



## **Supplementary Materials**

**Table S1.** IC50 values ( $\mu$ M or nM) for selected compounds obtained after 72 h treatment of 2008, C13\*, A2780, A2780/CP, and IGROV1 human ovarian cancer cell lines. Data indicate mean values and standard deviation from at least three experiments performed in duplicate.

Compounds	2008 cells	C13* cells	A2780 cells	A2780/CP cells	IGROV1 cells
5-FU	6.4±0.7	9.3±1	8.5±0.8	11.6±1.2	16.9±1
[DGln4]LR	6.2±0.3	9.1±0.8	8.7±0.6	10.2±0.5	5.2±0.4
cDDP	3.4±0.1	15.5±1.3	5.2±0.6	20.6±1.5	9.8±1.1
RTX (nM)	27.7±2.3	42.4±3.8	16.2±1.2	28.3±4.5	21.2±1.6
Pacli (nM)	37.4±4.6	52.2±8.3	9.2±0.8	19.4±1.2	22.5±3.3

**Table S2.** The effect of 72h-exposure to [DGln4]LR and 5-FU alone and in combination on the cell cycle phase distribution of 2008 and C13\* cell lines by cytofluorimetric analysis. The values are the mean of two experiments.

	2008 cells			C13* cells				
	$G_0G_1$	S	G2/M	hypodiploid	$G_0G_1$	S	G2/M	hypodiploid
Ctrl	75.19	9.78	10.83	1.25	67.58	11.71	11.62	0.37
[DGln4]LR	73.75	7.22	11.73	3.51	68.81	15.26	11.88	2.16
5FU 5-10uM	56.53	16.71	12.96	10.33	72.97	10.13	10.98	1.57
5FU 10-20uM	48.68	19.33	12.33	11.08	69.28	9.68	9.96	4.66
[DGln4]LR/ 5FU 5-10uM	53.22	13.93	8.05	19.07	69.62	11.05	11.33	2.78
[DGln4]LR/ 5FU 10-20uM	43.46	13.33	13.40	19.41	59.88	10.81	10.08	10.51

24 h after seeding, the cells were exposed to the drugs for 72h, then DNA content of untreated and treated cells was determined by flow cytometry after propidium iodide staining. Apoptotic cells are characterized by a lower DNA content (hypodiploid cells, having fewer than the diploid number of chromosomes) because of nuclear fragmentation.

	2008 cells				C13* cells				
48 hr	$G_0G_1$	S	G <sub>2</sub> /M	hypodiploid	$G_0G_1$	S	G2/M	Hypodiploid	
Ctrl	70.56	6.70	11.34	8.80	74.45	7.04	13.84	1.62	
[DGln4]LR	61.18	6.95	7.60	20.55	74.77	9.25	10.77	1.69	
cDDP	31.22	10.19	4.17	47.78	46.93	17.10	20.96	10.36	
RTX	24.98	5.51	2.19	63.21	43.01	9.67	8.18	32.68	
[DGln4]LR/ cDDP	23.18	9.67	2.87	58.72	29.52	15.31	30.54	20.33	
[DGln4]LR/ RTX	22.46	6.90	3.13	63.13	30.66	9.61	7.18	50.01	
	2008 cells					C13* cells			
72 hr	$G_0G_1$	S	G2/M	hypodiploid	$G_0G_1$	S	G2/M	hypodiploid	
Ctrl	62.90	9.96	14.37	2.56	62.61	13.30	11.87	8.72	
[DGln4]LR	65.01	9.75	11.18	3.01	56.94	9.80	8.14	7.01	
cDDP	31.24	8.82	9.61	41.87	17.73	12.18	45.83	18.44	
RTX	29.51	4.33	6.77	55.61	33.05	8.65	11.24	36.41	
[DGln4]LR/ cDDP	30.63	4.26	5.71	55.12	13.61	10.55	30.44	43.20	
[DGln4]LR/ RTX	24.05	2.54	4.51	66.61	37.87	6.34	6.66	48.98	

**Table 3.** The effect of 48h- and 72h-exposure to [DGln4]LR and cDDP alone and in combination on the cell cycle phase distribution of 2008 and C13\* cell lines by cytofluorimetric analysis. The values are the mean of two/three experiments.

24 h after seeding, the cells were exposed to the drugs or 48h or 72h, then DNA content of untreated and treated cells was determined by flow cytometry after propidium iodide staining. Apoptotic cells are characterized by a lower DNA content (hypodiploid cells, having fewer than the diploid number of chromosomes) because of nuclear fragmentation.



**Figure S1.** Cell cycle-related analysis of the 2008 cells after treatment with the indicated compounds. The variations of the cell distribution in the different phases of the cell cycle and especially in the sub-G0G1 area after the various treatments are shown. The results are representative of two/three independent assays.



**Figure S2.** Cell cycle-related analysis of the C13\* cells after treatment with the indicated compounds. The variations of the cell distribution in the different phases of the cell cycle and especially in the sub-G0G1 area after the various treatments are shown. The results are representative of two/three independent assays.



**Figure S3.** The effect of  $[DGln^4]LR$  and cDDP alone and in combination on the cell cycle phase distribution of 2008 and C13\* cells by cytofluorimetric analysis of the DNA content by PI staining. After 48-exposure to 5  $\mu$ M [DGln<sup>4</sup>]LR and 5 $\mu$ M (2008 cells) or 10 $\mu$ M (C13\* cells) cDDP, RTX 20nM alone and in concurrent combinations, cells were processed according to materials and methods. Inserted numbers indicate the % of cells in the different phases of the cell cycle. The values are the mean of two/three experiments. The error bars are omitted for a clearer visualization.