

Supporting Information

Indolinyl-thiazole based inhibitors of Scavenger Receptor-BI (SR-BI)-mediated lipid transport

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Contents

1. Additional SAR data: Table S1
2. Synthesis and characterization data for **17-11** (ML278)
3. Dose-response for [³H]cholesteryl ester uptake assay: Figure S1
4. Probe comparison: Table S2
5. Compound profiling protocols
6. Assay protocols
7. References
8. Off-target screening data for **17-11** (ML278)

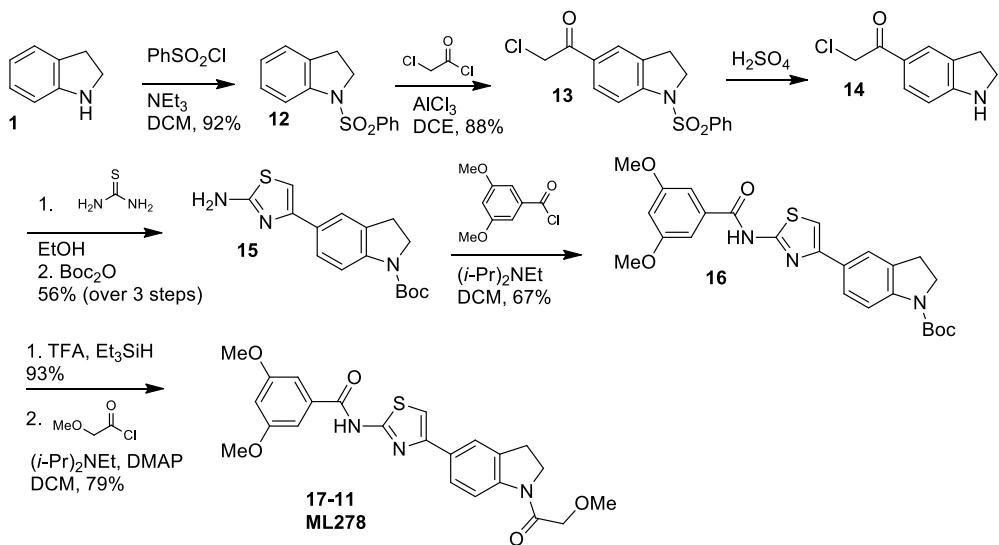
1. Additional SAR data

Table S1. Modifications to indoline ring^a

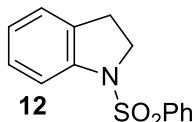
Cmp	Structure	IC ₅₀ (μM) ^a	Cmp	Structure	IC ₅₀ (μM) ^a
18		0.089	23		0.012
19		2.0	24		0.20
20		0.085 ± 0.02	25		0.09
21		0.2	26		0.40
22		0.020	27		0.07
			28		0.09

^a Average of at least two measurements in dI uptake assay, ± standard of mean when n > 2.

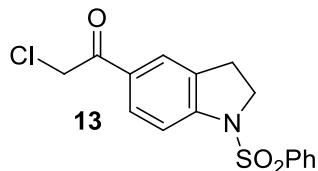
2. Synthesis and characterization data for 17-11 (ML278)



General Details. All reagents and solvents were purchased from commercial vendors and used as received. NMR spectra were recorded on a Bruker 300 MHz or Varian UNITY INOVA 500 MHz spectrometer as indicated. Proton and carbon chemical shifts are reported in parts per million (ppm; δ) relative to tetramethylsilane, CDCl₃ solvent, or d₆-DMSO (¹H δ 0, ¹³C δ 77.16, or ¹³C δ 39.5, respectively). NMR data are reported as follows: chemical shifts, multiplicity (obs. = obscured, app = apparent, br = broad, s = singlet, d = doublet, t = triplet, m = multiplet, comp = complex overlapping signals); coupling constant(s) in Hz; integration. Unless otherwise indicated, NMR data were collected at 25 °C. Flash chromatography was performed using 40-60 µm Silica Gel on a Teledyne Isco CombiFlash R_f system. Tandem liquid chromatography/mass spectrometry (LCMS) was performed on a Waters 2795 separations module and Waters 3100 mass detector. Analytical thin layer chromatography (TLC) was performed on EM Reagent 0.25 mm silica gel 60-F plates. Visualization was accomplished with UV light and aqueous potassium permanganate (KMnO₄) stain followed by heating. High-resolution mass spectra were obtained at the MIT Mass Spectrometry Facility with a Bruker Daltonics APEXIV 4.7 Tesla Fourier Transform Ion Cyclotron Resonance mass spectrometer. Compound purity and identity were determined by UPLC-MS (Waters, Milford, MA). Purity was measured by UV absorbance at 210 nm. Identity was determined on a SQ mass spectrometer by positive electrospray ionization. Mobile Phase A consisted of either 0.1% ammonium hydroxide or 0.1% trifluoroacetic acid in water, while mobile Phase B consisted of the same additives in acetonitrile. The gradient ran from 5% to 95% mobile Phase B over 0.8 minutes at 0.45 ml/min. An Acquity BEH C18, 1.7 µm, 1.0 x 50 mm column was used with column temperature maintained at 65 °C. Compounds were dissolved in DMSO at a nominal concentration of 1 mg/ml, and 0.25 µl of this solution was injected.

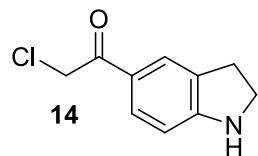


1-(Phenylsulfonyl)indoline (12): Indoline (1.4 ml, 12.6 mmol) was placed in a round-bottom flask equipped with a magnetic stirbar and dissolved in dichloromethane (42.0 ml). After cooling to 0 °C, triethylamine (3.5 ml, 25.2 mmol) was added to the solution, followed by benzenesulfonyl chloride (1.7 ml, 13.3 mmol). The reaction was stirred for 30 minutes while warming to room temperature. Saturated sodium bicarbonate solution (aqueous, 30 ml) was added to quench the reaction. The layers were separated, and the aqueous layer was extracted with dichloromethane (3 x 25 ml). The combined organics were washed with brine (25 ml), then shaken over magnesium sulfate, filtered, and concentrated under reduced pressure to give a light tan solid. The crude material was purified by column chromatography over silica gel (hexanes/ethyl acetate: 100/0 to 80/20) to give the title compound as a light pink solid (3.00 g, 92 %). **1H NMR (300 MHz, CDCl₃):** δ 7.82 – 7.77 (m, 2H), 7.66 (d, *J* = 8.0 Hz, 1H), 7.55 (t, *J* = 7.4 Hz, 1H), 7.44 (t, *J* = 7.9 Hz, 2H), 7.20 (t, *J* = 7.8 Hz, 1H), 7.08 (d, *J* = 7.4 Hz, 1H), 6.98 (t, *J* = 7.4 Hz, 1H), 3.93 (t, *J* = 8.4 Hz, 2H), 2.89 (t, *J* = 8.4 Hz, 2H); **MS (ESI⁺):** 260 (M+H).

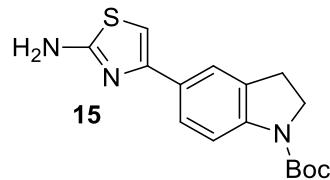


2-Chloro-1-(1-(phenylsulfonyl)indolin-5-yl)ethanone (13): A round-bottom flask was charged with a magnetic stirbar and anhydrous aluminum chloride (1.70 g, 12.7 mmol) then placed under nitrogen atmosphere. Anhydrous 1,2-dichloroethane (28 ml) was added to produce a pale yellow suspension. Chloroacetyl chloride (1.0 ml, 12.7 mmol) was added slowly by syringe. The mixture was stirred at room temperature for 30 minutes. A solution of 1-(phenylsulfonyl)indoline (3.00 g, 11.6 mmol) in anhydrous 1,2-dichloroethane (8.0 ml) was slowly added to the reaction by syringe. The syringe was rinsed twice with anhydrous 1,2-dichloroethane (3.0 ml), and the rinses were added to the reaction. The reaction, now a dark green mixture, was heated to 50 °C and stirred for 4 hours. Additional portions of aluminum chloride (1.30 g, 9.7 mmol) and chloroacetyl chloride (0.75 ml, 9.5 mmol) were dissolved in anhydrous 1,2-dichloroethane (10 ml) and added to the reaction. After stirring 2 more hours at 50 °C, 5 ml of anhydrous 1,2-dichloroethane containing another 0.8 g (6.0 mmol) of aluminum chloride and 0.5 ml (6.4 mmol) of chloroacetyl chloride was added. Stirring at 50 °C was continued for another hour to complete the reaction. The dark red mixture was slowly poured into ice water (approx. 250 ml) and further diluted with dichloromethane (200 ml). The resulting cloudy, orange mixture was stirred while warming to room temperature, after which the layers were separated and the aqueous phase was extracted with dichloromethane (3 x 100 ml). The combined organic layers were washed with water (100 ml) and brine (100 ml), then shaken over magnesium sulfate, filtered, and

concentrated under reduced pressure to give a brown solid (3.91 g). The crude material was purified by column chromatography over silica gel (hexanes/ethyl acetate: 100/0 to 0/100) to give the title compound as a tan solid (3.41 g, 88 %). **¹H NMR (300 MHz, CDCl₃):** δ 7.87 – 7.81 (m, 3H), 7.75 – 7.69 (m, 2H), 7.65 – 7.57 (m, 1H), 7.54 – 7.47 (m, 2H), 4.63 (s, 2H), 4.01 (t, *J* = 8.6 Hz, 2H), 3.05 (t, *J* = 8.6 Hz, 2H); **MS (ESI⁺):** 336 (M+H).

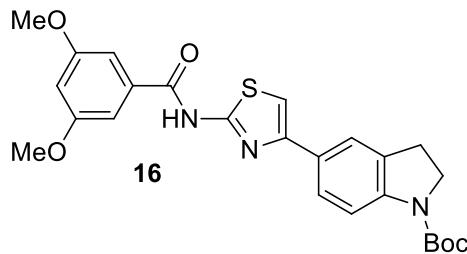


2-Chloro-1-(indolin-5-yl)ethanone (14): A microwave tube was charged with a magnetic stirbar and 2-chloro-1-(1-phenylsulfonyl)indolin-5-yl)ethanone (3.26 g, 9.54 mmol). Concentrated sulfuric acid (9.0 ml) was added, and the resulting suspension was microwaved for 10 minutes at 100 °C. The reaction was carefully poured into ice water (500 ml). The dark mixture was stirred while warming to room temperature then treated with 10% (w/v) aqueous sodium hydroxide (approximately 200 ml) until the pH >10. This mixture was then extracted with dichloromethane (3 x 250 ml). The combined extracts were washed with brine (200 ml), then shaken over magnesium sulfate, filtered, and concentrated under reduced pressure to give a brown solid (1.52 g). This material was used immediately without further purification. **¹H NMR (300 MHz, CDCl₃):** δ 7.72 (d, *J* = 1.2 Hz, 1H), 7.69 (dd, *J* = 8.3, 1.6 Hz, 1H), 6.54 (d, *J* = 8.2 Hz, 1H), 4.60 (s, 2H), 3.70 (t, *J* = 8.6 Hz, 2H), 3.09 (t, *J* = 8.6 Hz, 2H); **MS (ESI⁺):** 196 (M+H).

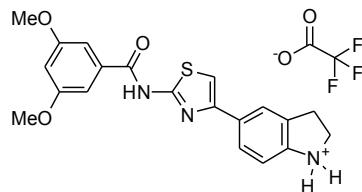


tert-Butyl 5-(2-aminothiazol-4-yl)indoline-1-carboxylate (15): 2-Chloro-1-(indolin-5-yl)ethanone (1.52 g, 7.75 mmol) was placed in a microwave vial and dissolved in anhydrous ethanol (30.0 mL) to give an opaque, black solution. Thiourea (0.66 g, 8.67 mmol) was added and the resulting mixture was microwaved for 30 minutes at 120 °C. 4-(*N,N*-Dimethylamino)pyridine (95.0 mg, 0.78 mmol) and *N,N*-diisopropylethylamine (1.5 ml, 9.3 mmol) were added to the reaction mixture. Neat di-*tert*-butyl dicarbonate (2.0 mL, 8.52 mmol) was added last, and the reaction was stirred at room temperature for 1 hour. The opaque, red-brown mixture was concentrated under reduced pressure to give a red-brown solid. This material was partitioned between water (50 ml) and ethyl acetate (75 ml) and stirred at room temperature until everything dissolved. The layers were separated and the aqueous phase was extracted with ethyl acetate (3 x 50 ml). The combined organic extracts were shaken over magnesium sulfate, filtered, and concentrated under reduced pressure to give an orange-brown solid. The crude material was purified by column chromatography over silica gel (hexanes/ethyl acetate: 100/0 to 40/60) to give the title compound as an orange solid (1.72 g, 56 % over three steps). **¹H NMR**

(300 MHz, d₆-DMSO): δ 7.61 (s, 1H), 7.60 (d, *J* = 8.1 Hz, 1H), 6.99 (s, 1H), 6.85 (s, 1H), 3.92 (t, *J* = 8.7 Hz, 2H), 3.07 (t, *J* = 8.7 Hz, 2H), 1.51 (s, 9H); **MS (ESI⁺):** 318 (M+H).

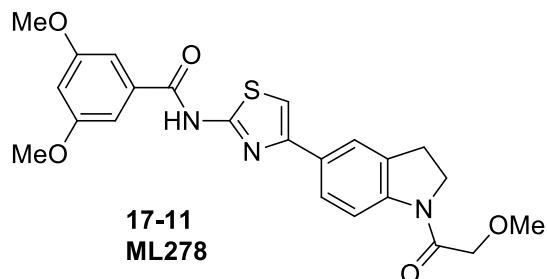


tert-Butyl 5-(2-(3,5-dimethoxybenzamido)thiazol-4-yl)indoline-1-carboxylate (16): In a round-bottom flask equipped with a magnetic stirbar, *tert*-butyl 5-(2-aminothiazol-4-yl)indoline-1-carboxylate (0.500 g, 1.58 mmol) was combined with 4-(*N,N*-dimethylamino)pyridine (19.0 mg, 0.16 mmol). Dichloromethane (4.0 ml) was added to produce an orange suspension that was cooled to 0 °C. *N,N*-Diisopropylethylamine (0.33 ml, 1.89 mmol) was added followed by a solution of 3,5-dimethoxybenzoyl chloride (0.35 g, 1.73 mmol) in dichloromethane (1.00 ml). The bright orange mixture was stirred for 1 hour while warming to room temperature. The resulting clear, red-brown solution was diluted with saturated sodium bicarbonate solution (aqueous, 10 ml) and dichloromethane (10 ml). The layers were separated, and the aqueous phase was extracted with dichloromethane (3 x 10 ml). The combined organics were washed with brine (10 ml) then shaken over magnesium sulfate, filtered, and concentrated under reduced pressure to give a thick, orange-brown oil. This material was purified by column chromatography over silica gel (hexanes/ethyl acetate: 100/0 to 70/30) to give the title compounds as a light yellow solid (0.51 g, 67%). **1H NMR (300 MHz, CDCl₃):** δ 9.84 (s, 1H), 7.60 (d, *J* = 12.2 Hz, 1H), 7.58 (s, 1H), 7.07 (s, 1H), 7.04 (s, 1H), 7.03 (s, 1H), 6.64 (t, *J* = 2.2 Hz, 1H), 4.01 (t, *J* = 8.7 Hz, 2H), 3.84 (s, 6H), 3.12 (t, *J* = 8.7 Hz, 2H), 1.58 (s, 9H); **MS (ESI⁺):** 482 (M+H).



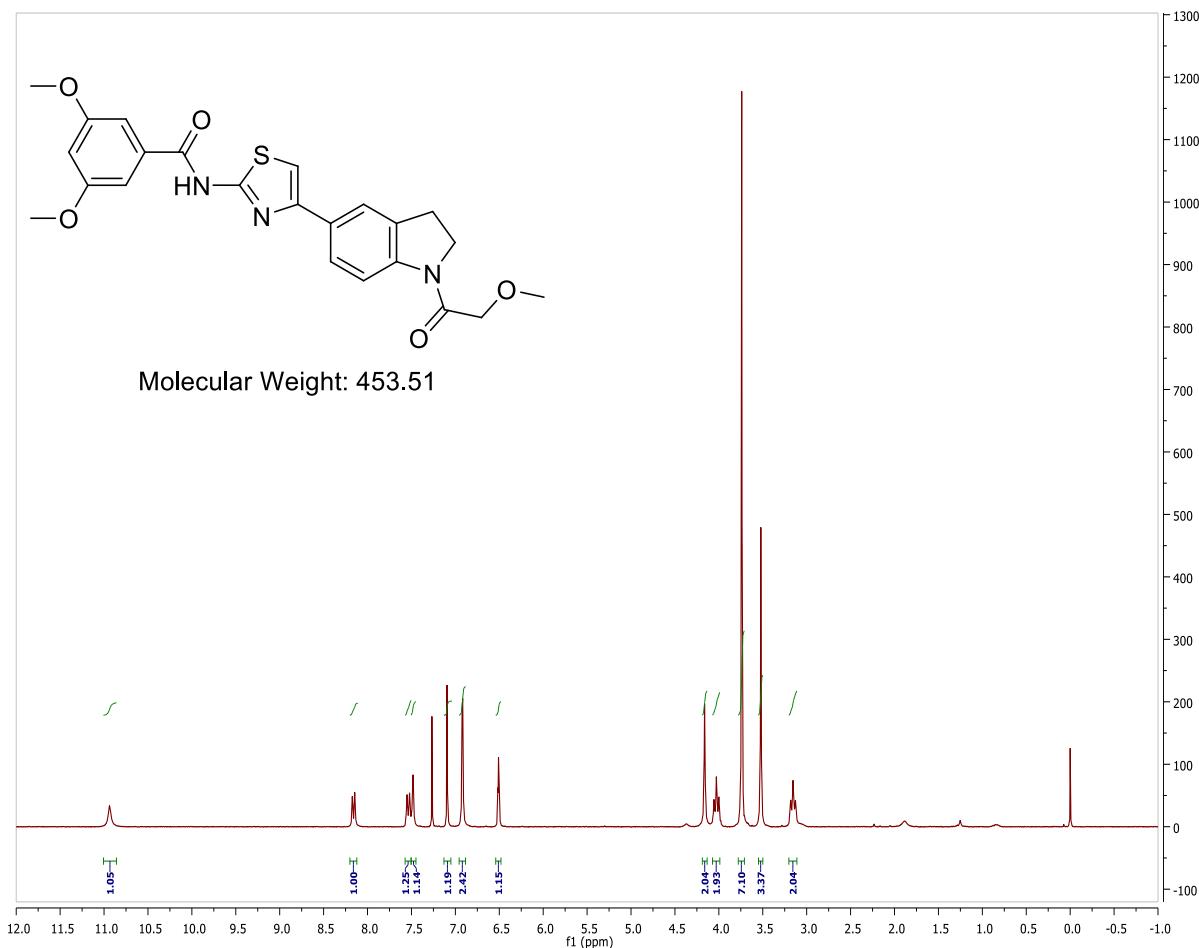
N-(4-(Indolin-5-yl)thiazol-2-yl)-3,5-dimethoxybenzamide, trifluoroacetic acid salt: *tert*-Butyl 5-(2-(3,5-dimethoxybenzamido)thiazol-4-yl)indoline-1-carboxylate (0.38 g, 0.79 mmol) was placed in a round-bottom flask with a magnetic stirbar and dissolved in dichloromethane (5.3 mL) to give a clear, yellow solution. Triethylsilane (1.3 mL, 7.93 mmol) was added followed by 2,2,2-trifluoroacetic acid (1.2 mL, 15.9 mmol), and the reaction quickly turned light orange and produced a gas. The reaction was stirred at room temperature for 1 hour. The light pink reaction was concentrated under reduced pressure to give a rose-colored solid. The crude material was suspended in diethyl ether (10 ml) and filtered. The collected solids were washed

with additional diethyl ether and air dried on the filter to give the title compound as light purple solid (0.36 g, 93 %). **¹H NMR (300 MHz, CD₃OD):** δ 8.05 (s, 1H), 7.99 (d, *J* = 10.0 Hz, 1H), 7.51 (s, 1H), 7.39 (d, *J* = 8.3 Hz, 1H), 7.20 (s, 1H), 7.20 (s, 1H), 6.74 (s, 1H), 3.87 (s, 6H), 3.85 (t, *J* = 7.8 Hz, 2H), 3.35 (t, *J* = 7.7 Hz, 2H); **MS (ESI⁺):** 382 (M+H).

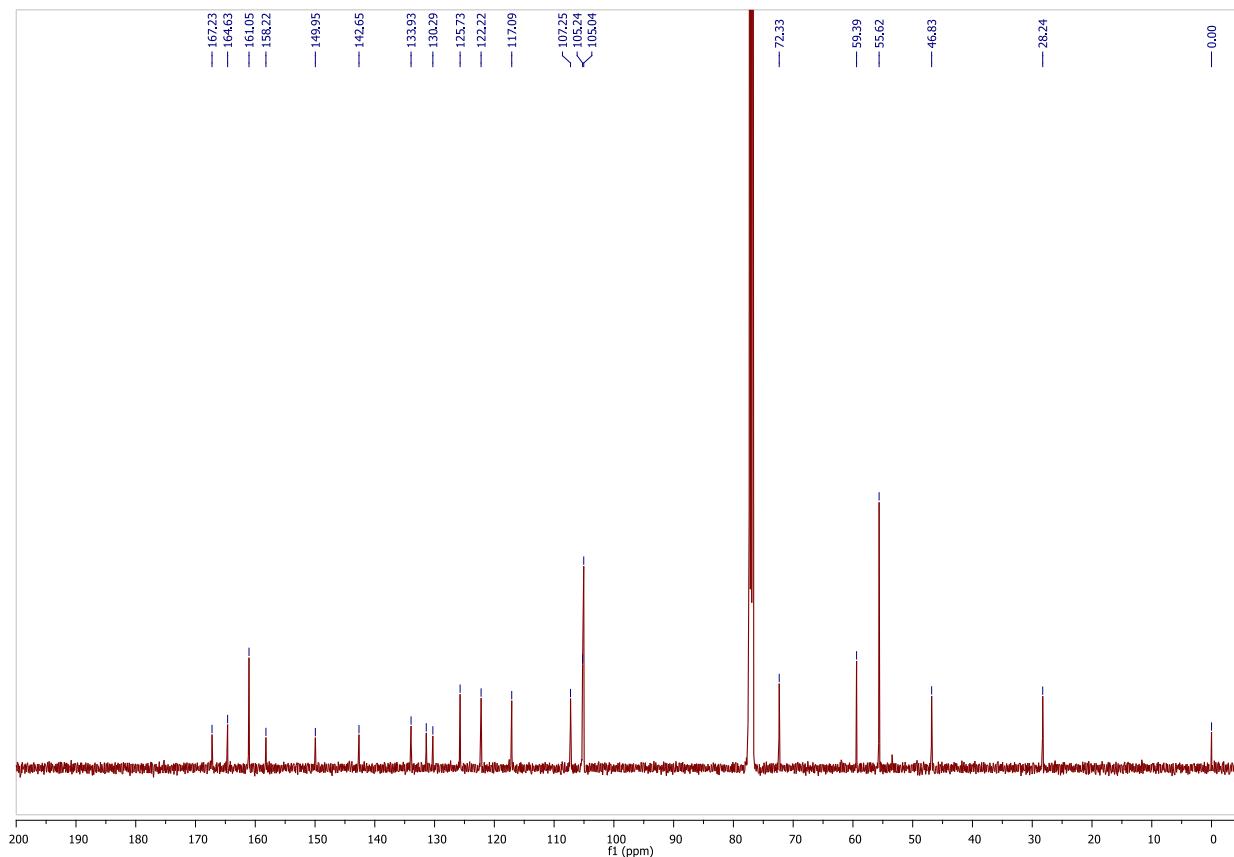


3,5-Dimethoxy-N-(4-(1-(2-methoxyacetyl)indolin-5-yl)thiazol-2-yl)benzamide (17-11, ML278): A round-bottom flask equipped with a magnetic stir bar was charged with 5-(2-(3,5-dimethoxybenzamido)thiazol-4-yl)indolinium 2,2,2-trifluoroacetate (75.0 mg, 0.15 mmol). Dichloromethane (1.5 ml) was added to produce a purple suspension that was cooled to 0 °C. 4-(*N,N*-Dimethylamino)pyridine (1.8 mg, 0.015 mmol) was added, followed by *N,N*-diisopropylethylamine (79 μl, 0.45 mmol) to give a clear, dark solution. 2-Methoxyacetyl chloride (15 μl, 0.17 mmol) was added last. The reaction was warmed to room temperature and stirred for 1 hour. The cloudy, tan mixture was diluted with dichloromethane (2 ml) and quenched with saturated sodium bicarbonate (aqueous, 2 ml). The layers were separated, and the aqueous portion was extracted with additional hot ethyl acetate (5 x 3 ml), with both layers heated to ~50 °C. The combined organic layers were shaken over magnesium sulfate, filtered, and concentrated under reduced pressure to give a light tan solid. The crude material was purified by column chromatography over silica gel (dichloromethane/methanol: 100/0 to 97/3) to give the title compound as an off-white solid (54.3 mg, 79 %). **¹H NMR (300 MHz, CDCl₃):** δ 10.14 (s, 1H), 8.23 (d, *J* = 8.3 Hz, 1H), 7.63 (d, *J* = 8.4 Hz, 1H), 7.60 (s, 1H), 7.11 (s, 1H), 7.01 (d, *J* = 1.9 Hz, 3H), 6.61 (t, *J* = 2.0 Hz, 1H), 4.18 (s, 2H), 4.07 (t, *J* = 8.3 Hz, 2H), 3.81 (s, 8H), 3.53 (s, 4H), 3.22 (t, *J* = 8.2 Hz, 2H). **¹³C NMR (125 MHz, CDCl₃):** δ 167.2, 164.6, 161.1, 158.2, 150.0, 142.7, 133.9, 131.4, 130.3, 125.7, 122.2, 117.1, 107.3, 105.2, 105.0, 72.3, 59.4, 55.6, 46.8, 28.2. **HRMS (ESI⁺):** calculated for C₂₃H₂₄N₃O₅S [M+H] 454.1437, found 454.1420.

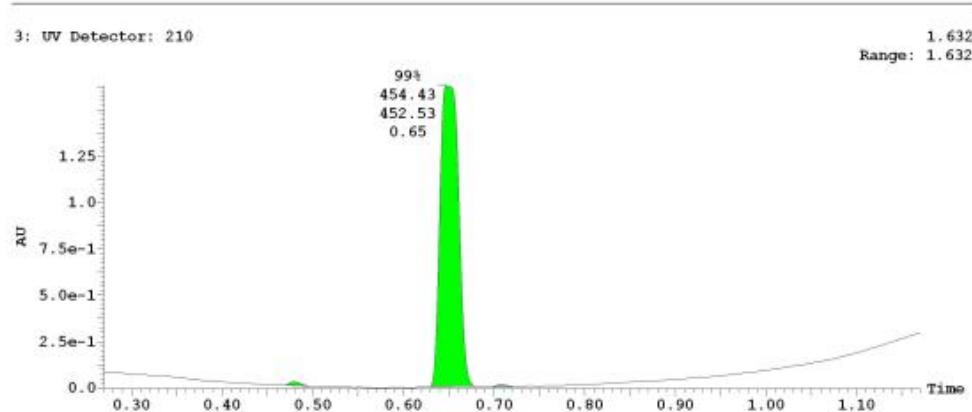
¹H NMR Spectrum (300 MHz, CDCl₃) of Probe 1 (ML278)



^{13}C NMR Spectrum (125 MHz, CDCl_3) of Probe 1 (ML278)

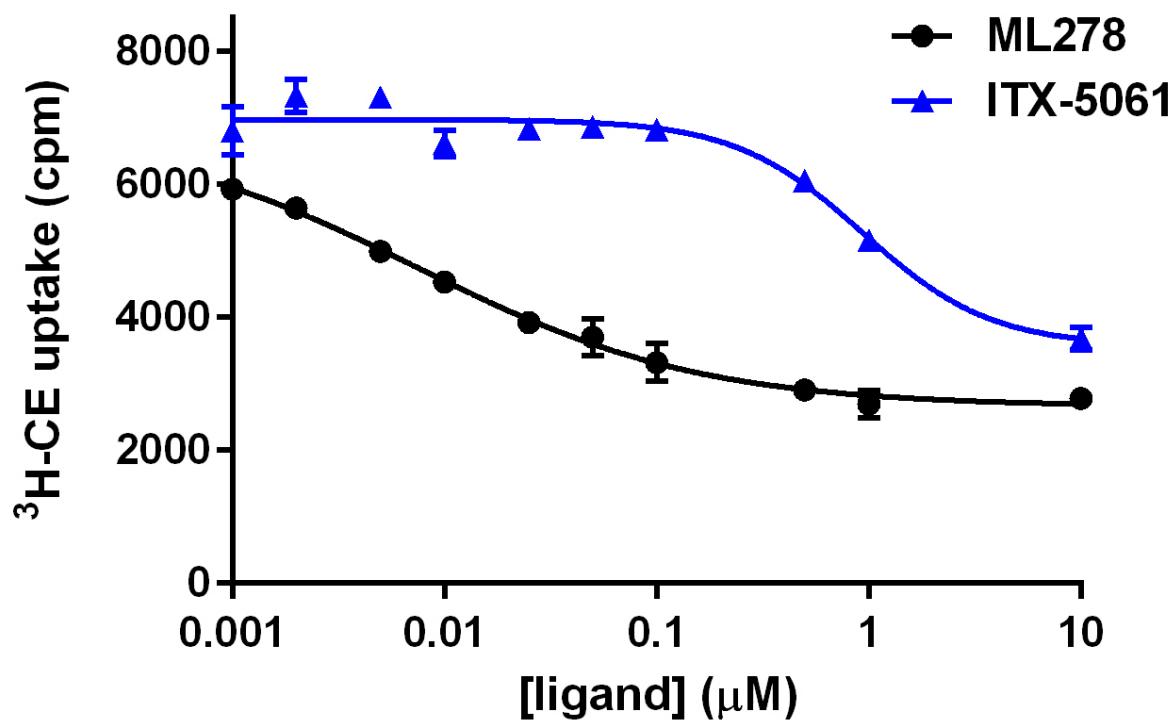


UPLC-MS Chromatogram of Probe 1 FW = 453.51 (ML278)



3. Dose-response for [³H]cholesteryl ester uptake assay

Figure S1. Dose-response for [³H]cholesteryl oleate uptake from HDL particles to ldlA[m-SR-BI] cells. Non-selective uptake (~5% of maximal uptake, see assay protocol) was quantified and subtracted from each uptake measurement prior to analysis. Points indicate the mean of $n = 3$, error bars indicate S.E.M. Uptake with no added inhibitor (ligand concentration = 0) was 7762 ± 302 cpm. IC₅₀s were determined (Table S2) and curves were plotted using non-linear regression analysis (4 parameters), using GraphPad Prism v. 6.



4. Probe comparison

Table S2. Comparison of SR-BI probes to prior art compounds^a

Property	BLT-1	ITX-5061	ML278 (17-11)
Inhibition of Dil-HDL uptake in IdIA[mSR-BI] cells: IC ₅₀	0.011±0.002 μM	0.254 μM	0.006±0.0003 μM
Inhibition of [³ H]CE-HDL uptake in IdIA[mSR-BI] cells: IC ₅₀	0.05 μM	0.94 μM	0.007 μM
Increase in Alexa488-HDL binding with IdIA[mSR-BI] cells: IC ₅₀	0.09 μM	NT	0.035 μM
Selectivity; receptor mediated endocytosis	No effect at up to 35 μM	No effect at up to 35 μM	No effect at up to 35 μM
Reversible	No ¹	NT	Yes
Cellular toxicity in IdIA[mSR-BI]	CC ₅₀ = 2.1 μM at 24 h	Non-toxic with CC ₅₀ >35 μM at 24 h	Non-toxic with CC ₅₀ >35 μM at 24 h
PBS Solubility (1% DMSO, pH 7.4, 23 °C)	1.3 μM	<1 μM	0.57 μM
Plasma protein binding, % bound (mouse, human)	NT	NT	>99%, 94%
Plasma stability, % remaining after 5 h (mouse, human)	NT	NT	>99%, >99%
Microsomal stability, % remaining after 1 h (mouse, human)	NT	NT	75%, 48%

^a Average of at least two measurements in Dil uptake assay, ± standard error of mean when n > 2. NT = not tested.

5. Compound profiling protocols

Solubility. Solubility was determined in phosphate buffered saline (PBS) pH 7.4 with 1% DMSO. Each compound was prepared in duplicate at 100 μM in both 100% DMSO and PBS with 1% DMSO. Compounds were allowed to equilibrate at room temperature with a 250 rpm orbital shake for 24 hours. After equilibration, samples were analyzed by UPLC-MS (Waters, Milford, MA) with compounds detected by SIR detection on a single quadrupole mass spectrometer. The DMSO samples were used to create a two-point calibration curve to which the response in PBS was fit.

Plasma Protein Binding. Plasma protein binding was determined by equilibrium dialysis using the Rapid Equilibrium Dialysis (RED) device (Pierce Biotechnology, Rockford, IL) for both human and mouse plasma. Each compound was prepared in duplicate at 5 μ M in plasma (0.95% acetonitrile, 0.05% DMSO) and added to one side of the membrane (200 μ l) with PBS pH 7.4 added to the other side (350 μ l). Compounds were incubated at 37 °C for 5 hours with a 250-rpm orbital shake. After incubation, samples were analyzed by UPLC-MS (Waters, Milford, MA) with compounds detected by SIR detection on a single quadrupole mass spectrometer.

Plasma Stability. Plasma stability was determined at 37 °C at 5 hours in both human and mouse plasma. Each compound was prepared in duplicate at 5 μ M in plasma diluted 50/50 (v/v) with PBS pH 7.4 (0.95% acetonitrile, 0.05% DMSO). Compounds were incubated at 37 °C for 5 hours with a 250-rpm orbital shake with time points taken at 0 hours and 5 hours. Samples were analyzed by UPLC-MS (Waters, Milford, MA) with compounds detected by SIR detection on a single quadrupole mass spectrometer.

Microsomal Stability. Microsomal stability was determined at 37 °C at 60 minutes in both human and mouse microsomes. Each compound was prepared in duplicate at 1 μ M with 0.3 mg/mL microsomes in PBS pH 7.4 (1% DMSO). Compounds were incubated at 37 °C for 60 minutes with a 350-rpm orbital shake with time points taken at 0 minutes and 60 minutes. Samples were analyzed by UPLC-MS (Waters, Milford, MA) with compounds detected by selected ion recording (SIR) on a single quadrupole mass spectrometer.

6. Assay protocols

Materials and Reagents

- DiI-HDL, custom purified HDL particles derived from human blood were prepared and labeled with 1,1'-dioctadecyl-3,3,3',3'- tetramethylindocarbocyanine perchlorate (DiI; Catalog No. D-282; Invitrogen, Carlsbad, CA). See section 5e) for details.
- Alexa 488 HDL, human HDL particles labeled with the Alexa Fluor® 488 Protein Labeling Kit (Catalog No. A-10235, Invitrogen; Carlsbad, CA) were purified and labeled by the Assay Provider.
- CellTiter-Glo® Luminescent Cell Viability Assay was purchased from Promega (Catalog No. G7573; Madison, WI).
- Radiolabeled cholesterol [$1,2^3\text{H}(\text{N})$]-, 1 mCi (37 MBq) was obtained from PerkinElmer – NEN (Catalog No. NET139001MC; Waltham, MA).
- Alexa Fluor-594 conjugated human transferrin (Catalog No. T-13343) was obtained from Invitrogen.

Cell Lines

The following cell lines were used in this study:

- ldlA[mSR-BI] is a Chinese hamster ovary (CHO) cell line that overexpresses murine SR-BI, isoform 1 (NP_058021) and lacks the LDL receptor was obtained from the Assay Provider (Krieger Laboratory). This cell line was used for the primary assay and several secondary assays. A variant of this cell line expressing mutant SR-BI, where a cysteine required for interaction with BLT-1 is mutated to serine (C384S SR-BI), was used in several secondary assays.
- [ldlA7] is the parental cell line to ldlA[mSR-BI] cells, and does not overexpresses SR-BI, and can be used to rule out compound activity independent of SR-BI.

a) DiI-HDL Uptake Assay

ldlA[mSR-BI] cells were plated into 384-well plates at 30 µl per well and incubated overnight. As a measurable surrogate for cholesterol uptake, human HDL particles were treated with the lipophilic fluorescent dye DiI and exposed to ldlA[mSR-BI] cells in lipoprotein-free media (Ham's F12/0.5% fatty acid-free Bovine Serum Albumin [BSA]/25 mM HEPES pH 7.4 plus 10 µg protein/ml DiI-HDL). Cells took up the Dil via SR-BI over 3 hours in the presence of compound. After significant uptake of the DiI, the cells became fluorescent. The level of fluorescence correlates with the amount of Dil uptake and can be measured with a standard plate reader. The uptake of lipid (represented by DiI) was inhibited by the compound BLT-1 or with an excess of HDL untreated with DiI.

The ldlA[mSR-BI] cells used in the assay were a CHO cell line lacking expression of the LDL receptors and overexpressing the Scavenger receptor (SR-BI). Inhibitors of SR-BI and HDL-mediated uptake lead to a reduction in fluorescence in this assay. Fluorescence was measured using a PerkinElmer EnVision plate reader. Primary HTS data were analyzed in Genedata Screener Assay Analyzer, and were normalized against DMSO and 1 µM BLT-1 (positive control). For the HTS, the average of two replicates was used to rank order activity and to choose compounds for retests. For dose studies, the curves were generated with Genedata Condeseo and showed percent (%) activity for the individual doses.

Detailed protocol

DiI-HDL Labeling Mix

2200 ml of Ham's/0.5% BSA/25 mM HEPES pH 7.4

DiI-HDL to 10 ug protein/ml final concentration (Stocks vary from 2.81-4 mg/mL)

Ham's/0.5% BSA/25 mM HEPES Assay Media (1 liter)

5 g BSA powder (Sigma fatty acid-free)

1 Liter Ham's F12 base media
25 ml 1 M HEPES pH 7.4

ldlA[mSR-BI] Chinese Hamster Ovary (CHO) cells were maintained in Ham's F12K (Cellgro Catalog No. 10-080-CV)/10% Fetal Calf Serum (Hyclone Catalog No. SH30071.03, Lot No. ATG32533)/ 1x Penicillin/streptomycin-L-Glutamine (Gibco Catalog No.10378-016)/200 ug/mL Geniticide (Gibco Catalog No.10131-027, Lot No.802967). Cells were fluid changed every 2 days and/or split upon reaching 80% confluence. For the primary HTS, cells were thawed at 6 million cells per Falcon T225 flask. After 2 days, the cells were passed to a Corning Hyperflask and plated after 3 days in the Hyperflask with a fluid change the day prior to plating.

On Day 0:

1. Plate 10,000 ldlA[mSR-BI] cells in 30 uL per well in Ham's F12K (Cellgro Catalog No. 10-080-CV)/5% Fetal Calf Serum (Hyclone Catalog No. SH30071.03, Lot No. ATG32533)/ 1x Penicillin/streptomycin-L-Glutamine (Gibco Catalog No. 10378-016) with a Thermo Combi Multi-drop fluid handler.
2. Use Aurora black 384-well, square, clear-bottomed, image-quality plates (Aurora Catalog No. 1022-11330) for the assay.

On Day 1:

1. Remove media with aspirator (Bio-Tek ELX405 plate washer).
2. Add 30 uL Ham's F12/0.5% Bovine Serum Albumin (fatty acid-free) /25 mM HEPES pH 7.4 (Invitrogen) + 10 ug protein/ml DiI-HDL with Thermo Combi fluid handler (slow speed setting).
3. Pin transfer 100 nL compounds and positive control (1 uM BLT-1). **It is recommended that low solubility compounds (such as ML278) be sonicated in DMSO to ensure complete dissolution prior to preparing stock solutions.**
4. Incubate 3 hours @ 37°C in humidified cell culture incubator.
5. Remove media with aspirator.
6. Wash twice with 30 uL PBS (+Ca and Mg) using the Thermo Combi on slow speed setting.
7. Analyze DiI-HDL uptake with Perkin-Elmer EnVision plate reader (Bodipy TMR mirror #405, excitation filter is Photometric 550 (#312) and emission filter is Cy3 595 (#229) with bottom read.

b) HDL Binding Assay

HDL binding was assessed using Alexa Fluor 488-labeled HDL particles. For this assay, the Alexa 488 dye was covalently bound to apolipoprotein components of the HDL particle via primary amines; thus, no transfer of the fluorophore to cell membranes occurs. In this manner, direct binding of the HDL particles to SR-BI can be measured. As a positive control, BLT-1 was used at 1 μ M, which is known to increase binding of HDL to SR-BI. It is possible that a compound can reduce binding of HDL to the receptor, and this would lead to a decrease in signal. This assay is used to characterize the mechanism of action of a particular compound; therefore, any outcome in the assay is acceptable. Data were normalized against DMSO and BLT-1 (positive control) wells in Genedata Assay Analyzer. Curves were generated with Genedata Condeseo and showed percent (%) activity for the individual doses.

Detailed protocol

ldlA[mSR-BI] Chinese Hamster Ovary (CHO) cells were maintained in Ham's F12K (Cellgro Catalog No. 10-080-CV)/10% Fetal Calf Serum (Hyclone Catalog No. SH30071.03, Lot No. ATG32533)/ 1x Penicillin-streptomycin-L-Glutamine (Gibco 10378-016)/200 ug/mL Geniticide (Gibco Catalog No.10131-027, Lot No.802967). Cells were fluid changed every 2 days and/or split upon reaching 80% confluence.

On Day 0:

1. Plate 10,000 ldlA[mSR-BI] cells in 30 uL per well in Ham's F12K (Cellgro Catalog No. 10-080-CV)/5% Fetal Calf Serum (Hyclone Catalog No. SH30071.03, Lot No. ATG32533)/ 1x Penicillin-streptomycin-L-Glutamine (Gibco 10378-016).
2. Use Aurora black, 384-well, square, clear-bottomed image-quality plates (Aurora Catalog No.1022-11330) for the assay.

On Day 1:

1. Remove media with aspirator (ELX405 plate washer).
2. Add 30 uL Ham's F12/0.5% Bovine Serum Albumin (fatty acid-free) (Sigma Catalog No. A8806-5G) /25 mM HEPES pH 7.4 (Invitrogen) plus 10 ug/mL Alexa 488-HDL with Thermo Combi fluid handler (slow speed setting).
3. Pin transfer 100 nL compounds and positive control (1 uM BLT-1).
4. Incubate 3 hours @ 37°C in a humidified cell culture incubator.
5. Remove media with aspirator.
6. Wash twice with 30 uL PBS (+Ca and Mg) using the Thermo Combi on slow speed setting.
7. Analyze Alexa-488-HDL binding with Perkin-Elmer EnVision plate reader (FITC mirror #403) with bottom read

c) Cholesterol Ester Uptake Assay

The goal of this assay is to verify compounds that disrupt the binding of HDL particles to SR-BI scavenger receptor using an alternative means of labeling cholestrylo esters and avoiding any type of fluorescence measurement. To measure this binding event, HDL particles are loaded with [³H]cholesteryl oleate ester and added to mSR-BI cells. Cells take up HDL and the [³H]cholesteryl ester via the SR-BI scavenger receptor in 2 to 3 hours. After significant uptake of the HDL, the radiolabel can be detected by liquid scintillation counting. The level of radioactivity correlates with the amount of [³H]cholesteryl ester uptake. The uptake can be inhibited by the compound BLT-1 or when co-treated with an excess of unlabeled HDL. The ldlA[mSR-BI] cells utilized in the assay are a CHO cell line lacking expression of the LDL receptors and overexpressing the scavenger receptor, SR-BI. Inhibitors of SR-BI and HDL uptake will have a reduction in liquid scintillation counts.

Detailed protocol

On Day 0:

1. Plate 50,000 ldlA[mSR-BI] cells in 1 mL per well in Ham's F12K/5% Fetal Calf Serum/ 1x Penicillin-streptomycin-L-Glutamine/G418.

On Day 2:

2. Wash cells with prewarmed (37°C) Ham's F12K medium plus 0.5% (wt/vol) fatty acid free BSA prior to adding radioactive lipoprotein.
3. Preincubate cells with 200 uL of inhibitors at indicated concentration or non-labeled HDL (at a 40 fold excess for measurement of non-specific activity) in Ham's F12/0.5% BSA/0.5%DMSO (vol/vol) for 1 hr at 37°C in a humidified cell culture incubator. **It is recommended that low solubility compounds (such as ML278) be sonicated in DMSO to ensure complete dissolution prior to preparing stock solutions.**
4. Add 50 uL Ham's F12/0.5% BSA/0.5%DMSO plus 50 ug/mL [³H]CE-HDL to reach a final concentration of 10 ug/mL and incubate 2 hours at 37°C in a humidified cell culture incubator
5. Remove the medium at the end of incubation at 4C
6. Wash the plates 2X with Tris-BSA buffer and 1X Tris-HCl buffer.
7. Add 1 ml isopropanol, and incubate for 20 min at 4C.
8. Collect 1 ml isopropanol/lysate into counting tube, add 4 ml liquid scintillation buffer, and prepare for counting.
9. Specific uptake is the difference between total and nonspecific activities.
10. Data were analyzed using Prism 6 (GraphPad Software).
11. All calculated errors represent standard errors of the mean.

e) HDL isolation and labeling

Human HDL was isolated from donors by density gradient ultracentrifugation and labeled with [³H]cholesteryl ester ([³H]CE, [³H]CE-HDL) as described previously.^{2,3,4,5,6}

7. References

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- ³ Chung, B. H.; Wilkinson, T.; Geer, J. C.; Segrest, J. P. Preparative and quantitative isolation of plasma lipoproteins: rapid, single discontinuous density gradient ultracentrifugation in a vertical rotor, *J Lipid Res.* **1980**, *21*, 284–291.
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- ⁵ Goldstein, J. L.; Basu, S. K.; Brown, M. S. Receptor-mediated endocytosis of low-density lipoprotein in cultured cells. *Methods Enzymol.* **1983**, *98*, 241–260.
- ⁶ Gwynne, J. T.; Mahaffee, D. D. Rat adrenal uptake and metabolism of high density lipoprotein cholestryl ester. *J. Biol. Chem.* **1989**, *264*, 8141–8150.

8. Off-target screening data for 17-11 (ML278)

Services Performed

LeadProfilingScreen (Total # of Assays: 67)

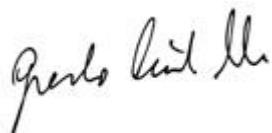
Study Objectives

To evaluate, in LeadProfilingScreen, the activity of test compound BRD-K83634925-001-01-1 (Ricerca # 1162315).

Study Signatures



Hua-Chou Hsing
Study Director for Binding Assays



Gonzalo Castillo, Ph.D.
Technical Director

"This study was conducted according to the procedures described in this report. All data presented are authentic, accurate and correct to the best of our knowledge."

Table of Contents

Report Section	Page
Cover Page	1
Study Signatures	2
Table of Contents	3
Summary	4
Summary of Significant Results	5
Experimental Results	6
Methods	9
Reference Compounds	26
Appendix	28

Summary

STUDY OBJECTIVE

To evaluate, in Radioligand Binding assays, the activity of compound BRD-K83634925-001-01-1 (BRD-9, PT# 1162315).

METHODS

Methods employed in this study have been adapted from the scientific literature to maximize reliability and reproducibility. Reference standards were run as an integral part of each assay to ensure the validity of the results obtained. Assays were performed under conditions described in the accompanying "Methods" section of this report.

Where presented, IC₅₀ values were determined by a non-linear, least squares regression analysis using MathIQ™ (ID Business Solutions Ltd., UK). Where inhibition constants (K_i) are presented, the K_i values were calculated using the equation of Cheng and Prusoff (Cheng, Y., Prusoff, W.H., Biochem. Pharmacol. 22:3099-3108, 1973) using the observed IC₅₀ of the tested compound, the concentration of radioligand employed in the assay, and the historical values for the K_D of the ligand (obtained experimentally at **Ricerca Biosciences, LLC**). Where presented, the Hill coefficient (n_H), defining the slope of the competitive binding curve, was calculated using MathIQ™. Hill coefficients significantly different than 1.0, may suggest that the binding displacement does not follow the laws of mass action with a single binding site. Where IC₅₀, K_i, and/or n_H data are presented without Standard Error of the Mean (SEM), data are insufficient to be quantitative, and the values presented (K_i, IC₅₀, n_H) should be interpreted with caution.

RESULTS

A summary of results meeting the significance criteria is presented in the following sections. Complete results are presented under the section labeled "Experimental Results". Individual responses, if requested, are presented in the section labeled "Individual Responses".

SUMMARY/CONCLUSION

Significant results are displayed in the following table(s) in rank order of potency for estimated IC₅₀ and/or K_i values.

Summary of Significant Results

Biochemical assay results are presented as the percent inhibition of specific binding or activity throughout the report. All other results are expressed in terms of that assay's quantitation method.

- For primary assays, only the lowest concentration with a significant response judged by the assays' criteria, is shown in this summary.
- Where applicable, either the secondary assay results with the lowest dose/concentration meeting the significance criteria or, if inactive, the highest dose/concentration that did not meet the significance criteria is shown.
- Unless otherwise requested, primary screening in duplicate with quantitative data (e.g., $IC_{50} \pm SEM$, $K_i \pm SEM$ and n_H) are shown where applicable for individual requested assays. In screening packages, primary screening in duplicate with semi-quantitative data (e.g., estimated IC_{50} , K_i and n_H) are shown where applicable (concentration range of 4 log units); available secondary functional assays are carried out (30 mM) and MEC or MIC determined only if active in primary assays >50% at 1 log unit below initial test concentration.

Significant responses ($\geq 50\%$ inhibition or stimulation for Biochemical assays) were noted in the primary assays listed below:

No significant results noted.

Experimental Results

Cat #	Assay Name	Batch*	Spec.	Rep.	Conc.	% Inh.	IC ₅₀ *	K _i	n _H	R
Compound: BRD-K83634925-001-01-1, PT #: 1162315										
200510	Adenosine A ₁	317535	hum	2	10 µM	20				
200610	Adenosine A _{2A}	317540	hum	2	10 µM	24				
200720	Adenosine A ₃	317577	hum	2	10 µM	43				
203100	Adrenergic α _{1A}	317536	rat	2	10 µM	3				
203200	Adrenergic α _{1B}	317547	rat	2	10 µM	13				
203400	Adrenergic α _{1D}	317537	hum	2	10 µM	5				
203620	Adrenergic α _{2A}	317534	hum	2	10 µM	2				
204010	Adrenergic β ₁	317532	hum	2	10 µM	2				
204110	Adrenergic β ₂	317533	hum	2	10 µM	4				
285010	Androgen (Testosterone) AR	317485	rat	2	10 µM	21				
212510	Bradykinin B ₁	317578	hum	2	10 µM	-7				
212620	Bradykinin B ₂	317579	hum	2	10 µM	3				
214510	Calcium Channel L-Type, Benzothiazepine	317482	rat	2	10 µM	16				
214600	Calcium Channel L-Type, Dihydropyridine	317531	rat	2	10 µM	20				
216000	Calcium Channel N-Type	317580	rat	2	10 µM	-17				
217030	Cannabinoid CB ₁	317529	hum	2	10 µM	8				
219500	Dopamine D ₁	317541	hum	2	10 µM	6				
219700	Dopamine D _{2S}	317543	hum	2	10 µM	-3				
219800	Dopamine D ₃	317544	hum	2	10 µM	7				
219900	Dopamine D _{4.2}	317552	hum	2	10 µM	-6				
224010	Endothelin ET _A	317641	hum	2	10 µM	12				
224110	Endothelin ET _B	317642	hum	2	10 µM	5				
225510	Epidermal Growth Factor (EGF)	317582	hum	2	10 µM	14				
226010	Estrogen ER _α	317583	hum	2	10 µM	6				
226600	GABA _A , Flunitrazepam, Central	317545	rat	2	10 µM	-5				
226500	GABA _A , Muscimol, Central	317556	rat	2	10 µM	8				
228610	GABA _{B1A}	317584	hum	2	10 µM	-12				
232030	Glucocorticoid	317476	hum	2	10 µM	-24				
232700	Glutamate, Kainate	317586	rat	2	10 µM	4				
232810	Glutamate, NMDA, Agonism	317587	rat	2	10 µM	-16				
232910	Glutamate, NMDA, Glycine	317588	rat	2	10 µM	-1				
233000	Glutamate, NMDA, Phencyclidine	317557	rat	2	10 µM	1				
239610	Histamine H ₁	317546	hum	2	10 µM	9				
239710	Histamine H ₂	317514	hum	2	10 µM	8				

Note: Items meeting criteria for significance ($\geq 50\%$ stimulation or inhibition) are highlighted.

* Batch: Represents compounds tested concurrently in the same assay(s).

ham=Hamster; hum=Human

Broad Institute

Study #: AB13141, Quote #: 30266-1, Compound Code: BRD-K83634925-001-01-1 (1162315)

Experimental Results

Cat #	Assay Name	Batch*	Spec.	Rep.	Conc.	% Inh.	IC ₅₀ *	K _i	n _H	R
239820	Histamine H ₃	317553	hum	2	10 µM	0				
241000	Imidazoline I ₂ , Central	317539	rat	2	10 µM	15				
243520	Interleukin IL-1	317484	mouse	2	10 µM	-10				
250460	Leukotriene, Cysteinyl CysLT ₁	317589	hum	2	10 µM	-6				
251600	Melatonin MT ₁	317513	hum	2	10 µM	-3				
252610	Muscarinic M ₁	317554	hum	2	10 µM	11				
252710	Muscarinic M ₂	317527	hum	2	10 µM	-9				
252810	Muscarinic M ₃	317528	hum	2	10 µM	0				
257010	Neuropeptide Y Y ₁	317468	hum	2	10 µM	-8				
257110	Neuropeptide Y Y ₂	317592	hum	2	10 µM	-1				
258590	Nicotinic Acetylcholine	317521	hum	2	10 µM	0				
258700	Nicotinic Acetylcholine α ₁ , Bungarotoxin	317522	hum	2	10 µM	-3				
260130	Opiate δ ₁ (OP1, DOP)	317525	hum	2	10 µM	-8				
260210	Opiate κ(OP2, KOP)	317524	hum	2	10 µM	2				
260410	Opiate μ(OP3, MOP)	317526	hum	2	10 µM	-15				
264500	Phorbol Ester	317558	mouse	2	10 µM	-4				
265010	Platelet Activating Factor (PAF)	317496	hum	2	10 µM	23				
265600	Potassium Channel [K _{ATP}]	317559	ham	2	10 µM	-1				
265900	Potassium Channel hERG	317560	hum	2	10 µM	-4				
268420	Prostanoid EP ₄	317561	hum	2	10 µM	2				
268700	Purinergic P _{2X}	317590	rabbit	2	10 µM	7				
268810	Purinergic P _{2Y}	317591	rat	2	10 µM	9				
270000	Rolipram	317562	rat	2	10 µM	8				
271110	Serotonin (5-Hydroxytryptamine) 5-HT _{1A}	317510	hum	2	10 µM	0				
271700	Serotonin (5-Hydroxytryptamine) 5-HT _{2B}	317551	hum	2	10 µM	24				
271910	Serotonin (5-Hydroxytryptamine) 5-HT ₃	317593	hum	2	10 µM	22				
278110	Sigma σ ₁	317523	hum	2	10 µM	-20				
255520	Tachykinin NK ₁	317474	hum	2	10 µM	7				
285900	Thyroid Hormone	317488	rat	2	10 µM	10				
220320	Transporter, Dopamine (DAT)	317469	hum	2	10 µM	-1				
226400	Transporter, GABA	317538	rat	2	10 µM	32				
204410	Transporter, Norepinephrine (NET)	317520	hum	2	10 µM	4				
274030	Transporter, Serotonin (5-Hydroxytryptamine) (SERT)	317549	hum	2	10 µM	9				

Note: Items meeting criteria for significance ($\geq 50\%$ stimulation or inhibition) are highlighted.

* Batch: Represents compounds tested concurrently in the same assay(s).

ham=Hamster; hum=Human

Broad Institute

Study #: AB13141, Quote #: 30266-1, Compound Code: BRD-K83634925-001-01-1 (1162315)

Tuesday, June 26, 2012

Page 7 of 28

Experimental Results

MDS has an exclusive, worldwide limited use license from Synaptic Pharmaceutical Corporation to perform these assays: Adrenergic Alpha 1D, Adrenergic Alpha 2B, and Dopamine D5 for safety and selectivity profiling. MDS' license excludes performing those assays in connection with drug discovery or development activities where the principal therapeutic mechanism of action of the test compound involves selective binding to a licensed receptor. Customers may contact Synaptic directly if they believe they need a broader license.

Note: Items meeting criteria for significance ($\geq 50\%$ stimulation or inhibition) are highlighted.

* Batch: Represents compounds tested concurrently in the same assay(s).

ham=Hamster; hum=Human

Methods

■ 200510 Adenosine A₁

Source:	Human recombinant CHO cells	Ligand:	1.0 nM [³ H] DPCPX
Vehicle:	1% DMSO	Non-Specific Ligand:	100 µM R(-)-PIA
Incubation Time/Temp:	90 minutes @ 25°C	Specific Binding:	85% *
Incubation Buffer:	20 mM HEPES, pH 7.4, 10 mM MgCl ₂ , 100 mM NaCl	Quantitation Method:	Radioligand Binding
Kd:	1.40 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	2.70 pmole/mg Protein *

■ 200610 Adenosine A_{2A}

Source:	Human recombinant HEK-293 cells	Ligand:	0.050 µM [³ H] CGS-21680
Vehicle:	1% DMSO	Non-Specific Ligand:	50.0 µM NECA
Incubation Time/Temp:	90 minutes @ 25°C	Specific Binding:	85% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 10 mM MgCl ₂ , 1 mM EDTA, 2 U/mL Adenosine Deaminase	Quantitation Method:	Radioligand Binding
Kd:	0.064 µM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	7.0 pmole/mg Protein *

■ 200720 Adenosine A₃

Source:	Human recombinant CHO-K1 cells	Ligand:	0.50 nM [¹²⁵ I] AB-MECA
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 µM IB-MECA
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	83% *
Incubation Buffer:	25 mM HEPES, pH 7.4, 5 mM MgCl ₂ , 1 mM CaCl ₂ , 0.1% BSA	Quantitation Method:	Radioligand Binding
Kd:	5.90 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	1.80 pmole/mg Protein *

■ 203100 Adrenergic α_{1A}

Source:	Wistar Rat submaxillary gland	Ligand:	0.25 nM [³ H] Prazosin
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Phentolamine
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 0.5 mM EDTA	Quantitation Method:	Radioligand Binding
Kd:	0.17 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.18 pmole/mg Protein *

* Historical Values

Methods

■ 203200 Adrenergic α_{1B}

Source:	Wistar Rat liver	Ligand:	0.25 nM [³ H] Prazosin
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 μ M Phentolamine
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 0.5 mM EDTA	Quantitation Method:	Radioligand Binding
Kd:	0.31 nM *	Significance Criteria:	\geq 50% of max stimulation or inhibition
		Bmax:	0.18 pmole/mg Protein *

■ 203400 Adrenergic α_{1D}

Source:	Human recombinant HEK-293 cells	Ligand:	0.60 nM [³ H] Prazosin
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 μ M Phentolamine
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	80% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4	Quantitation Method:	Radioligand Binding
Kd:	0.58 nM *	Significance Criteria:	\geq 50% of max stimulation or inhibition
		Bmax:	0.17 pmole/mg Protein *

■ 203620 Adrenergic α_{2A}

Source:	Human recombinant insect Sf9 cells	Ligand:	1.0 nM [³ H] MK-912
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 μ M WB-4101
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	95% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 12.5 mM MgCl ₂ , 2 mM EDTA	Quantitation Method:	Radioligand Binding
Kd:	0.60 nM *	Significance Criteria:	\geq 50% of max stimulation or inhibition
		Bmax:	4.60 pmole/mg Protein *

■ 204010 Adrenergic β_1

Source:	Human recombinant CHO-K1 cells	Ligand:	0.030 nM [¹²⁵ I] Cyanopindolol
Vehicle:	1% DMSO	Non-Specific Ligand:	100 μ M S(-)-Propranolol
Incubation Time/Temp:	2 hours @ 25°C	Specific Binding:	95% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 1.4 mM Ascorbic Acid, 0.001% BSA, 5 mM EDTA, 1.5 mM CaCl ₂ , 120 mM NaCl	Quantitation Method:	Radioligand Binding
Kd:	0.041 nM *	Significance Criteria:	\geq 50% of max stimulation or inhibition
		Bmax:	0.072 pmole/mg Protein *

* Historical Values

Methods

■ 204110 Adrenergic β_2

Source:	Human recombinant CHO cells	Ligand:	0.20 nM [³ H] CGP-12177
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 μ M ICI-118551
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	95% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 0.5 mM EDTA, 5.0 mM MgCl ₂ , 120 mM NaCl	Quantitation Method:	Radioligand Binding
Kd:	0.44 nM *	Significance Criteria:	\geq 50% of max stimulation or inhibition
		Bmax:	0.44 pmole/mg Protein *

■ 285010 Androgen (Testosterone) AR

Source:	Rat recombinant E. coli	Ligand:	1.50 nM [³ H] Mibolerone
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 μ M Mibolerone
Incubation Time/Temp:	4 hours @ 4°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 0.8 M NaCl, 10% Glycerol, 2 mM Dithiothreitol, 0.1% BSA, 2% EtOH	Quantitation Method:	Radioligand Binding
Kd:	3.0 nM *	Significance Criteria:	\geq 50% of max stimulation or inhibition
		Bmax:	930 pmole/mg Protein *

■ 212510 Bradykinin B₁

Source:	Human IMR-90 cells	Ligand:	0.50 nM [³ H] (Des-Arg ¹⁰)-Kallidin
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 μ M (Des-Arg ⁹ , Leu ⁸)-Bradykinin
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	80% *
Incubation Buffer:	20 mM HEPES, pH 7.4, 125 mM N-Methyl-D-glucamine, 5 mM KCl, 1 mM 1,10-Phenanthroline, 140 μ g/ml Bacitracin	Quantitation Method:	Radioligand Binding
Kd:	0.17 nM *	Significance Criteria:	\geq 50% of max stimulation or inhibition
		Bmax:	0.55 pmole/mg Protein *

■ 212620 Bradykinin B₂

Source:	Human recombinant Chem-1 cells	Ligand:	0.50 nM [³ H] Bradykinin
Vehicle:	1% DMSO	Non-Specific Ligand:	5.0 μ M Bradykinin
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM HEPES, pH 7.4, 0.2% BSA, 1 mM CaCl ₂ , 5 mM MgCl ₂	Quantitation Method:	Radioligand Binding
Kd:	0.85 nM *	Significance Criteria:	\geq 50% of max stimulation or inhibition
		Bmax:	9.40 pmole/mg Protein *

* Historical Values

Methods

■ 214510 Calcium Channel L-Type, Benzothiazepine

Source:	Wistar Rat brain	Ligand:	2.0 nM [³ H] Diltiazem
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Diltiazem
Incubation Time/Temp:	3 hours @ 4°C	Specific Binding:	73% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 0.1% BSA	Quantitation Method:	Radioligand Binding
Kd:	0.016 µM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.21 pmole/mg Protein *

■ 214600 Calcium Channel L-Type, Dihydropyridine

Source:	Wistar Rat cerebral cortex	Ligand:	0.10 nM [³ H] Nitrendipine
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 µM Nifedipine
Incubation Time/Temp:	90 minutes @ 25°C	Specific Binding:	91% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4	Quantitation Method:	Radioligand Binding
Kd:	0.18 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.23 pmole/mg Protein *

■ 216000 Calcium Channel N-Type

Source:	Wistar Rat frontal brain	Ligand:	10 pM [¹²⁵ I] ω-Conotoxin GVIA
Vehicle:	1% DMSO	Non-Specific Ligand:	0.10 µM ω-Conotoxin GVIA
Incubation Time/Temp:	30 minutes @ 4°C	Specific Binding:	96% *
Incubation Buffer:	20 mM Tris-HCl, pH 7.4, 0.5% BSA	Quantitation Method:	Radioligand Binding
Kd:	0.051 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.88 pmole/mg Protein *

■ 217030 Cannabinoid CB₁

Source:	Human recombinant Chem-1 cells	Ligand:	2.0 nM [³ H] SR141716A
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM R(+)-WIN-55,212-2
Incubation Time/Temp:	90 minutes @ 37°C	Specific Binding:	70% *
Incubation Buffer:	50 mM HEPES, pH 7.4, 5 mM MgCl ₂ , 1 mM CaCl ₂ , 0.2% BSA	Quantitation Method:	Radioligand Binding
Kd:	5.90 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	15.0 pmole/mg Protein *

* Historical Values

Methods

■ 219500 Dopamine D₁

Source:	Human recombinant CHO cells	Ligand:	1.40 nM [³ H] SCH-23390
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM (+)-Butaclamol
Incubation Time/Temp:	2 hours @ 37°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 1.4 mM Ascorbic Acid, 0.001% BSA, 150 mM NaCl	Quantitation Method:	Radioligand Binding
Kd:	1.40 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.63 pmole/mg Protein *

■ 219700 Dopamine D_{2S}

Source:	Human recombinant CHO cells	Ligand:	0.16 nM [³ H] Spiperone
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Haloperidol
Incubation Time/Temp:	2 hours @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 1.4 mM Ascorbic Acid, 0.001% BSA, 150 mM NaCl	Quantitation Method:	Radioligand Binding
Kd:	0.090 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	1.60 pmole/mg Protein *

■ 219800 Dopamine D₃

Source:	Human recombinant CHO cells	Ligand:	0.70 nM [³ H] Spiperone
Vehicle:	1% DMSO	Non-Specific Ligand:	25.0 µM S(-)-Sulpiride
Incubation Time/Temp:	2 hours @ 37°C	Specific Binding:	85% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 1.4 mM Ascorbic Acid, 0.001% BSA, 150 mM NaCl	Quantitation Method:	Radioligand Binding
Kd:	0.36 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	1.10 pmole/mg Protein *

■ 219900 Dopamine D_{4.2}

Source:	Human recombinant CHO-K1 cells	Ligand:	0.50 nM [³ H] Spiperone
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Haloperidol
Incubation Time/Temp:	2 hours @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 1.4 mM Ascorbic Acid, 0.001% BSA, 150 mM NaCl	Quantitation Method:	Radioligand Binding
Kd:	0.32 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.55 pmole/mg Protein *

* Historical Values

Methods

■ 224010 Endothelin ET_A

Source:	Human recombinant CHO-K1 cells	Ligand:	0.030 nM [¹²⁵ I] Endothelin-1
Vehicle:	1% DMSO	Non-Specific Ligand:	0.10 µM Endothelin-1
Incubation Time/Temp:	2 hours @ 37°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 0.1% BSA, 0.5 mM CaCl ₂ , 0.05% Tween-20	Quantitation Method:	Radioligand Binding
Kd:	0.048 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.35 pmole/mg Protein *

■ 224110 Endothelin ET_B

Source:	Human recombinant CHO-K1 cells	Ligand:	0.10 nM [¹²⁵ I] Endothelin-1
Vehicle:	1% DMSO	Non-Specific Ligand:	0.10 µM Endothelin-1
Incubation Time/Temp:	2 hours @ 25°C	Specific Binding:	75% *
Incubation Buffer:	50 mM HEPES, pH 7.4, 1 mM CaCl ₂ , 5 mM MgCl ₂ , 0.5% BSA	Quantitation Method:	Radioligand Binding
Kd:	0.085 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	4.30 pmole/mg Protein *

■ 225510 Epidermal Growth Factor (EGF)

Source:	Human A431 cells	Ligand:	0.080 nM [¹²⁵ I] EGF (human)
Vehicle:	1% DMSO	Non-Specific Ligand:	0.10 µM EGF (human)
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM HEPES, pH 7.7, 0.1% BSA, 1.2 mM CaCl ₂ , 5 mM KCl, 1.2 mM MgSO ₄ , 138 mM NaCl	Quantitation Method:	Radioligand Binding
Kd:	0.17 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	5.50 pmole/mg Protein *

■ 226010 Estrogen ER_A

Source:	Human recombinant insect Sf9 cells	Ligand:	0.50 nM [³ H] Estradiol
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 µM Diethylstilbestrol
Incubation Time/Temp:	2 hours @ 25°C	Specific Binding:	85% *
Incubation Buffer:	10 mM Tris-HCl, pH 7.4, 0.1% BSA, 10% Glycerol, 1 mM DTT	Quantitation Method:	Radioligand Binding
Kd:	0.20 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	1400 pmole/mg Protein *

* Historical Values

Methods

■ 226600 GABA_A, Flunitrazepam, Central

Source:	Wistar Rat brain (minus cerebellum)	Ligand:	1.0 nM [³ H] Flunitrazepam
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Diazepam
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	91% *
Incubation Buffer:	50 mM Phosphate Buffer, pH 7.4	Quantitation Method:	Radioligand Binding
Kd:	4.40 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	1.20 pmole/mg Protein *

■ 226500 GABA_A, Muscimol, Central

Source:	Wistar Rat brain (minus cerebellum)	Ligand:	1.0 nM [³ H] Muscimol
Vehicle:	1% DMSO	Non-Specific Ligand:	0.10 µM Muscimol
Incubation Time/Temp:	10 minutes @ 4°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4	Quantitation Method:	Radioligand Binding
Kd:	3.80 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	1.80 pmole/mg Protein *

■ 228610 GABA_{B1A}

Source:	Human recombinant CHO cells	Ligand:	4.0 nM [³ H] CGP-54626
Vehicle:	1% DMSO	Non-Specific Ligand:	3.0 mM GABA
Incubation Time/Temp:	3 hours @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 2.5 mM CaCl ₂ , 0.1% BSA	Quantitation Method:	Radioligand Binding
Kd:	3.30 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	48.0 pmole/mg Protein *

■ 232030 Glucocorticoid

Source:	Human recombinant Insect cells	Ligand:	5.0 nM [³ H] Dexamethasone
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Dexamethasone
Incubation Time/Temp:	1 day @ 4°C	Specific Binding:	97% *
Incubation Buffer:	5 mM KH ₂ PO ₄ , 8 mM Na ₂ HPO ₄ ·12H ₂ O, pH7.4, 137 mM NaCl, 2.7 mM KCl, 0.2% BSA	Quantitation Method:	Radioligand Binding
Kd:	4.60 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	1.0 pmole/mg *

* Historical Values

Methods

■ 232700 Glutamate, Kainate

Source:	Wistar Rat brain (minus cerebellum)	Ligand:	5.0 nM [³ H] Kainic acid
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 mM L-Glutamic acid
Incubation Time/Temp:	60 minutes @ 4°C	Specific Binding:	80% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4	Quantitation Method:	Radioligand Binding
Kd:	0.012 µM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.35 pmole/mg Protein *

■ 232810 Glutamate, NMDA, Agonism

Source:	Wistar Rat cerebral cortex	Ligand:	2.0 nM [³ H] CGP-39653
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 mM L-Glutamic acid
Incubation Time/Temp:	20 minutes @ 4°C	Specific Binding:	70% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4	Quantitation Method:	Radioligand Binding
Kd:	0.019 µM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	2.30 pmole/mg Protein *

■ 232910 Glutamate, NMDA, Glycine

Source:	Wistar Rat cerebral cortex	Ligand:	0.33 nM [³ H] MDL 105,519
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM MDL 105,519
Incubation Time/Temp:	30 minutes @ 4°C	Specific Binding:	85% *
Incubation Buffer:	50 mM HEPES, pH 7.7	Quantitation Method:	Radioligand Binding
Kd:	6.0 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	3.70 pmole/mg Protein *

■ 233000 Glutamate, NMDA, Phencyclidine

Source:	Wistar Rat cerebral cortex	Ligand:	4.0 nM [³ H] TCP
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 µM Dizocilpine ((+)-MK-801)
Incubation Time/Temp:	45 minutes @ 25°C	Specific Binding:	94% *
Incubation Buffer:	10 mM Tris-HCl, pH 7.4	Quantitation Method:	Radioligand Binding
Kd:	8.40 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.78 pmole/mg Protein *

* Historical Values

Methods

■ 239610 Histamine H₁

Source:	Human recombinant CHO-K1 cells	Ligand:	1.20 nM [³ H] Pyrilamine
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 μM Pyrilamine
Incubation Time/Temp:	3 hours @ 25°C	Specific Binding:	94% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 2 mM MgCl ₂ , 100 mM NaCl, 250 mM Sucrose	Quantitation Method:	Radioligand Binding
Kd:	1.10 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	6.70 pmole/mg Protein *

■ 239710 Histamine H₂

Source:	Human recombinant CHO-K1 cells	Ligand:	0.10 nM [¹²⁵ I] Aminopotentidine
Vehicle:	1% DMSO	Non-Specific Ligand:	3.0 μM Tiotidine
Incubation Time/Temp:	2 hours @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Phosphate, pH 7.4	Quantitation Method:	Radioligand Binding
Kd:	0.45 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	6.90 pmole/mg Protein *

■ 239820 Histamine H₃

Source:	Human recombinant CHO-K1 cells	Ligand:	0.40 nM [³ H] N-α-Methylhistamine (NAMH)
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 μM R(-)-α-Methylhistamine (RAMH)
Incubation Time/Temp:	2 hours @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 5 mM MgCl ₂ , 0.1% BSA	Quantitation Method:	Radioligand Binding
Kd:	0.38 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	2.0 pmole/mg Protein *

■ 241000 Imidazoline I₂, Central

Source:	Wistar Rat cerebral cortex	Ligand:	2.0 nM [³ H] Idazoxan
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 μM Idazoxan
Incubation Time/Temp:	30 minutes @ 25°C	Specific Binding:	85% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 0.5 mM EDTA	Quantitation Method:	Radioligand Binding
Kd:	4.0 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.14 pmole/mg Protein *

* Historical Values

Methods

■ 243520 Interleukin IL-1

Source:	Mouse 3T3-SWISS cells	Ligand:	0.10 nM [¹²⁵ I] Interleukin-1β
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Interleukin-1β
Incubation Time/Temp:	2 hours @ 37°C	Specific Binding:	80% *
Incubation Buffer:	RPMI 1640, 20 mM HEPES, pH 7.4, 0.1% Sodium Azide, 1% BSA	Quantitation Method:	Radioligand Binding
Kd:	0.25 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	820 R/cell Protein *

■ 250460 Leukotriene, Cysteinyl CysLT₁

Source:	Human recombinant CHO-K1 cells	Ligand:	0.30 nM [³ H] LTD ₄
Vehicle:	1% DMSO	Non-Specific Ligand:	0.30 µM LTD ₄
Incubation Time/Temp:	30 minutes @ 25°C	Specific Binding:	93% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 5 mM CaCl ₂ , 5 mM MgCl ₂ , 100 µg/ml Bacitracin, 1 mM Benzamidine, 0.1 mM PMSF	Quantitation Method:	Radioligand Binding
Kd:	0.21 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	3.0 pmole/mg Protein *

■ 251600 Melatonin MT₁

Source:	Human recombinant CHO-K1 cells	Ligand:	0.050 nM [¹²⁵ I] 2-lodomelatonin
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 µM 6-Chloromelatonin
Incubation Time/Temp:	3 hours @ 25°C	Specific Binding:	97% *
Incubation Buffer:	25 mM HEPES, pH 7.4, 5 mM MgCl ₂ , 1 mM CaCl ₂ , 0.5% BSA	Quantitation Method:	Radioligand Binding
Kd:	0.054 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	3.50 pmole/mg Protein *

■ 252610 Muscarinic M₁

Source:	Human recombinant CHO-K1 cells	Ligand:	0.80 nM [³ H] N-Methylscopolamine
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 µM Atropine
Incubation Time/Temp:	2 hours @ 25°C	Specific Binding:	95% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 10 mM MgCl ₂ , 1 mM EDTA	Quantitation Method:	Radioligand Binding
Kd:	0.26 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	2.0 pmole/mg Protein *

* Historical Values

Methods

■ 252710 Muscarinic M₂

Source:	Human recombinant CHO-K1 cells	Ligand:	0.80 nM [³ H] N-Methylscopolamine
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 μM Atropine
Incubation Time/Temp:	2 hours @ 25°C	Specific Binding:	95% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 10 mM MgCl ₂ , 1 mM EDTA	Quantitation Method:	Radioligand Binding
Kd:	0.58 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	5.10 pmole/mg Protein *

■ 252810 Muscarinic M₃

Source:	Human recombinant CHO-K1 cells	Ligand:	0.80 nM [³ H] N-Methylscopolamine
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 μM Atropine
Incubation Time/Temp:	2 hours @ 25°C	Specific Binding:	95% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 10 mM MgCl ₂ , 1 mM EDTA	Quantitation Method:	Radioligand Binding
Kd:	0.75 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	5.40 pmole/mg Protein *

■ 257010 Neuropeptide Y Y₁

Source:	Human SK-N-MC cells	Ligand:	0.015 nM [¹²⁵ I] Peptide YY
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 μM Neuropeptide Y (human, rat)
Incubation Time/Temp:	60 minutes @ 37°C	Specific Binding:	80% *
Incubation Buffer:	25 mM HEPES, pH 7.4, 1 mM MgCl ₂ , 2.5 mM CaCl ₂ , 0.1% BSA, 0.01% Bacitracin	Quantitation Method:	Radioligand Binding
Kd:	0.24 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.58 pmole/mg protein *

■ 257110 Neuropeptide Y Y₂

Source:	Human KAN-TS cells	Ligand:	10 pM [¹²⁵ I] Peptide YY
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 μM Neuropeptide Y (13-36) (porcine)
Incubation Time/Temp:	2 hours @ 37°C	Specific Binding:	90% *
Incubation Buffer:	25 mM HEPES, pH 7.4, 2.5 mM CaCl ₂ , 1 mM MgCl ₂ , 0.1% Bacitracin	Quantitation Method:	Radioligand Binding
Kd:	0.012 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.5 pmole/mg Protein *

* Historical Values

Methods

■ 258590 Nicotinic Acetylcholine

Source:	Human IMR-32 cells	Ligand:	0.10 nM [¹²⁵ I] Epibatidine
Vehicle:	1% DMSO	Non-Specific Ligand:	300 µM (-)-Nicotine
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	97% *
Incubation Buffer:	20 mM HEPES, pH 7.5, 150 mM NaCl, 1.5 mM KCl, 2 mM CaCl ₂ , 1 mM MgSO ₄ .	Quantitation Method:	Radioligand Binding
Kd:	0.22 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.46 pmole/mg Protein *

■ 258700 Nicotinic Acetylcholine α1, Bungarotoxin

Source:	Human RD cells	Ligand:	0.60 nM [¹²⁵ I] α-Bungarotoxin
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 µM α-Bungarotoxin
Incubation Time/Temp:	2 hours @ 25°C	Specific Binding:	85% *
Incubation Buffer:	150 mM NaCl, 4 mM KCl, 2.3 mM CaCl ₂	Quantitation Method:	Radioligand Binding
Kd:	1.10 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	1.0 pmole/mg Protein *

■ 260130 Opiate δ₁ (OP1, DOP)

Source:	Human recombinant HEK-293 cells	Ligand:	1.30 nM [³ H] Naltrindole
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 µM Naltrindole
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	95% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 1 mM EDTA, 10 mM MgCl ₂	Quantitation Method:	Radioligand Binding
Kd:	0.27 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	7.60 pmole/mg Protein *

■ 260210 Opiate κ(OP2, KOP)

Source:	Human recombinant HEK-293 cells	Ligand:	0.60 nM [³ H] Diprenorphine
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Naloxone
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4	Quantitation Method:	Radioligand Binding
Kd:	0.40 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	1.10 pmole/mg Protein *

* Historical Values

Methods

■ 260410 Opiate μ (OP3, MOP)

Source:	Human recombinant CHO-K1 cells	Ligand:	0.60 nM [3 H] Diprenorphine
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 μ M Naloxone
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4	Quantitation Method:	Radioligand Binding
Kd:	0.41 nM *	Significance Criteria:	\geq 50% of max stimulation or inhibition
		Bmax:	3.80 pmole/mg Protein *

■ 264500 Phorbol Ester

Source:	ICR Mouse brain	Ligand:	3.0 nM [3 H] PDBu
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 μ M PDBu
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	80% *
Incubation Buffer:	20 mM Tris-HCl, pH 7.4, 5 mM CaCl ₂	Quantitation Method:	Radioligand Binding
Kd:	8.70 nM *	Significance Criteria:	\geq 50% of max stimulation or inhibition
		Bmax:	26.0 pmole/mg Protein *

■ 265010 Platelet Activating Factor (PAF)

Source:	Human platelets	Ligand:	0.12 nM [3 H] PAF
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 μ M PAF
Incubation Time/Temp:	3 hours @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 100 mM KCl, 5 mM EDTA, 5 mM MgCl ₂ , 0.25% BSA	Quantitation Method:	Radioligand Binding
Kd:	0.13 nM *	Significance Criteria:	\geq 50% of max stimulation or inhibition
		Bmax:	120 R/cell *

■ 265600 Potassium Channel [K_{ATP}]

Source:	Hamster pancreatic HIT-T15 beta cells	Ligand:	5.0 nM [3 H] Glyburide
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 μ M Glyburide
Incubation Time/Temp:	2 hours @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM MOPS, pH 7.4, 0.1 mM CaCl ₂	Quantitation Method:	Radioligand Binding
Kd:	0.64 nM *	Significance Criteria:	\geq 50% of max stimulation or inhibition
		Bmax:	1.0 pmole/mg Protein *

* Historical Values

Methods

■ 265900 Potassium Channel hERG

Source:	Human recombinant HEK-293 cells	Ligand:	1.50 nM [³ H] Astemizole
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Astemizole
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	90% *
Incubation Buffer:	10 mM HEPES, pH 7.4, 0.1% BSA, 5 mM KCl, 0.8 mM MgCl ₂ , 130 mM NaCl, 1 mM EGTA, 10 mM Glucose	Quantitation Method:	Radioligand Binding
Kd:	6.80 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	6.30 pmole/mg Protein *

■ 268420 Prostanoid EP₄

Source:	Human recombinant Chem-1 cells	Ligand:	1.0 nM [³ H] Prostaglandin E ₂ (PGE ₂)
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Prostaglandin E ₂ (PGE ₂)
Incubation Time/Temp:	2 hours @ 25°C	Specific Binding:	90% *
Incubation Buffer:	10 mM MES, pH 6.0, 1 mM EDTA, 10 mM MgCl ₂	Quantitation Method:	Radioligand Binding
Kd:	0.69 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	4.30 pmole/mg Protein *

■ 268700 Purinergic P_{2X}

Source:	New Zealand derived albino Rabbit urinary bladder	Ligand:	8.0 nM [³ H] α, β-Methylene-ATP
Vehicle:	1% DMSO	Non-Specific Ligand:	100 µM β, γ-Methylene ATP
Incubation Time/Temp:	30 minutes @ 25°C	Specific Binding:	80% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4	Quantitation Method:	Radioligand Binding
Kd1:	2.20 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
Kd2:	2.20 µM *	Bmax1:	2.0 pmole/mg Protein *
		Bmax2:	790 pmole/mg Protein *

■ 268810 Purinergic P_{2Y}

Source:	Wistar Rat brain	Ligand:	0.10 nM [³⁵ S] ATP-αS
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM ADP-βS
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	87% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4	Quantitation Method:	Radioligand Binding
Kd:	0.015 µM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	16.0 pmole/mg Protein *

* Historical Values

Methods

■ 270000 Rolipram

Source:	Wistar Rat brain	Ligand:	1.80 nM [³ H] Rolipram
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Rolipram
Incubation Time/Temp:	60 minutes @ 4°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4	Quantitation Method:	Radioligand Binding
Kd:	1.0 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.31 pmole/mg Protein *

■ 271110 Serotonin (5-Hydroxytryptamine) 5-HT_{1A}

Source:	Human recombinant CHO-K1 cells	Ligand:	1.50 nM [³ H] 8-OH-DPAT
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Metergoline
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	75% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 0.1% Ascorbic Acid, 0.5 mM EDTA, 10 mM MgSO ₄	Quantitation Method:	Radioligand Binding
Kd:	2.0 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	1.30 pmole/mg Protein *

■ 271700 Serotonin (5-Hydroxytryptamine) 5-HT_{2B}

Source:	Human recombinant CHO-K1 cells	Ligand:	1.20 nM [³ H] Lysergic acid diethylamide (LSD)
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Serotonin (5-HT)
Incubation Time/Temp:	60 minutes @ 37°C	Specific Binding:	80% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 4 mM CaCl ₂ , 0.1% Ascorbic Acid	Quantitation Method:	Radioligand Binding
Kd:	2.10 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	1.10 pmole/mg Protein *

■ 271910 Serotonin (5-Hydroxytryptamine) 5-HT₃

Source:	Human recombinant HEK-293 cells	Ligand:	0.69 nM [³ H] GR-65630
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM MDL 72222
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 1 mM EDTA, 5 mM MgCl ₂	Quantitation Method:	Radioligand Binding
Kd:	0.20 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	11.0 pmole/mg Protein *

* Historical Values

Methods

■ 278110 Sigma σ₁

Source:	Human Jurkat cells	Ligand:	8.0 nM [³ H] Haloperidol
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 μM Haloperidol
Incubation Time/Temp:	4 hours @ 25°C	Specific Binding:	80% *
Incubation Buffer:	5 mM Potassium Phosphate, pH 7.5	Quantitation Method:	Radioligand Binding
Kd:	5.80 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.71 pmole/mg Protein *

■ 255520 Tachykinin NK₁

Source:	Human recombinant CHO cells	Ligand:	0.80 nM [³ H] Substance P
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 μM L-703,606
Incubation Time/Temp:	90 minutes @ 4°C	Specific Binding:	90% *
Incubation Buffer:	20 mM HEPES, pH 7.4, 1 mM MnCl ₂ , 0.1% BSA	Quantitation Method:	Radioligand Binding
Kd:	2.10 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	1.70 pmole/mg Protein *

■ 285900 Thyroid Hormone

Source:	Wistar Rat liver	Ligand:	0.030 nM [¹²⁵ I] Triiodothyronine
Vehicle:	1% DMSO	Non-Specific Ligand:	1.0 μM Triiodothyronine
Incubation Time/Temp:	18 hours @ 4°C	Specific Binding:	77% *
Incubation Buffer:	20 mM Tris-HCl, pH 7.6, 50 mM NaCl, 10% Glycerol, 2 mM EDTA, 5 mM DTT	Quantitation Method:	Radioligand Binding
Kd:	0.034 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.16 pmole/mg Protein *

■ 220320 Transporter, Dopamine (DAT)

Source:	Human recombinant CHO-K1 cells	Ligand:	0.15 nM [¹²⁵ I] RTI-55
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 μM Nomifensine
Incubation Time/Temp:	3 hours @ 4°C	Specific Binding:	90% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 100 mM NaCl, 1 μM Leupeptin, 10 μM PMSF	Quantitation Method:	Radioligand Binding
Kd:	0.58 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	0.047 pmole/mg Protein *

* Historical Values

Methods

■ 226400 Transporter, GABA

Source:	Wistar Rat cerebral cortex	Ligand:	6.0 nM [³ H] GABA
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM NO-711
Incubation Time/Temp:	20 minutes @ 25°C	Specific Binding:	80% *
Incubation Buffer:	10 mM HEPES, pH 7.5, 120 mM NaCl, 4 mM Ca(CH ₃ COO) ₂ , 10 µM Isoguvacine, 10 µM S(-)-Baclofen	Quantitation Method:	Radioligand Binding
Kd:	0.30 µM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	60.0 pmole/mg Protein *

■ 204410 Transporter, Norepinephrine (NET)

Source:	Human recombinant MDCK cells	Ligand:	0.20 nM [¹²⁵ I] RTI-55
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Desipramine
Incubation Time/Temp:	3 hours @ 4°C	Specific Binding:	75% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 100 mM NaCl, 1 µM Leupeptin, 10 µM PMSF	Quantitation Method:	Radioligand Binding
Kd:	0.024 µM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	2.50 pmole/mg Protein *

■ 274030 Transporter, Serotonin (5-Hydroxytryptamine) (SERT)

Source:	Human recombinant HEK-293 cells	Ligand:	0.40 nM [³ H] Paroxetine
Vehicle:	1% DMSO	Non-Specific Ligand:	10.0 µM Imipramine
Incubation Time/Temp:	60 minutes @ 25°C	Specific Binding:	95% *
Incubation Buffer:	50 mM Tris-HCl, pH 7.4, 120 mM NaCl, 5 mM KCl	Quantitation Method:	Radioligand Binding
Kd:	0.078 nM *	Significance Criteria:	≥50% of max stimulation or inhibition
		Bmax:	4.40 pmole/mg Protein *

* Historical Values

Reference Compounds

Cat #	Assay Name	Reference Compound	Historical			Concurrent	
			IC ₅₀ *	K _i	n _H	Batch *	IC ₅₀ *
200510	Adenosine A ₁	R(-)-PIA	0.83 μM	0.49 μM	0.90	317535	0.36 μM
200610	Adenosine A _{2A}	CGS-21680	0.13 μM	0.079 μM	1.0	317540	0.11 μM
200720	Adenosine A ₃	IB-MECA	0.78 nM	0.72 nM	0.80	317577	0.91 nM
203100	Adrenergic α _{1A}	Prazosin	0.69 nM	0.28 nM	0.90	317536	0.25 nM
203200	Adrenergic α _{1B}	Prazosin	0.27 nM	0.15 nM	1.0	317547	0.16 nM
203400	Adrenergic α _{1D}	Prazosin	0.88 nM	0.43 nM	0.70	317537	0.68 nM
203620	Adrenergic α _{2A}	Yohimbine	8.40 nM	3.10 nM	0.90	317534	4.48 nM
204010	Adrenergic β ₁	S(-)-Propranolol	2.50 nM	1.40 nM	0.80	317532	0.86 nM
204110	Adrenergic β ₂	S(-)-Propranolol	0.78 nM	0.54 nM	1.20	317533	0.28 nM
285010	Androgen (Testosterone) AR	Testosterone	6.50 nM	4.30 nM	1.0	317485	3.27 nM
212510	Bradykinin B ₁	(Des-Arg ¹⁰)-Kallidin	0.87 nM	0.22 nM	1.10	317578	0.42 nM
212620	Bradykinin B ₂	Bradykinin	1.80 nM	1.10 nM	1.0	317579	1.17 nM
214510	Calcium Channel L-Type, Benzothiazepine	Diltiazem	0.036 μM	0.032 μM	0.90	317482	0.021 μM
214600	Calcium Channel L-Type, Dihydropyridine	Nitrendipine	0.72 nM	0.46 nM	0.90	317531	0.26 nM
216000	Calcium Channel N-Type	ω-Conotoxin GVIA	0.034 nM	0.028 nM	1.60	317580	0.026 nM
217030	Cannabinoid CB ₁	R(+)-WIN-55,212-2	0.20 μM	0.15 μM	0.70	317529	0.26 μM
219500	Dopamine D ₁	R(+)-SCH-23390	1.40 nM	0.70 nM	0.90	317541	1.56 nM
219700	Dopamine D _{2S}	Spiperone	0.25 nM	0.089 nM	1.0	317543	0.086 nM
219800	Dopamine D ₃	Spiperone	0.36 nM	0.12 nM	0.90	317544	0.21 nM
219900	Dopamine D _{4.2}	Spiperone	0.50 nM	0.20 nM	0.90	317552	0.30 nM
224010	Endothelin ET _A	Endothelin-1	0.23 nM	0.14 nM	1.10	317641	0.17 nM
224110	Endothelin ET _B	Endothelin-1	0.13 nM	0.060 nM	0.90	317642	0.054 nM
225510	Epidermal Growth Factor (EGF)	EGF (human)	1.60 nM	1.10 nM	1.10	317582	0.97 nM
226010	Estrogen ER _α	Diethylstilbestrol	0.77 nM	0.22 nM	1.0	317583	0.81 nM
226600	GABA _A , Flunitrazepam, Central	Diazepam	0.016 μM	0.013 μM	0.80	317545	0.027 μM
226500	GABA _A , Muscimol, Central	GABA	0.032 μM	0.026 μM	0.90	317556	0.025 μM
228610	GABA _{B1A}	CGP-54626	6.40 nM	2.90 nM	1.0	317584	4.56 nM
232030	Glucocorticoid	Dexamethasone	3.80 nM	1.80 nM	0.90	317476	2.98 nM
232700	Glutamate, Kainate	L-Glutamic acid	0.24 μM	0.17 μM	0.80	317586	0.18 μM
232810	Glutamate, NMDA, Agonism	L-Glutamic acid	0.41 μM	0.37 μM	0.90	317587	0.18 μM
232910	Glutamate, NMDA, Glycine	MDL 105,519	0.022 μM	0.021 μM	0.60	317588	9.38 nM
233000	Glutamate, NMDA, Phencyclidine	Dizocilpine ((+)-MK-801)	5.10 nM	3.40 nM	0.70	317557	4.71 nM
239610	Histamine H ₁	Pyrilamine	3.30 nM	1.60 nM	1.0	317546	2.50 nM
239710	Histamine H ₂	Tiotidine	0.022 μM	0.018 μM	1.10	317514	0.020 μM
239820	Histamine H ₃	R(-)-α-Methylhistamine (RAMH)	2.30 nM	1.10 nM	1.10	317553	1.74 nM
241000	Imidazoline I ₂ , Central	Idazoxan	0.012 μM	8.0 nM	1.0	317539	7.50 nM
243520	Interleukin IL-1	IL-1β	0.19 nM	0.14 nM	1.30	317484	0.22 nM

* Batch: Represents compounds tested concurrently in the same assay(s).

Broad Institute

Study #: AB13141, Quote #: 30266-1, Compound Code: BRD-K83634925-001-01-1 (1162315)

Reference Compounds

Cat #	Assay Name	Reference Compound	Historical			Concurrent	
			IC ₅₀ *	K _i	n _H	Batch *	IC ₅₀ *
250460	Leukotriene, Cysteinyl CysLT ₁	LTD ₄	0.70 nM	0.29 nM	1.0	317589	1.29 nM
251600	Melatonin MT ₁	Melatonin	0.21 nM	0.11 nM	0.70	317513	0.45 nM
252610	Muscarinic M ₁	4-DAMP	4.50 nM	1.10 nM	1.0	317554	3.64 nM
252710	Muscarinic M ₂	4-DAMP	0.055 μM	0.023 μM	1.0	317527	0.024 μM
252810	Muscarinic M ₃	4-DAMP	5.10 nM	2.50 nM	1.10	317528	4.70 nM
257010	Neuropeptide Y Y ₁	Neuropeptide Y (human, rat)	0.22 nM	0.21 nM	1.10	317468	0.44 nM
257110	Neuropeptide Y Y ₂	Neuropeptide Y (13-36) (porcine)	0.21 nM	0.12 nM	0.90	317592	0.35 nM
258590	Nicotinic Acetylcholine	Epibatidine	0.076 nM	0.052 nM	0.90	317521	0.12 nM
258700	Nicotinic Acetylcholine α ₁ , Bungarotoxin	α-Bungarotoxin	1.10 nM	0.72 nM	1.10	317522	1.30 nM
260130	Opiate δ ₁ (OP1, DOP)	Naltrindole	0.91 nM	0.16 nM	1.0	317525	1.23 nM
260210	Opiate κ(OP2, KOP)	U-69593	0.016 μM	6.40 nM	0.5	317524	0.013 μM
260410	Opiate μ(OP3, MOP)	DAMGO	0.020 μM	8.10 nM	0.60	317526	0.019 μM
264500	Phorbol Ester	PMA	0.79 nM	0.59 nM	1.0	317558	0.69 nM
265010	Platelet Activating Factor (PAF)	PAF	0.28 nM	0.15 nM	0.90	317496	0.27 nM
265600	Potassium Channel [K _{ATP}]	Glyburide	5.70 nM	0.65 nM	0.80	317559	2.80 nM
265900	Potassium Channel hERG	Astemizole	2.60 nM	2.10 nM	1.10	317560	6.16 nM
268420	Prostanoid EP ₄	Prostaglandin E ₂ (PGE ₂)	1.10 nM	0.45 nM	0.90	317561	0.61 nM
268700	Purinergic P _{2X}	α, β-Methylene ATP	0.082 μM	0.018 μM	1.10	317590	0.029 μM
268810	Purinergic P _{2Y}	ATP	0.018 μM	0.018 μM	0.90	317591	0.038 μM
270000	Rolipram	Rolipram	5.70 nM	2.10 nM	1.0	317562	2.66 nM
271110	Serotonin (5-Hydroxytryptamine) 5-HT _{1A}	Metergoline	4.10 nM	2.30 nM	0.90	317510	2.45 nM
271700	Serotonin (5-Hydroxytryptamine) 5-HT _{2B}	Ketanserin	0.29 μM	0.18 μM	0.60	317551	0.53 μM
271910	Serotonin (5-Hydroxytryptamine) 5-HT ₃	MDL 72222	0.011 μM	2.50 nM	0.80	317593	0.018 μM
278110	Sigma σ ₁	Haloperidol	0.021 μM	8.80 nM	0.90	317523	8.84 nM
255520	Tachykinin NK ₁	L-703,606	3.60 nM	2.60 nM	1.0	317474	1.56 nM
285900	Thyroid Hormone	Triiodothyronine	0.034 nM	0.018 nM	1.0	317488	0.040 nM
220320	Transporter, Dopamine (DAT)	GBR-12909	1.70 nM	1.30 nM	0.90	317469	0.59 nM
226400	Transporter, GABA	NO-711	0.20 μM	0.20 μM	1.10	317538	0.21 μM
204410	Transporter, Norepinephrine (NET)	Desipramine	0.93 nM	0.92 nM	0.60	317520	0.44 nM
274030	Transporter, Serotonin (5-Hydroxytryptamine) (SERT)	Fluoxetine	8.60 nM	1.40 nM	0.90	317549	0.013 μM

* Batch: Represents compounds tested concurrently in the same assay(s).

Appendix Notes

The assay # 232020 Glucocorticoid (Ricerca Taiwan Ltd.) was replaced with # 232030 Glucocorticoid (Invitrogen #PV4690). The used material for membrane source was changed from in house to commercial supplier to simplify work process.

Assay #279510 sodium channel site 2 was removed due to backorder of radioligand from the supplier.