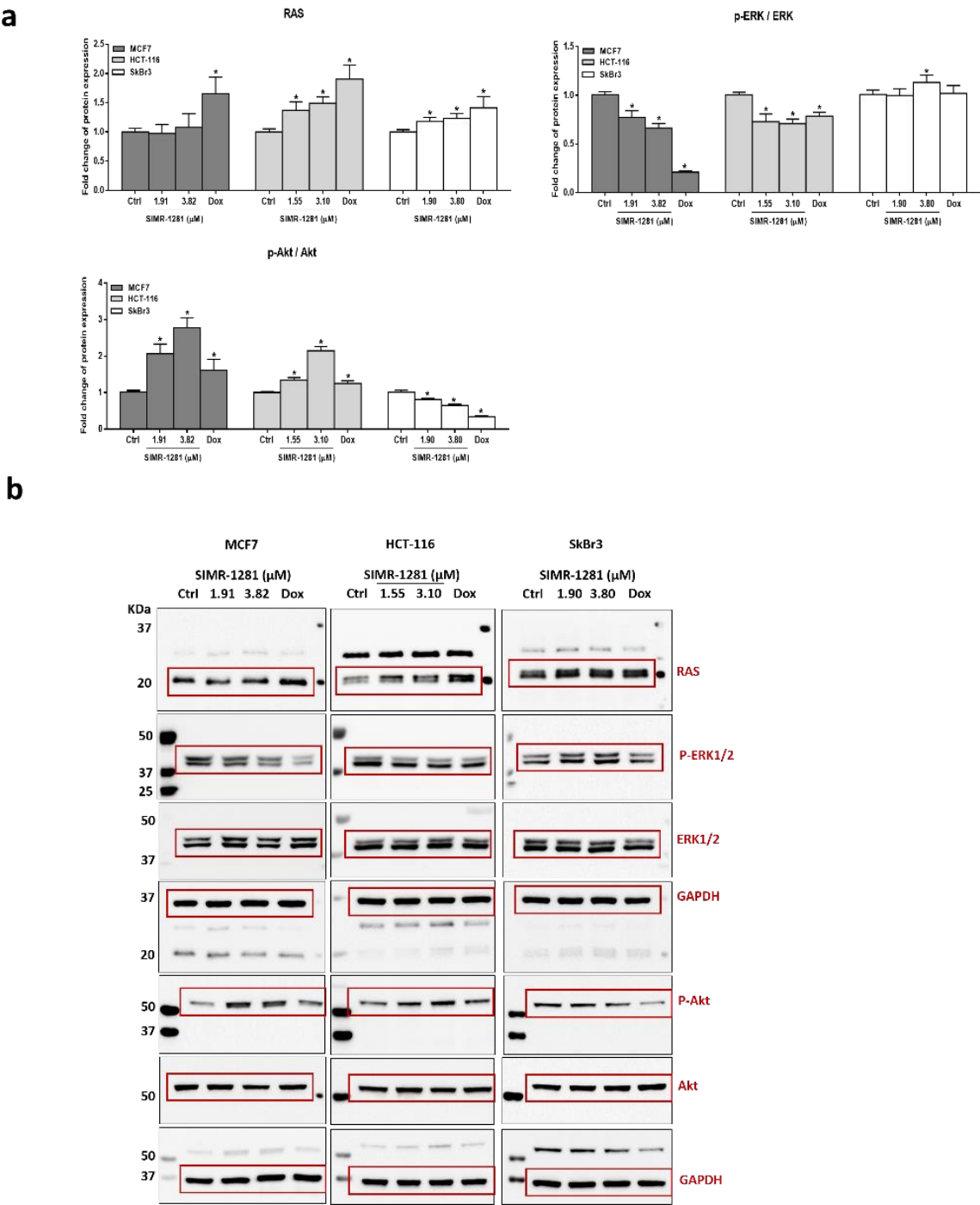
Supplementary Figure S3: Western blot analysis for γ -H2Ax/H2Ax, p-ATM/ATM and p-ATR/ATR proteins in MCF7, HCT116 and SKBR-3 cell lines after the treatment with SIMR-1281 and Dox for 24 h at the indicated concentrations (Whole blot corresponds to Fig3. a). **(a)** Quantification of the bands of γ -H2Ax, H2Ax, p-ATM, ATM, p-ATR and ATR proteins using Image lab software, Data expressed as mean \pm SEM ($n=3$) independent experiments. *indicates significant difference versus control at $p<0.05$ determined by two-tailed unpaired t -test. **(b)** Full blot Western blot of the indicated proteins.



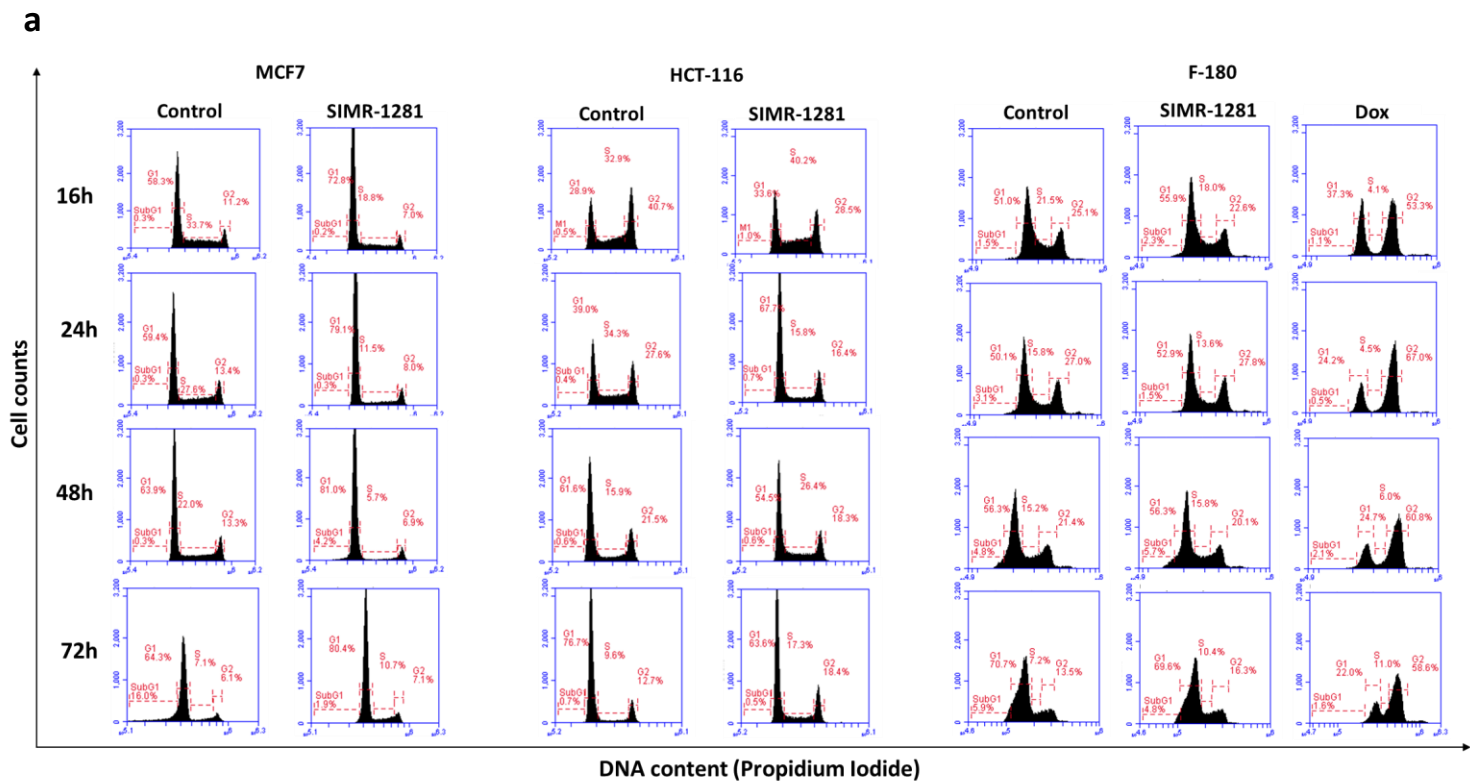
Supplementary Figure S4: Western blot analysis for p53, p21, p-Chk2/Chk2 and p-Chk1/Chk1 proteins in MCF7, HCT116 and SKBR-3 cell lines after the treatment with SIMR-1281 and Dox for 24 h at the indicated concentrations (Whole blot corresponds to Fig3. b). **(a)** Quantification of the bands of p53, p21, p-Chk2, Chk2, p-Chk1 and Chk1 proteins using Image lab software, Data expressed as mean \pm SEM ($n=3$) independent experiments. *indicates significant difference versus control at $p<0.05$ determined by two-tailed unpaired t -test. **(b)** Full blot Western blot of the indicated proteins



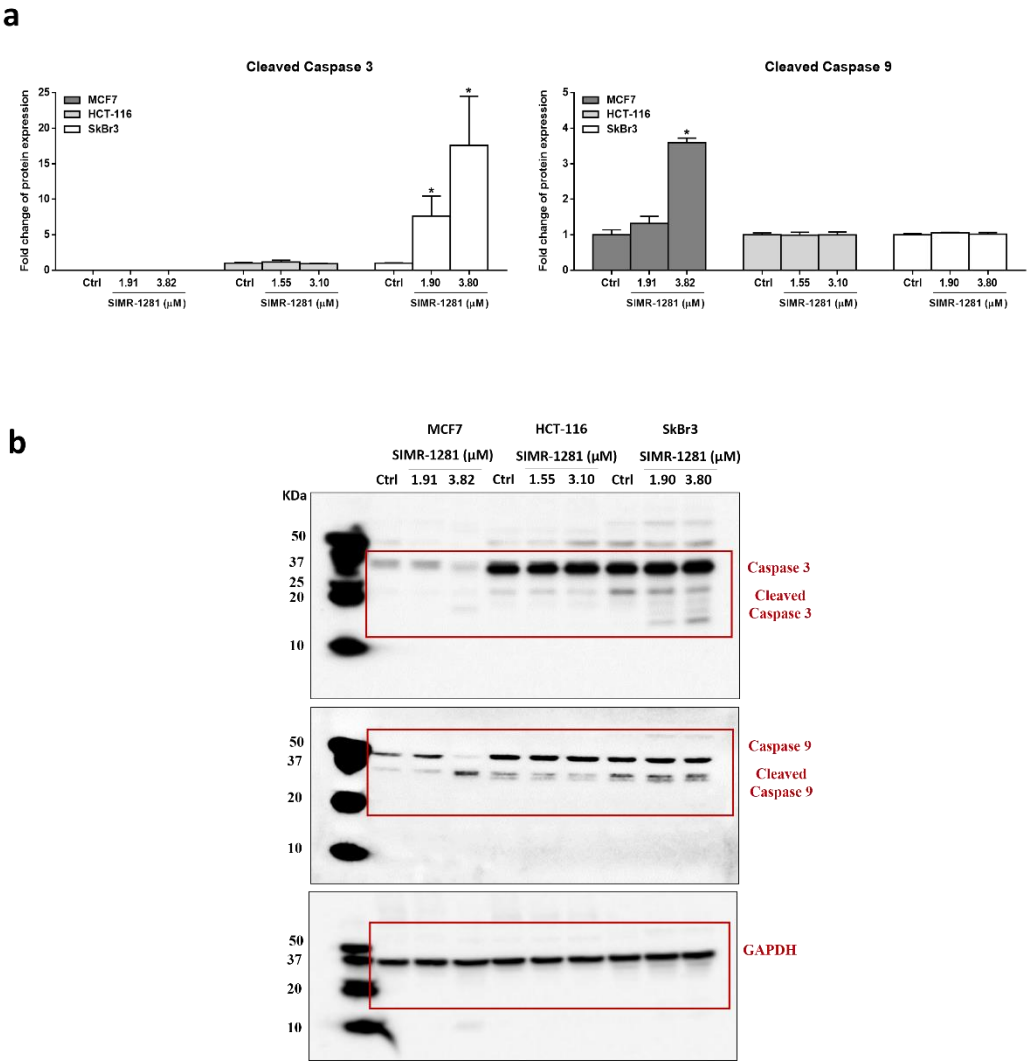
Supplementary Figure S5: Western blot analysis for Ras, p-ERK1/2, ERK1/2, p-Akt, Akt proteins in MCF7, HCT116 and SKBR-3 cell lines after the treatment with SIMR-1281 and Dox for 24 h at the indicated concentrations (Whole blot corresponds to Fig3. c). **(a)** Quantification of the bands of Ras, p-ERK1/2, ERK1/2, p-Akt, Akt proteins using Image lab software, Data expressed as mean \pm SEM ($n=3$) independent experiments. *indicates significant difference versus control at $p<0.05$ determined by two-tailed unpaired t -test. **(b)** Full blot Western blot of the indicated proteins



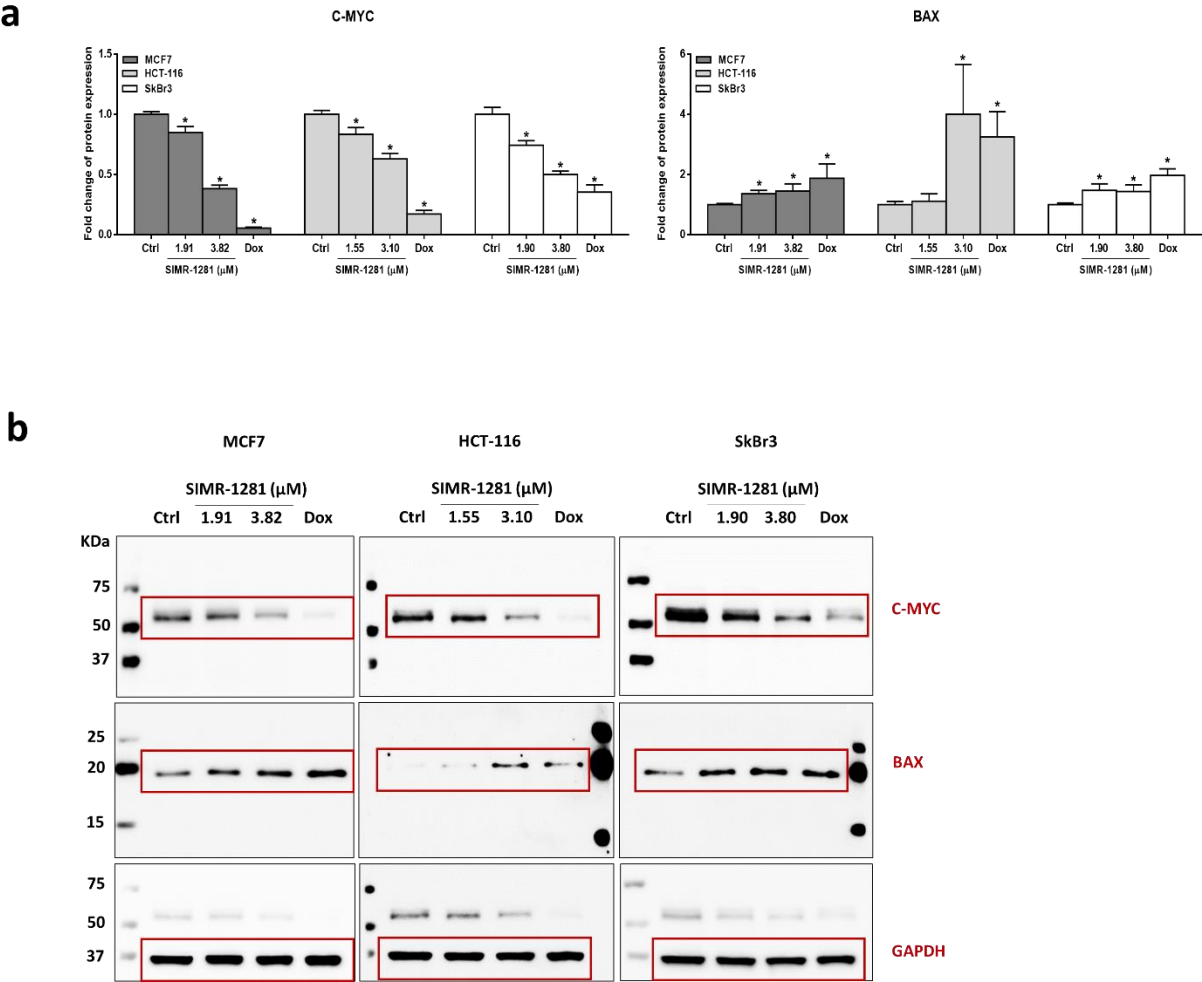
Supplementary Figure S6: Cell cycle analysis of MCF7, HCT116 and F-180 cells treated with SIMR-1281 at the indicated time points. **(a)** Histogram representation of the cell cycle distributions of MCF7, HCT116 and F-180 cells treated with SIMR-1281.



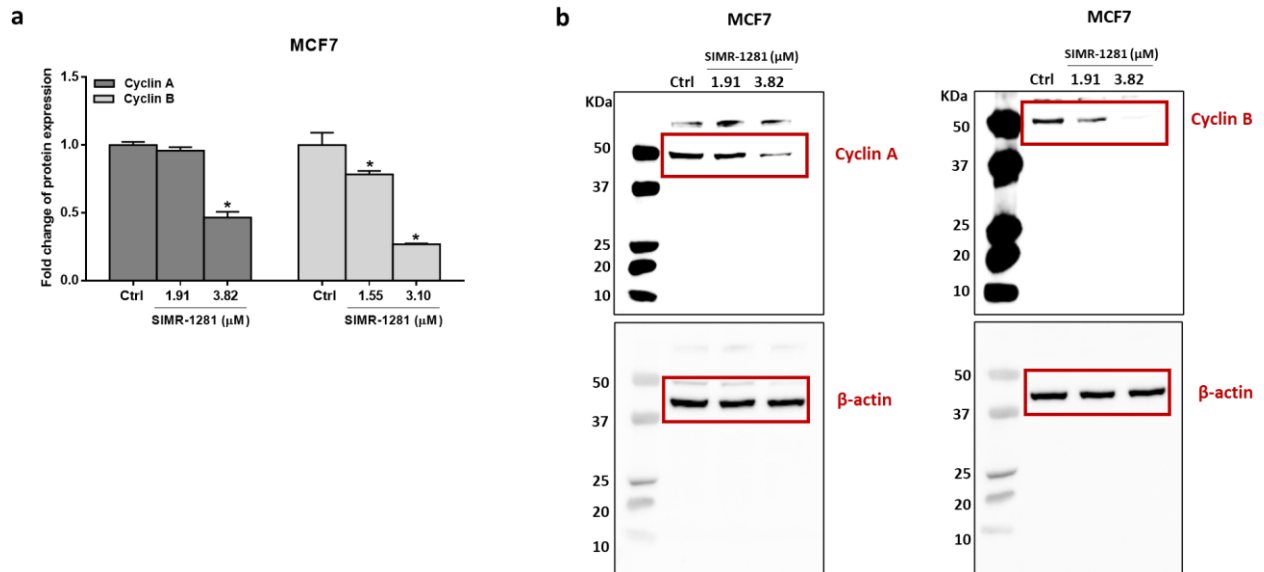
Supplementary Figure S7: Western blot analysis for caspase-3 and caspase-9 proteins in MCF7, HCT116 and SKBR-3 cell lines after the treatment with SIMR-1281 for 24 h at the indicated concentrations (Whole blot corresponds to Fig3. e). **(a)** Quantification of the cleavage bands of caspase 3 and caspase 9 proteins using Image lab software, Data expressed as mean \pm SEM ($n=3$) independent experiments. *indicates significant difference versus control at $p<0.05$ determined by two-tailed unpaired t -test. **(b)** Full blot Western blot of the caspases proteins



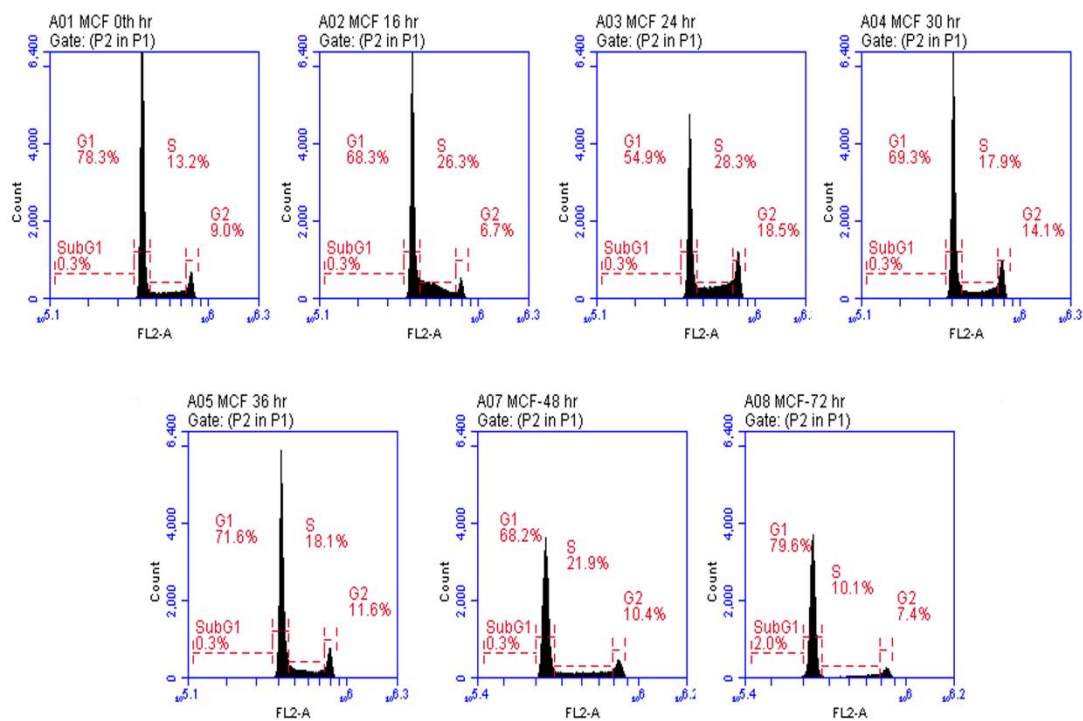
Supplementary Figure S8: Western blot analysis for apoptotic marker proteins in MCF7, HCT116 and SKBR-3 cell lines after the treatment with SIMR-1281 and Dox for 24 h at the indicated concentrations (Whole blot corresponds to Fig3. f). **(a)** Quantification of the apoptosis markers proteins bands using Image lab software, Data expressed as mean \pm SEM ($n=3$) independent experiments. *indicates significant difference versus control at $p<0.05$ determined by two-tailed unpaired t -test. **(b)** Full blot Western blot of the apoptosis markers proteins



Supplementary Figure S9: Western blot analysis for cyclin A and cyclin B proteins in MCF7 cell line after the treatment with SIMR-1281 for 24 h at the indicated concentrations, (Whole blot corresponds to Fig4. c). **(a)** Quantification of cyclin A and cyclin B proteins using Image lab software, Data expressed as mean \pm SEM ($n=3$) independent experiments. *indicates significant difference versus control at $p<0.05$ determined by two-tailed unpaired t -test. **(b)** Full blot Western blot of the cyclins proteins.



Supplementary Figure S10: Histogram representation of the cell cycle distributions of synchronized MCF7 cells at different time points.



Supplementary Figure S11: Hematology values^a and serum chemistry values^a from athymic nude mice bearing subcutaneous HCT116 xenograft tumors after 30 days of treatment with SIMR-1281. Values represent means \pm SD. ^a WBC, white blood cells; RBC, red blood cells; HCT, Hematocrit; MCV, mean corpuscular volume; MCH, Mean Corpuscular Hemoglobin; MCHC, Mean Corpuscular Hemoglobin Concentration; RDW, Red blood cell distribution width; RDWSD, Red blood cell distribution width; PLT, platelet; MPV, mean platelet volume; LY, leukocytes; MO, monocytes; NE, neutrophils; EO, eosinophils; BA, basophils. ^b ALT, Alanine aminotransferase; Aspartate Aminotransferase (AST); Alkaline Phosphatase (Alk Phos, ALP); Gamma Glutamyl Transpeptidase (GGT, GGTP or GTP); Significant difference versus control * **P<0.05** ****P<0.005** determined by two-tailed unpaired *t*-test.

Hematology ^b	Units	Vehicle	SIMR-1281 (50mg/kg)
WBC	$\times 10^3/\mu\text{l}$	2.44 \pm 0.85	2.22 \pm 1.15
RBC	$\times 10^3/\mu\text{l}$	7.18 \pm 0.31	6.72 \pm 0.45
Hemoglobin	g/dL	12.42 \pm 0.75	11.53 \pm 0.76
HCT	%	32.34 \pm 2.11	30.89 \pm 2.18
MCV	fL	44.99 \pm 1.35	45.96 \pm 1.10
MCH	pg	17.28 \pm 0.35	17.16 \pm 0.25
MCHC	g/dL	38.42 \pm 0.58	37.36 \pm 1.06
RDW	%	23.61 \pm 1.26	26.04* \pm 1.43
RDWSD	fL	26.14 \pm 0.5	29.46* \pm 3.12
PLT	$\times 10^3/\mu\text{l}$	853.63 \pm 95.68	718.09 \pm 371.26
MPV	fL	6.30 \pm 0.40	7.69 \pm 1.96
LY	%	90.28 \pm 0.40	86.64 \pm 7.46
MO	%	5.68 \pm 2.81	8.03 \pm 6.73
NE	%	3.16 \pm 1.72	4.25 \pm 1.77
EO	%	0.52 \pm 0.27	0.81 \pm 0.57
BA	%	0.36 \pm 0.21	0.27 \pm 0.14
Serum Chemistry^c			
Creatinine	mg/dL	1.04 \pm 0.89	1.176 \pm 0.398
ALT	iu/L	25.19 \pm 5.55	44.35 \pm 14.84
AST	iu/L	91.25 \pm 20.50	180.72 \pm 54.93
ALP	iu/L	33.42 \pm 19.21	25.722 \pm 13.25
GAMMA GT		11.88 \pm 9.20	16.45 \pm 2.041
Total Bilirubin	mg/dL	0.548 \pm 0.040	0.7 \pm 0.327
Total Protein	g/dL	7.136 \pm 0.157	10.124 \pm 4.383
Albumin	g/dL	3.277 \pm 0.168	2.660** \pm 0.2478
Urea		51.22 \pm 2.469	57.094 \pm 12.29
Glucose	mg/dL	283.55 \pm 20.00	176.59 \pm 50.93