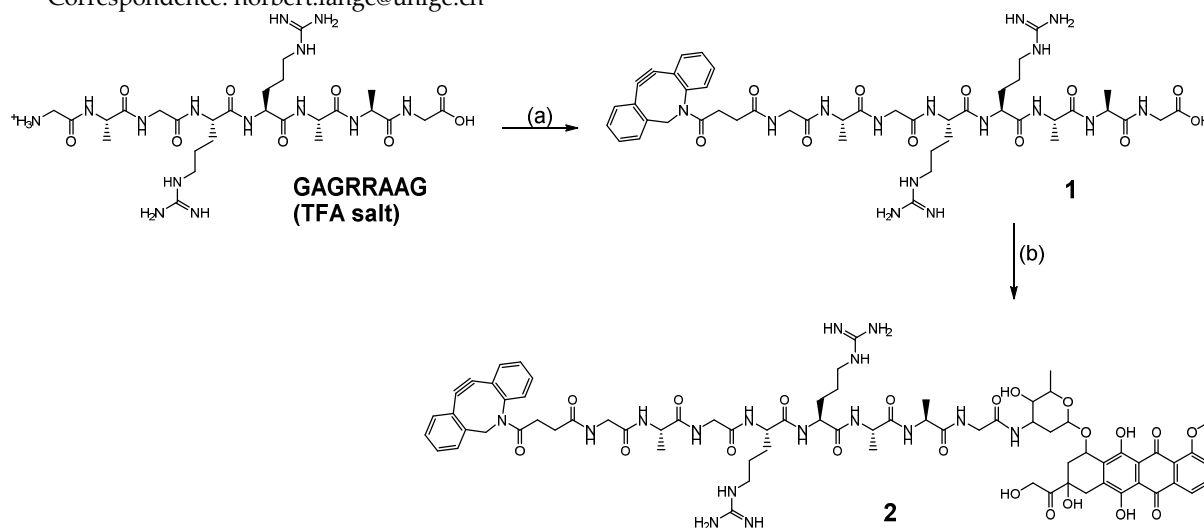


# Cathepsin B-cleavable Cyclopeptidic Chemotherapeutic Prodrugs

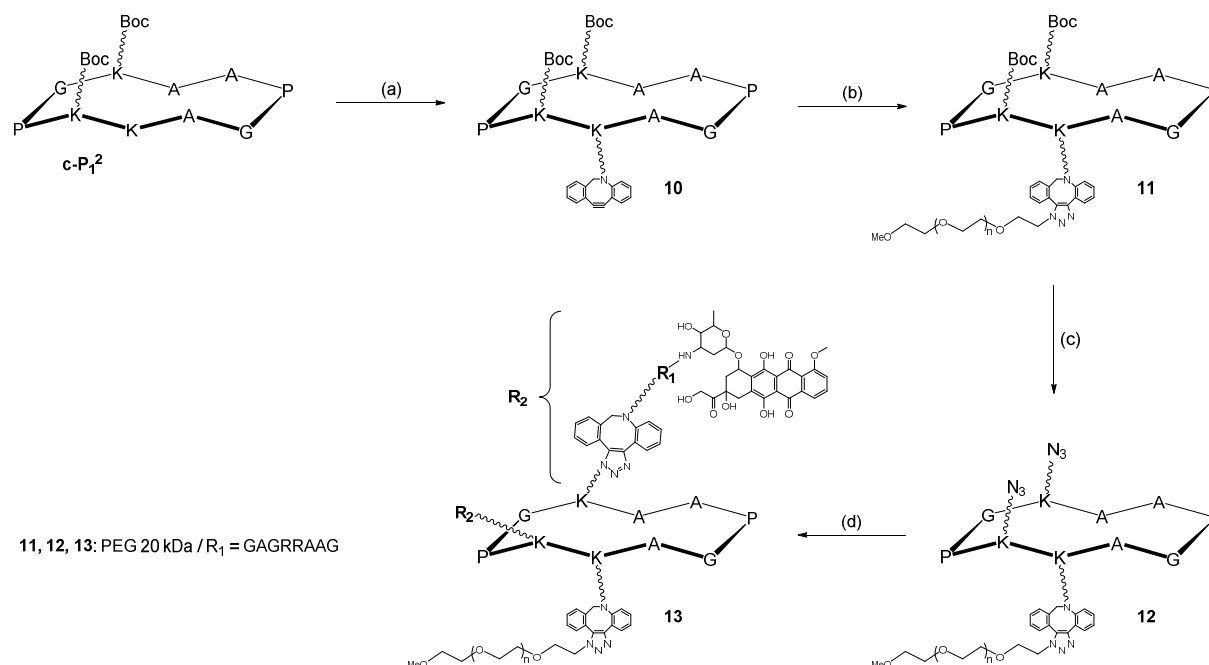
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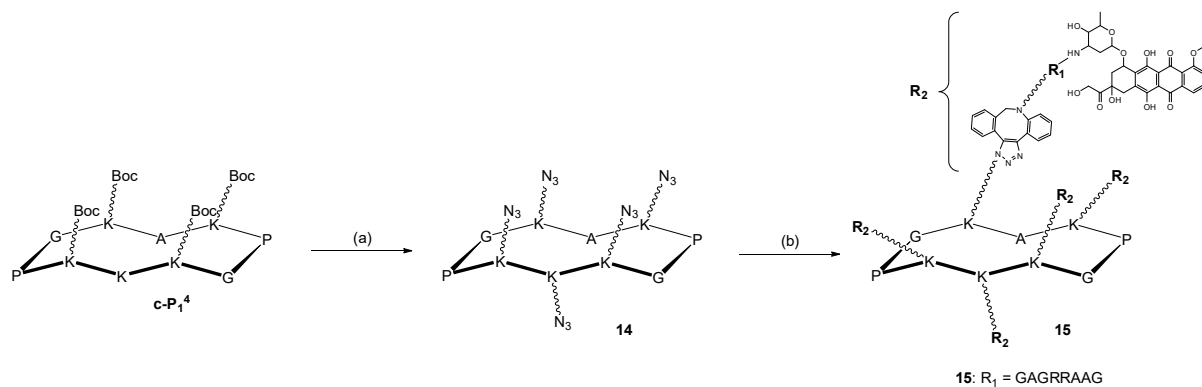
\* Correspondence: norbert.lange@unige.ch



Scheme S1 - Generic synthesis pathway for the DBCO-GAGRRAAG-DOX conjugate. *Reagents and conditions:* (a) DBCO-NHS, DIPEA, argon (ar.), dark, room temperature (r. t.), overnight (o/n); (b) DOX (HCl salt), HATU, DIPEA, ar., dark, r. t., o/n; (c) DBCO-NH<sub>2</sub>, HATU, DIPEA, ar., dark, r. t., o/n.



Scheme S2 - Generic synthesis pathway for the disubstituted-monoPEGylated conjugate cPCP<sub>2/20</sub>. *Reagents and conditions:* (a) DBCO-NHS, argon (ar.), room temperature (r. t.), overnight (o/n); (b)  $\alpha$ -azido- $\omega$ -methoxy-PEG (20 kDa), ar., r. t., o/n; (c) TFA/DCM (50:50), 30 min, r. t.; APA, DIPEA, HATU, ar., r. t., o/n; (d) **(2)**, ar., dark, r. t., o/n.



Scheme S3 - Generic synthesis pathway for the pentasubstituted conjugate cPCPs. *Reagents and conditions:* (a) TFA/DCM (50:50), 30 min, r. t.; azidoacetic acid-NHS, DIPEA, HATU, ar., r. t., o/n; (b) (2), ar., dark, r. t., o/n.

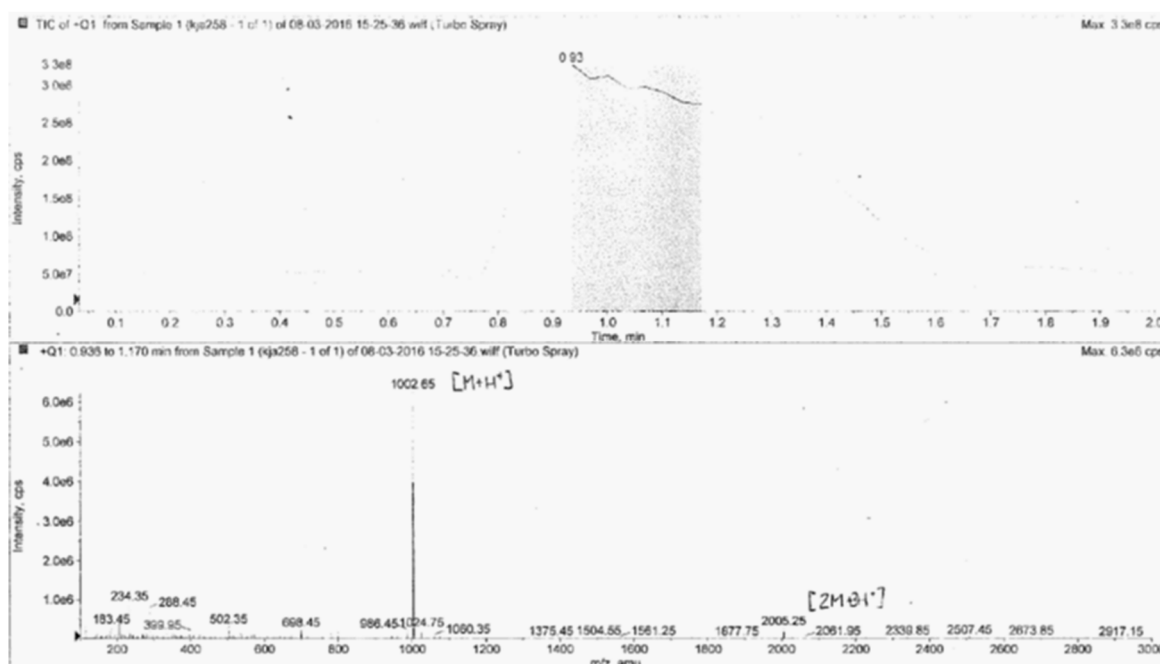


Figure S1 - ESI-MS analysis of DBCO-GAGRRAAG

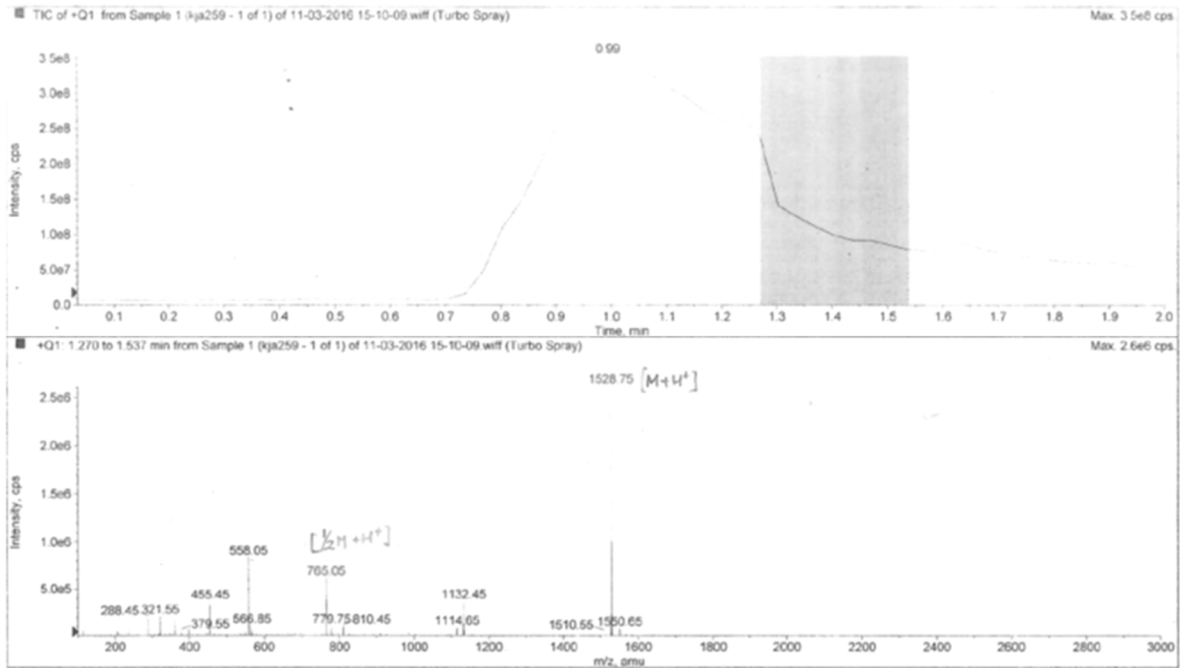


Figure S2 - ESI-MS analysis of DBCO-GAGRRAAG-DOX

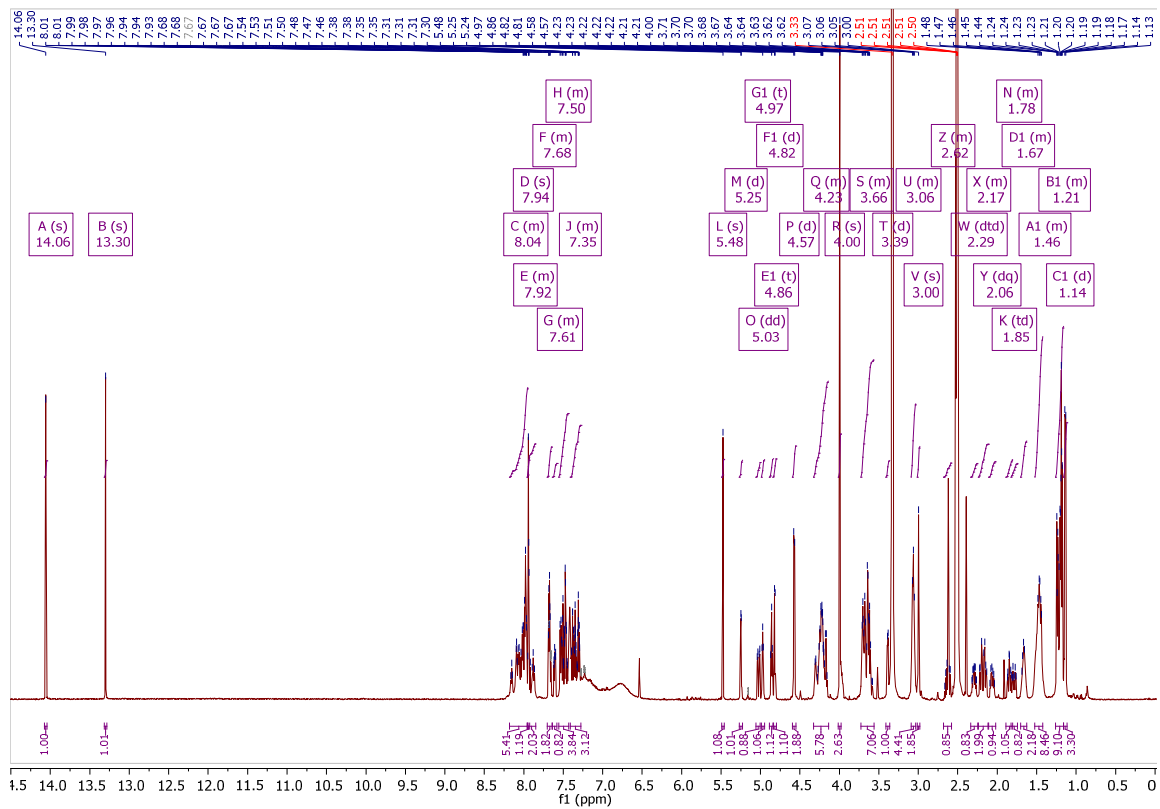


Figure S3 - <sup>1</sup>H NMR (DMSO-d<sub>6</sub>, 300 MHz) NMR analysis of DBCO-GAGRRAAG-DOX

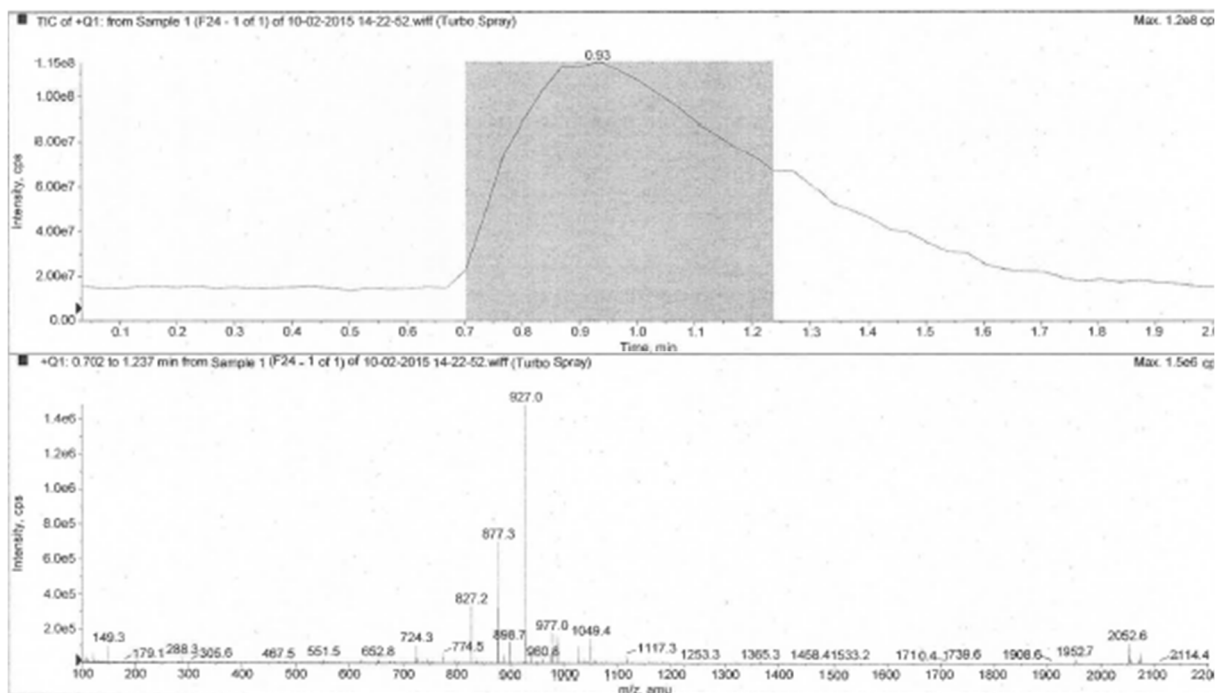


Figure S4 - ESI-MS analysis of (DBCO)<sub>2</sub>-cP<sub>24</sub>

Shimadzu Biotech Axima CFRplus 2.8.4.20081127: Mode Linear, Power: 85, P.Ext. @ 5000 (bin 114)

%Int. 0.6 mV[sum= 18 mV] Profiles 1-31 Smooth Av 10

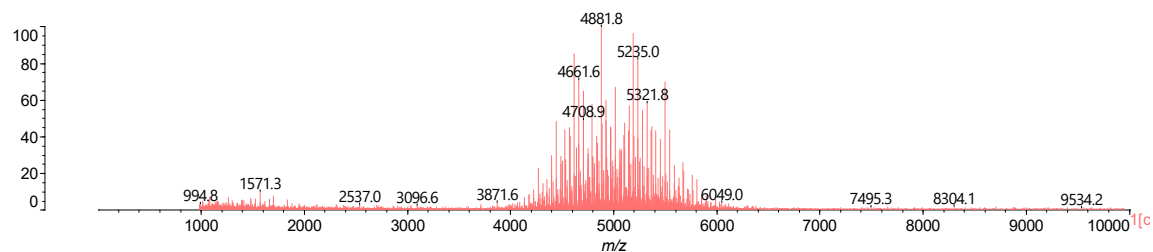
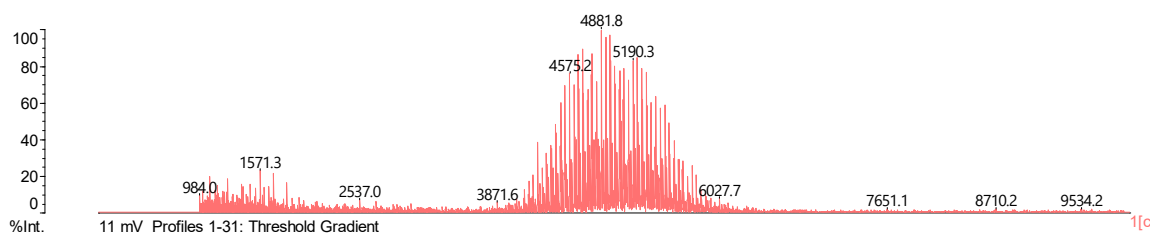


Figure S5 - MALDI-TOF analysis of MeO-PEG-N<sub>3</sub> 5 kDa. Mw expected according to the supplier: 5079 Da

Shimadzu Biotech Axima CFRplus 2.8.4.20081127: Mode Linear, Power: 85, P.Ext. @ 18000 (bin 192)

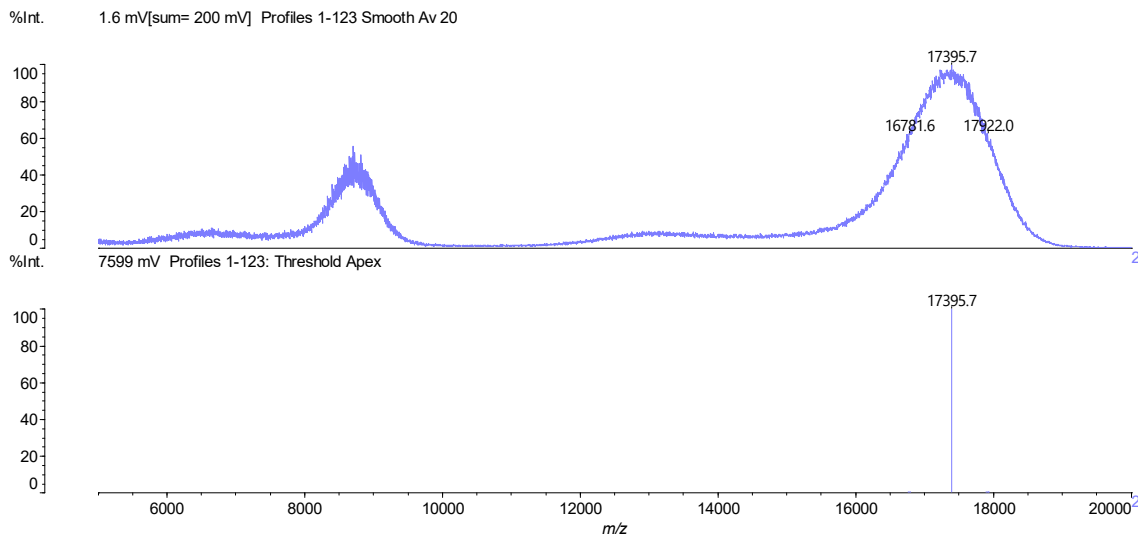


Figure S6 - MALDI-TOF analysis of cPCP<sub>4/5</sub><sup>2</sup>

Shimadzu Biotech Axima CFRplus 2.8.4.20081127: Mode Linear, Power: 90, P.Ext. @ 11000 (bin 150)

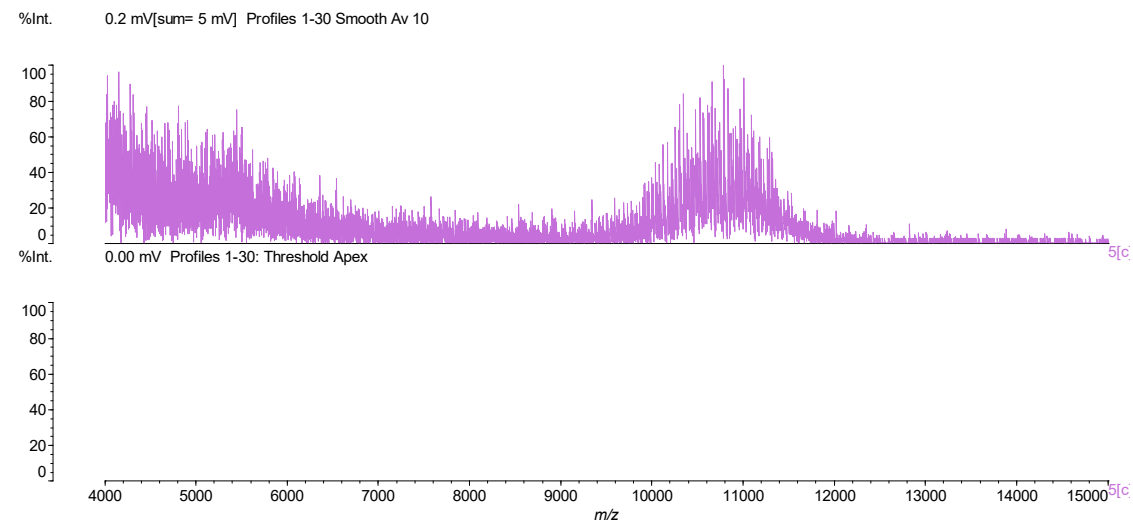


Figure S7 - MALDI-TOF analysis of MeO-PEG-N<sub>3</sub> 10 kDa. Mw expected according to the supplier: 11153 Da; found: ~10800 Da.

Shimadzu Biotech Axima CFRplus 2.8.4.20081127: Mode Linear, Power: 100, P.Ext. @ 30500 (bin 250)

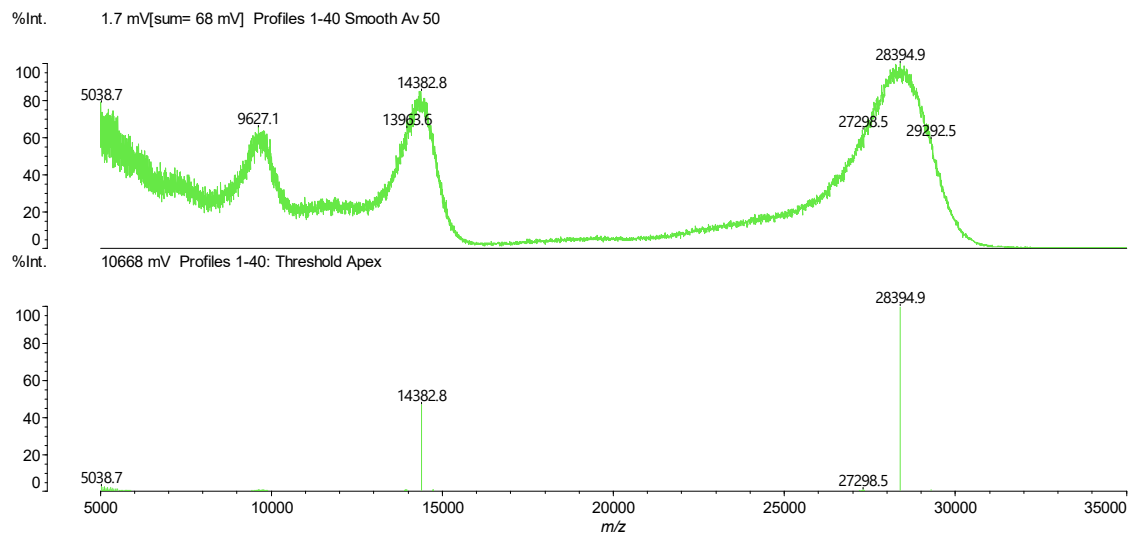
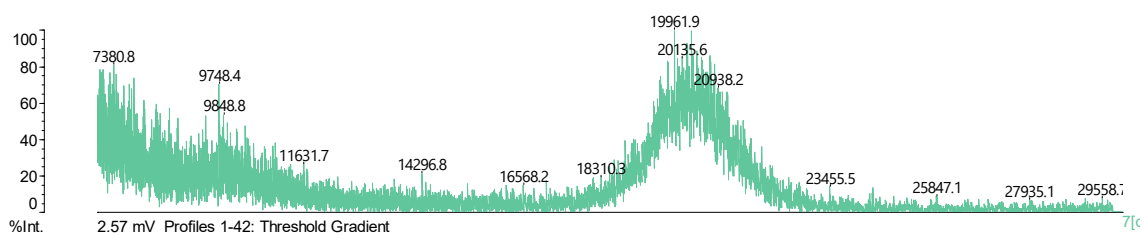


Figure S8 - MALDI-TOF analysis of cPCP<sub>4/10</sub><sup>2</sup>.

Shimadzu Biotech Axima CFRplus 2.8.4.20081127: Mode Linear, Power: 100, P.Ext. @ 25000 (bin 256)

%Int. 0.1 mV[sum= 3 mV] Profiles 1-42 Smooth Av 20



2.57 mV Profiles 1-42: Threshold Gradient

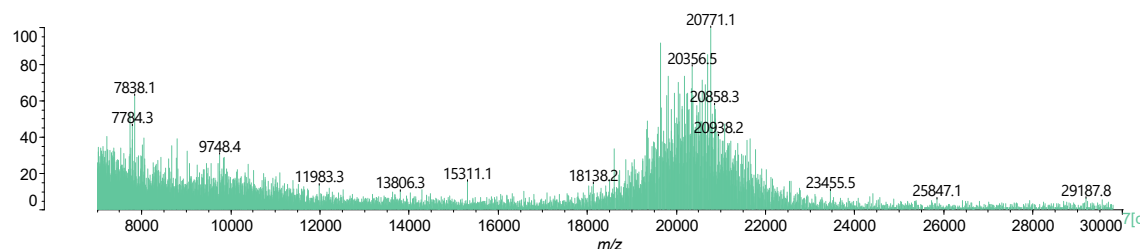
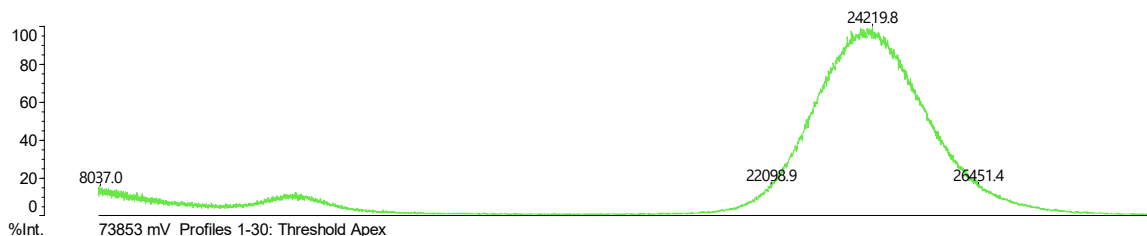


Figure S9 - MALDI-TOF of MeO-PEG-N<sub>3</sub> 20 kDa. Mw expected according to the supplier: 22271 Da; found: ~20400 Da

Shimadzu Biotech Axima CFRplus 2.8.4.20081127: Mode Linear, Power: 105, P.Ext. @ 26700 (bin 234)

%Int. 6.7 mV[sum= 201 mV] Profiles 1-30 Smooth Av 50



73853 mV Profiles 1-30: Threshold Apex

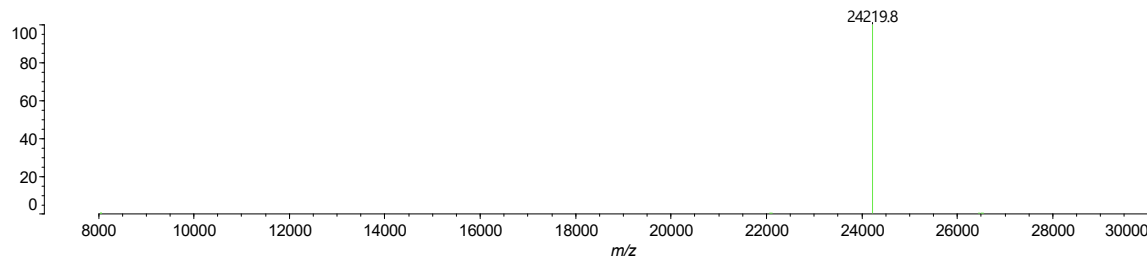


Figure S10 - MALDI-TOF analysis of cPCP<sub>2/20</sub>

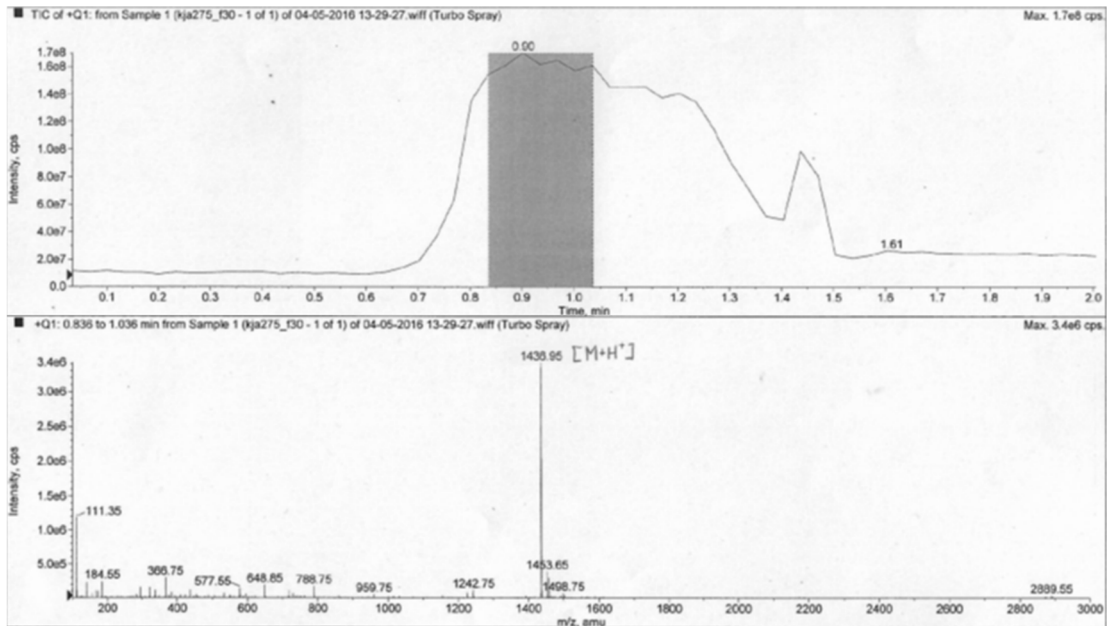


Figure S11 - ESI-MS analysis of cP<sub>1</sub><sup>4</sup> after Boc deprotection

Shimadzu Biotech Axima CFRplus 2.8.4.20081127: Mode Linear, Power: 95, P.Ext. @ 9070 (bin 136)

%Int. 6.2 mV[sum= 311 mV] Profiles 1-50 Smooth Av 20

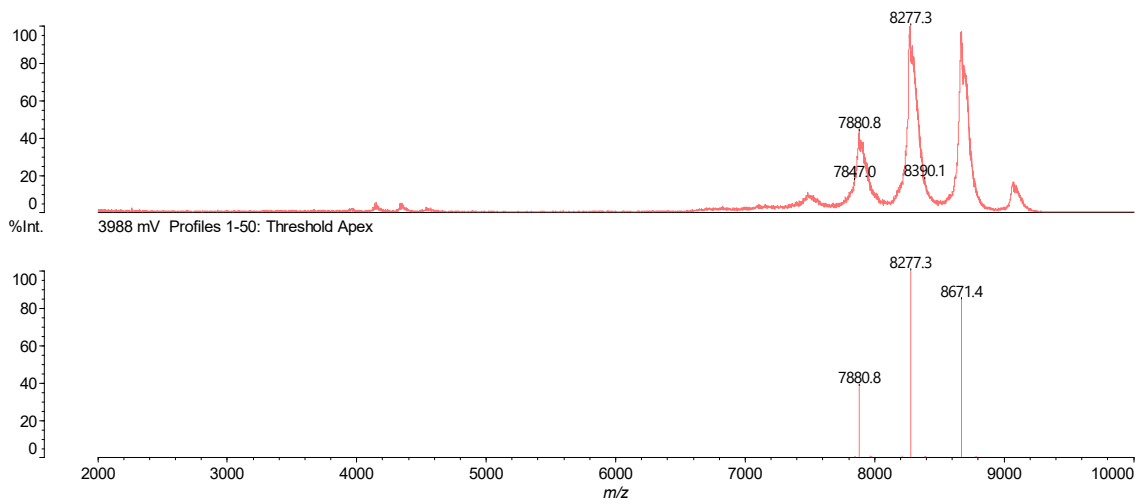


Figure S12 - MALDI-TOF analysis of cPCP<sub>5</sub>.

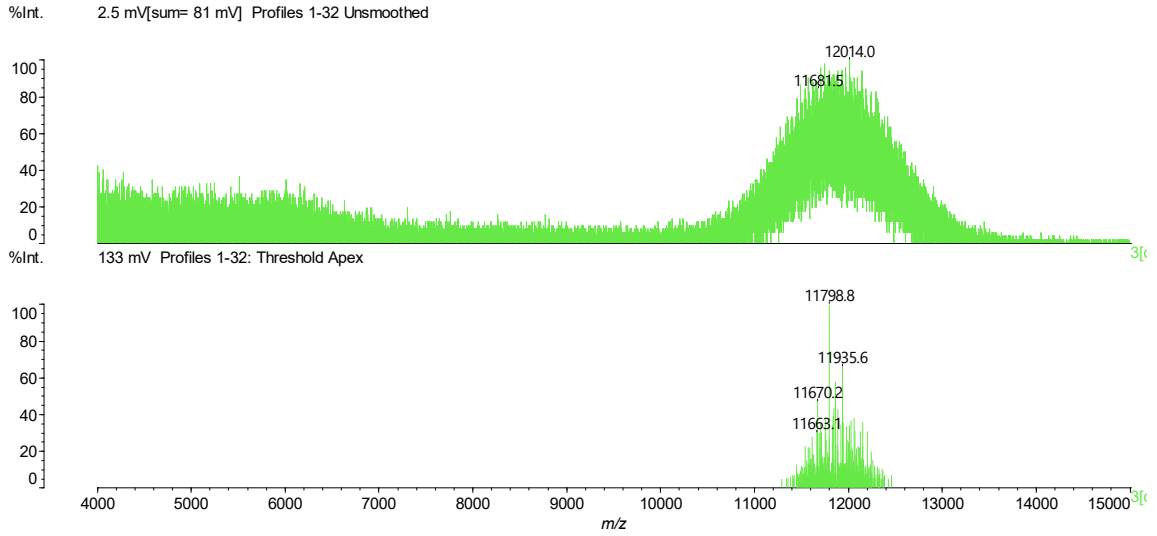


Figure S13 - MALDI-TOF analysis of PEG<sub>10</sub>-GAGRRAAG-DOX

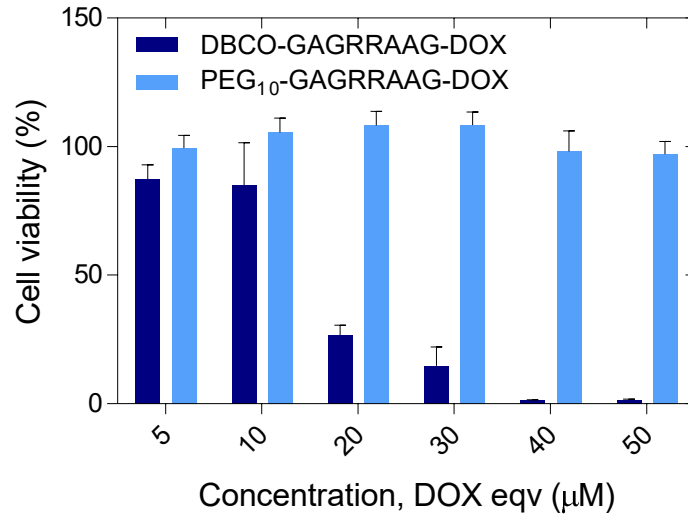


Figure S14 - WST-1 cell proliferation assay in HT1080 after 72 hours of incubation with PEG<sub>10</sub>-GAGRRAAG-DOX (n = 3).

[Dox] mM	0.01	0.05	0.10	0.15	0.2	0.4	0.5	0.6	0.8	1.0
SR (%)	86.4± 9.7	81.7±1 0.6	69.9± 9.7	53.7± 7.3	34.6± 4.7	6.9±1 .3	1.3±0 .4	0.4±0 .5	0±0 .7	0.4±0 .5

Table S1: WST-1 cell proliferation assay in HT1080 after 72 hours of incubation with DOX SR: Survival Rate



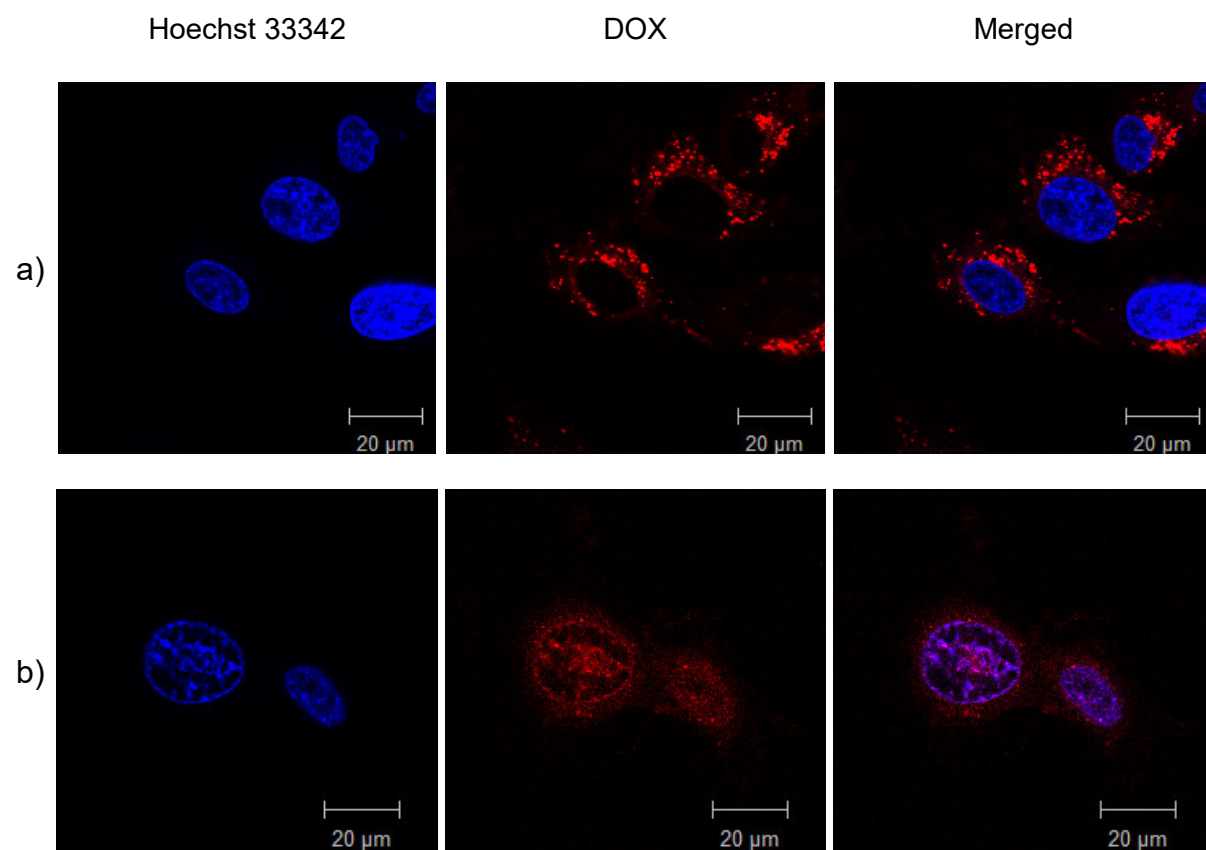


Figure S15 - HT1080 – fluorescence micrographs upon 24 hours of incubation with a) 30  $\mu\text{M}$  of  $\text{cPCP}_{4/5^2}$ , b) 0.3  $\mu\text{M}$  of free DOX. Magnification 63

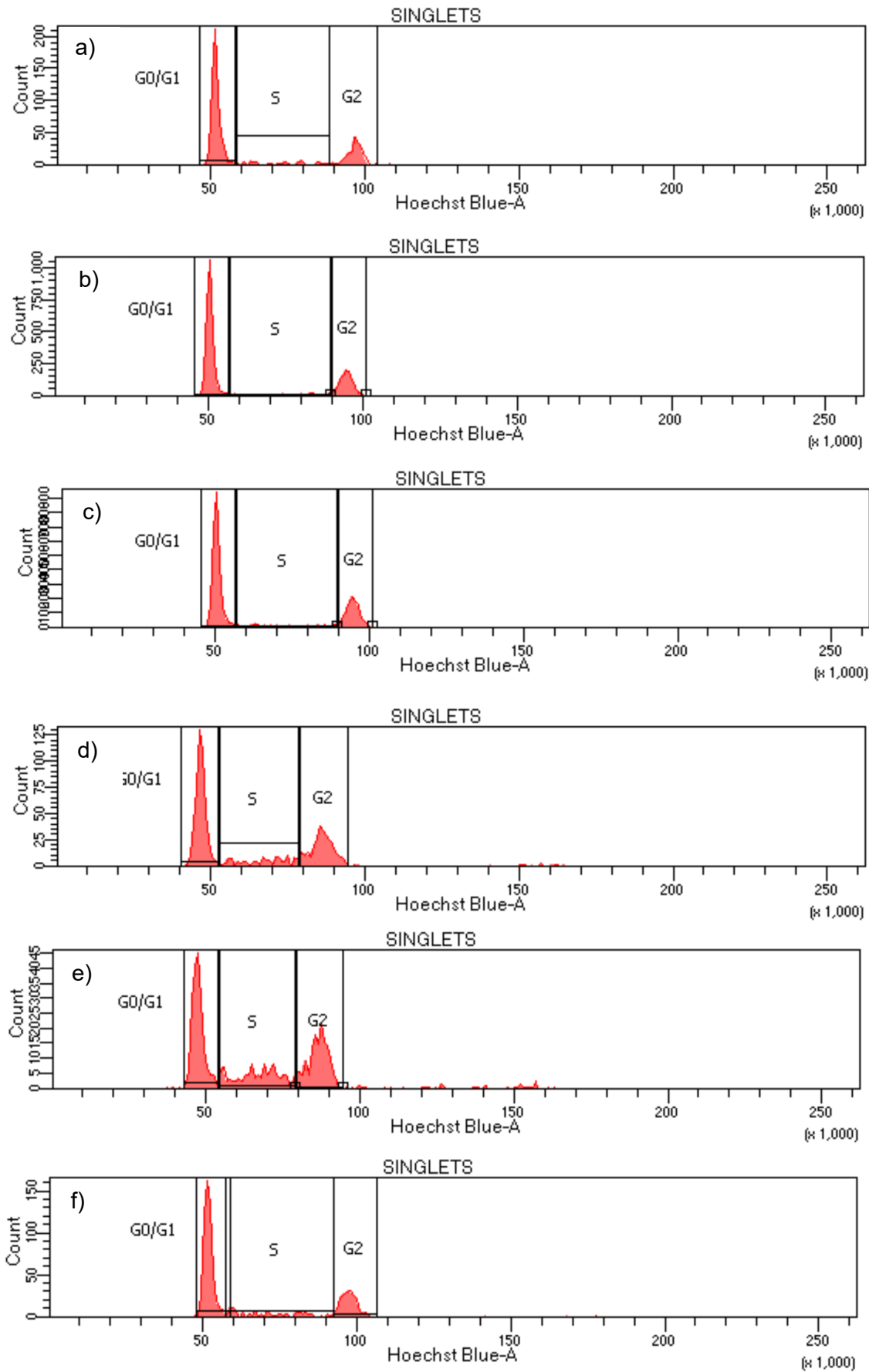


Figure S16 - HT1080-flow cytometry of cell cycle analysis after 24 hours of incubation with 0.3  $\mu\text{M}$  (DOX eq) of a) cPCP<sub>4/5</sub><sup>2</sup>, b) cPCP<sub>4/10</sub><sup>2</sup>, c) cPCP<sub>2/20</sub>, d) cPCP<sub>s</sub>, e) free DOX and f) control.

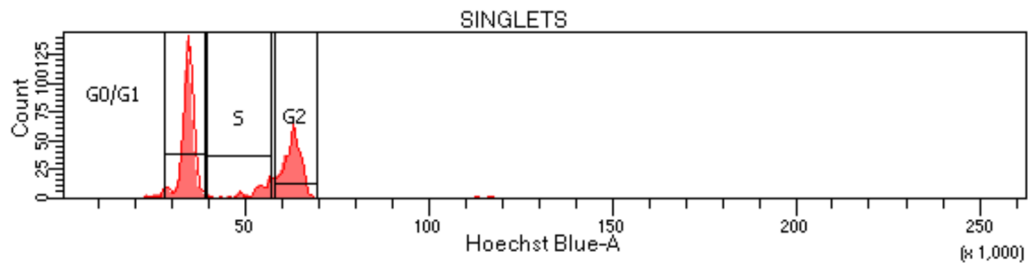
Table S2 - Cell cycle analysis of HT1080 cells after 48 hours of incubation with equimolar concentration (DOX eq) of free DOX, cPCP<sub>4/5</sub><sup>2</sup> and cPCP<sub>5</sub>.

<i>Product name</i>	<i>% G1</i>	<i>% S</i>	<i>% G2</i>
<i>CTR</i>	47.0 ± 1.7	25.3 ± 2.8	25.0 ± 1.7
<i>DOX</i>	29.2 ± 1.5	32.3 ± 4.4	33.6 ± 5.7
<i>cPCP<sub>4/5</sub><sup>2</sup></i>	50.2 ± 2.1	19.6 ± 10.4	30.2 ± 11.3
<i>cPCP<sub>5</sub></i>	35.45 ± 2.0	22.0 ± 4.9	39.9 ± 6.3

Table S3 - Cell cycle analysis of HT1080 cells after 72 hours of incubation with equimolar concentration (DOX eq) of free DOX, cPCP<sub>4/5</sub><sup>2</sup> and cPCP<sub>5</sub>.

<i>Product name</i>	<i>% G1</i>	<i>% S</i>	<i>% G2</i>
<i>CTR</i>	45.6 ± 0.5	28.5 ± 2.1	20.9 ± 2.0
<i>DOX</i>	32.6 ± 3.2	35.8 ± 6.7	27.3 ± 8.1
<i>cPCP<sub>4/5</sub><sup>2</sup></i>	46.8 ± 1.3	28.9 ± 3.4	18.6 ± 2.6
<i>cPCP<sub>5</sub></i>	38.1 ± 3.7	19.4 ± 11.2	39.3 ± 8.7

a)



b)

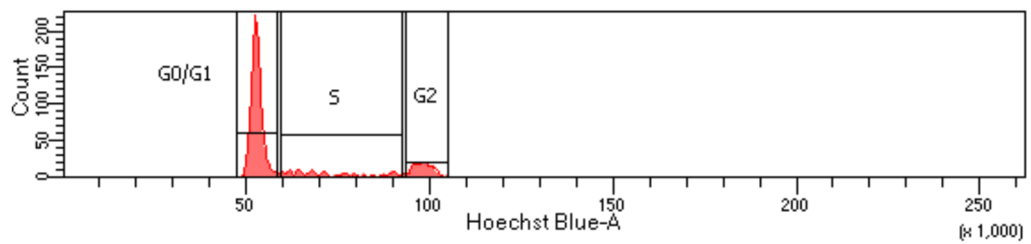


Figure S17 - HT1080 - flow cytometry of cell cycle analysis after 24 hours of incubation with A) 30 μM (DOX eq) of cPCP<sub>4/5</sub><sup>2</sup>, B) control cells.