Luminescent Iridium Complex-Peptide Hybrids (IPHs) for Therapeutics of Cancer: Design and Synthesis of IPHs for Detection of Cancer Cells and Induction of Their Necrosis-type Cell Death

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Figure S1. The results of MTT assay: cell viability of Jurkat cells (% of control at [Ir complex] = 0 μ M) in presence of TRAIL (150 ng/mL), **4**, **5** and **6** (75 μ M) after incubation at 37 °C for 16 h.



Figure S2. The results of competitive staining of Jurkat cells with **5** and **CP1** (luminescence microscopy images on Biorevo, BZ-9000, Keyence, x 40). (a-c) Jurkat cells were incubatied with **5** (10 μ M) at 37°C for 1 h. (d-f) Jurkat cells were incubated with **CP1** (100 μ M) at 37°C for 1 h and then with **5** (10 μ M) at 37°C for 1 h. Scale bar (white) = 10 μ m.



Figure S3. Flow cytometry analysis of competitive staining of Jurkat cells with **5** and **CP1**. The red histogram corresponds to Jurkat cells treated with **5** (5 μ M) at 37 °C for 1 h, the orange histogram corresponds to Jurkat cells treated with **CP1** (100 μ M) at 37 °C for 1 h and then **5** (5 μ M) at 37 °C for 1 h, the blue histogram corresponds to Jurkat cells treated with **CP1** (100 μ M) at 37 °C for 1 h and then **5** (2 μ M) at 37 °C for 1 h and the green histogram corresponds to the cells treated with **CP1** (100 μ M) at 37 °C for 1 h and then **5** (2 μ M) at 37 °C for 1 h and the green histogram corresponds to the cells treated with **CP1** (100 μ M) at 37 °C for 1 h and then **5** (2 μ M) at 37 °C for 1 h and the green histogram corresponds to the cells treated with **CP1** (100 μ M) at 37 °C for 1 h and then **5** (2 μ M) at 37 °C for 1 h.



Figure S4. Flow cytometry analysis of co-staining assay of anti-DR5 antibody and **5**. The blue histogram corresponds to the cells treated with anti-DR5 antibody (15 μ g/mL) at 4°C for 15 min, and the red histogram corresponds to the cells treated with anti-DR5 antibody (15 μ g/mL) at 4°C for 15 min and then with **5** (5 μ M) at 37°C for 1 h.



Figure S5. Flow cytometry analysis of co-staining of Jurkat cells with **5** and anti-DR5 antibody. The red histogram corresponds to the cells treated with only anti-DR5 antibody (15 μ g/mL) at 4 °C for 15 min (Method a of Chart 5 in text), the blue histogram corresponds to the cells treated with **5** (5 μ M) at 37 °C for 1 h and then anti-DR5 antibody (15 μ g/mL) at 4 °C for 15 min (Method c of Chart 5 in text), and the green histogram corresponds to the cells treated with **5** (10 μ M) at 37 °C for 1 h and then anti-DR5 antibody (15 μ g/mL) at 4 °C for 15 min (Method c of Chart 5 in text), and the green histogram corresponds to the cells treated with **5** (10 μ M) at 37 °C for 1 h and then anti-DR5 antibody (15 μ g/mL) at 4 °C for 15 min (Method c of Chart 5 in text).



Figure S6. Flow cytometry analysis of Jurkat cells incubated with **5** (5 μ M) and anti-DR5 antibody (15 μ g/mL). Red histograms correspond to the cells treated with **5** and anti-DR5 antibody, blue histograms correspond to the cells treated with only anti-DR5 antibody. (a) after 1 h incubation with **5** (Method c in Chart 5 (b) after 1 h incubation with **5** and then again 1 h incubation in fresh medium (Method d in Chart 5). (c) after 1 h incubation with **5** and then again 6 h incubation in fresh medium (Method d in Chart 5).



Figure S7. Luminescence microscopy images (Biorevo, BZ-9000, Keyence) of Jurkat cells, K562 cells and Molt-4 cells (x 40) stained with **5**. (a-c) Jurkat cells were incubated with **5** (10 μ M) at 37°C for 1 h. (d-f) K562 cells were incubated with **5** (10 μ M) at 37°C for 1 h. (g-i) Molt-4 cells were incubated with **5** (10 μ M) at 37°C for 1 h. (g-i) Molt-4 cells were incubated with **5** (10 μ M) at 37°C for 1 h. (g-i) Molt-4 cells were incubated with **5** (10 μ M) at 37°C for 1 h. Scale bar (white) = 10 μ m.



Figure S8. HPLC charts of (a,b) **5** after incubation at 37 °C in RPMI 1640 (MTG free) medium for 24 h. (c,d) only RPMI 1640 (MTG free) medium (without **5**) after incubation at 37 °C for 24 h. (Senshu Pak PEGASIL ODS 4.6 ϕ X 250 mm UV: 254 nm, Excitation: 366 nm, Emission: 520 nm, flow rate 1.0 mL/min with continuous gradient elution (20-50% solvent B, 0-30 min, linear) with solvent A (0.1% TFA in H₂O) and solvent B (0.1% TFA in MeCN).



Figure S9. The results of MTT assay of TRAIL (150 ng/mL) and **5** (75 μ M) in the presence and the absence of caspase inhibitor (Z-VAD-FMK), necroptosis inhibitor (Necrostatin-1) and oxidative stress induced necrosis inhibitor (IM-54): cell viability of Jurkat cells (% of control) after incubation at 37 °C for 24 h.



Figure S10. The results of flow cytometry assay of (a) TRAIL (150 ng/mL) and (b) **5** (75 μ M). The red histograms correspond to the Jurkat cells treated with TRAIL (150 ng/mL) or **5** (75 μ M) at 37 °C for 24 h and then with PI (30 μ M) at room temperature for 10-15 min. The blue histograms correspond to the Jurkat cells treated with caspase inhibitor (Z-VAD-FMK) (15 μ M) at 37 °C for 1h and then TRAIL (150 ng/mL) or **5** (75 μ M) at 37 °C for 24 h followed by the treatment with PI (30 μ M) at room temperature for 10-15 min. The green histograms correspond to the Jurkat cells not treated with TRAIL or **5**.



Figure S11. Luminescent microscopy images (Biorevo, BZ-9000, Keyence) of Jurkat cells (x 40) incubated with **5** (35 μ M) in presence of (aa-ad) 4-aminopyridine (10 μ M), (ba-bd) amiloride (100 μ M), (ca-cd) bafilomycin A1 (10 μ M), (da-dd) CCCP (40 μ M), (ea-ed) chloroquinine (10 μ M), (fa-fd)

phentolamine (10 μ M), (ga-gd) quinidine (100 μ M), (ha-hd) nicardipine (50 μ M), and (ia-id) verapamil (20 μ M) at 37 °C for 24 h. Scale bar (white) = 10 μ m.



Figure S12: ¹H NMR chart of NHS ester of Ir complex 7



Figure S13: ESI mass chart of NHS ester of Ir complex 7



Figure S14: ¹H NMR chart of Ir complex 8



Figure S15: ESI mass chart of Ir complex 8



Figure S16: ¹H NMR chart of NHS ester of Ir complex 8



Figure S17: ESI mass chart of NHS ester of Ir complex 8



Figure S19: ESI mass chart of Ir complex 9



Figure S20: HPLC analysis of CP1

Senshu Pak PEGASIL ODS 4.6 ϕ X 250 mm UV: 280 nm, flow rate 1.0 mL/min with continuous gradient elution (20-50% solvent B, 0-30 min, linear) with solvent A (0.1% TFA in H₂O) and solvent B (0.1% TFA in MeCN).





Figure S21: ¹H NMR chart of CP1



Figure S22: ESI Mass chart of CP1



Figure S23: HPLC analysis of CP2

Senshu Pak PEGASIL ODS 4.6 ϕ X 250 mm UV: 280 nm, flow rate 1.0 mL/min with continuous gradient elution (10-40% solvent B, 0-30 min, linear) with solvent A (0.1% TFA in H₂O) and solvent B (0.1% TFA in MeCN).

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Figure S24: ¹H NMR chart of CP2



Figure S25: ESI Mass chart of CP2



Senshu Pak PEGASIL ODS 4.6 ϕ X 250 mm UV: 280 nm, flow rate 1.0 mL/min with continuous gradient elution (10-40% solvent B, 0-30 min, linear) with solvent A (0.1% TFA in H₂O) and solvent B (0.1% TFA in MeCN).

Figure S26: HPLC analysis of CP3

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Figure S27: ¹H NMR chart of CP3



Figure S28: ESI Mass chart of CP3



Senshu Pak PEGASIL ODS 4.6 ϕ X 250 mm UV: 254 nm, Ex.: 366 nm, Em.: 520 nm flow rate 1.0 mL/min with continuous gradient elution (20-50% solvent B, 0-30 min, linear) with solvent A (0.1% TFA in H₂O) and solvent B (0.1% TFA in MeCN).

Figure S29: HPLC analysis of Ir complex-peptide hybrid 4



Figure S30: ¹H NMR chart of Ir complex-peptide hybrid 4



Figure S31: ESI Mass chart of Ir complex-peptide hybrid 4



Senshu Pak PEGASIL ODS 4.6 ϕ X 250 mm UV: 254 nm, Ex.: 366 nm, Em.: 520 nm flow rate 1.0 mL/min with continuous gradient elution (20-50% solvent B, 0-30 min, linear) with solvent A (0.1% TFA in H₂O) and solvent B (0.1% TFA in MeCN).

Figure S32: HPLC analysis of Ir complex-peptide hybrids 5



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Figure S33: ¹H NMR chart of Ir complex-peptide hybrids 5



Figure S34: ESI Mass chart of Ir complex-peptide hybrid 5



Senshu Pak PEGASIL ODS 4.6 ϕ X 250 mm UV: 254 nm, Ex.: 366 nm, Em.: 520 nm flow rate 1.0 mL/min with continuous gradient elution (10-40% solvent B, 0-30 min, linear) with solvent A (0.1% TFA in H₂O) and solvent B (0.1% TFA in MeCN).

Figure S35: HPLC analysis of Ir complex-peptide hybrids 6





Figure S36: ¹H NMR chart of Ir complex-peptide hybrids 6



Figure S37: ESI Mass chart of Ir complex-peptide hybrid 6