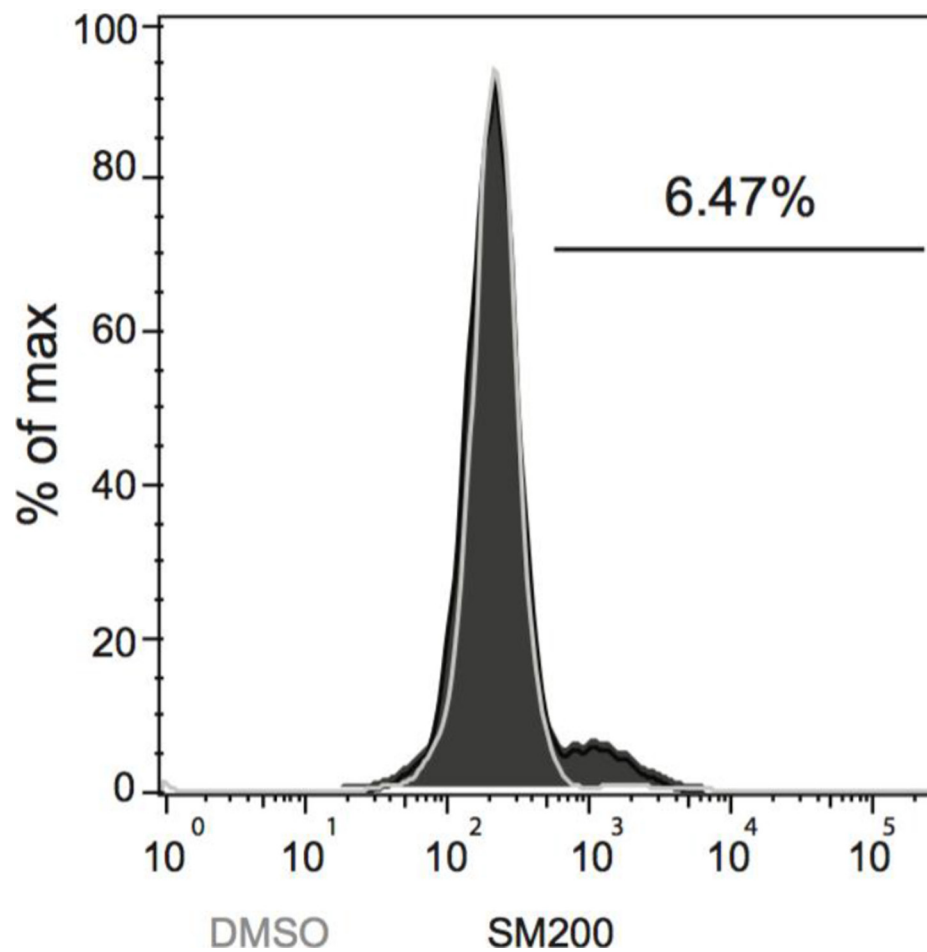
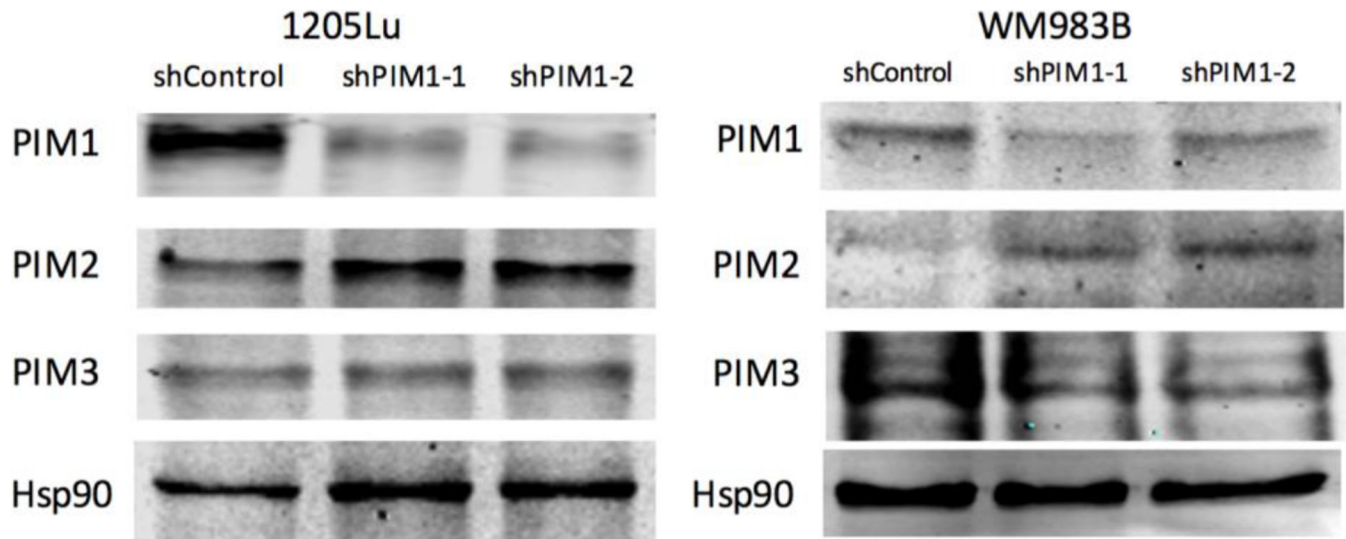


PIM kinases as therapeutic targets against advanced melanoma

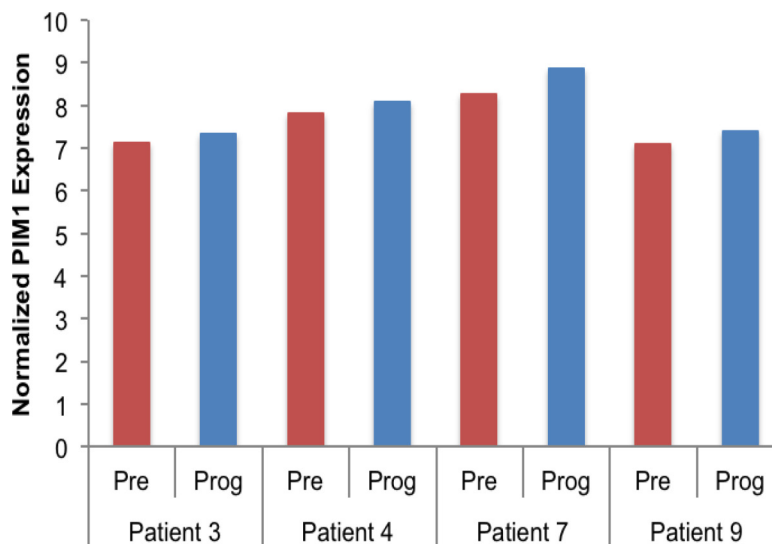
Supplementary Materials



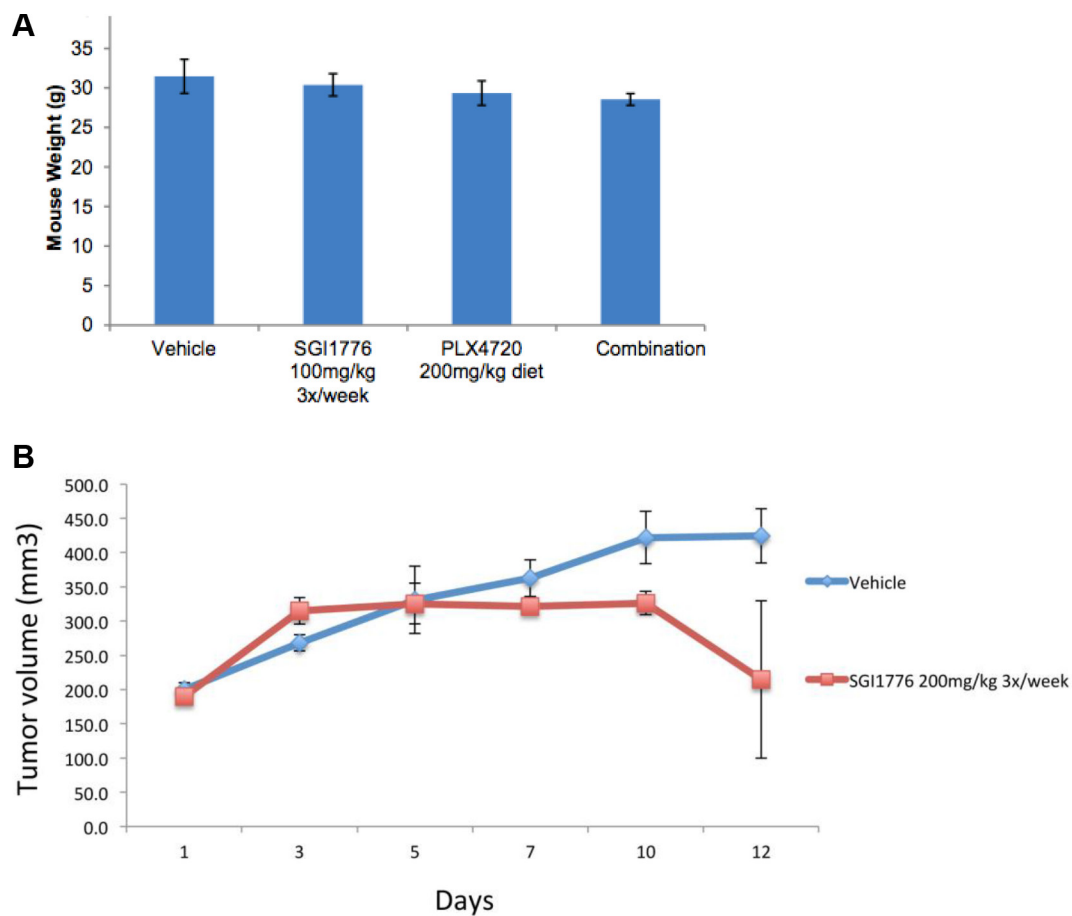
Supplementary Figure S1: Caspase-3 staining following SM200 treatment. Histograms represent caspase-3 activated staining in 1205Lu cells after DMSO (grey line) or SM200 treatment (black filled histogram). The percentage indicates the frequency of caspase3-positive cells found after 72 h SM200 (10 μ M) treatment. On X and Y axes, fluorescence intensity and % of max are respectively reported.



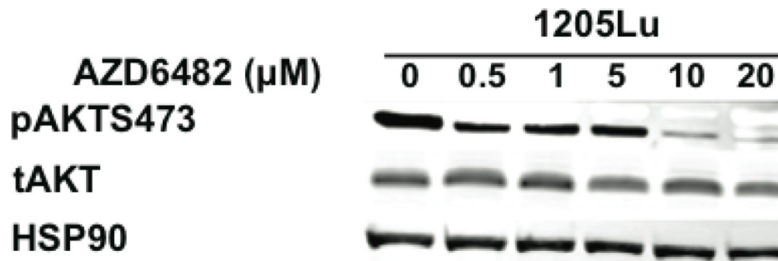
Supplementary Figure S2: Effects of PIM1 knockdown on melanoma cells. PIM1 was knocked down in 1205Lu and WM983B melanoma cells using two different shRNAs; expression of PIM isoforms was then assessed by western blot. These cell lines were independently established and are distinct from those shown in Figure 3A. Hsp90 served as loading control.



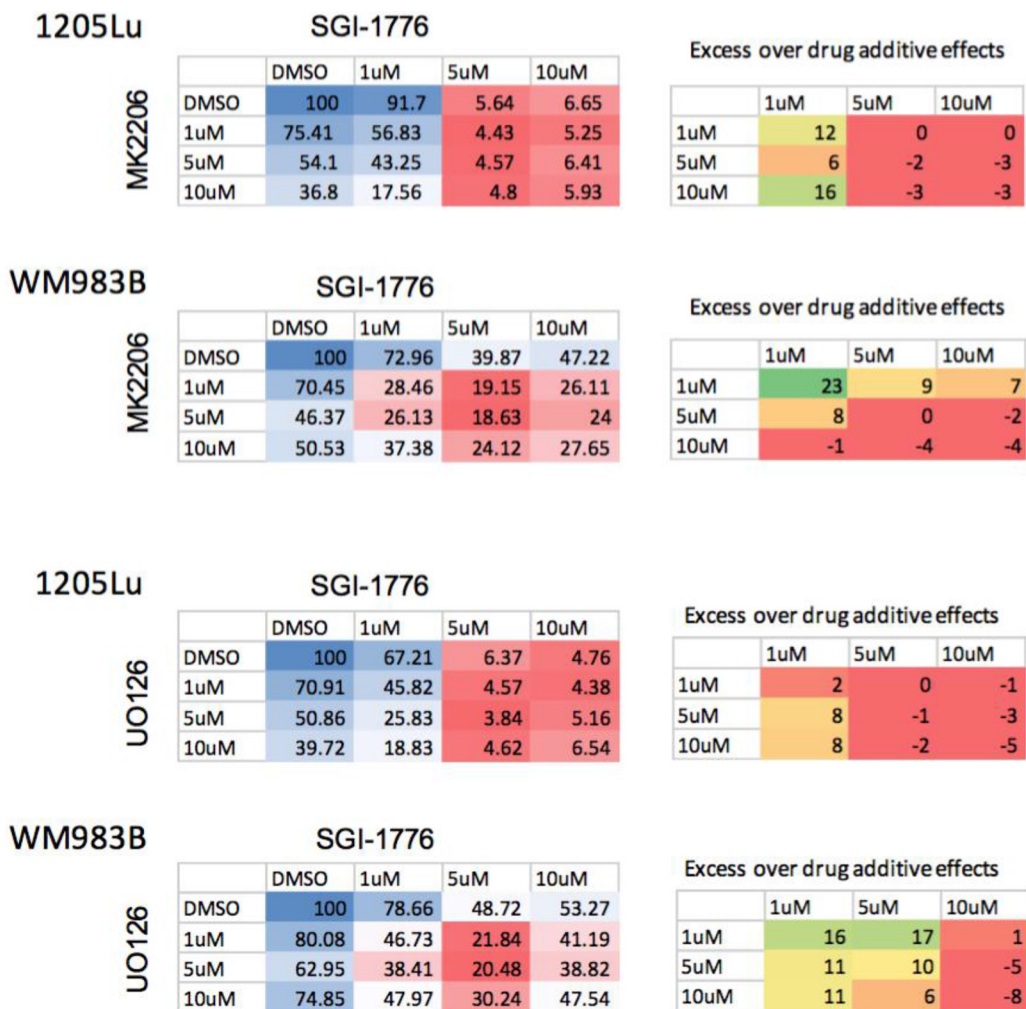
Supplementary Figure S3: PIM1 gene expression in melanoma tumors treated with a combination of dabrafenib and trametinib (pre-treatment and upon progression). RNA was isolated from fresh frozen melanoma tumors in 9 patients (GSE61992) using. The graph shows data for 4 patients with greater PIM1 gene expression in progressing tumor samples (Prog) than in pretreated samples (Pre). The y-axis is the normalized PIM1 gene expression level.



Supplementary Figure S4: SGI-1776 is effective as a single agent or in combination with PLX4720 *in vivo* melanoma models. (A) SGI-1776 has minimal effects on total mouse weight as a single agent or in combination with PLX4720. Overall mouse weights from the *in vivo* experiment shown in Figure 6A, measured on the last treatment day. Data are represented as mean \pm SEM. (B) SGI-1776 as a single agent is effective in reducing tumor volumes. NSG mice ($n = 7$ /group) were xenotransplanted with 1205Lu melanoma cells and tumors were allowed to grow to 200 mm³. Mice were then treated with the single agent SGI-1776 (200 mg/kg 3x/week). Tumor volumes were measured at the indicated time points and are represented as mean \pm SEM.



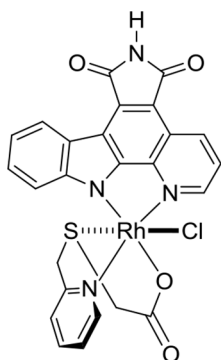
Supplementary Figure S5: AZD6482 target inhibition. Increasing doses of the PI3K β inhibitor AZD6482 were used to treat melanoma cells for 72 h before lysates were collected and probed for pAKTSer473 and total protein. Hsp90 was used as a loading control.



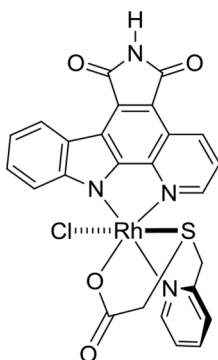
Supplementary Figure S6: SGI-1776 displays anti-melanoma activity in combination with an AKT or MEK inhibitor. Melanoma cells were treated for 72 h with SGI-1776 or the AKT inhibitor MK2206 or the MEK inhibitor UO126 as single agents or in combination. To facilitate statistical analysis of synergy effects, we used a grid-like design of constant ratio drug combinations. Viability was calculated relative to DMSO controls in each plate and the mean of three experiments provided. Synergy was calculated for combination experiments using the Bliss formula. The Bliss number gives the difference between predicted and observed inhibition values (excess over Bliss); a positive value indicates synergy, a negative value indicates antagonism and values near zero indicate an overlap of predicted and observed combination effects.

Screened organometallic compounds

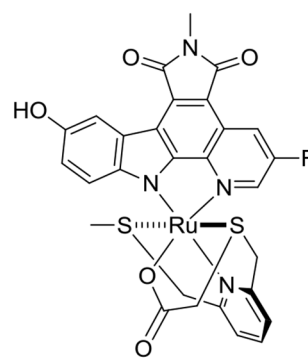
The compounds were synthesized according to published procedures [1–5].



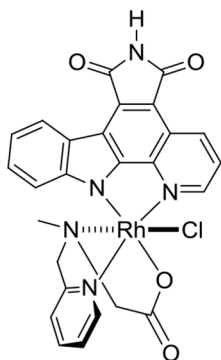
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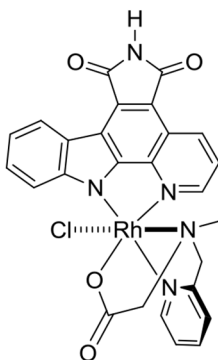
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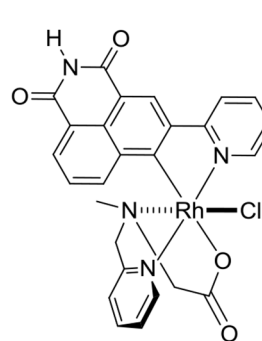
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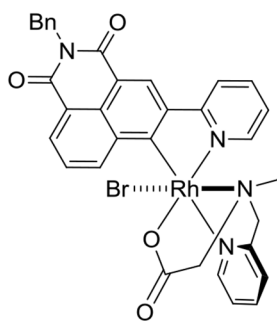
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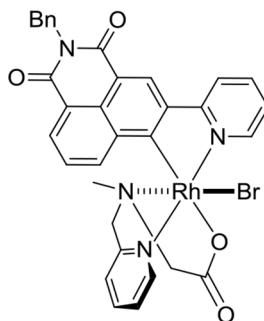
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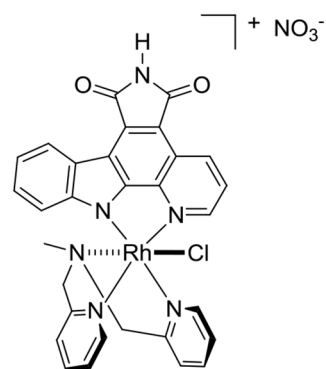
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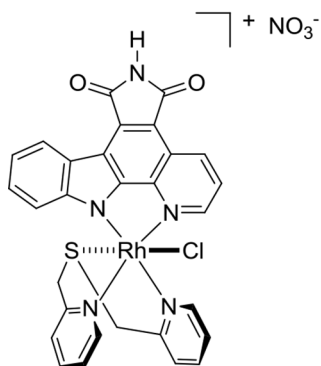
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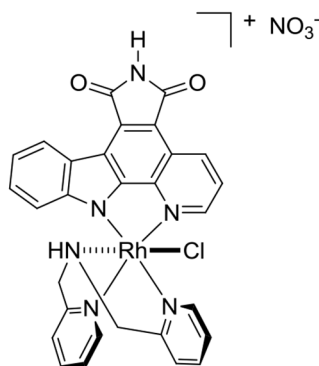
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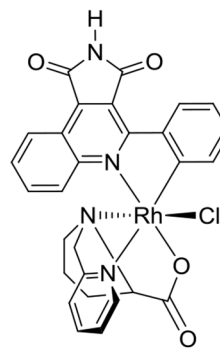
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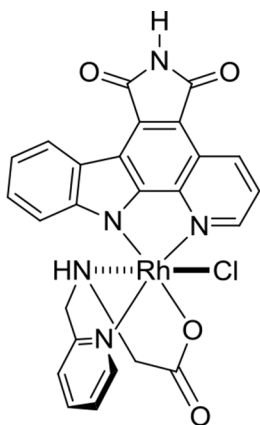
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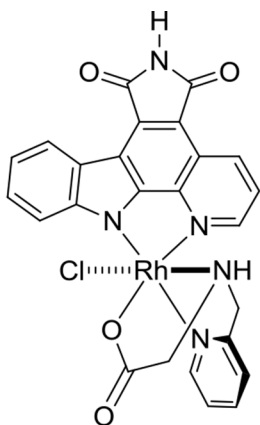
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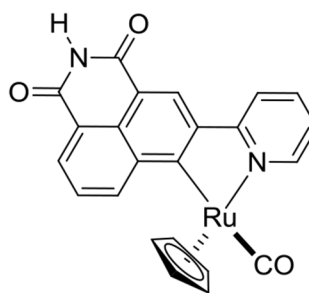
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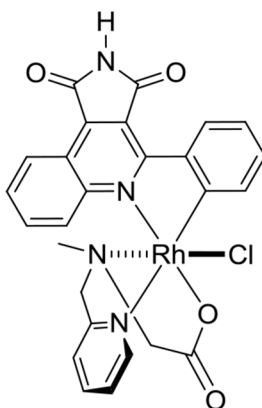
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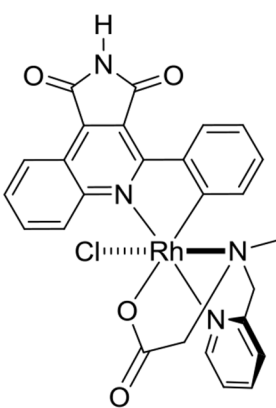
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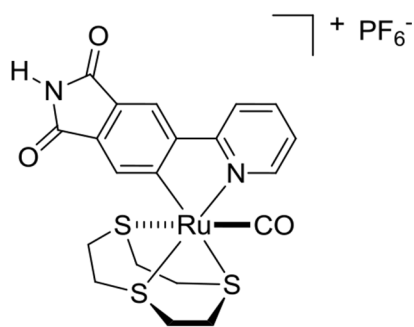
SB_TC-18



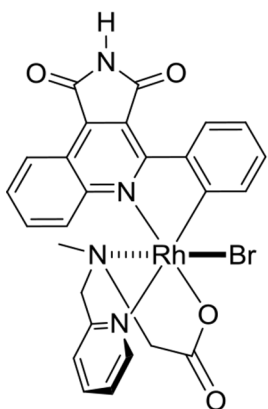
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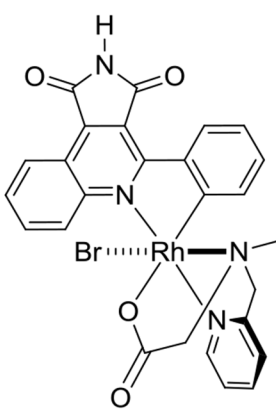
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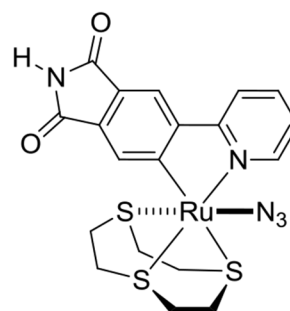
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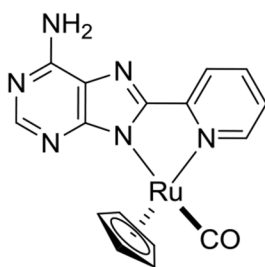
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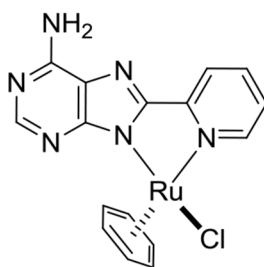
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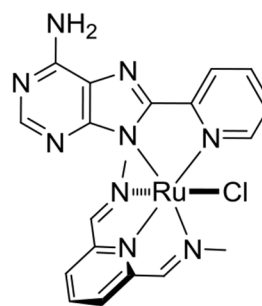
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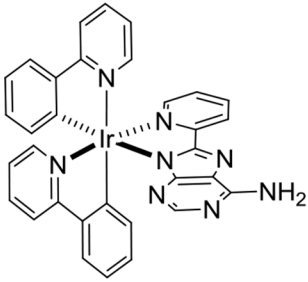
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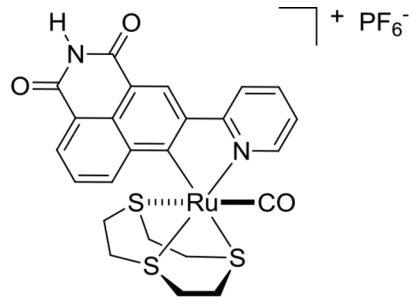
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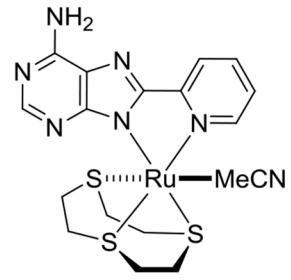
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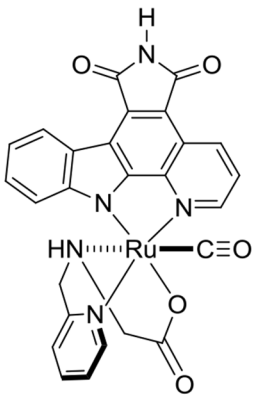
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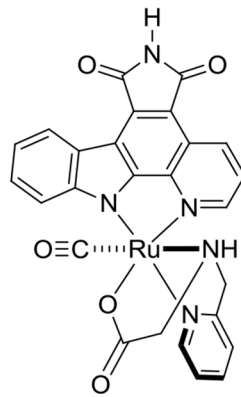
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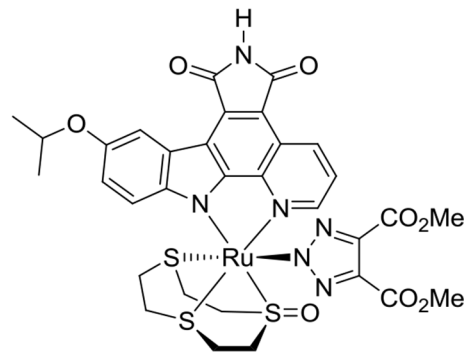
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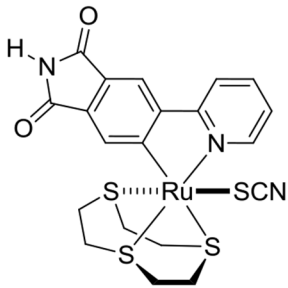
SM200-1 (SM200)



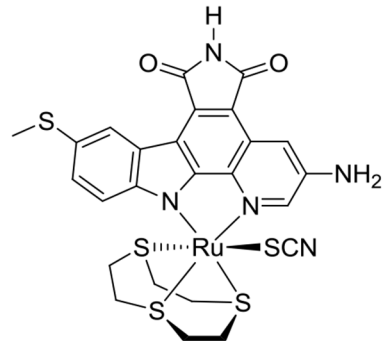
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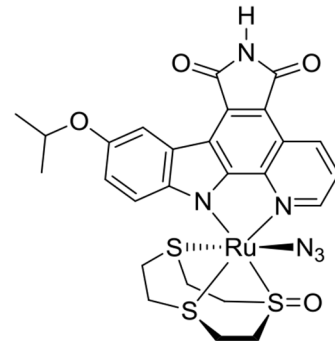
FL1543



SB_171



FL1334



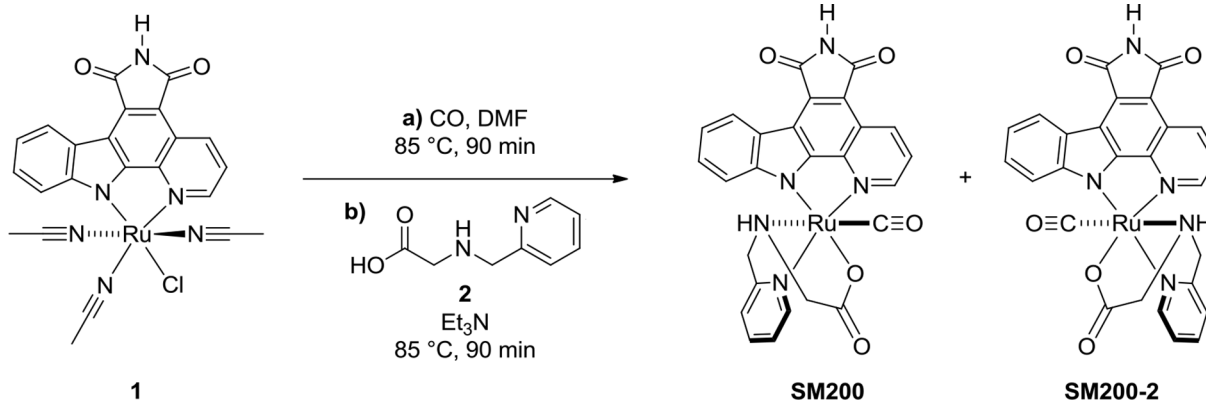
FL1528-1

Synthesis of SM200

The Ru(II)-precursor **1** [6] and *N*-(2-pyridylmethyl)glycine **2** [7] were synthesized according to reported methods. Et₃N was used without further purification and DMF was distilled over CaH₂ before usage. TLC plates for thin layer chromatography (*silica gel 60 F₂₅₄*) and silica gel for column chromatography (*silica gel 60, 40–63 μm*) were supplied by *Merck KGaA*. ¹H-NMR spectra were recorded using a *Bruker Avance 300* (300 MHz), for IR-spectra a *Bruker Alpha-P FT-IR* spectrometer and for mass spectra a *LTQ-FT Ultra* mass spectrometer from *Thermo Fisher Scientific*.

A solution of Ru(II)-precursor **1** (10.9 mg, 20.0 μmol) in 2.0 μL anhydrous DMF was heated to 85°C under an atmosphere of CO. After 90 min, Et₃N (8.4 mL, 60.0 μmol) and *N*-(2-pyridylmethyl)glycine **2** (3.7 mg, 20.0 μmol) were added to the red solution and stirred again at 85°C for 90 min. The red solution was cooled to room temperature, all volatile compounds were evaporated in vacuo and the mixture subjected to column chromatography (CH₂Cl₂/MeOH 50:1–30:1). After purification, 1.0 mg (1.72 μmol, 9%) of SM200 and 1.1 mg (1.89 μmol, 9%) of SM200-2 were isolated as red solids.

SM200: R_f = 0.18 (CH₂Cl₂/MeOH 20:1). ¹H-NMR (300 MHz, DMSO-*d*₆): δ (ppm) = 11.15 (s, 1H, CH_{ar}), 9.34



Kit), according to manufacturer's instructions. Cells were stained with primary Purified Rabbit anti-human caspase-3 activated antibody (Cell Signaling Technologies) diluted 1:200, 30 min at 4°C. This was followed by a secondary antibody incubation using anti-mouse Alexa488 (Life Technologies), 1:500 dilution, 20 min at 4°C. The percent

(d, J = 5.3 Hz, 1H, CH_{ar}), 9.20 (d, J = 7.3 Hz, 1H, CH_{ar}), 8.67–8.62 (m, 2H, CH_{ar}), 8.13 (td, J = 7.8 Hz, 1.4 Hz, 1H, CH_{ar}), 7.92 (dd, J = 8.4, 5.1 Hz, 1H, CH_{ar}), 7.72–7.68 (m, 2H, CH_{ar}), 7.25–7.15 (m, 2H, CH_{ar}), 6.55 (s, 1H, CH_{ar}), 5.95 (d, J = 6.5 Hz, 1H, CH_{ar}), 4.32 (d, J = 16.4 Hz, 1H, NCH₂), 4.00 (dd, J = 16.5, 5.6 Hz, 1H, NCH₂), 3.58 (dd, J = 17.2, 8.2 Hz, 1H, NCH₂), 3.18–3.13 (m, 1H, NCH₂). IR: n (cm⁻¹) 3415, 3241, 2924, 1948, 1747, 1701, 1612, 1422, 1346, 1231, 748, 643. HRMS (ESI(+)): C₂₆H₁₈N₅O₅Ru (M+H)⁺ calc. 582.0353, found 582.0342.

SM200-2: R_f = 0.13 (CH₂Cl₂/MeOH 20:1). ¹H-NMR (300 MHz, DMSO-*d*₆): δ (ppm) = 11.13 (s, 1H, CH_{ar}), 9.21 (d, J = 5.4 Hz, 1H, CH_{ar}), 9.11 (d, J = 8.4, 1.2 Hz, 1H, CH_{ar}), 8.70 (d, J = 7.9 Hz, 1H, CH_{ar}), 8.08–8.02 (m, 2H, CH_{ar}), 7.69–7.54 (m, 4H, CH_{ar}), 7.47 (td, J = 8.2, 1.3 Hz, 1H, CH_{ar}), 7.35–7.30 (m, 1H, CH_{ar}), 6.55 (s, 1H, CH_{ar}), 4.43–4.27 (m, 2H, NCH₂), 3.28–3.24 (m, 2H, NCH₂). IR: n (cm⁻¹) 3382, 1942, 1689, 1629, 1292, 1150, 1021, 817, 732, 703, 481, 424. HRMS (ESI(+)): C₂₆H₁₈N₅O₅Ru (M+H)⁺ calc. 582.0353, found 582.0341.

Flow cytometry staining

Caspase-3 activated intracellular staining was achieved after cell fixation and permeabilization (BD Cytotfix/Cytoperm™ Fixation/Permeabilization Solution

of positive cells after SM200 treatment are reported after subtraction of background from the control staining. Samples were acquired with a Becton Dickinson LSRII 14 color flow cytometer and analyzed with FlowJo software (Tree Star Inc. Ashland, OR, USA).

Supplementary Table S1: Melanoma cell line basic genetic information

Cell Line	Stage	BRAF	NRAS
WM3918	MET	WT	WT
WM983B WM983B-BR	MET	V600E	WT
1205Lu	MET	V600E	WT
451Lu 451Lu-BR	MET	V600E	WT
WM1361A	VGP	WT	Q61R

Stage and BRAF, NRAS mutation status is provided.

Abbreviations: MET, metastasis; WT, wild type; N/A, not available; BR, BRAF inhibitor resistant cell line.

Supplementary Table S2: Organometallic compound effects on melanoma cell lines

Compounds with no anti-melanoma activity	Compounds selective for melanoma cells (low dose)	Compounds selective for melanoma cells (higher dose)	Cytotoxic compounds
IC50 > 10 μ M	IC50 \leq 10 μ M	IC50 > 10 μ M	IC50 \geq 5 μ M
SB_178 SB_TC-18 MCS_C003 MCS_C010 SM250-2 SM406-1, -2 SM407-1, -2 SM415-1, -2 SM426-2 SM438-2, -3 SM439-2, -3 SM440-1, -3 SM449 SM450 SM451	SM200 SM200-2 FL1543	SB_179 SB_171 MCS_C002 MCS_C007	MCS_C004 SB_181 FL1528-01 FL1334

Compounds were tested on fibroblasts (control) and the melanoma cell lines 1205Lu, 451Lu, WM983B, and WM3918 over 72 h; effects were assessed using the MTS assay.

Supplementary Table S3: Kinase profiling of SM200. See Supplementary_Table_S3

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