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New Findings in the Signaling Pathways of cis and trans Platinum Iodido Complexes' Interaction with DNA of Cancer Cells

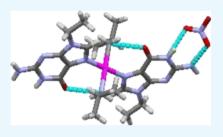
Adoración G. Quiroga,*^{,†}[©] Marta Cama,^{†,‡} Natalia Pajuelo-Lozano,[‡] Amparo Álvarez-Valdés,[†] and Isabel Sanchez Perez*,*

[†]Inorganic Chemistry Department, Universidad Autónoma de Madrid and IAdChem Universidad Autónoma de Madrid, Madrid 28049, Spain

[‡]Biochemistry Department, Faculty of Medicine, Instituto de Investigaciones Biomédicas Alberto Sols. CSIC-UAM, Madrid 28029, Spain

Supporting Information

ABSTRACT: We have selected a series of aliphatic amine platinum compounds bearing chloride and/or iodide as the leaving groups. The complexes' cytotoxicity and interaction with DNA indicated differences in the reactivity. Now, we are reporting on the analysis of their molecular mechanism of action on gastric cancer cells. Our data reveals differences between them. Chlorido drugs showed similar behavior to cisplatin; they both required p53 to induce apoptosis but only cis-ipa showed DNA damage requirement for apoptosis induction. On the contrary, cis and trans iodido induced cell death independent of p53 activity, and they induced cell death through Bid activation, so their toxicity could be enhanced in a combined treatment with novel



Bcl-2 protein family inhibitors. We also report the structural features of the DNA adduct for one of the complexes by X-ray diffraction. These findings represent a step forward in the search for new platinum-derived agents more specific and effective in the treatment of cancer.

1. INTRODUCTION

Cisplatin is a clinical drug used for the treatment of solid tumors; however, cisplatin can produce relevant disadvantages that are difficult to overcome, as for example the cellular development of resistance (intrinsic or acquired). In addition, treatment with cisplatin manifests systemic side effects, such as neurotoxicity among others. Therefore, obtaining metal derivatives of platinum that are equal or more effective than cisplatin will be of great benefit in the cure of the disease.

As we reported on the iodido complexes' potential antitumoral action possibly caused by their peculiar reactivity toward biological targets,¹ there has been a new research trend with reports trying to better understand iodido complexes' activity. The reevaluation of cis and trans iodido derivatives' cytotoxicity showed their efficiency and their interaction study versus model proteins indicate a possible different mechanism of action.² Further work proved this protein interaction of these complexes by X-ray diffraction.³

As DNA is cisplatin's main target, the studies of reevaluated iodide complexes with DNA are an important step to take into consideration. The basic models (cisplatin and its iodido analogues) were studied using oligo deoxyribo nucleotides as DNA models, indicating the same type of interaction toward guanine (leading to adducts containing $(Pt(NH_3)_2^{2+})$ with the iodido derivative always being more reactive than cisplatin.⁴ Another example is the work performed by Dvořák et al., varying the amine ligand and substituting for bulkier azaindoles.5 They studied the cytotoxic activity of iodide azaindole complexes and looked at the molecular level mechanism to find a decrease of tumor suppressor p53 amount, which can agree with our previous observations with the aliphatic amine iodido compounds.⁶

On putting this information together, it is clear that iodide complexes interact with DNA, but we need more data about the possible differences at molecular level and in particular, about the signaling pathways, to deeply analyze a broader spectrum of interactions before and after DNA damage. For this work, we will be using gastric cancer cell lines, with which we have recently reported the molecular processes involved in cisplatin-induced apoptosis.

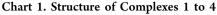
2. RESULTS

We selected some of the reported structures, and we synthesized and characterized the following series of compounds: cis-[Pt(ipa)₂Cl₂], 1, trans-[Pt(dma)(ipa)Cl₂], 2, cis-[Pt(ipa)₂I₂], 3, and trans-[Pt(ipa)₂I₂], 4 (depicted in Chart 1). All data collected with these complexes agreed with those previously reported,^{1,6,8} therefore we performed the following assays.

We began analyzing the viability in MNK45 gastric cancer cells after treatment with the selected series of platinum compounds and using cisplatin (CDDP) as a reference. The results are shown in Figure 1 we observed that the survival decreased in a dose-dependence manner with the following

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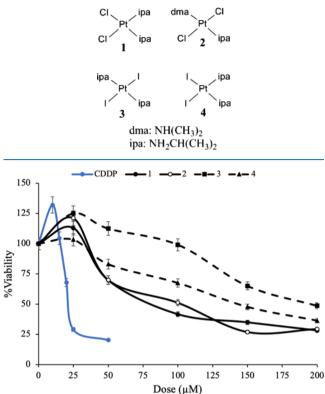


Figure 1. Cell viability results of cisplatin and complexes 1 to 4 for the MNK45 gastric cancer cells.

IC₅₀: CDDP (20 μ M), 1(80 μ M), 2 (100 μ M), 3 (180 μ M), and 4 (140 μ M). The compounds are less cytotoxic than cisplatin, but these complexes show good values in other type of cell lines, in particular, those which are not dependent on p53.⁶ Therefore, we analyzed the compound's effect in the HCT116 wild type (WT) and HCT116 p53 knock out (KO) cell lines.

Table 1 shows the IC_{50} values from HTC-116 cells (HCT-116 WT and p53KO) after treatment with the compounds.

Table 1. IC ₅₀ for Complexes 1–4						
	WT IC ₅₀ (μ M)	p53 ^{-/-} IC ₅₀ (µM)				
CDDP	15.34 ± 2.17	48.75 ± 4.80				
1	91.20 ± 9.90	172.18 ± 12.24				
2	66.37 ± 3.40	133.35 ± 15.21				
3	55.71 ± 3.22	85.50 ± 10.73				
4	28.64 ± 2.90	66.07 ± 5.38				

Our results show that in the absence of p53, HTC116 are more resistant to CDDP than the WT HCT116 cells. The treatment with compounds 1 and 2 shows an increase in the IC_{50} when cells lack p53. In contrast, cells treated with compounds 3 and 4 do not exhibit the big difference in viability that is shown by 1 and 2 independently of the genotype (Figure 2). Although compounds 3 and 4 show a difference within the range of cisplatin, these data have to be evaluated to find if the contribution of p53 in the cell death is partial and/or other pathways are involved in the process.

The cytotoxicity result in HTC-116 encouraged us to evaluate p53 activation in MNK45 after treatment with the

different compounds and CDDP as a control. To do that, we analyzed the phosphorylation of Ser15 in p53, which is indicative of activation by WB. The results indicate that p53 is activated in response to cisplatin and compound **2** in a persistent manner up to 48 h. In contrast, the treatment with compound **1** induced p53 phosphorylation, transitory with a maximum peak at 24 h (Figure 3). Surprisingly (however), no activation of p53 was observed after treatment with compound **3** and **4**.

Next, we analyzed the molecular markers that indicate apoptosis induction. We observed the activation of caspase 3 with the whole series of compounds. To corroborate these results, we analyzed the proteolysis of PARP (specific substrate of caspase 3) that shows a clear proteolysis, according to caspase 3 activation (Figure 4). This effect is also observed for cisplatin, as previously reported. In the end, apoptosis induction is corroborated.

CDDP activates apoptosis mainly through the intrinsic pathway, and then we explored the intrinsic apoptosis pathway activation by the compounds. First, we study the protein levels of different members of BcL2 family. We observed induction of the proapoptotic protein, Puma after treatment with CDDP and 1 but not 2. In fact compound 2 does not show differences at any time after treatment. In contrast, Puma protein is not induced after exposure to 3 and 4. The pro-apoptotic protein, Bax increases in response to CDDP and 1, whereas compound 2 slightly induces Bax. The iodide complex 3 instead provoked Bax induction, as we observed a small but clear level decrease at 48 h. Complex 4 does not modify Bax expression (Figure 5).

We observed induction of the proapoptotic protein, Puma after treatment with CDDP, 1, and 2 (Figure 5). However, 3 and 4 do not induce Puma. P53 contributes to Puma induction; therefore, this result is in agreement with the p53 levels previously observed (Figure 3). The pro-apoptotic protein, Bax increases in response to CDDP and 1, whereas compound 2 slightly induces Bax. The iodide complexes (3 and 4) do not modify Bax expression (Figure 5).

BID1 protein belongs to BH3-only Bcl2 subfamily; it is a cytoplasmic protein that once proteolyzed, translocates to the mitochondria membrane binding to BAX, which allows Cyt C release and caspase 9 and 3 activation. Then, BID1 is activated by CDDP, **1**, and **2** after 6 h of treatment; however, **3** and **4** provoke only a transitory activation, with a highest point after 6 h.

Finally, the antiapoptotic Mcl1 expression analysis after 6 h of treatment shows that CDDP produces clear degradation as it is described in the literature after 48 h.⁷ Compound 1 shows the same profile. No degradation is observed neither with 2 nor with 4, but a surprisingly marked degradation is observed with 3 h treatment.

It is known that JNK and p38 could phosphorylate MCL1 and induce its degradation.⁹⁻¹¹ We studied the MAKPs activation after treatment with the compounds; the results show a transient activation of p38 when treated with CDDP, 1, and 2, and a persistent activation with 3 and 4 (Figure 6). Only 2 and 3 show clear but transitory activation of JNK with the highest peak after 6 h. The entire compound series activates ERK, showing similar profile with the exception of 2 (which activates only in a transitory way). These results may indicate that MCL1 degradation is produced by p38 phosphorylation.

DNA is the main cellular target for platinum complexes, which leads to the activation of different checkpoints in the cell cycle. We then analyzed the cell cycle profile after treatment

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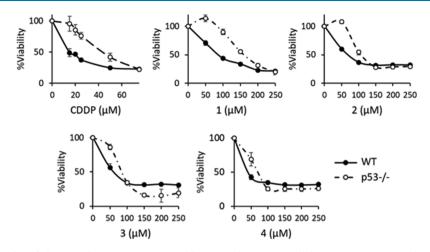


Figure 2. Cell viability results of the complexes in HCT116 wild type and p53 KO cell lines. CDDP was used as a reference. Cell survival percentage of HCT166^{WT} and HCT116 p53^{-/-} cells after 72 h of CDDP or compounds treatment. Cells were treated with increasing concentrations of CDDP ($0-75 \mu$ M) or compounds ($0-250 \mu$ M) The percentage of viable cells was quantified by the MTS assay. Data represent the mean values obtained in three experiments performed in quadruplicate.

MW (Kd)	CDDP 1	1	2	CDDP	3	4		
53					-	17 Aug	•	p-p53 (Ser15)
52							←	α-Tubulin
Time (h)	0 6 24 48 6 2	4 48	6 24 48	0 6 24 48 6	5 24 48	6 24 48		

Figure 3. Complex 1 and 2 induces p53 activation in MKN45 cells. Cells were treated with the complexes CDDP, 1, 2, 3, and 4 at IC50 dose for each compounds, and harvested at the indicated times. P53 total levels and phosphorylated were detected by Western blot using specific antibodies. Tubulin was used as a control.

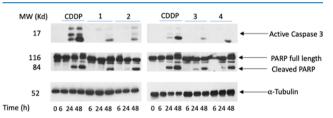


Figure 4. Caspase-3 activation and PARP proteolysis is detected after treatment with CDDP, 1, 2, 3, and 4 in MKN45 cells. Cells were treated as in Figure 3. Caspase 3 and PARP were detected by using specific antibodies.

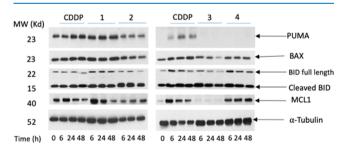


Figure 5. Bcl-2 protein family members analyzed by western blot from MKN45 cells treated with CDDP, 1, 2, 3, and 4 as in Figure 3. Proteins Puma, Bax, Bid, and MCL1 were detected by using specific antibodies. Tubulin was used as the control.

with the drugs by flow cytometry on the cancer cells. The results showed an increase in phase S (54.2%) for the cells treated with CDDP, which is indicative of intra S checkpoint activation. Compound 1 provokes an increase of cells in phase S (29%) and phase G2/M (31.3%), whereas the rest of the

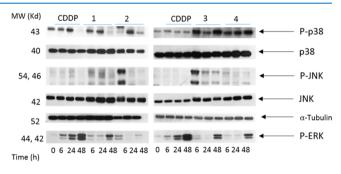


Figure 6. MAP Kinase members analyzed by western blot from MKN45 cells treated with CDDP, 1, 2, 3, and 4.

series produced such increase at the G2/M checkpoint (34.5% 2; 31.6%: 3; and 39.8%: 4) (Figure 7, Table 2).

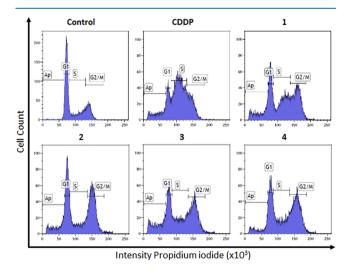


Figure 7. Cell cycle profile of MKN45 cells after treatment with the complexes. DNA content was assessed by flow cytometry, and cell cycle distribution was analyzed using Cell Quest Pro software. Table 2 shows the percentage of each cell phase, which is given as the mean of three experiments. G2/M: cells in G2 or mitosis; S: cells in phase of synthesis of DNA; G1: cells in G1, apoptosis.

 Table 2. Percentage of Cells in Each Cell Cycle Phase after

 Treatment with the Drugs

	apoptosis	G1	S	G2/M
control	2.8	55.6	21.5	19.4
CDDP	12.4	14	54.2	18.9
1	15.5	22.7	29.7	31.3
2	15.5	30.2	17.5	34.5
3	21.3	20.8	24.4	31.6
4	10	24.1	23.5	39.8

CDDP requires DNA damage during the S-phase to induce apoptosis. In order to know if the drugs share this requirement, we synchronized the cell in G1 by using Palbociclib (1 μ M) and then treated with the drugs (complexes 1 to 4). Our results showed that 1 and CDDP produce a lesion on phase S in order to induce cell death, whereas none of the compounds of the series causes such lesion. This result indicates that the apoptosis induction by 2, 3, and 4 could start at the phase G2/ M (Figure 8). As at the last step, we prove the reactivity of the complexes at the molecular level with a small model of DNA. The treatment of *trans*-[Pt(ipa)₂I₂] with two equivalents of AgNO₃ forced the aqua-complex formation which reacts with two equivalents of 9-EtG (9-ethyl guanine) to produce the complex *trans*-[Pt(9-EtG)₂(ipa)₂](NO₃)₂, where the iodide groups are replaced by the base.

The structure in Figure 9 shows a mononuclear adduct where the Pt atom is bound to two molecules of 9-EtG through N7 in a square planar geometry similarly to other examples of refs 12–14. Both 9-EtG molecules are coplanar, whereas isopropylamine ligands are 180° twisted giving the most stable conformation to the molecule. This conformation shows hydrogen bonding interaction between O₁ from the carbonyl group of 9EtG and hydrogen of the $-NH_2$ group from isopropylamine. This kind of interaction has been described to stabilize similar structures.^{15,16} The distance of 2.249 Å indicated strong intramolecular interaction which is the most significant data, as it might potentially hinder the 9-EtG ligands from rotating freely around Pt₁–N₂ when binding to any

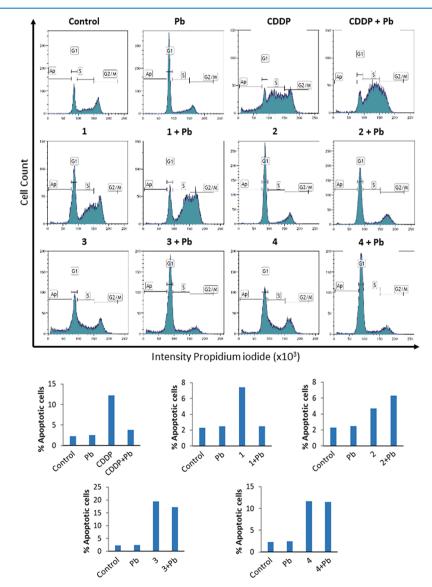


Figure 8. Apopstosis percentage of MKN45 cells after synchronizing with Palbociclib $(1 \ \mu M)$ for 18 h and the complexes. DNA content was assessed by flow cytometry, as in Figure 7. Results of the compounds in % synchronized culture cell by using Palbociclib. Ap: apoptosis.

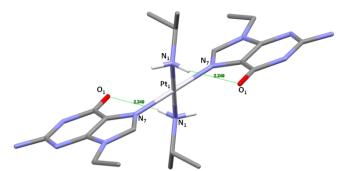


Figure 9. Structure of the complex *trans*-[Pt(9-EtG)₂(ipa)₂](NO₃)₂.

target.¹⁷ The crystal structure also shows intermolecular hydrogen bonds that involve the nitrate contraction linked by a series of NH O contacts. As can be seen in Figure 10, they are responsible for solid-state packaging.

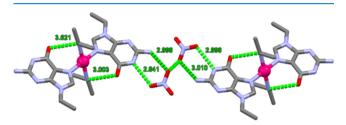


Figure 10. Molecules arrangement in complex *trans*-[Pt(9-EtG)₂(ipa)₂](NO₃)₂ crystal structure.

3. DISCUSSION

Based on the efficacy of CDDP (in spite of some of its side effects) and trying to find a clear signaling pathway for the iodide and chlorido compounds, we have analyzed the cytotoxic activity versus the gastric cancer cell line (MKN45) for four cis and trans selected complexes. The complexes show aliphatic amines in their structure and chloride and iodide as the leaving groups.

The compounds, together with CDDP, are capable to decrease the MKN45 cell survival, even though CDDP needs lower doses to produce the same percentage of cell death. It is well known that every cancer cell is different, and as we mentioned before, iodide complexes can be active in cancer cells from different tumors.² In fact, HCT116 cell lines results showed lower IC₅₀ values than those observed in MKN45.

p53 is a tumor suppressor protein which is mutated in more than the 50% of tumors; therefore, this finding is of a vital importance in the treatment of tumors p53 null. Mutation on the p53 gene is involved in CDDP resistance in cancer as a consequence of loss of the ability of p53 to transactivate Bax and Puma.^{18,19}

The bigger difference in the viability data of complexes 1 and 2 compared with cisplatin indicates a very clear p53 dependence, whereas complexes 3 and 4 did not show that difference. Moreover, complexes 1 and 2 induce apoptosis and p53 transcriptional dependence genes (Puma and Bax) as cisplatin does. However, the data indicate that the cell death produced by complexes 3 and 4 might only be dependent partially on p53 in spite of the 30 μ M difference in the genotype. Complexes 1 and 4 do not induce PUMA and BAX, which are the genes induced by 1, 2, and cisplatin.¹⁸ DNA is the target for chlorido and iodido complexes; based on the

differences observed in the reactivity, we thought that the iodido complexes could have some structural differences when binding to DNA. Nucleotides are of a common use in molecular biology studies, and they can also serve to achieve more information about the structural nature of the DNA adducts formed by platinum complexes, in particular 5'-GMP. In our case, we have used 9-EtG as a small DNA model base because this molecule is smaller than nucleotides or plasmidic DNA and more soluble in organic solvents than 5'-GMP, which is also a very convenient model to use for NMR monitoring studies. All these models of DNA help to evaluate the platinum compounds' reactivity and allow the final adducts characterization by different techniques and ideally by X-ray diffraction. The reactivity of trans-[Pt(ipa)₂I₂] with 9-EtG produces the formation of the adducts more rapidly than the cis counterpart,² leading to the final isolation of the bisadduct *trans*- $[Pt(9-EtG)_2(ipa)_2](NO_3)_2$ when forcing the conditions. In this article, we report the X-ray resolution of the final adduct, which supports the classical reactivity of the trans de Pt^{II}-I versus DNA hypothesized in our previous work.² We also report hydrogen bonding that can prevent the rotation of the molecule around the Pt1-N2 bond and be the direct structural feature produced by the platinum complex in the DNA molecule.

The analysis of the intrinsic apoptotic pathway induced by the compounds has revealed some important differences; compounds 1 and 3 generate an intrinsic apoptosis pathway similarly to CDDP inducing mitochondria perturbation. However, compound 4 seems to follow a different profile, and possibly more pathways are involved in its action. One possibility might be that compound 4 could induce cell death by the extrinsic pathway or by related receptors. Possibly the activation of caspase 8 will support this hypothesis and further studies will be carried out to explore this possibility. Therefore, compound 4 toxicity could be enhanced in a combined treatment with novel MCL1 inhibitors under development.² We have also demonstrated that compound 2, 3, and 4 induce cell death independently of replicative stress. Apoptosis is activated from G2/M, differently from CDDP (whose welldefined target is DNA), but cell death is induced by replicative stress along the S phase.

Compound **2** shows in general a midway behavior between CDDP and iodide complexes, and in particular the fact that JNK activates only in a transitory way could be related with a higher repair capacity of the DNA lesion.

Figure 11 shows a summary of the apoptotic pathway induced by the drug. We propose that iodide drugs increased cell death independent of p53, which is very promising for tumors with p53 mutated. Complex 2 could be used in combination with other drugs as a mimic of BH-only proteins to increase the activity. In summary, we have demonstrated that the synthetic variations on platinum complexes can modulate the interaction with these targets and the use of that information is very relevant for pharmaceutical purposes.

4. MATERIALS AND METHODS

4.1. Cell Lines. MKN45 and HCT116 human gastric adenocarcinoma cell lines were acquired from ATCC and cultured in RPMI medium or McCoy's 5a medium modified (Gibco), complemented with 10% FBS. Cultures were kept at 37 °C, 5% CO₂, and 95% humidity. MKN45 cells are wild type for TP53. Mycoplasma contamination tests are frequently run in our laboratory.

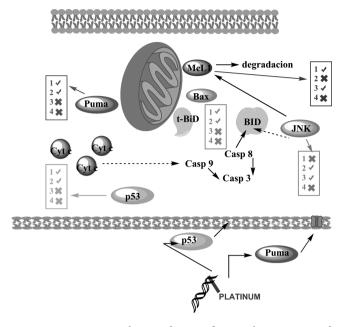


Figure 11. Apoptotic pathway and targets for complexes 1, 2, 3, and 4.

4.2. Chemicals. CDDP was kindly supplied by Ferrer FARMA. The compounds used for the assays (1, 2, 3, and 4) were synthesized following reported procedures, briefly described in the Supporting Information Section S1.²⁰ The solution for the MTS assay: CellTiter 96 AQueous One Solution Cell Proliferation Assay was acquired from Promega.

4.3. Reaction of trans-[Pt(ipa)2l2] with 9-EtG: trans- $[Pt(9-EtG)_2(ipa)_2](NO_3)_2$. A solution of AgNO₃ (16.5 mg) 0.097 mmol) in 0.2 mL of H₂O was added to a solution of complex 3 (27 mg, 0.048 mmol) in acetone and stirred at room temperature in the dark for 10 min. The mixture was filtered and the resulting solution mixed with a freshly prepared solution of 9 EtG (17.3 mg, 0.097 mmol) in 0.3 mL of a solution of acetone/H₂O (2:1), stirred at room temperature for 24 h, and allowed to stand until single crystals were formed and isolated. The final compound *trans*- $[Pt(9-EtG)_2(ipa)_2]$ - $(NO_3)_2$ was shiny white. Anal. Calc. for $PtC_{20}H_{36}N_{14}O_8 H_2O$: C, 29.50; H, 4.70; N, 24.64%. Found: C, 29.22; H, 5.00; N, 24.48. The structural features of *trans*- $[Pt(9-EtG)_2(ipa)_2]$ - $(NO_3)_2$ were unambiguously proven by X-ray diffraction. Data collection was performed on a Bruker Kappa Apex II (X8 APEX, Bruker, Rivas-Vaciamadrid, Madrid, Spain) area detector X-ray diffractometer using a graphite-monochromated Mo radiation. Crystal data are listed in Table SM1. Selected distances and angles are collected in Table SM2.

4.4. Cell Viability. Cell viability was assessed using a crystal violet-based staining method. The cells (3.5×10^5) were seeded in 0.5 mL of completed medium in 24 multiwell dishes, treated with different amounts of CDDP, Cl-1, Cl-3, I-5, and I-6 dissolved in the culture medium (RPMI) and fixed with 0.1% glutaraldehyde. After 30 min, they were washed with distilled water and stained with 0.1% crystal violet. A colorimetric ELISA-based assay set at 595 nm was used to estimate the number of cells per well. MTS protocol was performed according to the manufacturer's instructions. IC₅₀ were calculated by using the graph PAD prysme programme. We used nonlinear regression to fit the data to the log (inhibitor) versus response (variable slope).

4.5. Cell Cycle Analysis. MKN45 cells were seeded in 2.5 mL of RPMI 10% FBS in p60 plates and treated with solutions of the compounds following the established IC_{50} . The cells were fixed, 24 and 48 h after the treatment was applied, using 70% ethanol in phosphate-buffered saline (PBS). To perform the DNA content analysis, the cells were centrifuged, the supernatant was discarded and 200 μ l of propidium iodide was added, and the tubes were incubated for 15 min in darkness. Finally, 200 μ l of a solution of RNase in PBS (10 μ l/mL) was added to each tube. The cells were analyzed using a Becton Dickinson Flow Cytometer (Cowley, United Kingdom).

4.6. Western Blotting. The total protein extracts (WCE) were prepared using a lysis supplemented with phosphatase and protease inhibitors. WCE samples (20 μ g) were loaded in 15% (Bcl-2 family and caspase-3), 10% (JNK, p38 and ERK), and 8% (PARP-1, p53, MCL1) SDS-PAGE polyacrylamide gels. Once the electrophoresis was completed, they were transferred to polyvinylidene difluoride membranes and immunodetection was done using the correspondent antibodies. The following antibodies were purchased from Santa Cruz Technology: PARP-1 (sc-7150), Mcl-1 (sc-819), JNK (sc-827), p38 (sc-535), and ERK 1/2 (sc-154). The antibodies for Cleaved Caspase-3 (Asp175) (#9661), Bax, BID, PUMA (#12450), p-p38 (1:2000, #4631), p-p53 (Ser15) (#9284), and p53 (#9282) were acquired from Cell Signaling Technology. The antibody for p-JNK was bought from Promega Corporation, Spain. Last, the antibody for tubulin (1:10 000) was acquired from Sigma-Aldrich. The standard working dilution for the antibodies was 1:1000, unless a different one was specified above.

ASSOCIATED CONTENT

S Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acsomega.9b02831.

Synthesis and characterization of the platinum complexes, crystal data of trans- $[Pt(9-EtG)_2(ipa)_2](NO_3)_2$, and selected bond distances and angles for the structure of trans- $[Pt(9-EtG)_2(ipa)_2](NO_3)_2$ (PDF)

AUTHOR INFORMATION

Corresponding Authors

*E-mail: adoracion.gomez@uam.es (A.G.Q.). *E-mail: misanchez@iib.uam.es (I.S.P.).

ORCID 💿

Adoración G. Quiroga: 0000-0002-9261-9542 Isabel Sanchez Perez: 0000-0002-4829-201X

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Notes

The authors declare no competing financial interest.

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ABBREVIATIONS

CDDP, cisplatin; dma, dimethylamine; ipa, isopropylamine

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