Electronic Supporting Information

Anticancer application of half-sandwich iridium(III) ferrocene-

thiosemicarbazide Schiff base complexes

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Experimental Section

X-ray Crystallography

All diffraction data were obtained on a Bruker Smart Apex CCD diffractometer equipped with graphite-monochro mated Mo Ka radiation. Absorption corrections were applied using SADABS program. SQUEEZE option was used to remove the nonlocalized electron density at the final step of structure refinement. The structures were solved by direct methods using SHELXS (TREF) with additional light atoms found by Fourier methods. Complexes were refined against F2 using SHELXL, and hydrogen atoms were added at calculated positions and refined riding on their parent atoms. Xray crystallographic data for IrFc2 are available in Tables S1 and S2 and has been deposited in the Cambridge Crystallographic Data Centre under the accession numbers CCDC 2259306. IrFc2 was grown by slow diffusion of hexane into a saturated dichloromethane solution. X-ray crystallographic data in CIF format are available from the Cambridge Crystallographic Data Centre.

NMR Spectroscopy

¹H NMR spectra were acquired in 5 mm NMR tubes at 298K on Bruker DPX 500 (¹H =500.13 MHz) spectrometers. ¹H NMR chemical shifts were internally referenced to $(CHD_2)(CD_3)_2SO$ (2.50 ppm) for DMSO-d₆ and CHCl₃ (7.26 ppm) for chloroform-d₁. Additional, organic solvents were utilized in the synthesis and purification, which were maintained in ¹H NMR, including methanol (3.32 ppm), dichloromethane (5.30 ppm), and *n*-hexane (0.88 ppm) All data was carried out using XWIN-NMR version 3.6 (Bruker UK Ltd.).

UV-Vis Spectroscopy

The UV-Vis spectra of these compounds were recorded by TU-1901 UV spectrophotometer with 1 cm path-length quartz cuvettes (3 mL). Spectra were processed using UV Winlab software. Experiments were carried out at 298 K unless otherwise stated.

Stability Studies

The hydrolysis characteristics of IrFc1~IrFc6 (ca.250 μ M) in a 20% DMSO/80% H₂O (ν/ν) solution at 298 K for 8 h were evaluated by UV-Vis absorption spectra, and the presence of DMSO ensured the solubility of these complexes.

Reaction with NADH

The reaction of IrFc1~IrFc6 (ca. 1 μ M) with NADH (ca. 100 μ M) in 20% MeOH/80% H₂O (ν/ν) was monitored by UV-Vis at 298 K after various time intervals. TON was calculated from the difference in NADH concentration after 8 h divided by the concentration of iridium catalyst. The concentration of NADH was obtained using the extinction coefficient $\varepsilon_{339} = 6220 \text{ M}^{-1} \text{ cm}^{-1}$.

Cell Culture

Both human cancer cells (cervical carcinoma HeLa cells and lung cancer A549 cells) were obtained from Shanghai Institute of Biochemistry and Cell Biology (SIBCB) and were grown in Dubelco's Modified Eagle Medium (DMEM). All media were supplemented with 10% fetal bovine serum, and 1% penicillin-streptomycin solution. All cells were grown at 310 K in a humidified incubator under 5% CO₂ atmosphere.

Anti-proliferative assay (MTT assay)

After plating 5000 cells per well in 96-well plates, the cells were preincubated in drugfree media at 310 K for 24 h before adding different concentrations of the compounds to be tested. In order to prepare the stock solution of the drug, the solid compound was dissolved in DMSO. This stock was further diluted using cell culture medium until working concentrations were achieved. The drug exposure period was 24 h. Subsequently, 15 μ L of 5 mg mL⁻¹ MTT solution was added to form a purple formazan. Afterwards, 100 μ L of dimethyl sulfoxide (DMSO) was transferred into each well to dissolve the purple formazan, and results were measured using a microplate reader (DNM-9606, Perlong Medical, Beijing, China) at an absorbance of 570 nm. Each well was triplicated and each experiment repeated at least three times. IC₅₀ values quoted are mean ± SEM.

Measurement of Lipophilicity (logPo/w)

The octanol-water partition coefficients ($P_{a/w}$) of complexes were determined using a shake-flask method. Water (50 mL, distilled after milli-Q purification) and 1octanol (50 mL, vacuum distilled) were shaken together using a laboratory shaker, for 72 h to allow saturation of both phases. Stock solutions of IrFc7 and IrFc8 (50.0 μ M) were prepared in the aqueous phase and aliquots (5 mL) of each of these stock solutions were then added to an equal volume of the 1-octanol phase. The resultant biphasic solutions were mixed for 2 h and then centrifuged (3000×g, 5 min) to separate the phases. The concentrations of IrFc7 and IrFc8 in the organic and aqueous phases were then determined using UV-Vis. Log $P_{a/w}$ was defined as the logarithm of the ratio of the concentrations of the compound in the organic and aqueous phases (Values reported are the means of three separate determinations).

ROS Determination

Flow cytometry analysis of ROS generation in A549 cells caused by exposure to iridium compounds were carried out using the Reactive Oxygen Species Assay Kit (Beyotime Institute of Biotechnology, Shanghai, China) according to the supplier's instructions. Briefly, 1.5×10^6 A549 cancer cells per well were seeded in a six-well plate.

Cells were preincubated in drug-free media at 310 K for 24 h in a 5% CO₂ humidified atmosphere, and then drugs were added at concentrations of $1.0 \times IC_{50}$ and $2.0 \times IC_{50}$. After 24 h of drug exposure, cells were washed twice with PBS and then incubated with the DCFH-DA probe (10 μ M) at 310 K for 30 min, and then washed triple immediately with PBS. The fluorescence intensity was analyzed by flow cytometry (ACEA NovoCyte, Hangzhou, China). Data were processed using NovoExpressTM software.

Cell Cycle Analysis

A549 cancer cells at 1.5×10^6 per well were seeded in a six-well plate. Cells were preincubated in drug-free media at 310 K for 24 h, after which IrFc8 was added at concentrations of $0.5 \times IC_{50}$, $1.0 \times IC_{50}$ and $2.0 \times IC_{50}$ against A549 cells. After 24 h of exposure, supernatants were removed by suction and cells were washed with PBS. Finally, cells were harvested using trypsin-EDTA and fixed for 24 h using cold 70% ethanol. DNA staining was achieved by suspending the cell pellets in PBS containing propidium iodide (PI) and RNAse. Cell pellets were washed and suspended in PBS before being analyzed in a flow cytometer (ACEA NovoCyte, Hangzhou, China) using excitation of DNA-bound PI at 488 nm, with emission at 585 nm. Data were processed using NovoExpressTM software. The cell cycle distribution is shown as the percentage of cells containing G1, S and G2 DNA as identified by propidium iodide staining.

Induction of Apoptosis

Flow cytometry analysis of apoptotic populations of the cells caused by exposure to IrFc8 was carried out using the Annexin V-FITC Apoptosis Detection Kit (Beyotime Institute of Biotechnology, China) according to the supplier's instructions. Briefly, A549 cells $(1.5 \times 10^{6}/2 \text{ mL per well})$ were seeded in a six-well plate. Cells were preincubated in drug-free media at 310 K for 24 h, after which IrFc8 was added at concentrations of $1.0 \times IC_{50}$, $2.0 \times IC_{50}$ and $3.0 \times IC_{50}$ of IrFc8 against A549 cancer cells. After 24 h of drug exposure, cells were collected, washed once with PBS, and suspended in 195 μ L of Annexin V-FITC binding buffer which was then added to 5 μ L of Annexin V-FITC and 10 μ L of PI, and then incubated at room temperature in the dark for 15 min. Subsequently, the buffer placed in an ice bath in the dark. The samples

were analyzed by a flow cytometer (ACEA NovoCyte, Hangzhou, China).

Mitochondrial Membrane Assay

Analysis of the changes of mitochondrial potential in cells after exposure to IrFc8 was carried out using the mitochondrial membrane potential assay kit with JC-1 (Beyotime Institute of Biotechnology, Shanghai, China) according to the manufacturer's instructions. Briefly, 1.5×10^6 A549 cancer cells were seeded in six-well plates left to incubate for 24 h in drug-free medium at 310 K in a humidified atmosphere. Drug solutions, with the concentration changed from $0.5 \times IC_{50}$ to $3.0 \times IC_{50}$ of IrFc8 against A549 cells, were added in triplicate, and the cells were left to incubate for a further 24 h under similar conditions. Supernatants were removed by suction, and each well was washed with PBS before detaching the cells using trypsin-EDTA. Staining of the samples was done in flow cytometry tubes protected from light, incubating for 30 min at ambient temperature. The samples were immediately analyzed by a flow cytometer (ACEA NovoCyte, Hangzhou, China). For positive controls, the cells were exposed to carbonyl cyanide 3-chlorophenylhydrazone, CCCP (5 μ M), for 20 min. Data were processed using NovoExpressTM software.

Cellular Localization and Uptake Mechanism assay

Two Photon Laser Scanning Microscope (*/LSM/880NLO) is produced at Carl Zeiss AG, Germany. LTDR (Life Technologies, USA), MTDR (Life Technologies, USA), CCCP (Sigma Aldrich, USA), chloroquine (Sigma Aldrich, USA) was used as received. A549 cells were seeded into 35 mm dishes (Greiner, Germany) for confocal microscopy. After cultured overnight, the cells were incubated with IrFc8 ($1.0 \times IC_{50}$) for 1 h. The treated cells were observed immediately under a confocal microscope with excitation at 488 nm. For colocalization studies, the cells were incubated with IrFc8 ($1.0 \times IC_{50}$) for 1 h. Subsequently, the medium was replaced with staining medium containing MTDR/LDTR and stained for another 20 min and 1 h. The cells were washed twice with PBS, and then viewed immediately under a confocal microscope. Investigation of drug entry pattern: Cells were incubated with IrFc8 ($1.0 \times IC_{50}$) for 1 h, after which media was replaced with staining medium containing CCCP (10μ M, 1 h)

chloroquine (50 μ M, 1 h) and re-stained for 15 min. The cells were washed twice with PBS, and then viewed immediately under a confocal microscope.

Lysosomal damage assay

A549 cells seeded into six-well plate (Corning) were exposed to IrFc8 at the indicated concentrations for 12 h. The cells were then washed twice with PBS and incubated with AO (5 μ M) at 310 K for 15 min. The cells were washed twice with PBS and visualized by confocal microscopy (LSM/880NLO). Emission was collected at 510±20 nm (green) and 625 ± 20 nm (red) upon excitation at 488 nm.

Synthesis

Synthesis of $[(\eta^5 - Cp^*)Ir(\mu - Cl)Cl]_2$ (**Dimer1**).

IrCl₃·3H₂O (0.50 g, 1.7 mmol) was dissolved in methanol (20 mL) in a microwave vial, pentamethyl cyclopentadiene (0.69 g, 5.0 mmol) was added and reacted at 400K, 400 psi for 30 min in a microwave instrument. The reaction mixture was allowed to cool to ambient temperature and the dark green precipitate was filtered off. The volume of the dark red filtrate was reduced to ca. 15 mL on a rotary evaporator. Upon cooling to ambient temperature, an orange precipitate appeared and was collected by filtration. The product was washed with methanol and diethyl ether, dried in air, and pure Dimer 1 was obtained. Yield 0.40 g (65%). ¹H NMR (500 MHz, CDCl₃): δ 1.60 (s, *J* = 1.4 Hz, 30H, Cp-CH₃).

Synthesis of $[(\eta^5-Cp^{xph})Ir(\mu-Cl)Cl]_2$ (**Dimer2**).

Dimer 2 was synthesized using phenyltetramethyl cyclopentadiene (0.84 g, 5.0 mmol) and $IrCl_3 \cdot 3H_2O$ (0.50 g, 1.67 mmol) according to the method of Dimer 1. Yield 0.37 g (58.5%). ¹H NMR (500 MHz, CDCl₃): δ 7.58 (m, 4H, Ar-H), 7.35 (m, 6H, Ar-H), 1.72 (s, 12H, Cp-CH₃), 1.63 (s, 12H, Cp-CH₃).

Synthesis of ferrocene-unappended half-sandwich Ir^{III} TSC complexes (IrPh)



Scheme S1. Design of ferrocene-unappended half-sandwich Ir^{III} TSC complexes (IrPh)

Dimer 2 (92.0 mg) and 4-Phenylthiosemicarbazide ligands (33.4 mg) were added to a Schlenk flask and dissolved in 20 mL methanol. The reaction was carried out in nitrogen and room temperature for 24 h. Then ammonium hexafluorophosphate (130.4 mg, 0.80 mmol) was added and reacted for 6 h. The solvent was removed by vacuum distillation, 3 mL dichloromethane was added and filtered, then product was obtained after *n*-hexane added for diffusion. The ¹H NMR and ESI-MS of IrPh were presented in Figures S5. Yield 106.3 mg (75.8%). ¹H NMR (500 MHz, CDCl₃) δ 7.64 (s, 2H), 7.50 (d, *J* = 7.6 Hz, 4H, Ar-H), 7.42 (d, *J* = 8.2 Hz, 8H, Ar-H), 7.36 (t, *J* = 7.4 Hz, 4H, Ar-H), 7.26 (d, *J* = 6.9 Hz, 4H, Ar-H), 7.20 (t, *J* = 7.1 Hz, 2H), 2.10 (s, 6H, Cp-CH₃), 2.06 (s, 6H, Cp-CH₃), 2.05 (s, 6H, Cp-CH₃), 1.68 (s, 6H, Cp-CH₃). ESI-MS (*m*/*z*): calcd for C₄₄H₅₀N₆F₁₂P₂S₂Ir₂: C, 37.71; H, 3.60; N, 6.00%; Found C, 37.56; H, 3.98; N, 5.74%.

2. Figures





Figure S1 ¹H NMR spectra of L1- L4 in CDCl₃.





Figure S2. HR-MS of L1-L4.









Figure S3 ¹H NMR spectra of IrFc1~IrFc8 in CDCl₃/ DMSO-d₆.







Figure S4 HR-MS and ESI-MS of IrFc1~IrFc8

Figure S5 ESI-MS and ¹H NMR spectra in CDCl₃ of IrPh.

Figure S6 FT-IR of IrFc8 and L2.

Figure S7 The stability of IrFc1-IrFc6 in 20% DMSO/80% H₂O (*v/v*) solution at 298 K were recorded by UV-Vis spectra over 8 h.

gure S8 UV-Vis spectra of IrFc7 (a) and IrFc8 (b) in the 1-octanol and water solution.

Figure S9 UV-Vis (a) and fluorescence spectra (b) of IrFc8 (20 μ M) in DMSO solution.

3. Tables

| Complexes | IC ₅₀ (µM) | | | | |
|-----------|-----------------------|------------|--|--|--|
| | A549 cells | Hela cells | | | |
| IrPh | >100 | >100 | | | |
| L1 | 92.91±1.32 | >100 | | | |
| L2 | >100 | >100 | | | |
| L3 | >100 | >100 | | | |
| L4 | >100 | >100 | | | |

Table S1 IC_{50} values of L1-L4 and IrPh towards A549, Hela cells after 24 h of incubation.

Table S2 Crystallographic data and structure refinement for IrFc2.

| Empirical formula | $C_{56}H_{62}F_{12}Fe_2Ir_2N_6P_2S_2$ | Density (mg/m ³) | 3.349 | |
|---------------------------------------|---------------------------------------|------------------------------|--------------------------------|--|
| Formula weight | 373.11 | Z | 37 | |
| F(000) | 6105 | Cryst size (mm) | $0.46 \times 0.20 \times 0.04$ | |
| Volume (Å ³) | 6844.4(17) | Wavelength (Å) | 0.71073 | |
| Theta range for data collection (deg) | 2.023 to 26.000 | Crystal system | Monoclinic | |
| | -13≤h≤14, | | | |
| Limiting indices | -36≤k≤36, | Space group | P21/n | |
| | -24≤1≤24 | | | |
| Reflections | 13450/13450 | a (Å) | 11.6402(16) | |
| collected/unique | [R(int) = 0.0293] | a (A) | | |
| Data/restraints/paramete | ers 13450/2/715 | b (Å) | 29.210(4) | |
| Goodness-of-fit on F [^] | 2 1.064 | c (Å) | 20.134(3) | |
| Final R indices [I>2sigma(I)] | R1=0.0544, wR2=0.1650 | α (°) | 90 | |
| R indices (all data) | R1=0.0687, wR2=0.1791 | β (°) | 91.112(5) | |
| | | γ (°) | 90 | |
| | | | | |

| | | Population (%) | | | | | |
|---------|----------------------|----------------|-----------------|----------------|------------|--|--|
| | Concentration | Viable | Early apoptosis | Late apoptosis | Non-viable | | |
| Control | | 79.1±0.2 | 0.68±0.5 | 8.85±0.1 | 11.37±0.1 | | |
| | 1.0×IC ₅₀ | 70.7±0.9 | $0.78{\pm}0.2$ | 18.4±0.7 | 10.12±0.03 | | |
| IrFc8 | 2.0×IC ₅₀ | 62.8±3.3 | 1.33±0.3 | 31.0±2.0 | 4.87±0.07 | | |
| | 3.0×IC ₅₀ | 21.3±3.3 | 1.65±0.3 | 68.3±2.0 | 8.75±0.07 | | |

Table S3 Flow cytometry analysis to determine the percentages of apoptotic cells, using AnnexinV-FITC/PI staining, after exposing A549 cells to IrFc8.

 Table S4 Cell cycle analysis carried out by flow cytometry using PI staining after exposing A549 cells to IrFc8.

| | | Population (%) | | | | |
|---------|----------------------|----------------------|-----------|----------------------|--|--|
| | Concentration | G ₁ phase | S phase | G ₂ phase | | |
| Control | | 65.52±0.1 | 10.28±0.5 | 24.20±2.3 | | |
| IrFc8 | 0.5×IC ₅₀ | 65.91±2.7 | 10.61±0.8 | 23.48±1.3 | | |
| | 1.0×IC ₅₀ | 66.45±0.7 | 10.06±0.4 | 23.49±0.8 | | |
| | 2.0×IC ₅₀ | 77.32±1.3 | 6.47±0.1 | 16.20±0.7 | | |

Table S5 The mitochondrial membrane polarization of A549 cells induced by IrFc8.

| | | Population (%) | | | |
|------------------|----------------------|-----------------|---------------|--|--|
| | Concentration | JC-1 Aggregates | JC-1 Monomers | | |
| | 0.5×IC ₅₀ | 88.25±1.0 | 11.75±1.0 | | |
| | 1.0×IC ₅₀ | 85.68±1.5 | 14.31±0.8 | | |
| lrFc8 | 2.0×IC ₅₀ | 84.00±1.2 | 16.00±0.3 | | |
| | 3.0×IC ₅₀ | 82.99±0.9 | 17.01±0.7 | | |
| Negative Control | | 89.86±0.2 | 10.14±0.2 | | |
| Positive Control | | 27.67±1.4 | 72.33±2.4 | | |

| Group | Animals | 0d | 2d | 4d | 6d | 8d | 10d | 12d | 14d |
|-----------|---------|-------|-------|-------|-------|-------|-------|-------|-------|
| Cisplatin | 194 | 18.93 | 19.10 | 19.43 | 19.76 | 20.08 | 20.29 | 20.75 | 21.39 |
| | 195 | 17.82 | 18.33 | 18.75 | 19.06 | 19.44 | 19.80 | 20.43 | 20.92 |
| | 196 | 18.38 | 18.66 | 19.03 | 19.27 | 19.56 | 19.92 | 20.31 | 20.88 |
| IrFc8 | 189 | 18.30 | 18.67 | 19.05 | 19.72 | 20.24 | 20.68 | 21.03 | 21.35 |
| | 191 | 18.50 | 18.82 | 19.25 | 19.69 | 20.05 | 20.60 | 21.03 | 21.49 |
| | 192 | 17.87 | 18.05 | 18.34 | 18.95 | 19.33 | 19.56 | 20.01 | 20.43 |

 Table S6 14-Day body weight (g)

 Table S7 Tumor weight (mm³)

| Group | Animals | Tumor weight (g) |
|-----------|---------|------------------|
| | 195 | 1.4635 |
| Cisplatin | 196 | 1.0113 |
| 1 | 198 | 1.3462 |
| | 189 | 0.4671 |
| IrFc8 | 191 | 0.4483 |
| | 192 | 0.2615 |

 Table S8 14-day tumor volume (mm³)

| Group | Animal | s 0d | 2d | 4d | 6d | 8d | 10d | 12d | 14d |
|------------|--------|--------|--------|--------|--------|--------|---------|---------|---------|
| Cis-platin | 194 | 111.01 | 190.78 | 268.32 | 425.92 | 647.15 | 1164.06 | 1779.06 | 2572.80 |
| | 195 | 98.78 | 158.76 | 231.89 | 332.80 | 516.91 | 829.99 | 1094.40 | 1468.53 |
| | 196 | 90.40 | 138.60 | 183.79 | 255.79 | 404.60 | 569.92 | 896.72 | 1335.10 |
| IrFc8 | 189 | 98.84 | 143.65 | 181.30 | 241.88 | 305.94 | 400.95 | 497.66 | 622.91 |
| | 191 | 113.13 | 141.31 | 173.40 | 208.25 | 237.14 | 272.81 | 306.13 | 400.00 |
| | 192 | 91.85 | 111.01 | 129.60 | 159.74 | 205.80 | 247.50 | 294.40 | 349.27 |