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General Methods

Unless specifically stated otherwise, all the manipulations were carried in air in the fume hood in ambient environment. Oxygen- and moisture-sensitive manipulations were carried out under an inert atmosphere using either standard Schlenk techniques or a glove box. Chromatography for air-sensitive compounds was performed under an inert atmosphere using dry, degassed solvents and silica gel (240-300 mesh) which had been heated under vacuum in a 180 °C oil bath for 12 hours. Compound Febinac (4-biphenylacetic acid, CAS [5728-52-9]), **S6** (biphenyl-4-carboxylic acid, CAS [92-92-2]) and amine **S5** were purchased from Sigma-Aldrich. Reagents and catalyst were purchased from Sigma-Aldrich, TCI America, and Strem Chemicals and used without further purification. Anhydrous toluene, methylene chloride, pentane, THF, and ethyl ether were taken from a solvent purification system and used without further purification; the remaining solvents were dried over calcium hydride, distilled, and freeze-pump-thaw degassed before use. Additionally, all materials used under argon were either purged with argon (in the case of solids) or stirred over calcium hydride, distilled, then freeze-pump-thaw degassed (in the case of liquids). DMA solution for silica gel chromatography is a mixture of 80% CH₂Cl₂, 18% MeOH and 2% concentrated ammonium hydroxide.

¹H NMR, ¹³C NMR and ¹¹B NMR spectra were recorded on a Varian Unity/Inova 300, Varian Unity/Inova 400, Varian Unity/Inova 500 or Varian Unity/Inova 600 spectrometer at ambient temperature. ¹¹B NMR were externally referenced to $BF_3 \cdot Et_2O$ ($\delta = 0$). IR spectra were recorded on a Nicolet Magna 550 FT-IR.

High-resolution mass spectrometry data were obtained at the following facilities: 1) Boston College mass spectrometry center on a JEOL AccuTOF instrument (JEOL USA, Peabody, MA), equipped with a Direct Analysis in Real Time (DART) ion source (IonSense, Inc., Danvers, MA) in positive ion mode. 2) Small Molecule Mass Spectrometry, FAS-Center for Systems Biology at Harvard University.

Scheme S1. Synthesis of BN-biphenyl carboxylic acid S4



Reagents and conditions: (a) 2-(trimethylsilyl)ethanol, DMAP, DCC, CH_2Cl_2 , 98%; (b) $Pd(PPh_3)_4$, $(Me_3Sn)_2$, toluene, 100 °C, N₂, 82%; (c) 1-(tert-butyldimethylsilyl)-2-chloro-1,2-dihydro-1,2-azaborinine, $[Rh(C_2H_4)_2Cl]_2$, BIPHEP, toluene, 100 °C, N₂, 89%; (d) TBAF (1.0 M in THF), THF, -25 °C to RT, N₂, 84%.

2-(Trimethylsilyl)ethyl-4-bromobenzoate S1



To a solution of 4-bromobenzoic acid (5.10 g, 24.9 mmol) in CH_2Cl_2 (150 mL) was added 4dimethylaminopyridine (180 mg, 1.47 mmol), *N,N'*-dicyclohexylcarbodiimide (5.24 g, 25.4 mmol) and 2-(trimethylsilyl)ethanol (6.20 mL, 43.3 mmol) under ice-water bath. The reaction was stirred at room temperature for 14 hours. At the conclusion of the reaction, the mixture was passed through a glass frit, and the resulting filtrate was concentrated under reduced pressure. This crude material was purified by silica gel chromatography (5% EtOAc/hexane) (v/v) to provide the desired compound as colorless oil (7.52 g, yield 98%).

¹H NMR (300 MHz, CD₂Cl₂) δ 7.92 (d, J = 7.8 Hz, 2H), 7.62 (d, J = 7.8 Hz, 2H), 4.44 (t, J = 8.1 Hz, 2H), 1.16 (t, J = 4.8 Hz, 2H), 0.13 (s, 9H); ¹³C NMR (126 MHz, CD₂Cl₂) δ 165.7, 131.6, 131.0, 129.8, 127.6, 63.4, 17.3, -1.77; FTIR (thin film): \tilde{v} = 2953, 1708, 1588, 1483, 1398, 1274, 1247, 1174, 1103, 1064, 1009, 935, 829, 754, 681, 607, 459; HRMS (ESI) calcd for C₁₂H₁₇BrO₂SiNa (M+Na)⁺ 323.0073, found 323.0075.

2-(Trimethylsilyl)ethyl-4-(trimethylstannyl)benzoate S2



In the glovebox, to a 20 mL a pressure vessel, compound **S1** (100 mg, 0.330 mmol), tetrakis(triphenylphosphine)palladium (20 mg, 0.017 mmol), hexamethylditin (140 mg, 0.427 mmol) and toluene (4.0 mL) was added. The solution was heated to 100 °C for 15 hours, then allowed to cooled to room temperature. Volatiles were removed under reduced pressure, and the crude material was purified by silica gel chromatography (100% pentane then 10% EtOAc/hexane) (v/v) to afford the desired compound as a colorless oil (105 mg, yield 82 %).

¹H NMR (300 MHz, CD₂Cl₂) δ 7.99 (d, J = 4.5 Hz, 2H), 7.64 (d, J = 4.8 Hz, 2H), 4.44 (t, J = 5.1 Hz, 2H), 1.17 (t, J = 5.1 Hz, 2H), 0.36 (s, 9H), 0.13 (s, 9H); ¹³C NMR (126 MHz, CD₂Cl₂) δ 166.8, 149.4, 135.8, 130.4, 128.2, 63.0, 17.2, -1.8, -9.9; FTIR (thin film): $\tilde{\upsilon}$ = 1715, 1273, 1110, 1064, 858, 836, 752, 694, 528; HRMS (ESI-TOF) calcd for C₁₅H₂₆O₂SiSnNa (M+Na)⁺ 409.0618, found 409.0618.

Compound S3



In the glovebox, chlorobis(ethylene)rhodium dimer (52 mg, 0.13 mmol, 0.05 eq.), BIPHEP (140 mg, 0.26 mmol, 0.1 eq.) and toluene (5.0 mL) were added to a 20 mL vial. The solution was stirred for 30 min, then it was transferred to a pressure vessel containing 1-(tert-butyldimethylsilyl)-2-chloro-1,2-dihydro-1,2-azaborinine [CAS 1138164-75-6] (650 mg, 2.86 mmol, 1.1 eq.), compound **S2** (1.00 g, 2.59 mmol, 1.0 eq) and toluene (15 mL). The pressure vessel was sealed, and the reaction mixture was heated at 100 °C for 15 hours. At the conclusion of the reaction, the reaction mixture was allowed to cool to room temperature. Volatiles were removed under reduced pressure, and the resulting crude material was purified in the glovebox

by silica gel chromatography (10% diethyl ether/pentane) to provide the desired product as white solid (951mg, yield 89%).

¹H NMR (300 MHz, CD₂Cl₂) δ 8.00 (d, J = 8.1 Hz, 2H), 7.66-7.71 (m, 1H), 7.51-7.55 (m, 3H), 6.68 (d, J = 10.5 Hz, 1H), 6.54 (m, 1H), 4.49 (t, J = 4.2, Hz, 2H), 1.23 (t, J = 4.5 Hz, 2H), 0.99 (s, 9H), 0.18 (s, 9H), 0.11 (s, 6H); ¹³C NMR (126 MHz, CD₂Cl₂) δ 167.0, 143.2, 138.3, 132.0, 128.9, 127.5, 112.2, 62.9, 26.6, 18.8, 17.4, -1.7, -2.3 (the boron-bound carbono signals are not observed); ¹¹B NMR (96 MHz) δ 41.3; FTIR (thin film): \tilde{v} = 3282, 2931, 1733, 1613, 1542, 1456, 1370, 1233, 1137, 813, 731, 676, 644, 504, 457; HRMS (ESI) calcd for C₂₂H₃₇BNO₂ Si₂ (M+H)⁺ 414.2456, found 414.2468.

Compound S4



In the glovebox, compound **S3** (850 mg, 2.06 mmol) was dissolved into THF (12.0 mL) in a round-bottom flask and cooled in a -25 °C freezer for 30 min. The reaction flask was then taken out of the glovebox, and TBAF (5.0 mL, 1.0 M in THF, 5.0 mmol) was added slowly. The resulting yellow solution was stirred at room temperature for 3 hr. Volatiles were removed under reduced pressure, and the crude material was purified on silica gel chromatography using CH₂Cl₂/MeOH/AcOH 100:4:0.8 (v/v) as the eluent. The resulting off-white solid was recrystallized in CH₂Cl₂/hexane system to afford the desired compound (344 mg, yield 84%).

¹H NMR (300 MHz, acetone- d_6) δ 10.19 (br s, 1H), 7.89-8.07 (m, 4H), 7.81 (dd, J = 6.3, 11.1 Hz, 1H), 7.61 (t, J = 6.6 Hz, 1H), 7.22 (d, J = 11.1 Hz, 1H), 6.47 (t, J = 6.6 Hz, 1H) (COOH peak is not observed); ¹³C NMR (126 MHz, acetone- d_6) δ 167.2, 144.8, 135.2, 132.4, 130.6, 129.0, 111.2 (the boron-bound carbon signals are not observed); ¹¹B NMR (96 MHz) δ 33.6; FTIR (thin film): \tilde{v} = 1733, 1716, 1684, 1635, 1557, 1540, 1521, 1473, 1374, 724, 671, 419; HRMS (DART) calcd for C₁₁H₁₁BNO₂ (M+H)⁺ 200.08828, found 200.08758.

General experimental procedure to form BN-biarylcarboxamides

To the mixture of 2-chloro-4,6-dimethoxyl-1,3,5triazine (CDMT, 32.0 mg, 0.182 mmol) and BN Felbinac (38.0 mg, 0.176 mmol) or compound **S4** (35.0 mg, 0.176 mmol) in anhydrous CH_2Cl_2 (3.0 mL), *N*-methylmorpholine (NMM, 20.0 µL, 0.182 mmol) was added at 0 °C. The resulting clear solution was stirred at room temperature for 25 min. Then, the corresponding amine (1.1 equiv.) was added, and this mixture was stirred for 14 hours at room temperature (if this amine is in salt form, another 1 equiv. NMM is added). At the conclusion of the reaction, CH_2Cl_2 (50 mL) was added to the reaction mixture, and the resulting mixture was washed with saturated NaHCO₃. The organic phase was concentrated under reduced pressure. The resulting crude material was purified by silica gel chromatography to afford corresponding amide products. If necessary, recrystallization in CH_2Cl_2 /hexane or MeOH/CH₂Cl₂/hexane can provide desired amide compounds as an off-white solid.



Amine **S5** [CAS 21103-33-3] was purchased from Aldrich, and the bioactive amide **CC-1** was prepared according to literature procedures.¹ ¹H NMR and mass spectra of **CC-1** match with those reported in the literature.



¹H NMR (600 MHz, CDCl₃) δ 7.84-7.86 (m, 2H), 7.56-7.63 (m, 4H), 7.34-7.45 (m, 3H), 6.93-7.00 (m, 2H), 6.83-6.87 (m, 3H), 3.83 (s, 3H), 3.49-3.51 (m, 2H), 3.08-3.11 (m, 4H), 2.69-2.75 (m, 3H), 2.54-2.58 (m, 3H), 1.67-1.73 (m, 4H); HRMS (DART) calcd for C₂₈H₃₄N₃O₂ (M+H)⁺ 444.26510, found 444.26631.



Silica gel chromatography condition (CH₂Cl₂/MeOH), off-white solid (110 mg, yield 82%).

¹H NMR (500 MHz, CD₂Cl₂) δ 8.48 (br s, 1H), 7.79-7.86 (m, 5H), 7.47 (d, J = 6.5 Hz, 1H), 7.20 (d, J = 6.0 Hz, 1H), 6.86-6.97 (m, 5H), 6.46 (d, J = 6.0 Hz, 1H), 3.82 (s, 3H), 3.43-3.48 (m, 2H), 2.96-3.20 (m, 4H), 2.44-2.78 (m, 6H), 1.67-1.80 (m, br, 4H); ¹³C NMR (126 MHz, CD₂Cl₂) δ167.5, 152.4, 144.8, 141.5, 135.4, 128.4 (br), 126.5, 122.6, 120.9, 118.1, 111.4, 111.2, 57.9, 55.2, 50.2, 39.8, 31.6, 27.4, 24.2, 22.6, 13.9; ¹¹B NMR (160 MHz) δ 36.7; FTIR (thin film): \tilde{v} = 3553, 1699, 1612, 1541, 1499, 1462, 1364, 1310, 1239, 1180, 1117, 1088, 1022, 979, 923, 736, 677, 534; HRMS (DART) calcd for C₂₆H₃₄BN₄O₂ (M+H)⁺ 445.27748, found 445.27853.



Amine **S7** [CAS 1850290-88-8] and the bioactive amide **CC-2** were prepared according to literature procedures.² ¹H NMR and mass spectra signals of **CC-2** match with those reported in the literature.



¹H NMR (500 MHz, Acetone-*d*6) δ 9.07 (s, 1H), 8.51-8.53 (m, 1H), 8.34-8.36 (m, 1H), 7.84-7.91 (m, 3H), 7.72-7.74 (m, 4H), 7.44-7.54 (m, 3H), 3.89-3.97 (m, 4H); HRMS (DART) calcd for C₂₁H₁₈F₃N₂O₃S (M+H)⁺ 435.09902, found 435.09851.



After coupling reaction, the reaction mixture was directly purified by silica gel chromatography (DMA/CH₂Cl₂/hexane). Off-white solid (79 mg, yield 78%).

¹H NMR (500 MHz, DMSO-*d*₆) δ 10.67 (br s, 1H), 9.06 (s, 1H), 8.45-8.50 (m, 2H), 8.23 (d, J = 8.5 Hz, 1H), 7.85-7.90 (m, 2H), 7.63-7.72 (m, 1H), 7.58-7.60 (m, 2H), 7.48 (t, J = 7.5 Hz, 1H), 7.07 (d, J =11.0 Hz, 1H), 6.39 (t, J = 6.0 Hz, 1H), 3.84-3.93 (m, 2H), 3.60-3.78 (m, 2H); ¹³C NMR (126 MHz, DMSO-*d*₆) δ 166.7, 160.5, 147.7, 145.1, 137.3, 136.0, 133.7, 132.6, 127.7, 126.7, 122.7, 111.3, 50.9, 34.2 (boron-bound carbon and the CF₃ signals are not observed) ¹¹B NMR (160 MHz) δ 34.3; FTIR (thin film): \tilde{v} = 2924, 1734, 1642, 1540, 1463, 1361, 1300, 1141, 1097, 1072, 1012, 860, 814, 726, 612, 486; HRMS (DART) calcd for C₁₉H₁₈BF₃N₃O₃S (M+H)⁺ 436.11140, found 436.11278.



The synthesis of amine **S8** [CAS 326827-13-8]^{3a,3b} and the bioactive amide **CC-3**^{3c} was reported in the reference. ¹H NMR and mass spectra signals of **CC-3** match with those reported in the literature.

Compound BN-3 was prepared in a two-step procedure:

Amide coupling reaction: Amine **S8** (120 mg, 0.540 mmol) was dissovled in anhydrous CH₂Cl₂ (8.0 mL), then compound **BN Felbinac** (115mg, 0.54 mmol), 2-chloro-4,6-dimethoxyl-1,3,5-triazine (CDMT, 95 mg, 0.54 mmol), and N-methylmorpholine (NMM, 140 μ L, 1.08 mmol) was added at 0 °C. The resulting solution was stirred at room temperature for 16 hours. At the conclusion of the reaction, CH₂Cl₂ (100 mL) was added to the reaction mixture, and the resulting mixture was washed with saturated NaHCO₃. The organic phase was concentrated under reduced pressure and the resulting crude material was purified by silica gel chromatography (DMA/CH₂Cl₂/hexane) to afford the desired amide **S10** as an yellow oil (¹H NMR purity >90%) which was used directly for the next step.

Boc-deprotection procedure: To the amide **S10** obtained in the previous step, 10% TFA/CH₂Cl₂ (5.0 mL) was added at 0 °C. The mixture was then allowed to stir at room temperature for 2 hours. At the conclusion of the reaction, CH_2Cl_2 (100 mL) was added to the reaction mixture, and the resulting mixture was washed with saturated NaHCO₃. The organic phase was concentrated under reduced pressure. The resulting crude material was purified by silica gel chromatography (hexane/CH₂Cl₂ to DMA/CH₂Cl₂) to afford a solid, which was further recystalized from CH_2Cl_2 /hexane to yield pure product as off-white solid (104 mg, yield 61% for two steps).



¹H NMR (500 MHz, DMSO-*d*₆) δ 12.01 (s, 1H), 10.47 (s, 1H), 7.57-7.63 (m, 3H), 7.33-7.44 (m, 6H), 6.12 (s, 1H), 3.59 (s, 2H), 1.80-1.82 (m, 1H), 0.85-0.88 (m, 2H), 0.60-0.62 (m, 2H); HRMS (DART) calcd for C₂₀H₂₀N₃O (M+H)⁺ 318.16604, found 318.15979.



¹H NMR (500 MHz, DMSO-*d*₆) δ 12.04 (s, 1H), 10.58 (bs, 1H), 10.47 (s, 1H), 7.82 (d, J = 8.0 Hz, 2H), 7.67-7.70 (m, 1H), 7.47 (t, J = 7.0 Hz, 1H), 7.34 (d, J = 8.0 Hz, 2H), 7.06 (d, J = 11.0 Hz, 1H), 6.36 (t, J = 6.5 Hz, 1H), 6.15 (s, 1H), 3.59 (s, 2H), 1.76-1.83 (m, 1H), 0.88-0.90 (m, 2H), 0.62-0.65 (m, 2H); ¹³C NMR (150 MHz, DMSO-*d*₆) δ 168.5, 144.7, 137.1, 135.8, 132.9, 129.5, 129.1, 128.7, 127.7 (br), 126.9, 110.7, 43.0, 8.2, 7.3 (the exocyclic boron-bound carbon signal was not observed); ¹¹B NMR (160 MHz) δ 33.4; FTIR (thin film): $\tilde{\upsilon}$ = 3373, 1655, 1590, 1539, 1484, 994, 802, 731, 676, 597, 459; HRMS (DART) C₁₈H₂₀BN₄O (M+H)⁺ 319.17379, found 319.17413.

Air and water stability study of compound BN-1, BN-2 and BN-3

Air and water stability study procedures were adapted from a previously reported study with minor modifications.⁴

In the fume hood, a J-Young NMR tube was charged with compound **BN-1** (5.0 mg), CDCl₃ (0.5 mL), internal standard toluene (2 μ L), and H₂O (2 μ L). ¹H and ¹¹B NMR were taken at time 0 hr. Then the sample was vigorously shaken and immersed into a pre-heated 50 °C oil bath. ¹H and ¹¹B NMR were taken at indicated time points. ¹H and ¹¹B NMR analysis indicated no decomposition of **BN-1** after 24 hours.

In the fume hood, a J-Young NMR tube was charged with the individual compound **BN-2** or **BN-3** (5.0 mg), DMSO-d₆ (0.5 mL), internal standard toluene (2 μ L), and H₂O (2 μ L). ¹H and ¹¹B NMR were taken at time 0 hr. Then the sample was vigorously shaken and immersed into a pre-heated 50 °C oil bath. ¹H and ¹¹B NMR were taken at indicated time points. ¹H and ¹¹B NMR analysis indicated no decomposition of **BN-2** and **BN-3** after 24 hours.

Assays

Rat Liver Microsome assay

Performed according to standard literature methods (Obach, RS (1999) Drug Metab Dispos; 27:1350-1359). Liver microsomes from male Sprague-Dawley rats were obtained from BDGentest (catalog #452501), Briefly, test compound was dissolved in DMSO at 10 mM and diluted to 10 uM in 50 mM Pi, pH 7.4. 0.1 volume of this was added to a 0.5 mg/ml suspension of rat liver microsomes in 50 mM Pi, pH 7.4. Reaction is initiated by addition of 1 mM NADPH, 2 mM MgCl₂. Samples are withdrawn after 0, 5, 15, and 30 minutes incubation at 37 °C and mixed with 1 vol acetonitrile. Test compound remaining is determined by peak area in LC/MS chromatogram relative to t = 0 peak area. Clearance is calculated applying literature scaling factors to test compound elimination rate constant, which is determined by slope of ln[compound remaining] vs. time.

CYP3A4

Inhibition of CYP3A4 activity was performed according to literature methods (i.e. Kim, et al, (2005) Rapid Commun Mass Spectrom; 19:2650-2658 and Obach et al (2006) J Pharmacol Exp Ther; 316:336-348). Briefly, compounds at serially diluted concentrations between 0.1 and 20 uM (final) were included in incubations with human liver microsomes (0.05 mg/ml, BDGentest) at 37 °C. Midazolam was added at 1 uM as test substrate and reaction initiated by addition of 1 mM NADPH. Reaction was terminated after 10 minutes and amount of 1'-hydroxymidazolam (product formed from midazolam by CYP3A4) determined by LC/MS methods. Inhibition of 1'-hydroxymidazolam formation relative to buffer controls and calculation of IC50 was done by standard data analysis methods.

Rat in vivo PK

Male Sprague-Dawley rats, 9-11 weeks old and jugular vein cannulated, were obtained from Envigo. Test compound was dissolved in a formulation consisting of 15% PEG300, 7.5% Solutol and 7.5% CremophoreEL in water. **BN-3** was dosed to SD rat at 1 mg/kg by intravenous bolus injection (IV) through the jugular cannula. **CC-3** was dosed IV at 0.5 mg/kg. Blood samples were taken at times after dosing (up to 24 h). Similarly, **BN-3** and **CC-3** were dosed orally (PO, *per os*) at 5 mg/kg. Blood samples were taken up to 7 hours after dosing. Plasma was obtained by

centrifugation of the blood samples and analyzed for concentration of **BN-3** or **CC-3** by LC/MS methods. Non-compartmental analysis was performed on the concentration data according to standard methods to obtain area under the concentration-time curve (AUC), clearance (CL), bioavailability (F) and half-life (t $\frac{1}{2}$).

Time (h)	CC-3	BN-3	CC-3	BN-3
	0.5 mg/kg IV	1.0 mg/kg IV	5 mg/kg PO	5 mg/kg PO
	(nM)	(nM)	(nM)	(nM)
0.083	1177.5	2761.7		
0.25	552.1	2007.3	311.0	211.1
0.5	360.0	1066.4	691.6	239.7
1	90.7	514.0	448.6	485.6
2	11.1	69.4	341.8	685.4
4	9.1	30.4	75.8	546.8
7	4.2	16.2	31	117.4
24	4.8	7.6		

Concentration-time data (average of N=2):

hERG binding

Cell membranes were prepared from HEK-293 cells recombinantly expressing the KCNH2 (hERG) gene. Membranes (about 40 μ g protein) were incubated for 60 min at 22 °C with 3 nM [³H]dofetilide in the absence or presence of the test compound in a buffer containing 50 mM Tris-HCl (pH 7.4), 10 mM KCl and 1 mM MgCl₂, in an assay volume of 200 μ l in a 96-well plate. Non-specific binding was determined in the presence of 25 μ M terfenadine. Following the incubation, the samples were filtered rapidly under vacuum through glass fiber filters (GF/B) presoaked with 0.3% polyethyleneimine (PEI) and rinsed several times with ice-cold 50 mM Tris-HCl, 10 mM KCl and 1 mM MgCl₂ using a 96-sample cell harvester (Unifilter, Packard). The filters were then dried and counted for radioactivity in a scintillation counter (TopCount, PerkinElmer) using a scintillation cocktail (Microscint-O, PerkinElmer). Test compounds were dissolved in DMSO. The final concentration of DMSO in the assay was 1%.

PPAR (γ and δ) antagonist

LanthaScreen[™] TR-FRET PR Coactivator Assay Kits were purchased from Life Technologies (e.g. Cat. #PV4685 for PPARdelta and PV4548 for PPARgamma). A terbium-labeled anti-GST

antibody is used to indirectly label a nuclear receptor by binding to its GST tag. When an agonist binds to the receptor, a conformational change takes place allowing a fluorescein-labeled co-activator to bind with high affinity. When the terbium label on the anti-GST antibody is excited at 340 nm, energy is transferred to the fluorescein label on the bound co-activator peptide and detected as emission at 520 nm.

The assays were performed in a final volume of 50 μ L per well in a 384-well solid black plate (PerkinElmer, Cat. #6007279). The components of the PPAR δ assay were added in the following order: 0.25 μ L test compound in 100% DMSO, 50 μ L GST-PPAR δ -LBD, Tb-antiGST antibody and Fluorescein-C33 Peptide (final concentrations of 5 nM, 10 nM, 100 nM respectively). For antagonist activity an additional 0.25 μ L EC₈₀ agonist (GW0742, 10 nM final concentration) was added. The plates were incubated in the dark at room temperature for 2 hours and then read in a PerkinElmer Envision. The signal measured was the fluorescence ratio 520 nm/495 nm.

The components of the PPAR γ assay were added in the following order: 0.25 µL test compounds in 100% DMSO, 50 µL GST-PPAR γ -LBD, Tb-antiGST antibody and Fluorescein-TRAP220/DRIP-2 Peptide (final concentrations of 2.5 nM, 2.5 nM, 125 nM respectively). For antagonist activity an additional 0.25 µL EC₈₀ agonist (rosiglitazone, 50 nM final concentration) was added. The plates were incubated in the dark at room temperature for 2 hours and then read in a PerkinElmer Envision. The signal measured is the fluorescence ratio 520 nm/495 nm.

Dopamine D3 binding

Wheatgerm agglutinin (WGA) scintillation proximity assay (SPA) beads (RPNQ0001) were purchased lyophilized from PerkinElmer (Boston, MA, USA). One vial (500 mg) was reconstituted using 5 mL of distilled water to give a final concentration of 100 mg/mL. Reconstituted SPA beads were stored at 4°C and not frozen. 384-well white, clear bottom plates (Cat. #3706) were purchased from Corning (Lowell, MA, USA). The incubations were performed in a final volume of 50 μ L per well in 384-well white, clear bottom polystyrene plates. The components of the incubation were added to each well in the following order and volumes: 0.25 μ L test compound / NSB / reference in 100 % DMSO; 9.75 μ L distilled water; 20 μ L radioligand; 20 μ L SPA bead / membrane mixture. The final concentration of WGA beads was 4 mg/mL and that of D3 membranes (PerkinElmer Cat. #ES-173) was 10 μ g/mL. The final concentration of the radioligand [³H]-Methylspiperone was 0.4 nM. Total binding was determined by adding DMSO and non-specific binding (NSB) was determined by the addition of (+)Butaclamol, final concentration 10 μ M. The plates were then sealed and allowed to sit at room temperature for 5 hours of incubation time. Plates were counted in a PerkinElmer Microbeta Trilux reader for 90 seconds per well.

CDK2 kinase assay

Kinase reactions was performed in automated fashion for profiling with several kinases in parallel with higher throughput. All assays were performed in 384 well microtiter plates. Each assay plate contained 8-point serial dilutions for 40 test compounds, as well as two 16-point serial dilutions of staurosporine as reference compound, plus 16 high- and 16 low controls. Protocol: 0.1 µl Compound, 9 µl 2x peptide/ATP solution, 9 µl 2x enzyme solution, incubate for 60 min at 30 °C, 70 µl stop/run buffer. Independent of the kinase, all reactions were performed in 50 mM HEPES, pH 7.5, 1mM DTT, 0.02% (v/v) Tween20, 0.02% (w/v) BSA, 0.5-1% (v/v) DMSO, 10 mM beta-glycerophosphate, and 10µM sodium orthovanadate. CDK specific conditions: [CDK2] 8.2nM, [ATP] 25uM, [peptide] 2uM, [ATP KM] 25uM, [Mg] 8mM. Percent-inhibition values were calculated as the following: The relative amount of phosphorylated peptide r, was calculated using the heights of the substrate peak, s, and the product peak, p: r = p/(p+s). Percent inhibition was then determined as : %inhibition = $100 \cdot (1-(r-r_{low control})/(r_{high control} - r_{low control}))$

Equilibrium solubility:

Equilibrium solubility was determined using a miniaturized shake flask approach as described in Zhou et al.⁵ Aliquots of 10 mM DMSO compound solution were dispensed in triplicate in 96-well polypropylene plates. The DMSO was removed using a GeneVac HT4X evaporator for approximately one hour. Media (pH 4.0 buffer, pH 6.8 buffer, FaSSIF: Fasted State Simulated Intestinal Fluid) was added to each well to achieve a target concentration of 1 mM. The plate was sealed and shaken for a minimum of 16 hours, then centrifuged for phase separation. An aliquot of supernatant was transferred to a new plate, where it was further diluted for subsequent analysis. Quantification of solubility was performed using RapidFire/MS/MS and a four-point

calibration curve. Experimental variability was determined from approximately 300 duplicate measurements from different days and experimentalists, with a log standard deviation of 0.25.

HT logD:

Dried DMSO stock samples were incubated with buffer and 1-octanol and shaken for at least 4 hours. Sample preparation and phase separation were automated using centrifugation and liquid handling workstations. Both 1-octanol and buffer phases were quantified in triplicate and logD was derived from the ratio of peak area responses against an internal standard in a tandem mass spectrometer. To obtain logP, the buffer pH was selected where the compound was unionized, based on calculated pKa values.

PAMPA Assay:⁶

Permeation experiments were carried out in 96-well microtiter filter plates obtained from Millipore AG (Volketswil, Switzerland). Filter (isopore, polycarbonate) specifications were as follows: 3 um pore size, 9-10 um thickness, and 5-20% porosity. Each well of the filter plate was impregnated with 15 *u*L of 5% hexadecane dissolved in hexane (i.e., total amount of hexadecane: 0.75 *u*L) for at least 10 min to allow for complete evaporation of the hexane. Subsequently, the donor compartments were hydrated with 300 uL of 50 ug/ mL test compound in buffer, containing 5% DMSO and 100 mM KCl, and connected to a homemade Teflon acceptor plate which had been prefilled with buffer containing 5% DMSO. The resulting sandwich construct was incubated at room temperature under constant light shaking (50-100 rpm). After 5 h, the sandwich was disassembled and the solution in the acceptor was transferred to a disposable UVtransparent plate (Corning Costar, Corning, NY). UV absorption was measured with a SPECTRAmax190 microplate spectrophotometer (Molecular Devices Corp., Sunnyvale, CA) at absorption wavelengths between 260 and 290 nm. To ensure that the donor/acceptor fluxes were not due to porous or unstable hexadecane layers, the stability of the hexadecane membranes was tested at the end of the incubation period by electrical resistance measurements. Wells with barriers which displayed electrical resistance lower than 5 k Ω were discarded. Electrical resistance measurements were performed using a Keithley 6517A electrometer (Keithley Instruments S.A., Dubendorf, Switzerland) with Ag/AgCl electrodes from World Precision Instruments (Berlin, Germany). Ionization constants were measured by potentiometric titration in

0.15 M KCl at 25.0 °C using a GlpKa instrument (Sirius Analytical Instruments, Forest Row, U.K.). Partition and distribution coefficients were measured using the pH metric technique using a PCA101 automatic titrator (Sirius Analytical Instruments, Forest Row, U.K.).

At the end of the incubation period the sandwich was carefully disassembled, the acceptor plate measured with the UV microtiter plate spectrophotometer, and the donor plate submitted to current measurements to assess the integrity of the hexadecane membranes. The apparent permeability value Pa is determined from the ratio r of the absorbance of compound found in the acceptor chamber divided by the theoretical equilibrium absorbance (determined independently):

$$P_{\rm a} = -\frac{V_{\rm D}}{(V_{\rm D} + V_{\rm R})At} \cdot \ln(1 - t)$$
(1)

In this equation, $V_{\rm R}$ is the volume of the acceptor compartment (0.4 cm³), $V_{\rm D}$ is the donor volume (0.3 cm³), A is the accessible filter area (total filter area, 0.24 cm², multiplied by a porosity ratio of 20%), and t is the incubation time. Equation 1 is obtained from the differential equation (2) with cD(t) being the compound concentration in the donor compartment and cR(t) being the concentration in the acceptor compartment. In absence of membrane retention Pa is identical to Pe, the effective membrane permeability. When membrane retention occurs Pa can be converted to Pe using mass balance equations.

$$\frac{\mathrm{d}c_{\mathrm{R}}}{\mathrm{d}t} = \frac{P_{\mathrm{a}}A}{V_{\mathrm{R}}}(c_{\mathrm{D}} - c_{\mathrm{R}}) \tag{2}$$

Molecular modeling:

Docking poses were generated in a high resolution crystal structure of CDK2/cyclin A (PDB entry 1VYW) using Glide (version 6.6) from Schrödinger. The crystal structure was first prepared using the standard protein preparation wizard within the Maestro interface. Grid files were generated for docking after deleting the crystallographic water molecules. 3D-geometries of the ligands **BN-3** and **CC-3** were generated using ligprep. Compounds were docked in the pregenerated grid files in extra precision mode. Three docking poses were saved for each of the compounds and the docking results were analyzed visually. The robustness of the docking

protocol was confirmed by self-docking and cross-docking studies of the known CDK2 ligands from crystal structures. Geometry of the C-B bond in BN-3 was verified by comparing with the small molecule x-ray of 1,2-dihydro-2-phenyl-1,2-azaborine (CSD code WAJTEH).⁷

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		dmf 12361	dpwr 40	decwave w	dm nny	dof 0	dn H1	DECOUPLER	pw 11.750	tpwr 55	tof 4817.4	sfrq 160.382	tn B11	TRANSMITTER	ct 372	nt 4000	d1 0.100	bs 4	fb 17000	np 6410	at 0.100	sw 32051.3	ACQUISITION	71-28.fid	rsys/data/Peng/5-1~	file /home/Liu/vnm~	solvent cdcl3	date May 20 2015	SAMPLE	exp1 s2pu]	peng-103 stability 1
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