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Article

Roles of DNA Target in Cancer Cell-Selective Cytotoxicity by Dicopper Complexes with DNA Target/Ligand Conjugates

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ABSTRACT: The DNA target/ligand conjugates (HL^{X} , X = Pnand Mn, n = 1-3) were synthesized where various lengths of $-\text{CONH}(\text{CH}_2\text{CH}_2\text{O})_n\text{CH}_2\text{CH}_2\text{NHCO}-$ linkers with a 9-phenanthrenyl (P) or methyl (M) terminal as DNA targets replace the methyl group of 2,6-di(amide-tether cyclen)-*p*-cresol ligand (HL). DNA binding, DNA cleavage, cellular uptake, and cytotoxicity of $[\text{Cu}_2(\mu\text{-OH})(\text{L}^X)](\text{ClO}_4)_2$ ($\mathbf{1}^X$) are examined and compared with those of $[\text{Cu}_2(\mu\text{-OH})(\text{L})](\text{ClO}_4)_2$ (1) to clarify roles of DNA targets. Upon reaction of $\mathbf{1}^X$ with H_2O_2 , μ -1,1-O₂H complexes are formed for DNA cleavage. $\mathbf{1}^{P1}$, $\mathbf{1}^{P2}$, and $\mathbf{1}^{P3}$ are 22-, 11-, 3-fold more active for conversion of Form II to III in the cleavage of supercoiled plasmid DNA with H_2O_2 than 1, where the short Plinker may fix a dicopper moiety within a small number of base



pairs to facilitate DNA double-strand breaks (dsb). This enhances the proapoptotic activity of 1^{P1} , 1^{P2} , and 1^{P3} , which are 30-, 12-, and 9.9-fold cytotoxic against HeLa cells than 1. DNA dsb and cytotoxicity are 44% correlated in 1^{P1-3} but 5% in 1^{M1-3} , suggesting specific DNA binding of P-linkers and nonspecific binding of M-linkers in biological cells. 1^{P1-3} exert cancer cell-selective cytotoxicity against lung and pancreas cancer and normal cells where the short P-linker enhances the selectivity, but 1^{M1-3} do not. Intracellular visualization, apoptosis assay, and caspase activity assay clarify mitochondrial apoptosis caused by 1^{P1-3} . The highest cancer cell selectivity of 1^{P1} may be enabled by the short P-linker promoting dsb of mitochondrial DNA with H_2O_2 increased by mitochondrial dysfunction in cancer cells.

INTRODUCTION

Cancer is the most malignant disease, and the death rate due to cancer greatly increases to date. Cisplatin and related platinum complexes^{1,2} and iron bleomycin (Fe-BLM)^{3,4} are clinically used as anticancer agents. BLM forms an iron complex (Fe-BLM) after being dosed. These metal complexes induce apoptosis by binding and cutting DNA. These cytotoxic anticancer agents show high efficacy to a wide range of cancer disease, but there are problems of side effects by equally acting on cancer and normal cells.^{2,5,6} Therefore, the development of anticancer metal complexes to exert cancer cell-selective cytotoxicity is important to suppress the side effects.

DNA- and organelle-targeting metal complexes have attracted interests of chemists and medical scientists to develop anticancer reagents that are able to exert cancer cellselective cytotoxicity.^{7–10} Mitochondrial dysfunction and high metabolic activity increase H_2O_2 concentration in cancer cells as the specific microenvironments.^{11,12} So far, a variety of metal complexes that are able to target mitochondria for cancer therapy have been developed to induce mitochondrial apoptosis.^{9,13} For example, mitochondria-targeting metal complexes using Cu,^{7e} Ru,^{7f} Ir,^{7g,h} and Au⁷ⁱ were reported. The Cu complex induced mitochondrial apoptosis by DNA cleavage.^{7e} The Ru and Ir complexes were effective as a photosensitizer for photodynamic therapy.^{7fg} The phosphorescent cyclometalated Ir compounds caused reactive oxygen species (ROS) elevation, cellular ATP depletion, caspasedependent apoptosis through mitochondrial damage, and mitochondrial respiration inhibition.^{7h} The Au porphyrin complex that is highly stable in a biological environment directly targeted mitochondrial chaperone to inhibit the activity.⁷ⁱ Besides these complexes, DNA- and mitochondriatargeting metal complexes having a H₂O₂ activation site are promising candidates capable of enhancing cancer cell-selective cytotoxicity by inducing mitochondrial apoptosis by cleaving mitochondrial DNA.¹⁴

Moreover, a variety of copper complexes have been well investigated to make anticancer agents with low cytotoxicity.^{15,16} Copper complexes may be effective to exert cancer cell-selective cytotoxicity because copper is an endogenous

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Scheme 1. Chemical Structures of HL, HL^{X} , 1, and $1^{X} (X = P1-3 \text{ and } M1-3)^{a}$



^aDNA targets are highlighted by red frames.

metal and has antineoplastic characters. Moreover, copper complexes having DNA-targeting moieties have been developed to enhance the anticancer activity.¹⁶ However, the DNAtargeting dinuclear copper complexes have been less developed so far, and roles of DNA targets in the cancer cell-selective cytotoxicity have not been clarified yet.

Recently, we found that a μ -1,1-hydroperoxodicopper(II) complex is formed from the dicopper(II) complex of 2,6bis(amide-tether cyclen)-*p*-cresol (HL) $[Cu_2(\mu-OH)(L)]$ - $(ClO_4)_2$ (1) upon reaction with H_2O_2 and largely accelerates oxidative DNA cleavage.¹⁷ However, the cytotoxicity of 1 is too low to be used as an anticancer agent. To enhance the cytotoxicity, a phenanthrene/HL conjugate HL^{P2} was synthesized where a 9-phenanthrenyl terminal -CONH-(CH₂CH₂O)₂CH₂CH₂NHCO- linker (P-linker) is introduced into HL because phenanthrene can bind DNA as an intercalator.¹⁸ The P-linker drastically improved the DNA cleavage activity and cytotoxicity of the dicopper(II) complex of HL^{P2} [Cu₂(μ -OH)(L1^{P2})](ClO₄)₂ (1^{P2}). Moreover, 1^{P2} exerted relatively high cancer cell-selective cytotoxicity. The caspase activity assay and intracellular visualization of 1^{P2} by the phenathrenyl group as a fluorescent probe revealed that mitochondrial apoptosis plays a key role in cancer cell-selective cytotoxicity by $1^{\dot{P2}.18}$ In this stage, however, the systematic design of DNA target/ligand conjugates has not been addressed to optimize cancer cell-selective cytotoxicity.

In this study, a series of DNA target/ligand conjugates (HL^X, X = Pn and Mn, n = 1-3) were synthesized to clarify and optimize roles of the DNA target in cancer cell-selective cytotoxicity where 4-methyl of the original ligand HL is replaced by $-\text{CONH}(\text{CH}_2\text{CH}_2\text{O})_n\text{CH}_2\text{CH}_2\text{NHCO}-$ linkers having a 9-phenanthrenyl (P) or methyl (M) terminal as DNA targets, P-linkers, and M-linkers, respectively. HL^X form dicopper(II) complexes [Cu₂(μ -OH)(L^X)](ClO₄)₂ (1^X). The ligands and dicopper complexes used in this study are drawn in Scheme 1. DNA binding, DNA double-strand break (dsb) activity of plasmid DNA with H₂O₂, cellular uptake, and cytotoxicity against various normal and cancer cells of 1^{X} are examined and compared with 1 to clarify roles of the P- and M-linkers as DNA targets. The localization of 1^{P1-3} in mitochondria is visualized by means of the confocal microscopy. The apoptotic cell death by 1^{P1-3} is shown by fluorescence-activated cell sorting (FACS). Caspase activity assays clarified the intrinsic pathway of apoptosis by 1^{X} . These results clarify roles of the DNA targets in cancer-cell-selective cytotoxicity of 1^{X} .

EXPERIMENTAL SECTION

Materials. Unless otherwise noted, solvents and reagents were purchased and used as received. Calf thymus DNA (Type I, fibers) and supercoiled plasmid pUC19 DNA were purchased from Nippon Gene CO., Ltd., and Sigma-Aldrich, respectively. 3,5-Diformyl-4-hydroxybenzoic acid 3, 1-amino-6-azido-3-oxaoctane 4 (n = 1), 1-amino-11-azido-3,6,9-triox-aundecane 4 (n = 3), $N_iN'_iN''$ -4,7,10-triBOC-cyclen 5, phenanthrene-9-carbonyl chloride 6, acetyl chloride 7, 11 (n = 2), and dicopper complexes 1 and 1^{P2} were prepared as previously reported.¹⁷⁻²⁰

Measurements. A PerkinElmer Elemental Analyzer 2400 II was used for elemental analyses (C, H, and N). An Agilent 8454 UV spectroscopy system was used to measure the UVvis spectra. A HORIBA LAQUA electrode was used to measure pH, and a standard buffer solution was used. A SHIMADZU Single Reflection HATR IRAffinity-1 MIRacle 10 was used to obtain the infrared (IR) spectra. The ¹H NMR spectra were recorded on a JEOL ECA-500RX spectrometer using Me₄Si as an internal standard. A HITACHI Spectral fluorometer F-7000 was used to measure the fluorescence spectra. A JEOL JMS-T100CS spectrometer was used to measure cold spray ionization mass (CSI MS) and electrospray ionization mass (ESI MS). A SHIMADZU Inductively Coupled Plasma Mass Spectrometer ICP MS-2030 was used to determine copper content in cells. Confocal microscopic pictures were recorded on a confocal laser microscope A1 by

Nikon Instech Co., Ltd. A Thermo Scientific Multiskan FC was used to carry out MTT assay. Flow cytometry studies were conducted on an Attune NxT Focusing Cytometer (Invitrogen).

Preparations. *N*-(5-Azido-3-oxaoctyl)-3,5-diformyl-4-hydroxybenzamide (8 (n = 1)). A solution of 3 (2.95 g, 15.2 mmol), 4 (n=1) (6.60 g, 50.7 mmol), EDC·HCl (9.75 g, 50.8 mmol), and Et₃N (7.2 mL, 51.6 mmol) in CHCl₃ (200 mL) was stirred at 0 °C for 15 h under N₂. To the resultant solution was added 1 M aqueous HCl (300 mL) and stirred at 30 °C for 15 h. After the H₂O layer was separated using a separation funnel, the organic layer was washed with H₂O (3×30 mL) and dried over Na₂SO₄. After removal of Na₂SO₄ by filtration, the filtrate was concentrated by evaporation. The product was obtained as a brown solid. Yield: 3.37 g (11.0 mmol, 72%). S00 MHz ¹H NMR (CDCl₃): δ /ppm = 11.9 (s, H, OH), 10.3 (brs, 2H, CHO), 3.68–3.74 (m, 6H, CH₂), 3.42 (t, J = 5.1 Hz, 2H, CH₂).

N-(11-Azido-3,6,9-trioxaoctyl)-3,5-diformyl-4-hydroxybenzamide (8 (n = 3)). 8 (n = 3) was prepared from 3 and 4 (n = 3) in a similar method described for 8 (n = 1). Yield: 67%.500 MHz ¹H NMR (CDCl₃): δ /ppm = 11.9 (s, H, OH), 10.3 (brs, 2H, CHO), 8.52 (s, 2H, Ph), 3.62–3.75 (m, 14H, CH₂), 3.37 (t, J = 5.0 Hz, 2H, CH₂).

5-((5-Azido-3-oxaoctyl)carbamoyl)-2-hydroxyisophthalic Acid (9 (n = 1)). In a 200 mL round-bottom flask, 8 (n= 1) (3.37 g, 11.0 mmol) and Ag₂O (7.89 g, 34.1 mmol) were placed and suspended in aqueous NaOH (3.64 g, 90.9 mmol, 70 mL). The mixture was stirred for 15 h at 60 °C. The insoluble materials were filtered and washed with a small amount of hot H₂O. The filtrate was adjusted to pH 1 using 12 M aqueous HCl. The product precipitated as a white solid and was collected by filtration. Yield: 3.23 g (9.56 mmol, 87%). 500 MHz ¹H NMR (DMSO-d₆): δ /ppm = 8.56 (t, J = 5.2 Hz, H, NH), 8.48 (s, 2H, CH), 3.62 (t, J = 4.6, 5.2 Hz, 2H, CH₂), 3.56 (t, J = 6.3 Hz, 2H, CH₂), 3.40 (t, J = 4.6, 5.2 Hz, 4H, CH₂).

5-((11-Azido-3,6,9-trioxaoctyl)carbamoyl)-2-hydroxyisophthalic Acid (9 (n = 3)). 9 (n = 3) was obtained from 8 (n = 3) similarly to the preparation of 9 (n = 1). Yield: 69%. 500 MHz ¹H NMR (DMSO- d_6): δ /ppm = 8.53 (t, J = 5.2 Hz, H, NH), 8.47 (s, 2H, CH), 3.60 (t, J = 5.2 Hz, 2H, CH₂), 3.54– 3.58 (m, 4H, CH₂), 3.52 (t, J = 6.3 Hz, 2H, CH₂), 3.39 (t, J = 6.3 Hz, 2H, CH₂), 3.37 (t, J = 6.3 Hz, 2H, CH₂).

Compound 10 (n = 1). To a DMF (20 mL) solution of DIPEA (1.7 mL, 9.90 mmol) and 9 (n= 1) (2.40 g, 5.08 mmol) was added a DMF (10 mL) solution of 5 (530 mg, 1.57 mol) and TBTU (1.50 g, 4.67 mmol) and stirred under N₂ for 15 h. The reaction mixture was concentrated in vacuo. To the residue was added H₂O (60 mL). The product was extracted with CHCl₃ (4 × 60 mL), purified by silica gel column chromatography (gradient from EtOAc to EtOAc/MeOH, 10/1), and obtained as a brown solid. Yield: 1.39 g (1.11 mmol, 71%). 500 MHz ¹H NMR (CDCl₃): δ /ppm = 7.86 (s, 2H, Ph), 3.33–3.70 (m, 38H, CH₂), 3.36 (t, *J* = 5.2, 4.6 Hz, 2H, CH₂), 1.28–1.52 (m, 54H, CH₃). ESI MS (positive mode) (MeOH): *m*/*z* = 646.1 (646.4: calcd for [10 (*n*= 1) + 2Na]²⁺) and 1269.1 (1269.7: calcd for [10 (*n*= 1) + Na]⁺).

Compound **10** (*n* = **3**). **10** (*n* = **3**) was prepared from **9** (*n* = **3**) and **5** by a similar method described for **10** (*n* = **1**). Yield: 49%. 500 MHz ¹H NMR (CDCl₃): δ /ppm = 7.87 (s, 2H, Ph), 3.21–3.83 (m, 48H, CH₂), 1.26–1.53 (m, 54H, CH₃). ESI MS (positive mode) (MeOH): *m*/*z* = 690.2 (690.4: calcd for [**10**

 $(n = 3) + 2Na]^{2+}$ and 1357.4 (1357.8: calcd for $[10 (n = 3) + Na]^{+}$).

Compound 11 (n = 1). To a solution of 10 (n = 1) (301 mg, 0.241 mmol) in MeOH (15 mL) was added 10% Pd-C (275 mg) under N₂. The gas phase was replaced by evacuation and refilling with H₂. The mixture was stirred at room temperature for 15 h. Pd-C was removed by filtration. After the concentration of the filtrate, the product was obtained as a pale yellow solid. Yield: 266 mg (0.217 mmol, 90%). ESI MS (positive mode) (MeOH): m/z = 1221.1 (1221.7: calcd for [11 (n = 1) + H]⁺).

Compound 11 (n = 3). 11 (n = 3) was prepared from 10 (n = 3) by a similar method described for 11 (n = 1). Yield: 93%. ESI MS (positive mode) (MeOH): m/z = 1309.8 (1309.8: calcd for $[11 (n = 3) + H]^+$).

Compound 12 (n = 1). To a THF (10 mL) solution of 6 (34.0 mg, 0.141 mmol) was added a THF (10 mL) solution of 11 (n=1) (169 mg, 0.138 mmol) and Et₃N (0.086 mL, 0.617 mol). The mixture was stirred at 0 °C for 1 h and at room temperature for 15 h under N2 and then concentrated to dryness. The product was extracted with H₂O (15 mL) and $CHCl_3$ (3 × 30 mL). The organic layers were combined and dried over Na2SO4. After Na2SO4 was filtered off, the filtrate was concentrated to dryness. The crude product was purified by silica gel column chromatography (gradient from CHCl₃ to $CHCl_3/MeOH$, 10/1). After the concentration of the fractions, the product was obtained as a pale-yellow solid. Yield: 115 mg (0.0805 mmol, 58%). 500 MHz ¹H NMR $(CDCl_3): \delta/ppm = 8.71 (d, I = 8.0 Hz, 1H, Phen5), 8.67 (d, I)$ = 8.4 Hz, 1H, Phen4), 8.35 (d, J = 8.4 Hz, 1H, Phen1), 7.90 (s, 1H, Phen10), 7.89 (d, J = 7.3 Hz, 1H, Phen8), 7.82 (s, 2H, Ph), 7.60-7.72 (m, 4H, Phen2, Phen3, Phen6, Phen7), 3.30-3.81 (m, 40H, CH₂), 1.26-1.52 (m, 54H, CH₃). ESI MS (positive mode) (MeOH): m/z = 735.1 (735.4: calcd for [12] $(n=1) + 2Na^{2+}$ and 1447.2 (1447.8: calcd for [12 (n=1) + $Na^{+}).$

Compound 12 (n = 3). 12 (n = 3) was prepared using 11 (n = 3) and 6 by a similar procedure to that described for 12 (n = 1). Yield: 81%. 500 MHz ¹H NMR (CDCl₃): δ /ppm = 8.71 (d, J = 8.6 Hz, 1H, Phen4), 8.68 (d, J = 8.0 Hz, 1H, Phen5), 8.36 (d, J = 8.0 Hz, 1H, Phen4), 7.92 (s, 1H, Phen10), 7.90 (d, J = 8.0 Hz, 1H, Phen1), 7.83 (s, 2H, Ph), 7.58–7.74 (m, 4H, Phen2, Phen3, Phen6, Phen7), 3.06–3.87 (m, 48H, CH₂), 1.26–1.53 (m, 54H, CH₃). ESI MS (positive mode) (MeOH): m/z = 779.0 (779.4: calcd for [12 (n = 3) + 2Na]²⁺), 1535.5 (1535.8: calcd for [12 (n = 3) + Na]⁺).

Compound 13 (n = 1). To a CH_2Cl_2 (5 mL) suspension of 11 (n= 1) (200 mg, 0.164 mmol) and K₂CO₃ (70.0 mg, 0.506 mol) was added a CH₂Cl₂ (5 mL) solution of acetyl chloride (35.0 μ L, 0.493 mmol). The mixture was stirred at 0 °C for 1 h and at room temperature for 15 h under N2. The solid generated was filtered off. The filtrate was placed on a separation funnel and washed with H_2O (3 × 10 mL). The organic layer was separated and dried over Na2SO4. The crude product was purified by silica gel column chromatography (gradient from EtOAc to EtOAc/MeOH, 10/1). The product was obtained after the concentration of fractions as a pale yellow solid. Yield: 103 mg (0.0819 mmol, 50%). 500 MHz 1 H NMR (CDCl₃): δ /ppm = 7.86 (s, 2H, Ph), 3.41–3.56 (m, 40H, CH₂), 2.01 (s, 3H, CH₃), 1.26–1.67 (m, 54H, CH₃). ESI MS (positive mode) (MeOH): m/z = 654.0 (654.3: calcd for $[13 (n=1) + 2Na]^{2+}$, 1285.5 (1285.7: calcd for [13 (n=1) + $Na|^+$).

Compound 13 (n = 2). 13 (n = 2) was prepared from 11 (n = 2) and acetyl chloride by a similar method described for 13 (n = 1). Yield: 45%. 500 MHz ¹H NMR (CDCl₃): δ /ppm = 7.88 (s, 2H, Ph), 3.35-3.66 (m, 44H, CH₂), 1.95 (s, 3H, CH₃), 1.30-1.54 (m, 54H, CH₃). ESI MS (positive mode) (MeOH): m/z = 676.0 (676.4: calcd for [13 (n = 2) + 2Na]²⁺), 1329.5 (1329.7: calcd for [13 (n = 2) + Na]⁺).

Compound 13 (n = 3). 13 (n = 3) was prepared using 11 (n = 3) and acetyl chloride by a similar procedure to that described for 13 (n = 1). Yield: 48%. 500 MHz ¹H NMR (CDCl₃): δ /ppm = 7.86 (s, 2H, Ph), 3.36–3.63 (m, 48H, CH₂), 1.95 (s, 3H, CH₃), 1.25–1.49 (m, 54H, CH₃). ESI MS (positive mode) (MeOH): m/z = 698.0 (698.4: calcd for [13 (n = 3) + 2Na]²⁺), 1373.5 (1373.8: calcd for [13 (n = 3) + Na]⁺).

HL^{P1}.6*HCl.* To a EtOH (6 mL) solution of **12** (*n*= **1**) (699 mg, 0.490 mmol) was added 12 M HCl (2 mL) dropwise at 0 °C. After being stirred at room temperature for 15 h, the mixture was concentrated to dryness. The white solid obtained was washed by a small amount of dry EtOH. Yield: 449 mg (0.430 mmol, 88%). 500 MHz ¹H NMR (D₂O): δ /ppm = 8.88 (d, *J* = 8.0 Hz, 1H, Phen4), 8.84 (d, *J* = 8.0 Hz, 1H, Phen5), 8.11 (d, *J* = 8.0 Hz, 1H, Phen8), 8.08 (d, *J* = 8.0 Hz, 1H, Phen1), 7.93 (s, 1H, Phen10), 7.90 (s, 2H, Ph), 7.64–7.86 (m, 4H, Phen2, Phen3, Phen6, Phen7), 2.65–3.93 (m, 40H, CH₂). ESI MS (positive mode) (H₂O): *m*/*z* = 413.0 (413.2: calcd for $[HL^{P1} + 2H]^{2+}$), 825.1 (825.5: calcd for $[HL^{P1} + H]^+$).

HL^{P3.}6*HCl*. **HL**^{P3.}6*HCl* was prepared from **12** (*n* = **3**) by a similar method described for **HL**^{P1.}6*HCl*. Yield: 85%. 500 MHz ¹H NMR (D₂O): δ /ppm = 8.87 (d, *J* = 8.0 Hz, 1H, Phen4), 8.82 (d, *J* = 8.0 Hz, 1H, Phen5), 8.17 (d, *J* = 8.0 Hz, 1H, Phen4), 8.05 (d, *J* = 8.0 Hz, 1H, Phen5), 8.17 (d, *J* = 8.0 Hz, 1H, Phen10), 7.75 (s, 2H, Ph), 7.69–7.83 (m, 4H, Phen2, Phen3, Phen6, Phen7), 2.90–3.90 (m, 48H, CH₂). ESI MS (positive mode) (H₂O): *m*/*z* = 457.3 (457.3: calcd for [**HL**^{P3} + 2H]²⁺), 913.5 (913.5: calcd for [**HL**^{P3} + H]⁺).

HL^{M1}·6*HCl.* **HL**^{M1}·6*HCl* was prepared from 13 (n = 1) by a similar method described for **HL**^{P1}·6*HCl.* Yield: 90%. 500 MHz ¹H NMR (D₂O): δ/ppm = 7.86 (s, 2H, Ph), 3.07–3.90 (m, 40H, CH₂), 1.96 (s, 3H, CH₃). ESI MS (positive mode) (H₂O): m/z = 332.0 (332.2: calcd for [**HL**^{M1} + 2H]²⁺), 663.2 (663.4: calcd for [**HL**^{M1} + H]⁺).

HL^{M2}·6*HCl.* **HL**^{M2}·6*HCl* was prepared from **13** (*n* = **2**) by a similar method described for **HL**^{P1}·6*HCl.* Yield: 95%. 500 MHz ¹H NMR (D₂O): δ /ppm = 7.96 (s, 2H, Ph), 3.04–3.94 (m, 44H, CH₂), 1.99 (s, 3H, CH₃). ESI MS (positive mode) (H₂O): *m*/*z* = 354.0 (354.2: calcd for [**HL**^{M2} + 2H]²⁺), 707.3 (707.5: calcd for [**HL**^{M2} + H]⁺). *HL*^{M3}·6*HCl.* **HL**^{M3}·6*HCl* was prepared from **13** (*n* = **3**) by a

HL^{M3}·6*HCl.* **HL**^{M3}·6*HCl* was prepared from 13 (n = 3) by a similar method described for **HL**^{P1}·6*HCl.* Yield: 98%. 500 MHz ¹H NMR (D₂O): δ /ppm = 7.90 (s, 2H, Ph), 3.09–3.95 (m, 48H, CH₂), 1.99 (s, 3H, CH₃). ESI MS (positive mode) (H₂O): m/z = 376.0 (376.2: calcd for [**HL**^{M3} + 2H]²⁺), 751.3 (751.5: calcd for [**HL**^{M3} + H]⁺).

Dicopper(II) Complex 1^{P1} . The HCl salt HL^{P1} ·6HCl (103 mg, 0.0990 mmol) was treated with aqueous NaOH (1 M, 600 μ L). After the mixture was concentrated and dried in vacuo, to the residue was added CH_2Cl_2 , and HL^{P1} was extracted. After concentration, HL^{P1} was obtained as a pale-yellow solid (57.2 mg, 0.0693 mmol; yield: 70%). To a H_2O (200 μ L) solution of HL^{P1} (57.2 mg, 0.0693 mmol) was added a H_2O (200 μ L) solution of $Cu(ClO_4)_2$ ·6H₂O (36.4 mg, 0.139 mmol) under N₂. The mixture was adjusted to pH 8 with aqueous NaOH (1

M). The mixture was concentrated in vacuo. $[Cu_2(\mu - OH)(L^{P1})](ClO_4)_2$ (1^{P1}) was obtained as a green solid and recrystallized from MeOH/Et₂O. Yield: 65.5 mg (0.0561 mmol, 81%). Anal. calcd for $[Cu_2(\mu - OH)(L^{P1})](ClO_4)_2$. NaClO₄·4H₂O: C, 38.82; H, 5.03; N, 10.29. Found: C, 38.96; H, 4.96; N, 9.81. ESI MS (positive mode) (H₂O): m/z = 474.0 (474.2: calcd for $[1^{P1} - OH - 2ClO_4 - H]^{2+}$), 483.1 (483.2: calcd for $[1^{P1} - 2ClO_4]^{2+}$), and 1065.0 (1065.3: calcd for $[1^{P1} - ClO_4]^+$). IR (KBr): $\tilde{\nu}/cm^{-1}$ = 2870, 1628, 1533, 1443, 1362, 1300, 1273, 1090, 953, 932, 837, 810, 781, 756, 729, 691, 654, 623. Other dicopper(II) complexes 1^{P3} and 1^{M1-M3} were synthesized by the almost same method as 1^{P1} where HL^{P3} and HL^{M1-M3} were used instead of HL^{P1}. Data of each complex are shown below.

Dicopper(II) Complex 1^{P3} . 1^{P3} was prepared from HL^{P3}. 6HCl by a similar method described for 1^{P1} . Yield: 75%. Anal. calcd for $[Cu_2(\mu-OH)(L^{P3})](ClO_4)_2\cdot 8H_2O: C, 41.20; H, 6.05; N, 10.01. Found: C, 41.39; H, 5.59; N, 10.2. ESI MS (positive mode) (H₂O): <math>m/z = 527.1$ (527.2: calcd for $[1^{P3} - 2ClO_4]^{2+}$), 1153.32 (1153.31: $[1^{P3} - ClO_4]^+$). IR (KBr): $\tilde{\nu}/$ cm⁻¹ = 3744–3134, 2928, 2878, 2020, 1726, 1632, 1537, 1443, 1360, 1300, 1273, 1246, 1076, 1001, 988, 953, 930, 839, 812, 786, 756, 731, 700, 656, 623. *Dicopper(II) Complex* 1^{M1} . 1^{M1} was prepared from HL^{M1}.

Dicopper(II) Complex 1^{M1} . 1^{M1} was prepared from HL^{M1}. 6HCl by a similar method described for 1^{P1} . Yield: 90%. Anal. calcd for $[Cu_2(\mu-OH)(L^{M1})](ClO_4)_2 \cdot 6H_2O$: C, 33.46; H, 5.98; N, 12.59. Found: C, 33.70; H, 6.23; N, 12.85. ESI MS (positive mode) (H₂O): m/z = 402.0 (402.1: calcd for $[1^{M1} - 2ClO_4]^{2+}$), 903.0 (903.2: calcd for $[1^{M1} - ClO_4]^+$). IR (KBr): $\tilde{\nu}/cm^{-1} = 2908$, 2029, 1624, 1539, 1437, 1364, 1302, 1271, 1082, 988, 930, 880, 850, 837, 812, 790, 766, 692, 667, 648, 625.

Dicopper(II) Complex 1^{M2} . 1^{M2} was prepared from HL^{M2}. 6HCl by a similar method described for 1^{P1} . Yield: 73%. Anal. calcd for $[Cu_2(\mu-OH)(L^{M2})](ClO_4)_2\cdot 4H_2O$: C, 35.36; H, 5.93; N, 12.50. Found: C, 35.08; H, 6.00; N, 12.25. ESI MS (positive mode) (H_2O) : m/z = 414.9 (415.1: calcd for $[1^{M2} - OH - 2ClO_4 - H]^{2+}$), 424.0 (424.2: calcd for $[1^{M2} - 2ClO_4]^{2+}$), 947.0 (947.3: calcd for $[1^{M2} - ClO_4]^+$). IR (KBr): $\tilde{\nu}/cm^{-1} = 2875$, 2054, 1628, 1547, 1445, 1368, 1302, 1275, 1083, 950, 932, 840, 810, 785, 772, 690, 654, 620.

Dicopper(II) Complex 1^{M3}. 1^{M3} was prepared from HL^{M3}. 6HCl by a similar method described for 1^{P1}. Yield: 80%. Anal. calcd for $[Cu_2(\mu-OH)(L^{M3})](ClO_4)_2\cdot 3H_2O$: C, 36.65; H, 5.98; N, 12.21. Found: C, 36.55; H, 6.02; N, 12.30. ESI MS (positive mode) (H₂O): m/z = 437.0 (437.2: calcd for $[1^{M3} - OH - 2ClO_4 - H]^{2+}$), 446.0 (446.2: calcd for $[1^{M3} - 2ClO_4]^{2+}$), 991.0 (991.3: calcd for $[1^{M3} - ClO_4]^{+}$). IR (KBr): $\tilde{\nu}/cm^{-1} = 2905$, 2054, 1628, 1553, 1445, 1360, 1275, 1085, 935, 885, 850, 805, 700, 654, 613.

Ethidium Bromide Method.²¹ To a buffer (pH 6.0 (MES, 10 mM)) solution of ct-DNA (20 μ M bp) and EtBr (3.3 μ M) containing NaCl (10 mM) was added a buffer (pH 6.0 (MES, 10 mM)) solution of 1^X containing NaCl (10 mM) in 2 μ M intervals until it reached 400 μ M. The fluorescence spectra with excitation at 510 nm were measured in 520–800 nm at 37 °C. The apparent binding constants K_{app} (M⁻¹) were obtained using eq 1 where [complex] = (the concentrations of 1^{P1,3} and 1^{M1-3} at a 50% reduction of the fluorescence intensity of EtBr), $K_{FtBr} = 1.1 \times 10^7 \text{ M}^{-1}$, and [EtBr] = 1.65 μ M were used.

$$K_{\text{EtBr}}[\text{EtBr}] = K_{\text{app}}[\text{complex}]$$
(1)

Measurements of DNA Cleavage. To evaluate the DNA cleavage activity of 1^{x} , pUC19 DNA was used as a substrate. 1^{X} (0-50 μ M) was dissolved in an aqueous solution containing pUC19 DNA (50 µM bp), buffer (10 mM, pH 6.0 (MES)), and NaCl (10 mM). To the mixture was added H_2O_2 (0-500 μ M) at 37 °C in the dark. The reaction was followed by taking an aliquot at 0, 10, 20, 30, 40, 60, 120, 180, and 300 min and quenched by addition of a loading dye (0.025% bromophenol blue, 0.025% xylene cyanol FF, 1.0 mM EDTA, and 30% glycerol). Each sample was subjected to electrophoresis on a 1% agarose gel in a TAE (tris/acetate/ EDTA) buffer for 1 h at 100 V. The gels were stained with EtBr (0.5 μ g μ L⁻¹) for 1 h. The DNA bands were visualized on a UV transilluminator, photographed by a Vilber Lourmat ECX-20-M. For the reduced stain uptake of Form I, a correction factor of 1.06 was used to quantify.

Spectroscopic Identification of μ -1,1-Hydroperoxodicopper(II) Complexes. To a MeCN solution of 1^{P1} or 1^{P3} (0.25 mM, 2 mL) in a quartz cell at -30 ± 0.2 °C (or rt) was added a MeCN solution of H₂O₂ (10 equiv). The formation of μ -1,1-hydroperoxodicopper(II) complexes was monitored by the electronic absorption spectral change in the range of 290–850 nm and also observed by the CSI MS spectra in H2O using not only H₂¹⁶O2 but also H₂¹⁸O2.

Cell Culture. Prof. Kitagishi provided HeLa (human cervical cancer) cells used in this study (Doshisha University, Kyoto, Japan). The RIKEN Bioresource Center provided WI-38 (human lung fibroblast normal), A549 (human lung cancer), PK-59 (human cancer), and 2C6 (human embryonic pancreas normal) cells used in this study (Ibaraki, Japan). HeLa, A549, WI-38, and PK-59 cells were maintained in Dulbecco's modified Eagle's medium (D-MEM, high glucose) (FUJIFILM Wako Pure Chemical Corporation, Osaka, Japan), Dulbecco's modified Eagle's medium (DMEM, low glucose) (Thermo Fisher Scientific, Tokyo, Japan), minimum essential medium Eagle (MEM) (Sigma-Aldrich Japan, Tokyo, Japan), or Roswell Park Memorial Institute 1640 Medium (RPMI 1640) (Thermo Fisher Scientific, Tokyo, Japan), respectively. These cell culture media were supplemented with fetal bovine serum (10%) (Biowest, Nuaillé, France) and penicillin (100 units mL⁻¹) and streptomycin (100 μ g mL⁻¹) (FUJIFILM Wako Pure Chemical Corporation, Osaka, Japan) at 37 °C in a humidified atmosphere of 5% CO₂. 2C6 cells were maintained in Mixture F-10 Ham (HamF10) (Sigma-Aldrich Japan, Tokyo, Japan) supplemented with newborn calf serum (10%) (Sigma-Aldrich Japan, Tokyo, Japan), fetal bovine serum (5%) (Biowest, Nuaillé, France), horse serum (2.5%) (Sigma-Aldrich Japan, Tokyo, Japan), and penicillin (100 units mL⁻¹) and streptomycin (100 μ g mL⁻¹) (FUJIFILM Wako Pure Chemical Corporation, Osaka, Japan) at 37 °C in a humidified atmosphere of 5% CO₂.

Cellular Uptake Measurements. HeLa cells were seeded onto dishes (φ 60 mm) in a culture medium (2.5 × 10⁵ cells per mL, 4 mL) and grown at 37 °C in a 5% CO₂ incubator for 24 h. After rinsing twice with PBS(-), 4 mL of 1^X (25 μ M) was added to each dish containing a complete medium and incubated for 48 h. In the control experiment, 1^X was not added. Cellular uptake was measured one time for each of 1^X. The cells were peeled off with trypsin (0.25%) after rinsing twice with PBS(-) and suspended in PBS(-) (500 μ L) after being centrifuged and washed with PBS(-) (1 mL). The cell suspension (1 × 10⁶ cells) was lyophilized and treated with 70% HNO₃ (250 μ L) for 3 h at 60 °C. The solution diluted with H₂O (250 μ L) was used for ICP MS.

Lipophilicity Measurements.²² The lipophilicity of $\mathbf{1}^{X}$ was estimated from distribution coefficients in an octanol/water system determined by the standard shake-flask method. Each measurement was repeated three times. Concentrations of $\mathbf{1}^{X}$ in organic and aqueous layers were determined separately by measuring the absorbance on a UV–vis spectrophotometer after proper dilution where their respective molar extinction coefficients were used. The distribution coefficients were calculated from the equation of log Pow = log[$\mathbf{1}^{X}$]o/[$\mathbf{1}^{X}$]w.

Cytotoxicity Measurements.^{23–25} MTT Cell Proliferation Assay Kit (Cayman Chemical Company, Michigan, USA) was used to assess the cytotoxicity of $\mathbf{1}^{\mathbf{X}}$ and cisplatin by a standard 3-(4,5-dimethylthiazole)-2,5-diphenyl-tetrazolium bromide (MTT) assay. In 96-well microassay culture plates (1×10^4) cells per well), the cells were placed and grown in a humidified atmosphere of a 5% CO₂ incubator for 24 or 48 h at 37 °C. To each well was added 1^{Pn} at 1 to 400 μ M, 1^{Mn} at 1 to 1000 μ M, or cisplatin at 0.1 to 20 μ M. Only a culture medium was added to the blank and control well. Each well was rinsed twice with PBS(-) after incubation. A MTT dye solution (100 μ L, 0.5 mg mL⁻¹) was added, and after 3 h, DMSO (100 μ L) was added to solubilize MTT Formazan. The half maximal inhibitory concentrations (IC_{50}) were determined as previously reported.¹⁸ All experiments were repeated at least three times and expressed as the mean \pm standard deviation.

Fluorescence Imaging.²⁶ Onto observation dishes (φ 14 mm), HeLa cells in a culture medium (600 μ L) were seeded and grown in a humidified atmosphere of 5% CO₂ at 37 °C for 24 h. The dishes were rinsed twice with PBS(-). A solution of 1^{P1} , 1^{P2} , or 1^{P3} (200 μ M, 600 μ L) in the culture medium was added to the dishes and incubated at 37 °C for 1 h. After rinsing twice with PBS(-), to the dishes was added a solution of Mito Tracker Deep Red FM (Thermo Fisher) (50 nM, 600 μ L) in a complete culture medium for mitochondrial staining. The dishes were incubated at 37 °C for 10 min and observed on the confocal fluorescence microscope to obtain pictures. A Nikon NIS-Elements Ar was used for image analysis. The cellular uptake of 1^{P1-3} to HeLa cells was examined at 4 and 37 °C as follows. Onto observation dishes (φ 14 mm) were seeded HeLa cells in a culture medium (600 μ L) and grown at 37 °C for 24 h. A 600 μ L solution of 1^{P1} (200 μ M), 1^{P2} (500 μ M), or 1^{P3} (800 μ M) in the culture medium was added to the dishes after rinsing twice with PBS(-). The dishes were incubated at 37 or 4 °C for 1 h and observed on the confocal fluorescence microscope.

Apoptosis Assay.²⁷ Onto six-well microassay culture plates $(3 \times 10^5$ cells per well) were seeded HeLa cells in a culture medium (2 mL) and grown in a humidified atmosphere of a 5% CO₂ incubator at 37 °C for 24 h. A solution (2 mL) of 1^{P1} , $1^{P3}\text{, or }1^{M1-3}$ (each IC_{50} value at 24 h) was added to each well and incubated for 1 or 12 h. The cells were peeled off with trypsin (0.25%) after rinsing twice with PBS(-), washed with PBS(-) (2 mL), and collected by centrifugation. A solution (100 μ L) of 1 × 10⁶ cells per mL was prepared with 1× BB. To the solution was added Annexin V-FITC solution (5 μ L) and incubated at rt for 10 min. The solution was centrifuged and aspirated. To the cells was added $1 \times BB$ (500 μL), suspended, and collected by centrifugation. To the cells was added $1 \times BB$ (200 μ L). To the solution was added PI (5 μ L) and suspended. The cell suspension (10,000 cells) was observed using channel BL1 (filter, 530/30; filter range, 515/545) and

Scheme 2. Synthetic Route of HL^{X} (X = P1-3 and M1-3)



channel BL2 (filter, 574/26; filter range, 561–587) of blue (488 nm) under an Attune NxT Acoustic Focusing Cytometer. *Caspase-9 Activity Assay*.^{28,29} CaspGLOW Fluorescein

Caspase-9 Activity Assay.^{28,29} CaspGLOW Fluorescein Active Caspase-9 Kit (Thermo Fisher Scientific, Tokyo, Japan) was used for the measurement. HeLa cells were grown with the six-well micro assay culture plates as described above. To each well, 1 or 1^X (2 mL) was added to the final concentration of 97.3 μ M (IC₅₀ of 1^{P1} at 24 h) and incubated at 37 °C for 1 or 12 h. The cells were peeled off with trypsin (0.25%) after rinsing twice with PBS(-). The cells were collected by centrifugation. The cell suspension (1 × 10⁶ cells per mL, 300 μ L) was prepared with a culture medium. To the suspension was added a solution of FITC-LEHD-FMK (FITC-Leu-Glu-His-Asp-fluoromethyl ketone) (1 μ L) and incubated for 60 min. After being collected by centrifugation, the cells were washed with buffer (500 μ L). The cell suspension (10,000 cells) was observed using channel BL1 (filter, 530/30; filter range, 515/545) of blue (488 nm) under an Attune NxT Acoustic Focusing Cytometer. The cells were gated to FSC and SSC.

Caspase-3/7 Activity Assay.³⁰ The measurements were carried out using CellEvent Caspase-3/7 Green Flow Cytometry Assay Kit (Thermo Fisher Scientific, Tokyo, Japan). HeLa cells were prepared as described in Caspase-9 Activity Assay. 1 or 1^{X} (2 mL) was added to each well to the final concentration of 97.3 μ M (IC₅₀ of 1^{P1} at 24 h). The cell suspension (1 × 10⁶ cells per mL, 1 mL) was prepared with PBS(-). To the suspension was added CellEvent Caspase-3/7 Green Detection Reagent (1 μ L) and incubated at 37 °C for 25 min. After being washed and collected by centrifugation, the cells were suspended with PBS(-) (1 mL). The cell suspension (10,000 cells) was observed using channel BL1

(filter, 530/30; filter range, 515/545) and channel BL3 (filter, 595/40; filter range, 675–715) of blue (488 nm) under an Attune NxT Acoustic Focusing Cytometer.

RESULTS AND DISCUSSION

Preparations of DNA Target/Ligand Conjugates HL^X and Their Copper(II) Complexes 1^x. The synthetic routes of the DNA target/ligand conjugates $[HL^X (X = Pn \text{ and } Mn, n]$ = 1-3] are shown in Scheme 2. In the beginning, three different lengths of PEG-linkers (n = 1-3) were attached to 2,6-diformyl-4-caroboxyphenol through the amide bond. In the next steps, tri-Boc cyclen and terminal phenanthrenyl (P) or methyl (M) groups were introduced stepwise. Boc groups were removed by acidic solvolysis in the final stage. The resultant DNA target/ligand conjugates HL^X form dicopper(II) complexes $[Cu_2(\mu-OH)(L^X)](ClO_4)_2$ $[1^X, X = P1 (1^{P_1}), P2 (1^{P_2}), P3 (1^{P_3}), M1 (1^{M_1}), M2 (1^{M_2}), and M3 (1^{M_3})]$ upon reaction with $Cu(ClO_4)_2$ at pH 8.0 adjusted by aqueous NaOH (1 M). The isolated solid was purified by recrystallization from MeOH/Et₂O. The synthetic details, elemental analysis, electronic absorption, IR, and ESI MS data of HL^X and 1^X are shown in Experimental Section and the Supporting Information, except for HLP2 and 1P2 reported previously.

Structures of Dicopper Complexes 1^X. Structures of $[Cu_2(\mu-OH)(L^X)](ClO_4)_2$ (1^X) were examined by the electronic absorption spectra in Figure S1, and the ESI MS spectra are shown in Figures S2–S6. Structures of 1 and 1^{P2} were previously reported.^{15,16} In the electronic spectra of 1^X, two bands appeared at 380 and 640 nm assignable to PhO⁻ to Cu(II) LMCT and d–d transition bands (see Experimental Section), close to those of 1 and 1^{P2} previously reported.^{17,18} The PhO⁻ to Cu(II) LMCT bands are similar to the bands at 340–400 nm reported for the μ -OH- μ -OPh dicopper(II) complexes.^{17,18,31,32} The moleculer ion peaks of 1^{P1}, 1^{P3}, 1^{M1}, 1^{M2}, and 1^{M3} in the ESI MS spectra appear at m/z 1065.17, 1153.32, 903.28, 947.32, and 991.38 corresponding to $[L^X + 2Cu(II) + OH + ClO_4]^+$, respectively. All spectral data demonstrate that a common μ -OH- μ -OPh dicopper(II) core structure of 1^X is kept in MeCN and aqueous solutions.

DNA Binding Constants of 1^X. Apparent binding constants K_{app} (M⁻¹) of 1^X with calf thymus DNA (ct-DNA) were estimated by an ethidium bromide method.^{21,22} The fluorescence emission intensity decreases upon addition of 1^X as shown in Figure S7. Figure 1 shows plots of intensity at 610 nm vs concentrations of 1^{P1-3} and DNA binding constants K_{app} (M⁻¹) of 1^X in comparison with 1. The DNA binding constants of 1^{P1-3} and 1^{M1-3} are 6.5–9.1 and 3.5–4.6 times larger than that of 1, respectively, showing that the P-linkers are two-fold more effective as DNA targets than the M-linkers. The DNA binding ability of 1^X is enhanced by a phenanthrenyl group that may serve as an intercalator and slightly by PEG-linkers. The DNA binding abilities of 1^{P2} and 1^{M2} having the PEG-linker (n = 2) may be slightly enhanced by antiorientation of the dicopper moiety and phenathrenyl or methyl group in the transoid conformation of the PEG-linker.

Spectroscopic Identification of μ -1,1-Hydroperoxo Complexes. Previously, we reported that μ -1,1-hydroperoxo complexes $[Cu_2(\mu$ -O₂H)(L)]^{2+} (2) and $[Cu_2(\mu$ -O₂H)(L^{P2})]^{2+} (2^{P2}) are formed from 1 and 1^{P2} upon reaction with H₂O₂ to accelerate the DNA cleavage and identified by the resonance Raman, cold-spray ionization (CSI) MS, and electronic absorption spectra.^{17,18}



Figure 1. (A) Plots of fluorescence emission intensity at 601 nm vs concentrations of 1 (red), 1^{P1} (light green), 1^{P2} (light blue), and 1^{P3} (purple). (B) DNA binding constants K_{app} (×10⁶ M⁻¹) estimated from these data. Conditions: [complex] = 0-400 μ M, [ct-DNA] = 20 μ M bp, [EtBr] = 3.3 μ M, [NaCl] = 10 mM, [buffer] = 10 mM (pH 6.0, MES), $\lambda_{ex} = 510$ nm at 37°C.

In this study, μ -1,1-hydroperoxo complexes [Cu₂(μ -O₂H)-(L^X)]²⁺ (2^X, X = P1 and P3) (Scheme 3) are identified by the

Scheme 3. Chemical Structures of 2^{X} (X = P1 and P3)



electronic absorption (Figure S8) and CSI MS spectra (Figures S9–S12). The electronic absorption spectra of 2^{P1} and 2^{P3} in MeCN at -30 °C show two bands at 340 ($\varepsilon = 5800$) and 400 ($\varepsilon = 3100$) nm and at 340 ($\varepsilon = 5600$) and 400 ($\varepsilon = 3200 \text{ M}^{-1} \text{ cm}^{-1}$) nm, respectively. These are almost the same as 340 ($\varepsilon = 5600$) and 398 ($\varepsilon = 4800 \text{ M}^{-1} \text{ cm}^{-1}$) nm of **2**. The CSI MS spectra of 2^{P1} and 2^{P3} in H₂O at 0 °C show two major peaks at m/z 491 and 1081 corresponding to $[L^{P1} + 2Cu(II) + O_2H]^{2+}$ and $[L^{P1} + 2Cu(II) + O_2H + ClO_4]^+$ and at m/z 535 and 1169 corresponding to $[L^{P3} + 2Cu(II) + O_2H]^{2+}$ and $[L^{P3} + 2Cu(II) + O_2H + ClO_4]^+$, respectively. When ¹⁸O-labeled H₂¹⁸O₂ was used instead of H₂¹⁶O₂, these peaks shifted by four mass units (Figures S10 and S12), showing that two O-atoms of 2^{P1} and 2^{P3} come from H₂O₂. These show that the common μ -1,1-hydroperoxo species are formed from 1^X upon reaction with H₂O₂ for the DNA cleavage.

DNA Cleavage by 1^x. Using supercoiled plasmid pUC19 DNA (Form I) as a substrate, the DNA cleavage activity of 1^x with H_2O_2 was examined at pH 6.0 at 37 °C under the reaction conditions shown in Experimental Section and compared with that of a dicopper complex 1 to clarify roles of the DNA targets. Forms II and III, nicked circular and linear doublestrand DNA, respectively, were produced from Form I.³³ These are analyzed by agarose gel electrophoresis. The time courses for the decrease of Form I, gel patterns, and rates (%) of Forms I, II, and III are shown in Figures S13, S14, and S19 and Table S1, respectively. Time courses for the increase of Form III (%) in the DNA cleavage by 1^{P1-3} and 1^{M1-3} are shown in comparison with 1 in Figure 2A,B, respectively.



Figure 2. Time courses for the increase of Form III (%) in the reaction of pUC19 DNA (50 μ M bp) with the complex (50 μ M) and H₂O₂ (0.5 mM) at pH 6.0 (MES, 10 mM) at 37°C. (A) 1^{P1} (right green), 1^{P2} (right blue), 1^{P3} (purple), and 1 (red) and (B) 1^{M1} (right green), 1^{M2} (right blue), 1^{M3} (purple), and 1 (red). Experiments were repeated at least three times.

The Form I decay rates obey the first-order kinetics as shown in Figure S15. The pseudo-first-order rate constants of 3.2, 7.1, 3.5, and 2.5×10^{-2} min⁻¹ for 1^{P1} , 1^{P2} , 1^{M1} , and 1^{M2} , respectively, are slightly larger than 2.3×10^{-2} min⁻¹ of 1, but 2.3 and 1.5×10^{-2} min⁻¹ of 1^{P3} and 1^{M3} are smaller. These data show that conversion of Form I to II is only slightly accelerated by short linkers but not by long ones. Meanwhile, as shown in Figure 2, Form III production is greatly accelerated by 1^{X} and by shortening the linker. It accompanies a large induction decay because conversion of Form I to II is much faster due to the distorted supercoiled structure of Form I than that of Form II to III. Therefore, short linkers specifically accelerate the slow conversion of Form II to III.

Form III is produced via DNA double-strand breaks (dsb) of opposed strands within the 16 base pairs.³⁴ Thus, to clarify roles of DNA targets in the DNA dsb by 1^{X} , the yield of Form III (%) in the 3 h reaction by 1^{X} or 1 and enhancement rates $(1^{X}/1)$ in the DNA dsb are shown in Table 1. P-linkers two-

Table 1. Yield of Form III (%) in the 3 h Reaction by 1^{X} or 1 and Enhancement Rates $(1^{X}/1)$ in the DNA dsb

complex	Form III (%) in 3 h	enhancement rate (1 ^X /1) in DNA dsb	complex	Form III (%) in 3 h	enhancement rate (1 ^X /1) in DNA dsb
1 ^{P1}	47 ± 1	22	1 ^{M1}	23 ± 3	11
1 ^{P2}	23 ± 4	11	1 ^{M2}	9 ± 2	4
1 ^{P3}	7 ± 1	3	1 ^{M3}	4 ± 1	2
1	2 ± 1	1	1	2 ± 1	1

fold enhance DNA dsb than M-linkers in the same linker size. This is consistent with the fact that DNA binding constants of $1^{P_{1-3}}$ are two-fold larger than those of $1^{M_{1-3}}$, showing that DNA dsb by 1^{X} is accelerated by DNA binding. Moreover, $1^{P_{1}}$ gives the highest yield 47% of Form III in the 3 h reaction in 1^{X} , and enhancement rates $(1^{X}/1)$ in the DNA dsb increase with shortening of the linkers. Thus, the short P-linker of $1^{P_{1}}$ is the optimized DNA target for the DNA dsb. These results suggest that the short P-linker of $1^{P_{1}}$ fixes the dicopper moiety within a small number of base pairs via intercalation of the phenanthrenyl group to facilitate the second break of a closely opposed strand to the first break in the DNA dsb.

To gain some insights into the active species, we carried out inhibition experiments for the DNA cleavage by 1^{P1} using KI (H₂O₂ scavenger) and DMSO (HO• scavenger). The results are shown in Figure S21. Either the decrease in Form I or the increase in Form III was not inhibited in the presence of DMSO (1.0, 5.0, 10 mM). Thus, a diffusible HO• is not involved in the DNA cleavage. Meanwhile, KI (1.0, 5.0, and 10 mM) retarded both the decrease in Form I and the increase in Form III. This is caused by reduction of H₂O₂ by KI. Thus, HO• and H₂O₂ are not direct oxidants for the DNA cleavage. Therefore, a Cu-bound peroxide species may directly attack DNA.

Cellular Uptake and Lipophilicity of 1^X. The intracellular toxic ability of 1^x depends on the cellular uptake. Thus, the cellular uptake of 1^{X} (25 μ M) to HeLa cells (2.5 \times 10⁵ cells/mL) after 48 h incubation was determined by measuring Cu contents in cells and the culture medium with ICP MS³ and compared with that of 1 to clarify roles of the DNA targets in the cytotoxicity of 1^{X} . Rates of [Cu in cells]/[Cu in culture medium] for 1, 1^{P1} , 1^{P2} , 1^{P3} , 1^{M1} , 1^{M2} , and 1^{M3} are 2.04, 4.92, 4.38, 3.70, 3.55, 1.72, and 1.25%, showing that enhancement rates $(1^{X}/1)$ in the cellular uptake of 1^{P1} , 1^{P2} , 1^{P3} , 1^{M1} , 1^{M2} , and 1^{M3} are 2.41, 2.15, 1.81, 1.74, 0.84, and 0.61, respectively. The cellular uptake of 1^{P1-3} is higher than that of 1^{M1-3} . The former decreases only to 75% by elongating linkers where the phenanthrenyl group is dominant but the latter to 35% where the linker size is dominant. Since lipophilicity may influence cellular uptake, the lipophilicity of $\mathbf{1}^{\hat{\mathbf{X}}}$ was estimated from the distribution coefficients (log P_{ow})²² in an octanol/water system. The log P_{ow} values of 1, 1^{P1}, 1^{P2}, 1^{P3}, 1^{M1}, 1^{M2}, and 1^{M3} are -2.38, -1.22, -1.14, -0.98, -1.39, -1.22, and -1.11, respectively, showing that the lipophilicity of $\mathbf{1}^{X}$ is enhanced by the P- and M-linkers and slightly by elongating the linker. These suggest that the cellular uptake of 1^{P_1-3} is enhanced by the membrane permeability of the phenanthrenyl group but not by the lipophilicity.

Table 2. In Vitro Cytotoxicity of 1, 1^x, and Cisplatin against Various Cells in MTT Assay (48 h) (Mean ± SD)

	IC_{50} (μ M) (mean ± SD)							
	cervical	lung		pancreas				
complex	HeLa (cancer)	A549 (cancer)	WI-38 (normal)	SF	PK-59 (cancer)	2C6 (normal)	SF	
1	660 ± 28	284 ± 16	838 ± 82	3.0	241 ± 4	1140 ± 50	4.7	
1 ^{P1}	21.8 ± 6.5	20.9 ± 0.6	215 ± 20	10.3	11.8 ± 2.5	90.4 ± 2.8	7.7	
1 ^{P2}	56.6 ± 1.8	23.8 ± 2.4	218 ± 3	9.2	18.4 ± 1.6	104 ± 8	5.7	
1 ^{P3}	66.4 ± 2.7	71.7 ± 1.4	268 ± 3	3.8	57.4 ± 2.0	128 ± 1	2.2	
1 ^{M1}	199 ± 11	206 ± 7	431 ± 32	2.1	147 ± 1	552 ± 5	3.8	
1 ^{M2}	257 ± 1	247 ± 18	527 ± 9	2.1	152 ± 9	657 ± 12	4.3	
1 ^{M3}	283 ± 22	279 ± 8	636 ± 16	2.3	161 ± 4	675 ± 5	4.2	
cisplatin	0.93 ± 0.01	1.83 ± 0.03	4.55 ± 0.03	2.5	2.01 ± 0.02	1.15 ± 0.08	0.6	

Cytotoxicity of 1^x. The cytotoxicity of 1^x against various cells was examined by MTT assay.²³⁻²⁵ IC_{50} values of 1^x determined by plots of cell viability vs $\log[1^X]$ after 24 and 48 h incubations are shown in Table S2 and Table 2, respectively. The IC₅₀ data show that cytotoxicity is kept intact after 48 h. Moreover, the stability of the dicopper complexes 1^{P1} and 1^{M1} under the incubation conditions used in this study was monitored by the electronic absorption spectra in 72 h as shown in Figure S20. The spectra of 1^{P1} and 1^{M1} did not change significantly, showing that these complexes are kept intact during the incubation. IC₅₀ data (24 h) of HL^X against all cell lines used in this study are larger than 1000 μ M as shown in Table S2, showing that HL^X are much less cytotoxic than the dicopper complexes. Moreover, we carried out MTT assay in the presence of HL^{P1} and $Cu(ClO_4)_2$. The IC_{50} data (24 h) against all cell lines used in this study are larger than 1000 μ M as shown in Table S2. Therefore, HL^{P1} plus Cu(II) ion are much less cytotoxic than the dicopper complexes. IC_{50} data (48 h) of 1^{P1}, 1^{P2}, 1^{P3}, and 1 against HeLa cells are 21.8, 56.6, 66.4, and 660 μ M, respectively, showing that enhancement rates $(1^{X}/1)$ in the cytotoxicity of 1^{P1} , 1^{P2} , 1^{P3} are 30, 12, and 9.9, respectively. These are much larger than enhancement rates in cellular uptake. Thus, the cytotoxicity of $\mathbf{1}^{X}$ is greatly enhanced by the P-linkers and by shortening the linker. This may be correlated to enhancement rates $(1^{X}/1)$ in DNA dsb enhanced by shortening the linker. However, IC₅₀ data (48 h) of 199, 257, and 283 μ M of 1^{M1}, 1^{M2}, and 1^{M3}, respectively, show that the cytotoxicity of 1^{M1-3} against HeLa cells is only 2to 3-fold enhanced than that of 1 and almost independent of the linker size.

To clarify roles of P- and M-linkers in DNA dsb by $\mathbf{1}^{X}$ in HeLa cells, expected enhancement rates $(\mathbf{1}^{X}/\mathbf{1})$ in DNA dsb in HeLa cells (A) × (C) are estimated from enhancement rates $(\mathbf{1}^{X}/\mathbf{1})$ in DNA dsb (A) and enhancement rates $(\mathbf{1}^{X}/\mathbf{1})$ in cellular uptake to HeLa cells (C) as shown in Table 3. Plots of

Table 3. Data for the Plot of Expected Enhancement Rates $(1^X/1)$ in DNA dsb in HeLa Cells (A) × (C) vs Enhancement Rates $(1^X/1)$ in Cytotoxicity against HeLa Cells (B)^{*a*}

complex	(A) enhancement rate (1 ^x /1) in DNA dsb	 (B) enhancement rate (1^X/1) in cytotoxicity against HeLa cells 	(C) enhancement rate $(1^{X}/1)$ in cellular uptake	$(A) \times (C)$		
1	1	1	1	1		
1 ^{P1}	22	30	2.4	52.8		
1 ^{P2}	11	12	2.1	23.1		
1 ^{P3}	3	9.9	1.8	5.4		
1 ^{M1}	11	3.3	1.7	18.7		
1 ^{M2}	4	2.6	0.84	3.36		
1 ^{M3}	2	2.3	0.61	1.22		
^{<i>a</i>} Enhancement rates $(1^{X}/1)$ in DNA dsb (A) and enhancement rates						

 $(1^{X}/1)$ in cellular uptake (C) are used to estimate (A) × (C).

the expected enhancement rates $(1^{X}/1)$ in DNA dsb in HeLa cells (A) × (C) vs enhancement rates in cytotoxicity against HeLa cells (B) are made for 1^{P1-3} and 1^{M1-3} in Figure 3. The plots for 1^{P1-3} and 1^{M1-3} give straight lines with slopes 0.44 and 0.053, respectively. These values show that cytotoxicity and DNA dsb are 44% correlated for 1^{P1-3} but only 5% for 1^{M1-3} . Thus, P-linkers proportionally enhance DNA dsb and cytotoxicity whereas M-linkers only DNA dsb but not cytotoxicity, suggesting that P-linkers promote DNA dsb via



Figure 3. Plots of expected enhancement rates in DNA dsb in HeLa cells (A) \times (C) shown in Table 3 vs enhancement rates in cytotoxicity against HeLa cells (B) for $1^{P_{1-3}}$ (red line) and $1M^{1-3}$ (purple line). The slopes of the red and blue lines are 0.44 and 0.053, respectively.

specific DNA binding by intercalation of the phenanthrenyl group in the cells to enhance cytotoxicity, but M-linkers do not due to nonspecific binding.

The cytotoxicity of 1^X against A549 (lung cancer), WI-38 (lung normal), PK-59 (pancreas cancer), and 2C6 (pancreas normal) is shown by the IC_{50} data (48 h) in Table 2. IC_{50} data (48 h) of 1, 1^{P1} , 1^{P2} , and 1^{P3} against A549 lung cancer cells are 284, 20.9, 23.8, and 71.7 μ M, respectively, showing that the cytotoxicity of 1^{P1-3} against the cancer cells is largely enhanced by P-linkers and by shortening the linker. However, based on IC_{50} data (48 h) of 215, 218, and 268 μ M for 1^{P1} , 1^{P2} , and 1^{P3} against WI-38 lung normal cells, respectively, the cytotoxicity of 1^{P1-3} against normal cells is not so greatly enhanced by Plinkers and independent of the linker size. These results show that 1^{P1-3} exert cancer cell-selective cytotoxicity. Moreover, cancer cell-selective cytotoxicity is evaluated by selectivity factor (SF) = $(IC_{50} \text{ of normal cells})/(IC_{50} \text{ of cancer cells})$ shown in Table 2. SF values of 1^{P1}, 1^{P2}, and 1^{P3} for A549 lung cancer and WI-38 lung normal cells are 10.3, 9.2, and 3.8, respectively, which are much higher than 2.3-2.1 of 1^{M1-3} and 2.5 of cisplatin. Thus, 1^{P1-3} exert cancer cell-selective cytotoxicity, but either 1^{M1-3} or cisplatin do not. The short P-linker enhances cancer cell-selective cytotoxicity in 1^{P1-3}, but there is no linker size effect in 1^{M1-3} . Similarly, 1^{P1-3} are cancer cell-selective in cytotoxicity against PK-59 pancreas cancer and 2C6 pancreas normal cells, but 1^{M1-3} are not (see Table 2). SF values of 1^{P1} for lung and pancreas cancer and normal cells are 10.3 and 7.7, respectively, showing that 1^{P1} exerts the highest cancer cell-selective cytotoxicity in 1^x. Thus, the short P-linker is the DNA target optimized for cancer cell-selective cytotoxicity.

Cytoplasmic Localization of 1^{P1-3}. To gain some insight into the mechanism of the cancer cell-selective cytotoxicity of 1^{P1-3}, their cytoplasmic localization was examined by means of confocal microscopy. After 1^{P1-3} (200 μ M) were added to HeLa cells in a culture medium and incubated for 1 h, and the cells were treated with Mito Tracker Deep Red (50 nM) as the mitochondrial staining dye, the cytoplasmic localization was observed from the merged image. The confocal microscopic images are shown in Figure 4. The red and blue colors show stained mitochondria and the presence of 1^{P1-3} in the cell, respectively. The merged image clearly shows that 1^{P1-3} are localized at mitochondria as shown by the purple color. The image processing using ImageJ shows that 17% of area in mitochondria is occupied by 1^{P1-3} as shown in Table S6.



Figure 4. Confocal microscopic images of $1^{P_{1-3}}$ (200 μ M) in HeLa cells in the dark on 1 h incubation. Bright-field images (A, E, I). Blue fluorescence indicates the fluorescence of 1^{P_1} (B), 1^{P_2} (F), and 1^{P_3} (J) (λ_{ex} = 405 nm). Red fluorescence is mitochondrial staining by Mito Tracker Deep Red FM (50 nM) (Thermo Fisher) (C, G, K) (λ_{ex} = 640 nm). (D, H, L) Overlay images of panels (A)–(C), (E)–(G), and (I)–(K), respectively. The scale bar is 20 μ m.

Moreover, the measurements on confocal microscopy were carried out at 4 and 37 °C as shown in Figure S16. The blue color of 1^{P1-3} was observed in mitochondria at 4 °C similarly to the fluorescence imaging observed at 37 °C. This suggests that the cellular uptake of 1^{P1-3} occurs mainly via direct membrane permeation but not via endocytosis. This is consistent with the fact that the cellular uptake of 1^{P1-3} is promoted by the membrane permeability of the P-linkers as described above. For the intracellular toxicity, direct membrane permeation is more favorable than endocytosis.

Apoptosis Assay. Fluorescence-activated cell sorting was carried out using Annexin V-FITC/PI²⁷ by flow cytometry to determine the mode of cell death. To quantify the apoptotic cells by flow cytometry, 1^{P1} , 1^{P3} , or 1^{M1-3} (each IC₅₀ value at 24 h) were added to HeLa cells and incubated for 1 and 12 h. Assay was performed using staining solutions of Annexin V-FITC and PI. Dot plots and rates of Q1–Q4 obtained are shown in Figure S17 and Tables S3–S5. As previously reported, in the presence of 1^{P2} , 15.8% of cells were dead in 1 to 12 h incubation, and 85% were early apoptosis.³⁶ In the case of 1^{P1} and 1^{P3} , 27.1 and 18.8% of cells were dead in 1 to 12 h incubation, where nearly 100 and 89% were early apoptosis induction rates, in which 1^{P1} exerts the highest cytotoxicity induced by 100% apoptosis. 1^{M1-3} , however, show lower cytotoxicity than 1^{P1-3} based on the Q1–Q4 rates shown in Table S5.

Caspase Activity Assay. Since 1^{P1-3} localize at mitochondria to induce apoptotic cell death as described above, caspase activity assays of 1^{X} were carried out for the initiator caspase- $9^{37,38}$ and effector caspase- $3/7^{37,38}$ to clarify whether the apoptosis proceeds via an intrinsic pathway. A Leu-Glu-His-Asp (LEHD) sequence is recognized by the activated caspase- 9^{28} as a cleavage site of proteins, and thus, a nontoxic and cell-permeable FITC-LEHD-FMK (FITC-Leu-Glu-His-Asp-fluoromethyl ketone) can be used to evaluate the activity of caspase- $9^{.29}$ HeLa cells were incubated with 1 or 1^{X} (97.3 μ M, IC₅₀ of 1^{P1} at 24 h) for 1 and 12 h. After that, the cells were separated and washed and incubated with FITC-LEHD-FMK

for 30 min. As described in Experimental Section, the caspase-9 activity was measured. Moreover, the CellEvent Caspase-3/7 Green Detection Reagent³⁰ of a peptide sequence Asp-Glu-Val-Asp was used to examine caspase-3/7 activity.³⁹ It emits a strong fluorescence via DNA binding when the sequence is cleaved. Thus, the caspase-3/7 activity is estimated from the fluorescence intensity. The activities of caspase-9 and caspase-3/7 are shown in Figure S18. In incubation for 12 h, 1^{P1-3} showed higher activity for caspase-9 and caspase-3/7 than 1^{M1-3} , and the activity 1^{P1} is the highest. Thus, 1^{P1} efficiently induces mitochondrial apoptosis by the intrinsic pathway.^{37,40,41} As shown above, 1^{P1} exerts the highest activity in the DNA dsb with H₂O₂ and the highest cancer cell-selective cytotoxicity. Therefore, it is proposed that the short P-linker of 1^{P1} enhances cancer cell-selective cytotoxicity by promoting the dsb of mitochondrial DNA with H2O2 increased by mitochondrial dysfunction in cancer cells. This means that $\mathbf{1}^{P_1'}$ induces mitochondrial apoptosis by the dsb of mitochondrial DNA in cancer cells but not in normal cells because of relatively low concentration of H₂O₂ in normal cells, leading to the cancer cell-selective cytotoxicity.

Comparison in Cancer Cell-Selective Cytotoxicity between 1^{Pn} and a Related Mitochondria-Targeting Copper Complex. A mononuclear copper complex with a terpyridine (ttpy) ligand attaching a tetraphenyl phosphonium (tpp) group as a mitochondrial target [Cu(ttpy-tpp)Br₂]Br (14) was reported to exert high cytotoxicity against cancer cells via mitochondrial apoptosis.^{7e} 14 exerted cytotoxicity to cancer cells by the oxidative DNA cleavage via ROS generation. The cancer cell-selective cytotoxicity of 14 is attained by high cellular uptake and mitochondrial localization. On the other hand, 1^{Pn} exerted the cancer cell-selective cytotoxicity by activating H_2O_2 because the H_2O_2 concentration is slightly higher in cancer cells due to the mitochondrial dysfunction than in normal cells. In fact, 1^{P1} exerted 10.3-fold higher cytotoxicity against lung cancer cells (A549) than lung normal cells (WI-38). Thus, 1^{Pn} showed promising cancer cell-selectivity, although 1^{Pn} was lower in cytotoxicity than 14. Moreover, the DNA cleavage activity and cytotoxicity of 1^{Pn} are proportionally enhanced by shortening the PEG tether, where the cancer cell-selective cytotoxicity of 1^{Pn} is also enhanced. This clearly shows that the cancer cell-selective cytotoxicity of 1^{Pn} can be controlled by the tether length. Therefore, the cancer cell-selective cytotoxicity of our dicopper complex has a chance to be further enhanced by optimizing the tether length and DNA target in future works.

CONCLUSIONS

A series of DNA target/ligand conjugates (HL^X, X = Pn and Mn, n = 1-3) and their dicopper complexes $[Cu_2(\mu - OH)(L^X)](ClO_4)_2$ (1^X) were synthesized. Roles of DNA targets in DNA binding, DNA cleavage, cellular uptake, and cytotoxicity were assessed. Conversion of Form II to III in DNA double-strand breaks (dsb) of 1^X was greatly promoted by short linkers fixing a dicopper moiety within a small number of base pairs. The cytotoxicity of 1^X was enhanced by P-linkers and by shortening the linker but not by M-linkers. DNA dsb and cytotoxicity are 44% correlated in 1^{P1-3} but only 5% in 1^{M1-3}. This suggested that P-linkers promote DNA dsb via specific DNA binding in the cells to enhance cytotoxicity, but M-linkers do not due to nonspecific binding. 1^{P1} exerted the highest cancer cell-selective cytotoxicity in 1^X for lung and pancreas cancer and normal cells. Intracellular visualization,

apoptosis assay, and caspase activity assay revealed that $1^{\rm P1-3}$ induced mitochondrial apoptosis. It is concluded that the short P-linker of $1^{\rm P1}$ enhances cancer cell-selective cytotoxicity via mitochondrial apoptosis. This is probably because the short P-linker promotes the dsb of mitochondrial DNA with $\rm H_2O_2$ increased by mitochondrial dysfunction in cancer cells. The clarification of roles of the DNA target may shed light on the development of new anticancer agents to reduce the heavy side effects.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acsomega.3c03387.

Detailed experimental procedures, Figures S1–S21, and Tables S1–S6 (PDF)

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M.H. mainly conducted experimental work and contributed to make Schemes and Figures. J.U. worked on measuring conformal microscopic images. Y.H. contributed discussion. The manuscript was written through contributions of all authors. All authors have given approval to the final version of the manuscript.

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Notes

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