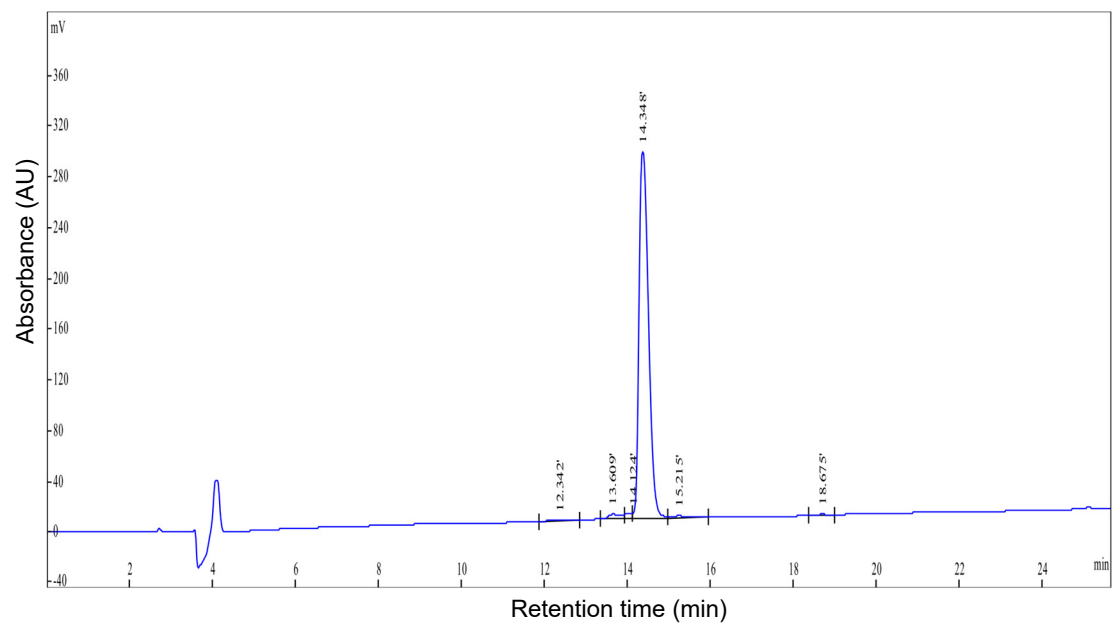


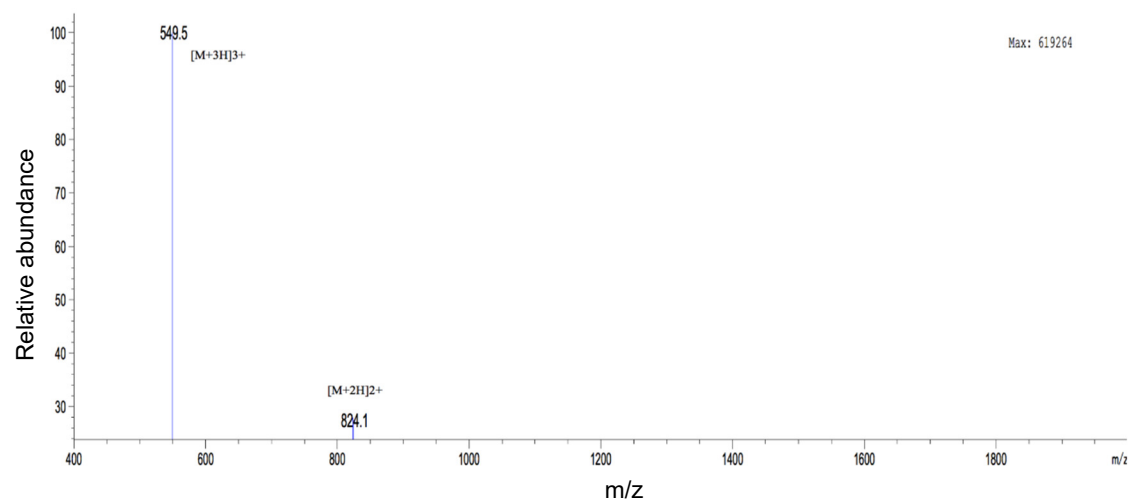
3-NAntC: A Potent Crotoxin B-Derived Peptide against the Triple-Negative MDA-MB-231 Breast Cancer Cell Line

Supplementary results.

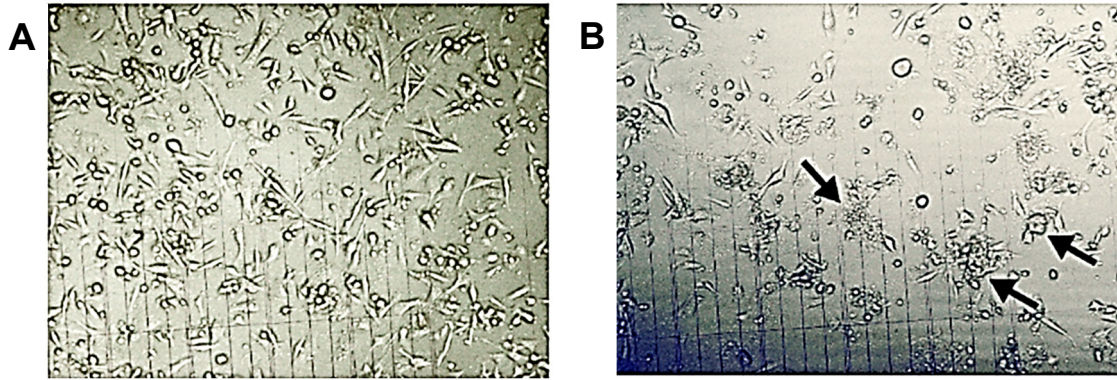


Peak No.	Ret Time	Height	Area	Conc.
1	12.342	1289	35239	0.7983
2	13.609	3409	71174	1.612
3	14.124	4266	44560	1.009
4	14.348	288295	4223309	95.67
5	15.215	1754	32466	0.7354
6	18.675	1101	7769	0.176
Total				100.0000

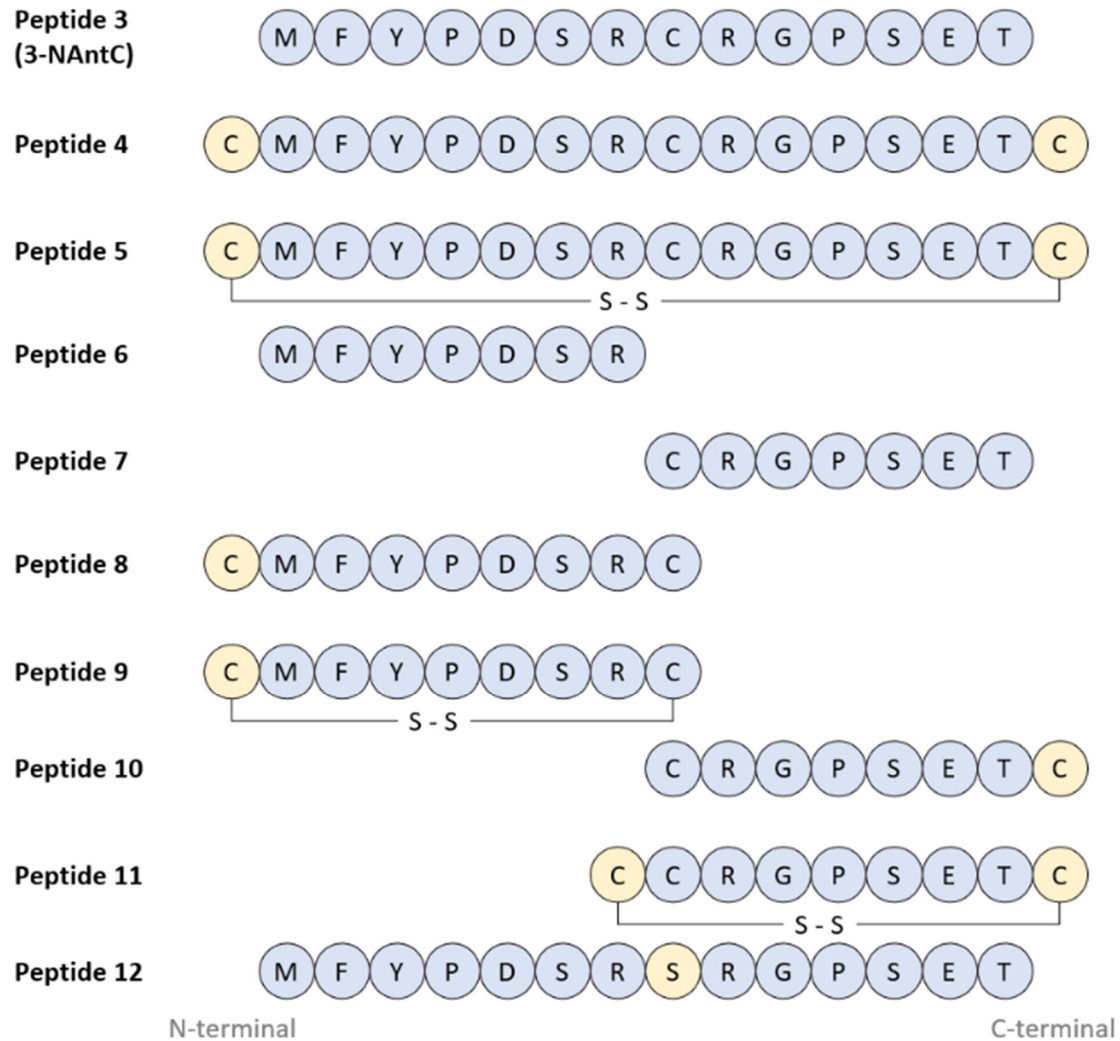
Supplementary Figure S1. HPLC chromatogram of 3-NantC. The peptide had a purity higher than 95%. Absorbance at the wavelength of 220 nm.



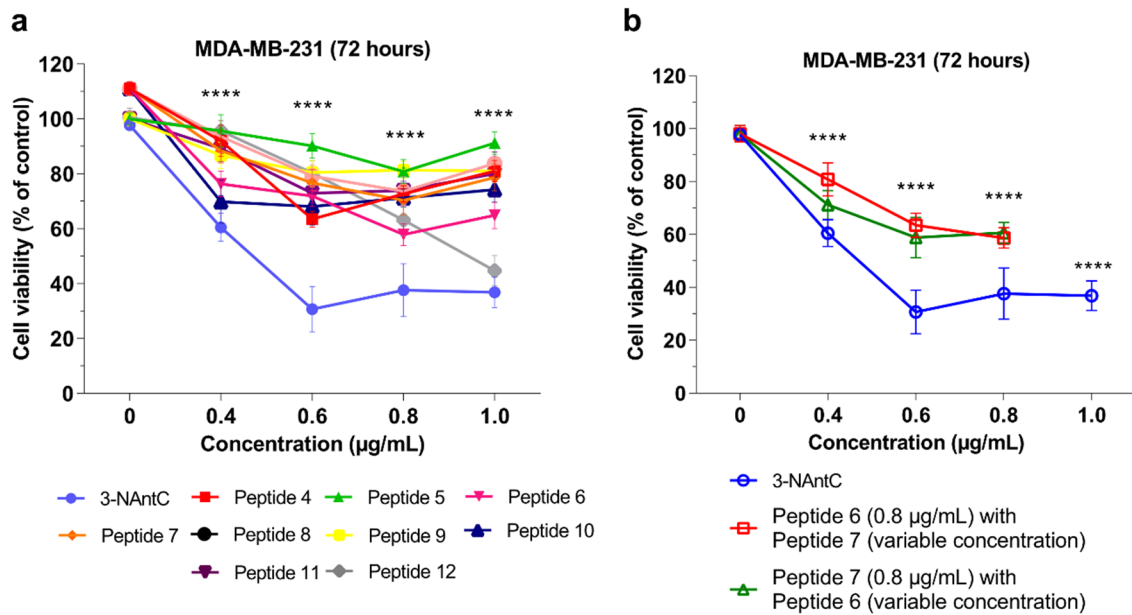
Supplementary Figure S2. MS spectrum of 3-NantC. The mass spectrometry characterization with ESI ionization confirmed the peptide sequence [H]-MFYPDSRCRGPSET-[OH] (Molecular weight = 1,645.95 g/mol).



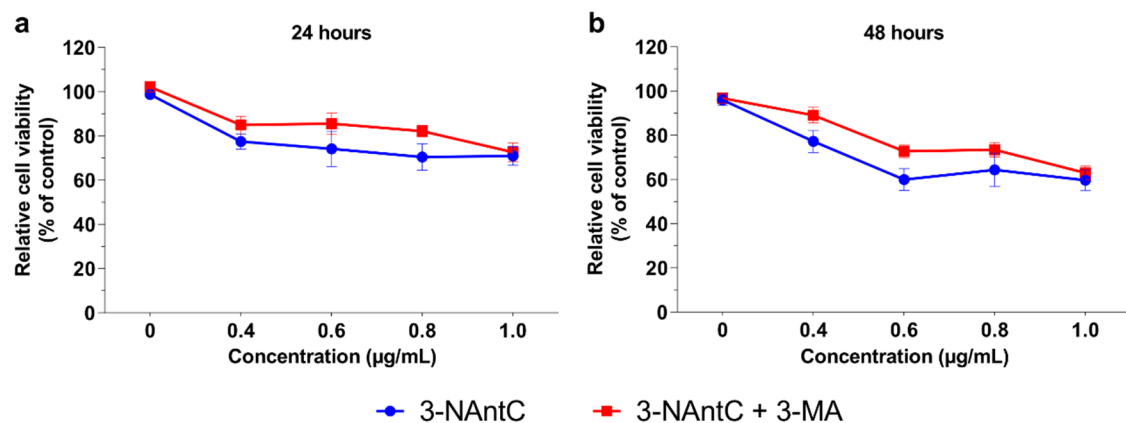
Supplementary Figure S3. Antitumor effect of the peptide 3-NantC against triple-negative breast cancer. MDA-MB-231 cells were incubated in presence and absence of 3-NantC (1 µg/mL) for 24 hours. (A) Untreated triple negative tumor cells exhibiting a large number of cells. (B). Tumor cells after being treated with the peptide 3-NantC. The arrows indicate the tumor cells (MDA-MB-231), which are shown in smaller numbers and with vesicles compatible with apoptotic bodies when compared with the untreated tumor cells.



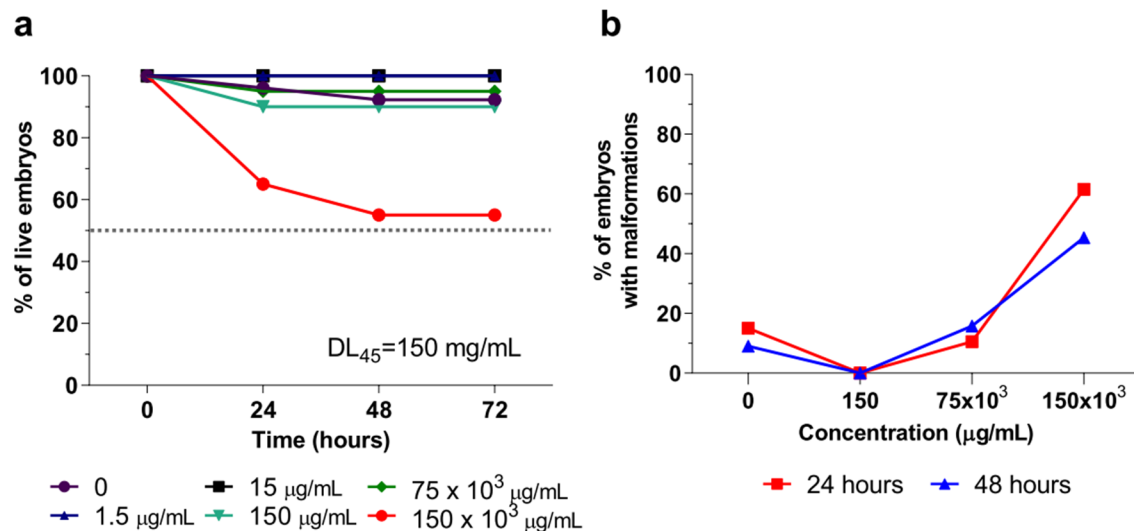
Supplementary Figure S4. Amino acid sequences of the 3-NAntC-derived peptides. The amino acids matching the original 3-NAntC sequence are shown in blue, while the additional amino acids are shown in light orange. Intramolecular disulfide bonds are denoted as 'S-S'. The sequences are shown from the N- to the C-terminal regions.



Supplementary Figure S5. Effect of 3-NAntC-derived peptides on TNBC cells (MDA-MB-231) viability. (A) MDA-MB-231 cells were treated with 3-NAntC or peptides 4 – 12 for 72 hours at the concentration range of 0.4 – 1.0 $\mu\text{g/mL}$. (B) Antitumor effect of 3-NAntC in comparison with the combination of the peptides 6 and 7. Cellular viability assay was conducted using the MTT method for the TNBC cell line. Data are shown as mean \pm SEM of at least three independent assays in triplicate. Treatment groups for each peptide were compared to the respective negative control (PBS 0.02%) using the two-way ANOVA and the Bonferroni post hoc test. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, and **** $p < 0.0001$ denote statistically significant differences for all the groups at a specific concentration.

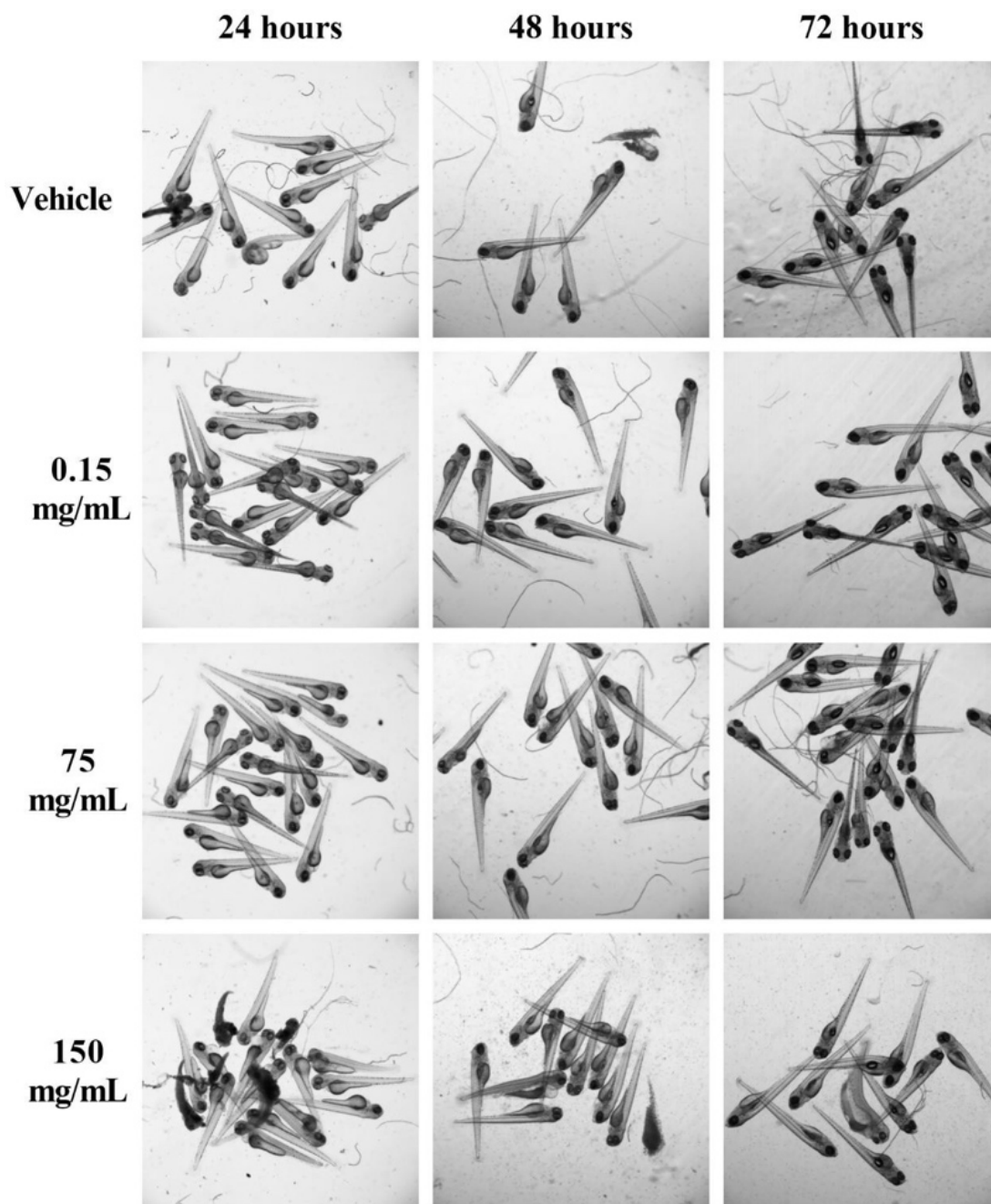


Supplementary Figure S6. Effect of 3-methyladenine (3-MA) (an autophagy inhibitor) on the decrease of the MDA-MB-231 cell viability mediated by 3-NAntC. MDA-MB-231 cells were treated with increasing concentrations of 3-NAntC in the presence or not of 3-methyladenine (3-MA) for 24 (A) or 48 hours (B). Cell viability was measured using the MTT assay. Data are shown as mean \pm SEM of at least three independent assays in triplicate. Groups treated with 3-NAntC alone were compared to those treated with 3-NAntC + 3-MA (the autophagy inhibitor) using the one-way ANOVA and the Bonferroni posthoc test. No statistically significant differences were observed between the groups, $p > 0.05$.



Supplementary Figure S7. 3-NAntC safety evaluation using zebrafish embryos.

Embryo survival (A) and malformation in live embryos (B) were assessed following the exposure to 3-NAntC (1.5 – 150 $\times 10^3 \mu\text{g/mL}$) for up to 72 hours. 3-NAntC exhibits low toxicity and malformation in zebrafish embryos up to 75 mg/mL . Data is shown as the percentage of live embryos and percentage of live embryos with malformation after the treatment with 3-NAntC (n=20 embryos/group).



Supplementary Figure S8. Representative images of the tolerability test of 3-NantC in zebrafish embryos. Zebrafish embryos were incubated with vehicle (control) or 3-NantC peptide (0.15, 75, and 150 mg/mL) for 24, 48, or 72 hours.

Supplementary Table S1. Normality score for each data set used in the present study.

Data set			n	Shapiro-Wilk	
				score	p-value
Figure 2	24h	MCF10A Peptide 1	40	0.9527	0.7543
		MCF10A Peptide 2	51	0.8985	0.3217
		MCF10A Peptide 3	46	0.9157	0.4365
		MDA-MB-231 Peptide 1	29	0.9402	0.0514

		MDA-MB-231 Peptide 2	44	0.9541	0.0785
		MDA-MB-231 Peptide 3	37	0.9579	0.1737
Figure 2	48h	MCF10A Peptide 1	40	0.9278	0.5324
		MCF10A Peptide 2	57	0.9838	0.8265
		MCF10A Peptide 3	51	0.9609	0.6406
		MDA-MB-231 Peptide 1	21	0.9366	0.1865
		MDA-MB-231 Peptide 2	37	0.9589	0.1857
		MDA-MB-231 Peptide 3	39	0.9633	0.2291
Figure 3	24h	MCF10A	23	0.9283	0.1004
		HMEC	22	0.9638	0.1892
		HDFa	49	0.973	0.3173
		MDA-MB-231	58	0.9727	0.2155
		MCF7	28	0.9304	0.1633
Figure 3	48h	MCF10A	26	0.8954	0.1237
		HMEC	29	0.9451	0.1363
		HDFa	52	0.9566	0.5580
		MDA-MB-231	57	0.9821	0.5590
		MCF7	24	0.9409	0.1705
Figure 3	72h	MCF10A	28	0.9409	0.1164
		HMEC	37	0.951	0.1243
		HDFa	47	0.9619	0.1279
		MDA-MB-231	64	0.9382	0.0514
		MCF7	20	0.9724	0.8051
Figure 4	24h	MDA-MB-231 3-NAntC	60	0.9477	0.1210
		MDA-MB-231 Doxorubicin	68	0.9579	0.2192
		MDA-MB-231 Cisplatin	54	0.9635	0.2985
		HDFa 3-NAntC	50	0.9739	0.3317
		HDFa Doxorubicin	35	0.9243	0.1903
		HDFa Cisplatin	35	0.9532	0.2146
Figure 4	48h	MDA-MB-231 3-NAntC	52	0.9607	0.3843
		MDA-MB-231 Doxorubicin	66	0.9035	0.0858
		MDA-MB-231 Cisplatin	54	0.9795	0.4784
		HDFa 3-NAntC	54	0.9515	0.3790
		HDFa Doxorubicin	32	0.9286	0.3588

		HDFa Cisplatin	21	0.8565	0.05515
Figure 4	72h	MDA-MB-231 3-NAntC	83	0.8868	0.2450
		MDA-MB-231 Doxorubicin	69	0.8537	0.2010
		MDA-MB-231 Cisplatin	51	0.9732	0.3997
		HDFa 3-NAntC	53	0.9668	0.3470
		HDFa Doxorubicin	30	0.8243	0.1877
		HDFa Cisplatin	29	0.9127	0.3003
Figure 5	24h	MDA-MB-231	74	0.9587	0.3165
	48h	MDA-MB-231	74	0.9306	0.1608
		MDA-MB-231 G0/G1	19	0.8936	0.0574
		MDA-MB-231 S	23	0.9378	0.1649
		MDA-MB-231 G2/M	21	0.9696	0.4241
	72h	MDA-MB-231	67	0.9201	0.0585
		HDFa	35	0.9699	0.4405
Figure 6	24h	MDA-MB-231 Apoptosis	14	0.9511	0.5773
		MDA-MB-231 Necrosis	18	0.8226	0.0583
		MDA-MB-231 LDH	13	0.9251	0.2938
	48h	MDA-MB-231 Apoptosis	16	0.7916	0.2309
		MDA-MB-231 Necrosis	12	0.9499	0.6362
		MDA-MB-231 LDH	16	0.7745	0.2127
	72h	MDA-MB-231 Apoptosis	77	0.9294	0.3584
		MDA-MB-231 Necrosis	85	0.951	0.4746
		MDA-MB-231 LDH	72	0.9703	0.5867