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Linker design for the modular assembly of multifunctional and targeted platinum(II)-containing anticancer agents†

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Abstract

A versatile and efficient modular synthetic platform was developed for assembling multifunctional conjugates and targeted forms of platinum-(benz)acridines, a class of highly cytotoxic DNAtargeted hybrid agents. The synthetic strategy involved amide coupling beween succinyl estermodified platinum compounds (P1, P2) and a set of 11 biologically relevant primary and secondary amines (N1–N11). To demonstrate the feasibility and versatility of the approach, a structurally and functionally diverse range of amines was introduced. These include biologically active molecules, such as rucaparib (a PARP inhibitor), E/Z-endoxifen (an estrogen receptor antagonist), and a quinazoline-based thyrosine kinase inhibitor. Micro-scale reactions in Eppendorf tubes or on 96-well plates were used to screen for optimal coupling conditions in DMF solution with carbodiimide-, uronium-, and phosphonium-based compounds, as well as other common coupling reagents. Reactions with the phosphonium-based coupling reagent PyBOP produced the highest yields and gave the cleanest conversions. Furthermore, it was demonstrated that the chemistry can also be performed in aqueous media and is amenable to parallel synthesis based on multiple consecutive reactions in DMF in a "one-tube" format. In-line LC-MS was used to assess the stability of the conjugates in physiologically relevant buffers. Hydrolysis of the conjugates occurs at the ester moiety and is facilitated by the aquated metal moiety under lowchloride ion conditions. The rate of ester cleavage greatly depends on the nature of the amine component. Potential applications of the linker technology are discussed.

Introduction

Platinum-based cytotoxics have a wide range of applications as curative and palliative oncology drugs.¹ Despite their overall clinical success, these agents suffer from unfavorable ADME (absorption, distribution, metabolism, excretion) properties and high systemic toxicities.² In addition to overcoming the dose-limiting toxicities of these drugs, improving the response of notoriously chemoresistant forms of cancer to platinum-based therapeutics remain high-priority goals.³ To address the latter drawback, several classes of non-classical

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Several strategies are being pursued to improve the pharmacological properties (systemic circulation and clearance, target-selective action) of chemotherapeutic agents. These include delivery of a cytotoxic payload in form of prodrugs, receptor-targeted conjugates, and nanoformulations, as well as co-delivery of agents that sensitize the affected tissue to the cytotoxic agent.⁹ Polymer-based delivery platforms and (liposomal) nanoencapsulation are currently under (pre)clinical investigation with the goal of improving the pharmacological properties of cisplatin and its second-generation derivatives.¹⁰⁻¹³ The majority of targeted nanodelivery vehicles contain the kinetically more inert platinum(IV) forms of the clinical platinum drugs.¹⁴ Release of active drug from these carriers is achieved by bio-reductive loss of axial ligation in the Pt(IV) precursors, while activation/release of Pt(II)-based formulations is achieved by simple, or acidic pH-promoted, ligand substitution.^{10, 14} In addition to serving as attachment points for passive or receptor-mediated delivery, the axial positions in Pt(IV) prodrugs have also been exploited to generate "dual-threat" conjugates containing bioactive ligands, such as chemosensitizers, or a second cytotoxic component.¹⁴

To allow incorporation of our Pt(II)-based platinum–acridines as cytotoxic "warheads" into multifunctional scaffolds, we have designed analogues containing conjugatable functional groups on the nonleaving intercalator moiety (e.g., OH and COOH in R_2 and R_3 in compounds **2a–c**, Fig. 1).⁸ We have previously demonstrated that ester derivatives of compound **2c** (e.g., **2d**, Fig. 1) are activated by platinum-promoted ester hydrolysis.¹⁵ The mechanism takes advantage of the concentration differential that exists between serum and intracellular chloride (Scheme 1), and the rate of cleavage depends on the length of the linker and the nature of R_3 . Derivatives such as valproic ester **2e** (Fig. 1),¹⁵ which are chemically robust but selectively cleaved by carboxylesterase-2 (CES-2), a prodrugactivating enzyme expressed at high levels in the liver, gastrointestinal tract, and certain solid tumors,¹⁶⁻¹⁸ can also be generated.

The goal of the current study was to develop versatile modular chemistry that combines our ester technology with amide coupling chemistry under conditions compatible with the reactive Pt(II) moiety in platinum–acridines. To demonstrate the utility of this strategy, a diverse set of multifunctional model pharmacophores was generated and the linker stability tested under physiologically relevant conditions.

Results and discussion

Development and optimization of coupling chemistry

Initial attempts to directly couple carboxylic acid-modified derivative **2a** (Fig. 1) with biologically relevant amines were unsuccessful. No amide bond formation was observed in

the presence of coupling reagents routinely used in peptide synthesis (unpublished data). We attributed this lack of reactivity to the steric hindrance produced by the platinum and acridine moieties adjacent to residue R_2 (see Fig. 1). To reduce the steric bulk of R_2 , which also improves the biological activity of the DNA-targeted agent,^{8, 15} we chose R_1 on the amidine carbon as the point of linker attachment. Previously, we identified compound 2d as a butyric ester-masked lipophilic prodrug of the highly cytotoxic hydroxyl-modified platinum–acridine **2c** (Fig. 1).¹⁵ To enable amide coupling chemistry at R_1 , we introduced a carboxylic acid functionality in 2c by extending the terminal hydroxyl group with a succinyl group. Two derivatives, acridine-based P1 and benz[c]acridine-based¹⁹ P2, were generated (Fig. 2). Both analogues were used as the carboxylic acid-containing components in coupling reactions with 11 different primary and secondary amines, N1-N11 (Fig. 2). P1 and P2 were generated in a simple three-step "one-pot" synthesis in DMF. The reactions involved substitution of one chlorido ligand in silver ion-activated $[Pt(pn)Cl_2]$ (pn = propane-1,3-diamine) with succinyl-modified nitrile (4-(3-cyanopropoxy)-4-oxobutanoic acid, 3) and subsequent addition of N^1 -(acridin-9-yl)- N^2 -methylethane-1,2-diamine) (A1) and N^{1} -(benzo[c]acridin-7-yl)- N^{2} -methylethane-1,2-diamine (**B1**), respectively (see the Experimental Section and the ESI for details).

Using carboxyl-functionalized building block **P1** and the simple primary amine 2azidoethan-1-amine (N1, Fig. 2), initially 14 solution-phase micro-scale reactions were assembled to screen a wide range of coupling reagents under varying conditions. Conversion to the desired amide in the mixtures was monitored by in-line LC-MS (see the ESI), and the yield of each reaction was determined from HPLC traces (Table 1, entries 1-14). Surprisingly, the carbodiimide-based reagents, DCC[±] and EDC,²⁰⁻²² completely failed to mediate amide bond formation. Likewise, no product was observed when coupling reactions with these reagents were performed in the presence of base (diisopropylethylamine, DIPEA, "Hünig's base") or the carboxylic acid-activating reagent N-hydroxysuccinimide (NHS). By contrast, the desired amide was produced with conversion yields of greater than 90% when uronium-based HBTU and phosphonium-based PyBOP were used. This outcome is consistent with the rapid formation of the activated benzotriazolyl esters by these coupling reagents under mild reaction conditions.²¹ On the other hand, COMU, another uroniumbased coupling reagent, which usually performs better than HBTU in solution-phase and solid-phase peptide synthesis,²² was not as efficient under the same conditions in our system. Finally, good conversion was also achieved in the presence of excess CDI (10 equiv.), which required removal of unreacted coupling reagent prior to adding the amine component. While cost-efficient, this additional step can be considered a major disadvantage when assembling libraries of conjugates for high-throughput screening.

To test if the above coupling chemistry can be applied to secondary amino groups in bulkier amines, 6 additional reactions were assembled with (E/Z)-4-hydroxy-N-desmethyltamoxifen (endoxifen, N6, Fig. 2). Similar to the primary amine N1, N6 showed the best conversion in the presence of PyBOP and HBTU (> 90%) (Table 1, entries 15–20). Since HBTU and other uronium-based coupling reagents also react with amines, resulting in guanidinium side

[‡]For a complete list of acronyms and systematic names of coupling reagents used in this study and their chemical structures, see the ESI.

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products,^{20, 23} PyBOP was selected as the coupling reagent of choice for assembling the final, extended library of conjugates. Overall, the highest conversion yields were observed for reaction mixtures containing 1.0 equiv. of carboxylic acid, 1.1 equiv. of amine, 1.0 equiv. of base, 1.5 equiv. of PyBOP, and when incubations were run for 16 h at room temperature at a concentration of **P1** of 5 mM or higher.

Assembly of an extended library in a 96-well plate format

With the optimized coupling conditions we set forth to validate the modular fragment-based conjugate assembly for the 2 functionalized platinum precursors and 10 amine components (P1, P2, N1–N10, Fig. 2). Micro-scale coupling reactions (200 µL) were performed in parallel and analyzed by fully automated LC-MS. To demonstrate the scope and utility of the amide/ester-based conjugation chemistry, a set of structurally and functionally diverse amino group containing, biologically relevant molecules were studied. Coupling of azido-amine N1 introduces a handle into the conjugate allowing additional chemical ligation (click chemistry).⁸ Lipophilic hexadecylamine (N2) and the phospholipid 1,2-distearoyl-snglycero-3-phosphoethanolamine (DSPE, N3) were chosen to turn the cationic, highly water soluble platinum-(benz)acridines into amphiphilic molecules. Such conjugates have the potential to self-assemble into micellar structures²⁴ or liposomal vesicles,²⁵ which are known to passively accumulate in tumor tissue.²⁶ The arginine derivative N4 was introduced as a model of amino acids and oligopeptides that might potentially facilitate uptake of platinum-(benz)acridines into cancer cells.²⁷ Multifunctional conjugates containing other clinically approved drugs or their key pharmacophores targeted at specific forms of cancer were also generated. The amine components introduced include (i) rucaparib²⁸ (Clovis Oncology) (N5), a member of the class of poly-ADP-ribose polymerase-1 (PARP-1) inhibitors, which have been shown to sensitize cancer cells to DNA-targeted chemotherapies, 29 (ii) endoxifen (N6), the active form of the estrogen receptor (ER) antagonist and breast cancer treatment tamoxifen,³⁰ and (iii) the epidermal growth factor receptor (EGFR) kinase-targeted quinazoline-based inhibitor N7,³¹ a derivative of the lung cancer therapy gefitinib (Astra Zeneca).³² Finally, amide coupling of P1 and P2 with mitochondria-targeted triphenylphosphonium (TPP) salts³³ (N8, N9) and with a poly(amidoamine) dendrimer (PAMAM-G3-NH₂)³⁴ (N10) was also studied.

Using the previously optimized conditions for PyBOP-mediated amide coupling, 20 reactions were assembled in Eppendorf tubes and analyzed by LC-MS (see the ESI). The majority of reactions went virtually to completion (14 out of 20) without producing undesired side products. The outcome of coupling reactions deemed successful (yields > 95%) and the structures of the resulting conjugates (confirmed by high-resolution mass spectrometry, HR-MS) are summarized in the ESI. A comparison of the results obtained for the TPP salts, **N8** and **N9**, shows that PyBOP coupling is limited by the steric hindrance of the amine component. While yields for **N8** did not exceed 70%, introduction of an extended 11-aminoundecanoic acid linker in **N9** to separate the bulky TPP moiety from the reactive primary amino group led to quantitative conversion under the same conditions. No reaction of **P1/P2** with the dendrimer–amine (**N10**) and DSPE (**N3**) was observed, which is most likely due to steric crowding and lack of solubility in DMF, respectively.

To demonstrate that the coupling reactions can be performed on a useful preparative scale, we resynthesized a representative conjugate, **P1-N7**, and isolated the pure product with a yield of 83%. To confirm formation of the desired conjugate by establishing scalar connectivity within the linker, **P1-N7** was fully characterized by 2-D NMR spectroscopy (see the ESI).

Towards aqueous coupling chemistry

While the coupling chemistry in DMF solvent provides a versatile synthetic platform, methodology would be desirable by which amide bonds with P1 and P2 can be formed in aqueous buffers or biological media. One potential application would be the attachment of platinum-(benz)acridines to cancer-targeted peptides or proteins that are insoluble in DMF. As a model reaction we chose amide coupling between compound P2 and 8-aminooctanoic acid (N11, Fig. 2). In peptide synthesis, this scenario typically requires protecting group chemistry to avoid cross-reactions between unprotected carboxyl group and coupling reagent. Because the conditions used for the deprotection steps would result in the decomposition of P2, a two-step synthesis was designed (Scheme 2). The reaction involves conversion of the carboxylic acid group in P2 with 1.5 equiv. of TSTU, a uronium-based coupling reagent,³⁵ to produce an NHS-activated ester, which is stable against hydrolysis, but reactive with amines in aqueous solution. The resulting intermediate, P2-NHS, was isolated by ether precipitation and reacted with N11 in phosphate-buffered saline (PBS). The aqueous reaction step was conducted at a high chloride ion concentration to avoid aquation of the Pt(II) moiety and prevent coordination of the primary amino group in N11 with the metal. **P2-NHS** and **N11** cleanly converted (yields > 70%) to the desired conjugate, **P2-N11**, after the mixture was incubated for 16 h at room temperature (see LC-MS analysis, ESI). It is noteworthy to mention that no hydrolysis of the succinyl ester was observed during the incubation in PBS. Thus, P2-NHS appears to be a chemically robust reagent that should allow efficient modification of primary amines, such as lysine residues in peptides or proteins,³⁶ with platinum–(benz)acridines.

Multistep "one-tube" assembly of conjugates for combinatorial library design

We have previously used a combinatorial approach for generating and screening modular libraries of functionalized platinum–acridines.⁸ We are now devising an efficient four-step reaction scheme that allows assembly of ester–amide linked multifunctional conjugates in DMF solution starting from simple diam(m)inedichlorido platinum precursor. The synthesis involves two simple precipitation/centrifugation steps, but requires no additional purification of intermediates. To demonstrate the feasibility of the proposed "one-tube" assembly, conjugate **P1-N5** was generated. The reaction sequence along with a schematic diagram is shown in Fig. 3. Briefly, [PtCl₂(pn)] was reacted with one equiv. of AgNO₃ (Step 1), AgCl was removed by centrifugation, and the activated complex was treated with excess 4-(3-cyanopropoxy)-4-oxobutanoic acid (**3**). To the resulting platinum–nitrile intermediate was then added acridine–amine **A1** to generate **P1**, which was recovered quantitatively and separated from unreacted **A1** by ether precipitation prior to the amide coupling step. LC-MS analysis performed after steps 1–3 showed that **P1** had been generated with a conversion yield of 85% (Fig. 4A). Finally, a coupling reaction using PyBOP was assembled that was

able to convert 72% of **P1** into the desired rucaparib (**N5**) conjugate, **P1-N5** (Fig. 4B). The only minor side product detected in the LC-MS traces recorded after steps 3 and 4 (Fig. 4C) resulted from direct reaction of **A1** with residual [PtCl(NO₃)pn]. With further optimization of the procedure it should be possible to generate even higher yields and purities that will allow for direct screening of the resulting library mixtures in cell-based assays. For applications requiring removal of unreacted precursor molecules, pure batches of the conjugates can be generated by passing the final reaction mixtures through gel filtration columns (Sephadex LH-20, methanol:acetonitrile, 1:1).

Conjugate stability and hydrolytic linker cleavage

Depending on the mechanism of action of a particular conjugate, the timing of linker cleavage is a critical parameter that assures maximum target selectivity and/or synergistic activity of its components. Suitable linkers are stable in circulation but undergo cleavage in target tissue as the result of a chemical or biological stimulus.³⁶ Pendant ester groups in platinum–acridines have been demonstrated to be chemically significantly more reactive than amide and carbamate groups.^{8, 15, 37} To test the hydrolytic stability of the conjugates, 7 representative derivatives were incubated in physiologically relevant buffers, and linker cleavage was monitored by LC-MS (see the ESI for a detailed product analysis). All conjugates show selective cleavage at the ester bond resulting in hydroxyl-modified platinum–acridine **2c** (Fig. 1) or the analogous benz[c]acridine hybrid, which undergo partial exchange of the chlorido ligand by water or formate ion under the conditions of this assay (see Fig. 5 for **P1-N1** as an example). Decomposition of conjugates unrelated to linker cleavage was observed after incubation periods of longer than 48 h, but only to a minor extent and only for conjugates otherwise stable to hydrolysis (see the ESI).

The time course of ester cleavage was monitored by determining the composition of reaction mixtures after 3, 6, 12, 24, and 48 h of continuous incubation (Fig. 6). Surprisingly, while all tested conjugates contain the same succinic ester linkage, -(CH₂)₃OC(O)(CH₂)₂C(O)NH-, major differences are observed in their hydrolysis rates. While cleavage in some cases is rapid and virtually complete within 12 h (P1-N1, P2-N1), other conjugates are remarkably stable (e.g., P1-N6) with only a minor amount of cleaved ester observed after 48 h. In general, the sterically least hindered alkyl amines were most reactive, whereas amines with charged (P1-N4) or bulkier groups (P1-N6, P2-N6, P1-N7) were more stable in buffered solution. A significant decrease in rate of ester hydrolysis of up to ~4-fold is observed in phosphate buffer supplemented with chloride (PBS) relative to chloride-free buffer. This observation confirms the previously established mechanism (Scheme 1) in which aquated platinum facilitates Lewis-acid promoted ester cleavage.¹⁵ Although the rate of cleavage primarily depends on the nature of the coupled amine, replacing acridine in P1-N1 and P1-N6 with benz[c]acridine in P2-N1 and P2-N6 also alters the rate of ester hydrolysis. From the LC-MS profiles (see the ESI) it appears that the extent of aquation of platinum in the conjugates correlates well with the rate of linker hydrolysis, further corroborating the proposed metal-assisted cleavage mechanism.

It should be noted that incubations of several of the chemically more stable new conjugates with CES-2 did not lead to enhanced linker cleavage (unreported data). Whether the

conjugates are susceptible to cleavage by other ubiquitous serum esterases remains to be determined. Likewise, potential proteolytic cleavage of the amide linkage in biological media remains to be investigated.

Conclusions

In conclusion, we have expanded the toolbox of linker and conjugation chemistries for platinum–acridines, a class of highly cytotoxic anticancer agents. Amide coupling appears to be a versatile and efficient platform for assembling multifunctional conjugates containing a Pt(II) moiety. The method is amenable to high-throughput library-based synthesis and is compatible with applications that require conjugation in aqueous media. Combining the coupling chemistry with cleavable linkers of variable reactivity has the potential to lead to the discovery of new multifunctional anticancer drugs with novel, synergistic modes of action. The methodology will also enable conjugation of otherwise too toxic platinum–acridines to targeted carriers, such as monoclonal antibodies. Whether the particular amide–ester type linker used in this study will be sufficiently stable for use in long-circulating, immuno-targeted antibody–drug conjugates (ADCs)³⁶ remains to be established. The linker chemistry developed in this study, in combination with previously introduced carbamate-coupled and azide–alkyne clickable analogues,^{8,37} should also have broad applicability in other clinical and experimental platinum and non-platinum metallodrugs.

Experimental section

Materials and procedures

The platinum complex [PtCl₂(pn)]⁸ and acridine precursors A1⁸ and B1¹⁹ were synthesized as described in the literature. 4-(3-Cyanopropoxy)-4-oxobutanoic acid (3)³⁸ and compounds N1,⁸ N4,³⁹ N6,³⁷ N7,³¹ and N8⁴⁰ were synthesized according to published procedures. Compound N9 was generated by HBTU/DIPEA coupling of N8 with 11-aminoundecanoic acid. Amines N2, N3, N5 (Cayman Chem. Comp., Ann Arbor, MI), N10 (Dendritech Inc., Midland, MI), and N11 were commercially available from common vendors or the provider of specialty chemicals indicated and used as supplied. All other reagents (incl. coupling reagents) were obtained from common vendors and used without further purification.

¹H NMR spectra of newly synthesized target compounds and intermediates were recorded on Bruker Advance DRX-500 and 300 MHz instruments. ¹³C {¹H} NMR spectra were recorded on a Bruker DRX-500 instrument operating at 125.8 MHz. 2-D COSY and ¹H-¹³C gradient-selected Heteronuclear Multiple Bond Coherence (gs-HMBC) data sets were acquired on a Bruker DRX-500 instrument equipped with a TBI probe. HMBC spectra were collected at 25 °C with 2048 pts in t₂ (sw = 6510 Hz), 256 pts in t₁ (sw = 27670 Hz), 128 scans per t₁ increment, and a recycle delay (*d*1) of 1.5 s. Chemical shifts (δ) are reported in parts per million (ppm) relative to internal standard tetramethylsilane (TMS). ¹H NMR data is reported in the conventional format including chemical shifts (δ , ppm), multiplicity (s = singlet, d = doublet, t = triplet, q = quartet, m = multiplet, br = broad), coupling constants (*J*, Hz), and integrated signal intensities. ¹³C NMR data are reported as chemical shift listings (δ , ppm). The NMR spectra were processed and analyzed using the MNova software package (Mestrelab Research, version 9.0.0).

LC-MS analysis was performed on an Agilent 1100LC/MSD ion trap mass spectrometer equipped with an atmospheric pressure electrospray (ES) source. HPLC-grade solvents were used for all HPLC separations in mass spectrometry experiments. Eluent nebulization was achieved with a Na pressure of 50 psi and solvent evaporation was assisted by a flow of Na

achieved with a N₂ pressure of 50 psi and solvent evaporation was assisted by a flow of N₂ drying gas (350 °C). Mass spectra in positive-ion mode were recorded with a capillary voltage of 2800 V and a mass-to-charge (m/z) scan range of 150 to 2200. HR-MS was performed on a Thermo Scientific LTQ Orbitrap XL equipped with an electrospray source, and the data was processed with Xcalibur 2.1 (Thermo Scientific). To determine the purity of target compounds, samples were diluted in methanol containing 0.1 % formic acid and separated using a 4.6 mm × 150 mm reverse-phase Agilent ZORBAX SB-C18 (5 mm) analytical column at 25 °C, using the following solvent system: solvent A, optima water, and solvent B, methanol/0.1 % formic acid, with a flow rate of 0.5 mL/min and a gradient of 95 % A to 5 % A over 30 minutes. HPLC traces were recorded with a monitoring wavelength range of 363-463 nm. Peak integration was done using the LC/MSD Trap Control 4.0 data analysis software (Agilent).

Synthetic procedures and product characterization

Synthesis of [PtClC₂₇H₃₈N₆O₄](NO₃) (P1)-P1 was synthesized using a two-step, onepot approach. A mixture of 0.678 g (2.00 mmol) of [PtCl₂(pn)] and 0.338 g (2.00 mmol) of AgNO₃ in 10 mL of anhydrous DMF was stirred at room temperature in the dark for 16 h. Precipitated AgCl was filtered off through a syringe filter. 2.96 g (16 mmol) of nitrile 3 was added to the filtrate, and the suspension was stirred at 65 °C for 4 h in the dark. The reaction mixture was then cooled to 4 °C in an ice bath. Acridine–amine A1 (0.51 g, 2.02 mmol) was added to the solution, and the suspension was stirred at 4 °C for another 48 h. To quench the reaction, the mixture was added dropwise into 200 mL of anhydrous diethyl ether, and the resulting yellow slurry was vigorously stirred for 30 min. The precipitate was recovered by membrane filtration and recrystallized from hot ethanol, affording P1 as yellow a solid. Yield: 704 mg (43 %). ¹H NMR (300 MHz, MeOH- d_4) δ 8.56 (d, J = 8.7 Hz, 2H), 7.95 (t, J = 7.5 Hz, 2H), 7.81 (d, J = 8.4 Hz, 2H), 7.57 (t, J = 7.6 Hz, 2H), 5.08 (br s, 2H), 4.80 (br s, 2H), 4.38 (t, J = 6.7 Hz, 2H), 4.32–4.22 (m, 3H), 3.94 (t, J = 6.6 Hz, 3H), 3.13–2.96 (m, 5H), 2.82–2.46 (m, 8H), 2.15 (s, 2H), 1.81 (s, 2H). ¹³C NMR (75 MHz, MeOH-d₄) δ 174.33, 172.68, 168.67, 161.75, 161.75, 154.16, 142.51, 131.29, 126.77, 121.35, 119.47, 117.88, 65.29, 63.64, 44.43, 43.58, 33.34, 32.57, 31.97, 28.99, 14.91. HR-MS (positive-ion mode): m/z for C₂₇H₃₈ClN₆O₄Pt ([M]⁺), Calculated 740.2291; found, 740.2297 (tolerance: 0.81 ppm). Analytical purity > 98% by LC-MS (see the ESI).

Synthesis of [PtClC₃₁H₄₀N₆O₄](NO₃) (P2)—P2 was synthesized by the same procedure as **P1** from 0.678 g (2.00 mmol) of [PtCl₂(pn)] and benzo[c]acridine–amine **B1** (0.61 g, 2.02 mmol). Yield: 782 mg (46 %). ¹H NMR (300 MHz, MeOH- d_4) & 9.06 (d, J = 7.9 Hz, 1H), 8.55 (d, J = 8.6 Hz, 1H), 8.27 (t, J = 9.3 Hz, 2H), 8.14–7.76 (m, 5H), 7.69 (t, J = 7.7 Hz, 1H), 5.08 (br s, 2H), 4.80 (br s, 2H), 4.35 (t, J = 6.5 Hz, 2H), 4.20 (d, J = 5.9 Hz, 2H), 3.93–3.89 (m, 2H), 3.19 – 2.90 (m, 5H), 2.88–2.45 (m, 8H), 2.08 (s, 2H), 1.79 (s, 2H). ¹³C NMR (75 MHz, MeOH- d_4) & 175.59, 173.30, 168.88, 154.31, 150.58, 143.04, 135.22, 132.17, 131.75, 130.20, 128.42, 127.56, 126.32, 125.37, 123.09, 122.94, 122.07, 116.70, 113.97, 64.04, 44.13, 43.48, 32.77, 31.75, 28. 93, 27.50, 15.44. HR-MS (positive-

ion mode): m/z for C₂₇H₃₈ClN₆O₄Pt ([M]⁺), Calculated 790.2447; found, 790.2463 (tolerance: 2.02 ppm). Analytical purity > 98% by LC-MS (see the ESI).

Synthesis of [PtCl(pn)C₄₃H₄₇ClFN₉O₄](NO₃)₂ (P1-N7)—100 mg of P1 (0.122 mmol), 71 mg of N7 (0.183 mmol), 127 mg of PyBOP (0.244 mmol) and 31 mg of DIPEA were mixed in 3 mL of anhydrous DMF. The reaction mixture was incubated at room temperature for 16 h and then poured into 200 mL of vigorously stirred anhydrous ethyl ether. The precipitate was recovered by membrane filtration and recrystallized from the mixture of ethanol and ethyl acetate (3:1) to give 121 mg of conjugate P1-N7 (yield: 83%) as a yellow solid which was dried in a vacuum. ¹H NMR (500 MHz, DMF) δ 13.91 (br s, 1H), 10.05 (br s, 1H), 8.91–8.87 (m, 1H), 8.74–8.68 (m, 2H), 8.17–8.12 (m, 1H), 8.03–7.96 (m, 5H), 7.92-7.84 (m, 1H), 7.68-7.51 (m, 4H), 6.72–6.27 (m, 2H), 5.49 (br s, 2H), 5.05 (br s, 2H), 4.58–4.50 (m, 2H), 4.37–3.99 (m, 6H), 3.86–3.74 (m, 2H), 3.50–3.47(m, 2H), 3.28– 3.13 (m, 8H), 2.83–2.53 (m, 8H), 2.30–2.17 (m, 2H), 1.90–1.80 (m, 2H), 1.56–1.43 (m, 3H). ¹³C NMR (126 MHz, DMF) δ 172.94, 172.91, 172.74, 171.31, 168.33, 158.56, 158.31, 157.43, 157.22, 156.23, 154.28, 154.08, 146.39, 140.75, 139.82, 135.30, 135.16, 126.27, 125.99, 124.58, 123.82, 119.75, 119.60, 118.95, 116.93, 116.76, 109.17, 109.03, 99.03, 98.59, 97.20, 96.77, 65.45, 63.53, 56.69, 47.53, 45.99, 43.45, 42.82, 41.57, 41.08, 33.05, 28.15, 27.87, 14.92, 13.91. HR-ESMS (positive-ion mode): *m/z* for C₄₆H₅₇Cl₂FN₁₁O₄Pt $([M]^+)$, Calculated 1111.3604; found, 1111.3615 (tolerance: 0.99 ppm). Analytical purity > 98% by LC-MS (see the ESI).

Micro-scale reactions and library assembly

Stock solutions of the carboxylic acid-modified platinum–(benz)acridines (**P1**, **P2**) (25 mM), coupling reagent PyBOP (25 mM), amines to be linked via amide bond formation (**N1–N10**) (25 mM), and DIPEA (25 mM) were prepared in anhydrous DMF. Micro-scale coupling reactions were carried out in 1.5-mL Eppendorf tubes or 96-well plates by mixing all the reaction components using the following volumes: **P1** or **P2** (40 μ L), PyBOP (60 μ L), DIPEA (40 μ L), and amines **N1–N10** (44 μ L). Finally, 16 μ L of dry DMF were added to produce a total incubation volume of 200 μ L and final platinum concentration of 5 mM. Mixtures were incubated at room temperature for 16 hours in a vortex shaker. To characterize conjugates and to determine conversion yields, 5- μ L samples are removed from the solutions and diluted with 995 μ L of methanol containing 0.1% formic acid prior to LC-MS analysis. Chromatographic separations and product quantification were performed as described above.

Chemical hydrolysis assay

Selected conjugates were generated using the microscale reactions as described before. To purify the conjugates, the reaction mixtures were precipitated in ethyl ether and separated by size exclusion chromatography (Sephadex LH-20, 10×1 cm column) using methanol:acetonitrile (1:1, v/v) as eluent. Ester hydrolysis was carried out by incubating 0.1 mM of each test compound in 10 mM phosphate buffer (pH 7.4) or 1× PBS containing 150 mM NaCl (pH 7.4) at 37 °C. Samples were withdrawn from the reaction mixture at various time points and analyzed by in-line LC-MS using the method and settings described above.

Supplementary Material

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 $\begin{array}{l} \textbf{2d:} \ R_3 = C_3 H_7 \ (butyric \ ester) \\ \textbf{2e:} \ R_3 = C H (C_3 H_7)_2 \ (valproic \ ester) \end{array}$

Fig. 1.

Structures of the most cytotoxic platinum–acridine agent (1), functionalized derivatives (2a– c), and ester-modified precursors (2d,e).





Fig. 2.

(A) Structures of platinum (**P**) and amine (**N**) components used for methodology development and library assembly. (B) Schematic representation of the coupling strategy for amide–ester linkers. The benz[c]acridine derivative in (A) contains an additional fused ring highlighted with dashed bonds.



Figure 3.

Synthetic scheme (A) and schematic representation (B) of four-step "one-tube" assembly of conjugate **P1-N5**.



Figure 4.

"One-tube" coupling of **P1** with **N5** monitored by LC-MS. HPLC traces recorded after steps 3 (A) and 4 (B) and mass spectra recorded in positive-ion mode with molecular ions $[M]^+$ and $[M]^{2+}$ labeled (C) are shown.



Figure 5.

Product analysis by LC-MS for ester cleavage in conjugate **P1-N1** (12 h, 37 °C, pH 7.4). (A) HPLC traces, mass spectra recorded in positive-ion mode with molecular ions $[M]^+$ and $[M]^{2+}$ labeled (C), and the corresponding product structures (C) are shown.



Figure 6.

Progress of ester hydrolysis in selected conjugates monitored by LC-MS in phosphate buffer (blue columns) and phosphate-buffered saline (PBS, red columns). The percentages of cleaved ester are plotted for 3, 6, 12, 24, and 48 h of continuous incubation. Data represent averages of three incubations. Numbers on the last column in each data set indicate final % cleavage after 48 h.



Scheme 1. Platinum-promoted ester cleavage in a low-chloride environment (X = chlorido or aqua ligand)



Scheme 2. Aqueous amide coupling using NHS-activated ester

1) Conversion $b(\%) \pm S.D.$		no reaction	no reaction	no reaction	no reaction	no reaction	no reaction	76 ± 2	98 ± 2	94 ± 3	99 ± 1	83 ± 1	22 ± 3	17 ± 2	18 ± 1		67 ± 3	92 ± 1	
Pt concentration (mN		N	5	S	5	5	5	5	5	S	5	S	0.5	0.5	0.5		5	5	
Molar ratio of reactants (Pt:amine:coupling reagent:base)	HNUH2 HNUH2 HNU N H H H H HZN Southing Reagent HRN N H HZN HZN Southing Reagent N HZN HZN HZN HZN HZN HZN HZN HZN HZN HZ	1:1.2:1.2:0	1:1.2:1.2:1	1:1.2:1.2(DCC):1.2(NHS):1	1:1.2:1.2:0	1:1.2:1.2:1	1:1.2:1.2(EDC):1.2(NHS):1	1:1.2:1.2:0	1:10:1.2:0	1:1.1:1.1:1	1:1.1:1.1:1	1:1.1:1.1:1	1:1.1:1.1:1	1:11:11:1	1:1:1:1:1	the state of the s	1:10:1.2:0	1:1.1:1.1:1	
Base N	HO HO HO HO HO HO HO HO HO HO HO HO HO H	:	DIPEA	DIPEA	;	DIPEA	DIPEA	1	1	DIPEA	DIPEA	DIPEA	DIPEA	DIPEA	DIPEA		1	DIPEA	
Coupling reagent		DCC	DCC	DCC/NHS	EDC	EDC	EDC/NHS	CDI	CDI	HBTU	PyBOP	COMU	HBTU	PyBOP	COMU		CDI	HBTU	
Entry ^a		-	2	ю	4	5	9	7	80	6	10	11	12	13	14		150	16	

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Entry ^a	Coupling reagent	Base	Molar ratio of reactants (Pt:amine:coupling reagent:base)	Pt concentration (mM)	Conversion ^{b} (%) \pm S.D.
18	COMU	DIPEA	ELELET	5	73 ± 1
19	HBTU	DIPEA	1:1.1:1.5:1	5	88 ± 1
20	P_{yBOP}	DIPEA	1:1.1:1.5:1	5	99 ± 1

 $\frac{a}{2}$ Each reaction was performed in triplicate and reaction mixtures were characterized by LC-MS without purification.

 $b_{
m The}$ conversion yields were determined from HPLC traces recorded in an acridine-specific wavelength range.

^cCDI was pre-incubated with Pt complexes for 5 h before the intermediate was precipitated with diethyl ether to remove unreacted CDI before mixing with amine.