



## Supplemental Material

# Liposomal formulations of a new zinc(II) complex exhibiting high therapeutic potential in a murine colon cancer model

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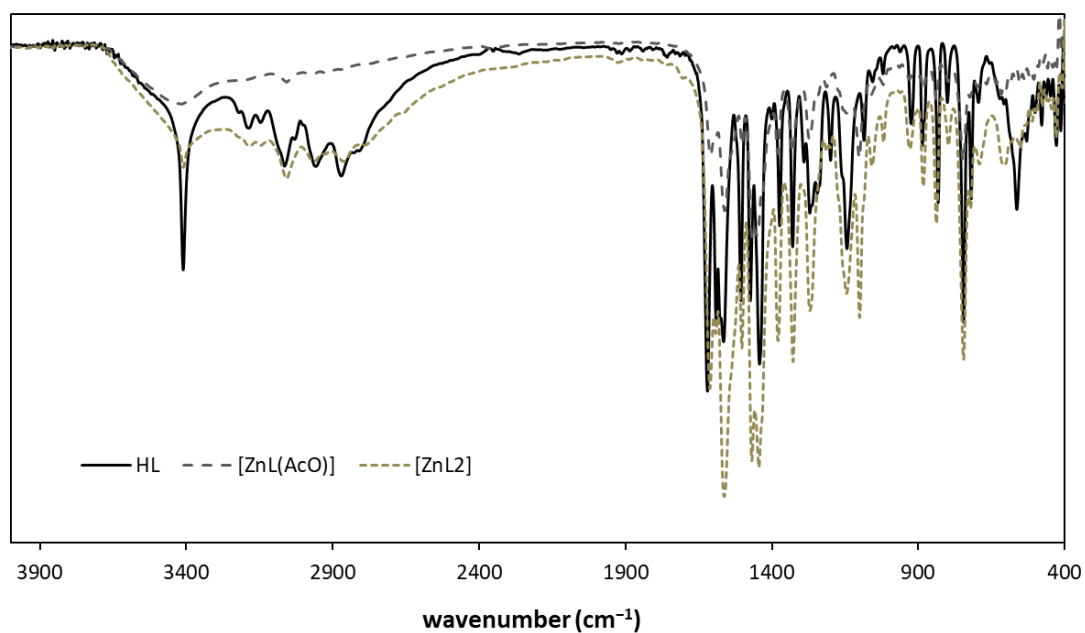
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## S1 - Characterization of the Zn(II) complexes

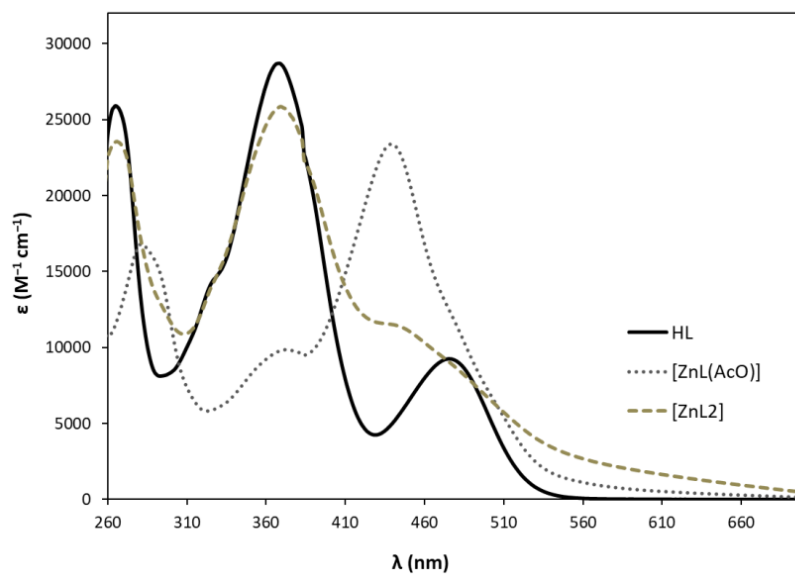
Despite several attempts, it was not possible to obtain single crystals suitable for X-ray diffraction. Thus, spectroscopic features of the coordinating groups were used to complement the elemental analysis and mass spectrometry results in order to provide a description of the Zn(II) complexes synthesized in this work. The main results are summarized in Table S1, while Figures S1, S2 and S3 show the spectra for FTIR, UV-Vis absorption and <sup>1</sup>H NMR, respectively.

**Table S1.** Hydroxyl and imine functions FTIR (in cm<sup>-1</sup>) and NMR signals (in ppm) in the ligand precursor and Zn(II) complexes.

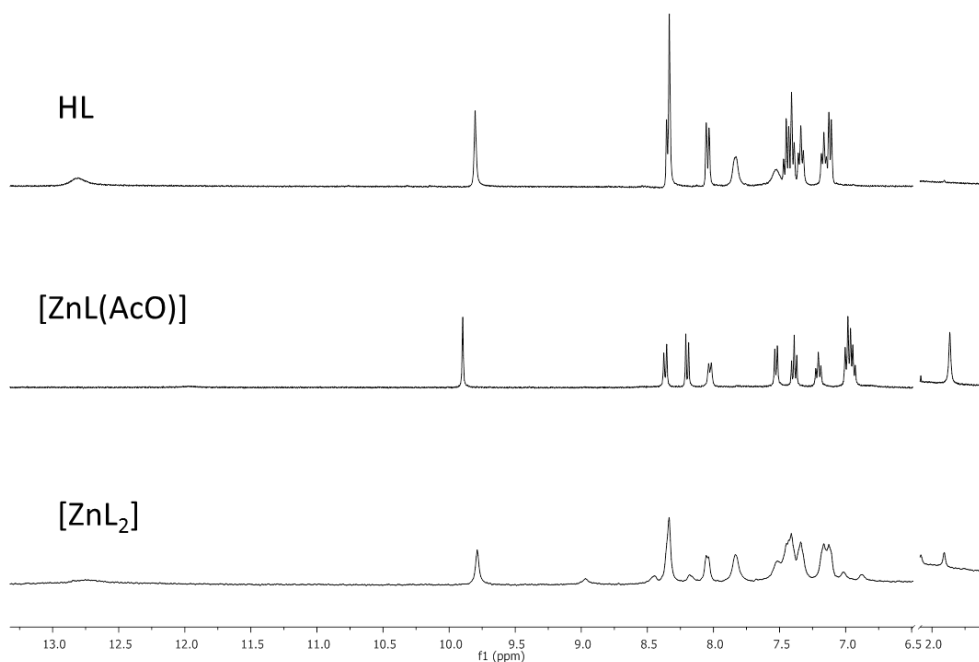
	$\nu_{\text{OH}}$ (cm <sup>-1</sup> )	$\nu_{\text{C=N}}$ (cm <sup>-1</sup> )	$\delta_{\text{H}}^{\text{OH}}$ (ppm)	$\delta_{\text{H}}^{\text{C=N}}$ (ppm)	$\delta_{\text{C}}^{\text{C=N}}$ (ppm)
HL	3411	1621	9.82	8.34	144.1
[ZnL(AcO)]	—	1610	—	9.90	144.9
[ZnL <sub>2</sub> ]	3411	1561	9.79	8.34	143.6



**Figure S1.** Infra-Red spectra of HL, [ZnL(AcO)] and [ZnL<sub>2</sub>] obtained as KBr pellets.



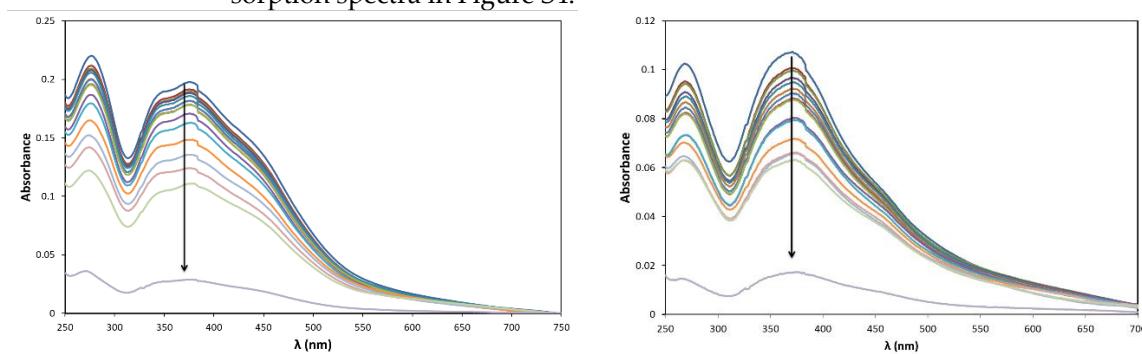
**Figure S2.** UV-Vis absorption spectra of HL, [ZnL(AcO)] and [ZnL<sub>2</sub>] in DMSO at *ca.* 35  $\mu$ M.



**Figure S3.**  $^1\text{H}$  NMR spectra of HL,  $[\text{ZnL}(\text{AcO})]$  and  $[\text{ZnL}_2]$  in  $\text{DMSO-}d_6$  in a 400 MHz apparatus.

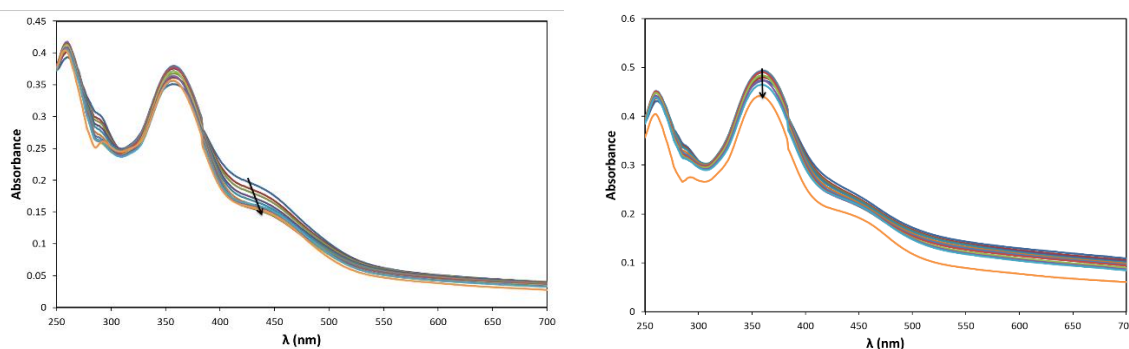
## S2 - Studies of interactions with BSA

A compound to be used as a drug has to possess water solubility and stability to some extent. The zinc complexes synthesized in this work present very low water solubility and, furthermore, even when dissolved in an organic solvent, they start to precipitate when in contact with an aqueous medium. This effect can be observed in the UV-Vis absorption spectra in Figure S4.



**Figure S4.** UV-Vis spectrum of a solution  $[\text{ZnL}(\text{AcO})] = 28 \mu\text{M}$  (left) and  $[\text{ZnL}_2] = 14 \mu\text{M}$  (right), in 0.6% DMSO/ PBS buffer (pH 7.4; 0.01 M phosphate; 0.138 M NaCl; 0.0027 M KCl); arrows show the changes occurring during 24 h of experiment.

A way to circumvent this effect is to protect them in a hydrophobic environment. When traveling inside a living organism, several of these exogenous compounds bind to carrier proteins, such as albumins. In fact, as we can see in Figure S5, when the complexes are bound to BSA there is a substantial increase in their stability and solubility in the aqueous medium.



**Figure S5.** UV-Vis spectrum of a solution  $[\text{ZnL}(\text{AcO})] = 24 \mu\text{M}$  (left) and  $[\text{ZnL}_2] = 22 \mu\text{M}$  (right) with equimolar concentration of BSA, in 0.6% DMSO/ PBS buffer (pH 7.4; 0.01 M phosphate; 0.138 M NaCl; 0.0027 M KCl); arrows show the changes occurring during 24 h of experiment.

Spectrophotometric techniques are easy and accessible methods to perform interaction studies with proteins and calculated the binding constant. Bovine serum albumin (BSA) exhibits fluorescence emission due to the presence of tryptophan (Trp) residues in its structure. When a substance interacts with BSA in a way that affects the Trp surroundings, a decrease in the intensity of the fluorescence emission is observed, in a process named quenching. The process can be analysed with the Stern-Volmer equation, which is related to the efficiency of the quenching process, providing clues on the type of involved phenomenon [1]:

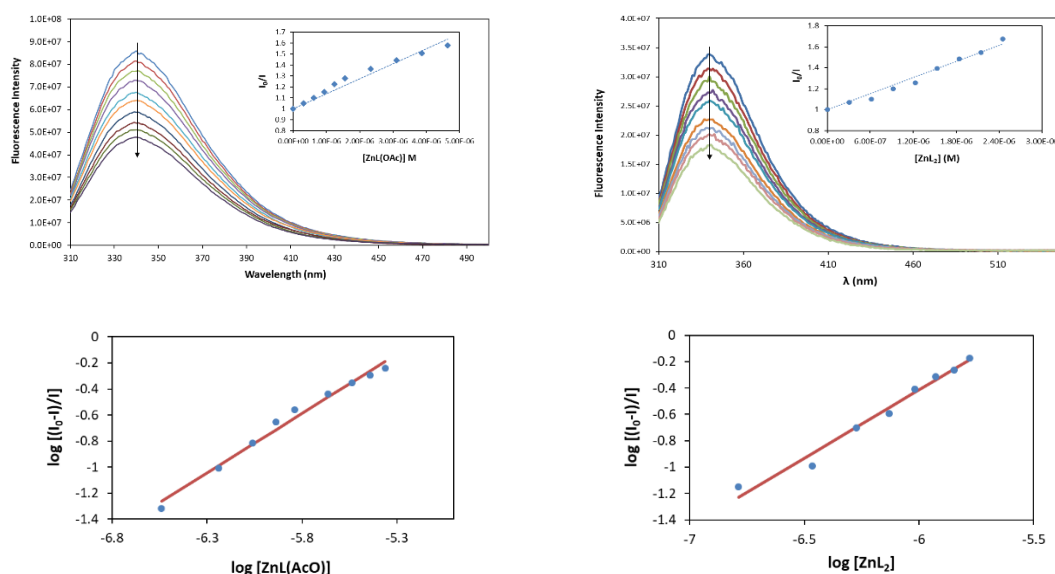
$$I_0/I = 1 + k_q\tau_0[Q] = 1 + K_{SV}[Q]$$

here  $I_0$  and  $I$  are the fluorescence intensities in the absence and presence of quencher, respectively, corrected for reabsorption and inner filter effects [2,3],  $k_q$  is the bimolecular quenching constant,  $\tau_0$  is the fluorophore average fluorescence lifetime,  $[Q]$  is the total quencher concentration and  $K_{SV}$  is the Stern-Volmer quenching constant. If the quenching process is solely due to diffusive encounters (collisions) between the compound and albumin, then  $K_{SV}$  is the dynamic constant ( $K_D$ ) and  $k_q$  should have a value lower than the limiting diffusion rate in water ( $3 \times 10^{10} \text{ M}^{-1} \text{ s}^{-1}$ ) [4]. The obtained Stern-Volmer constants are  $1.7 \times 10^5 \text{ M}^{-1}$  for  $[\text{ZnL}(\text{AcO})]$  and  $2.6 \times 10^5 \text{ M}^{-1}$  for  $[\text{ZnL}_2]$ . Since the Trp emission in albumins has  $\tau_0$  in the nanoseconds range, this means that the calculated  $k_q$  are above the collisional limit and, therefore, the quenching process has a static component, from the formation of a non-fluorescent ground complex between the compounds and albumin.

When small molecules bind to BSA, the association process can be analysed with the Scatchard equation [5]:

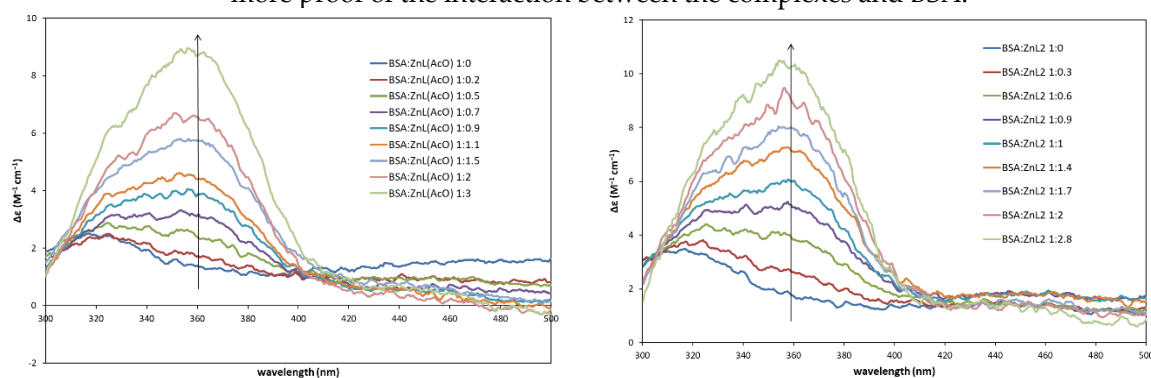
$$\log[(I_0 - I)/I] = \log K + n \log[Q]_f$$

where  $I_0$  and  $I$  have the same meaning as in the Stern-Volmer equation,  $K$  is the association constant between the two species,  $n$  is the number of binding sites and  $[Q]_f$  is the concentration of the free quencher [6]. Application of this analysis to both systems yields an association constant of  $5.0 \times 10^4 \text{ M}^{-1}$  for the 1:1 complex, while  $[\text{ZnL}_2]$  shows a constant more than an order of magnitude higher,  $K = 6.4 \times 10^5 \text{ M}^{-1}$ . In both cases one binding site is obtained, with  $n = 0.91$  for  $[\text{ZnL}(\text{AcO})]$  and 1.04 for  $[\text{ZnL}_2]$ . These results are in good agreement with previously published data for 8-hydroxyquinoline derivatives [7], and other metal complexes bearing hydrazone ligands [8]. Association constants in the range  $10^4 - 10^5 \text{ M}^{-1}$  are considered to represent moderate to strong binders, which may be delivered by the protein to their biological targets.



**Figure S6.** Top: Fluorescence emission of BSA (1.6 μM) in the presence of [ZnL(AcO)] (0 – 1.6 μM) and in the presence of [ZnL<sub>2</sub>] (0 – 2.5 μM) with the Stern-Volmer plots as insets. Down: Scatchard plots corresponding to the above titrations.

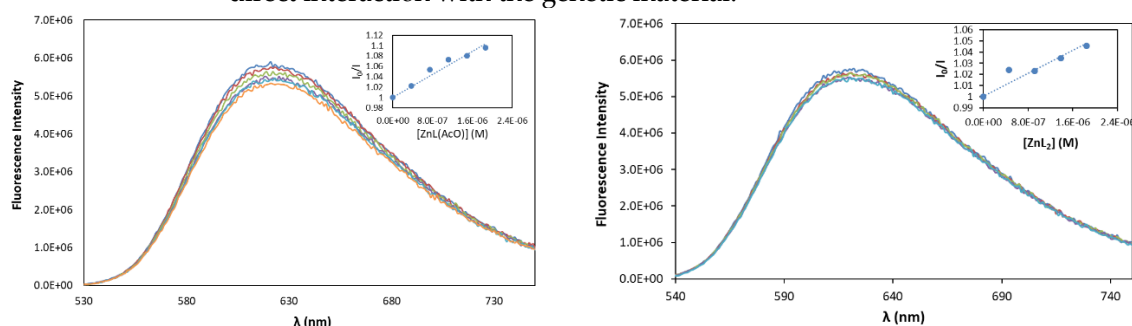
Another useful spectroscopic technique to study these interactions is circular dichroism (CD) that may be used to detect protein conformational changes or the development of induced CD signals [9,10]. Although limited by its low sensitivity and requirement of relatively high amounts of chiral species, induced circular dichroism (ICD) signals provide valuable information on the interaction process. BSA is chiral but does not possess absorption bands above 300 nm (it is a non-absorbing host in this spectral range), while the complexes present absorption bands in the region 310 – 600 nm but they are the non-chiral (absorbing guest molecules). Upon binding, the symmetry of the non-chiral guest is perturbed by the chiral environment of the host resulting in the generation of non-zero rotational strengths [11]. Titrations of a fixed concentration of BSA (24 μM) with increasing amounts of each complex, revealed the appearance of ICD signals providing once more proof of the interaction between the complexes and BSA.



**Figure S7.** Titration of BSA (24 μM) with [ZnL(AcO)] (0 – 74 μM) and with [ZnL<sub>2</sub>] (0 – 68 μM) followed by circular dichroism spectroscopy showing the appearance of ICD bands in the absorption region of the complexes. The experiments were done in PBS buffer, with a maximum content of 2% (v/v) DMSO; the induced CD spectra were obtained by subtracting the BSA spectrum measured under the same conditions.

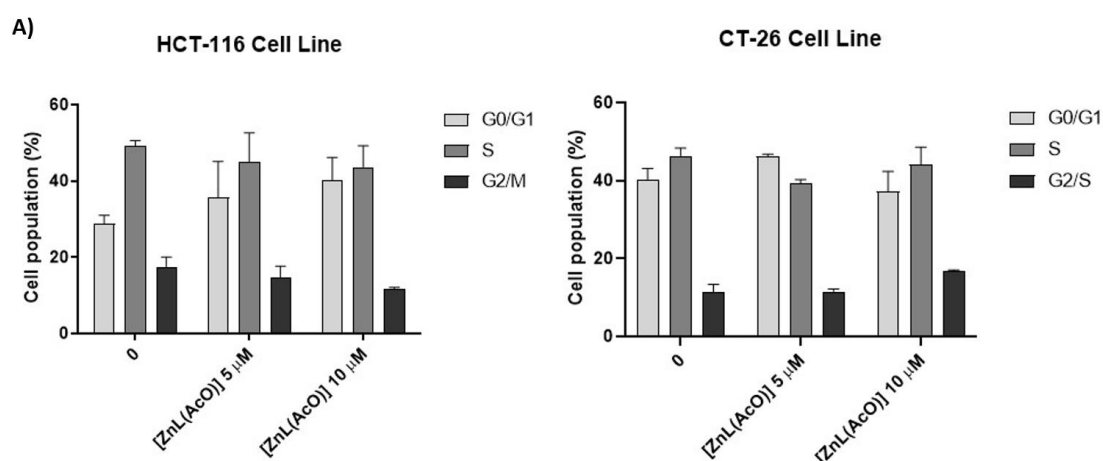
### S3 - Studies of interactions with ctDNA

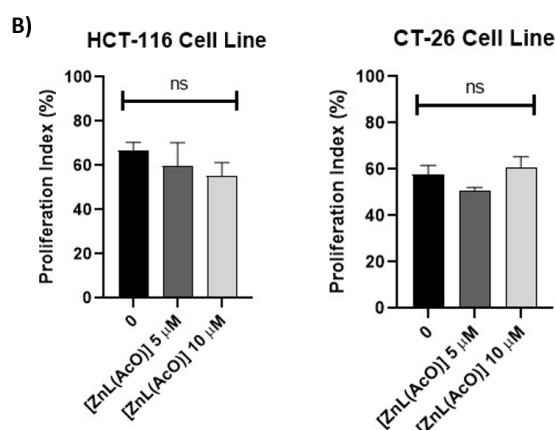
The evaluation of the interaction of each complex with the genetic material (DNA) was also done using spectroscopic techniques, as used for BSA. However, no induced circular dichroism bands were observed, and the complexes did not show ability to displace the classical intercalator ethidium bromide (Figure S8). Additionally, the very poor solubility in water precluded the use of UV-Vis absorption titrations. Overall, the set of results obtained with DNA makes us believe in a cell death mechanism that does not include a direct interaction with the genetic material.



**Figure S8.** Titration of the system EB:DNA (1.85  $\mu$ M: 2.27  $\mu$ M) with  $[\text{ZnL}(\text{AcO})]$  (0 – 3.7  $\mu$ M, left) and with  $[\text{ZnL}_2]$  (0 – 2.3  $\mu$ M, right) followed by fluorescence emission spectroscopy, using  $\lambda_{\text{ex}} = 510$  nm and slits 5/5 nm. The experiments were done in HEPES buffer, pH 7.4 (0.1 M HEPES; 0.15 M KCl), with a maximum content of 0.8% DMSO (p/v).

### S4 – Cell death and cell cycle analysis





**Figure S9.** A) Percentage of HCT-116 and CT-26 cells cultured in a 2D setting in G0/G1, S, and G2/M cell cycle phases after a 24 h incubation with then Zn-complexes at 5 and 10  $\mu$ M and with complete medium (0), obtained by flow cytometry analysis. B) Proliferation indexes for the complexes determined according to cell cycle distribution of exponentially growing populations of HCT-116 and CT-26 as the sum of percentage of cells in S and G2/M phases. Results correspond to the cell population expressed as mean percentage (%)  $\pm$  SD of three independent experiments. Ns: non-significant.

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