Supporting Information

Discovery of a Pyrrolopyrimidine (JH-II-127), a Highly Potent, Selective and Brain Penetrant LRRK2 Inhibitor

John M. Hatcher^{a,b,d}, Jinwei Zhang ^{c,d}, Hwan Geun Choi^a, Genta Ito^c, Dario R. Alessi^c,* and Nathanael S. Gray^{a,b},*

- aDepartment of Cancer Biology, Dana-Farber Cancer Institute, Boston, MA 02115, USA.
- *bDepartment of Biological Chemistry & Molecular Pharmacology, Harvard Medical School, 250 Longwood Ave, SGM 628, Boston, MA 02115, USA.*
- *CMRC Protein Phosphorylation Unit, College of Life Sciences, University of Dundee, Dow Street, Dundee DD1 5EH, Scotland, United Kingdom.*dThese authors contributed equally to this work.
- * To whom correspondence should be addressed. For N.S.G.: phone, 1-617-582-8590; fax, 1-617-582-8615; E-mail, Nathanael Gray@dfci.harvard.edu, for D.R.A.: phone, 44-1382-385602; fax, 44-1382-223778; E-mail, d.r.alessi@dundee.ac.uk

Experimental Procedures and Compounds Characterization

General

All reactions were monitored by thin layer chromatography (TLC) with 0.25 mm E. Merck pre-coated silica gel plates (60 F₂₅₄) and Waters LCMS system (Waters 2489 UV/Visible Detector, Waters 3100 Mass, Waters 515 HPLC pump, Waters 2545 Binary Gradient Module, Waters Reagent Manager, Waters 2767 Sample Manager) using SunFireTM C18 column (4.6 x 50 mm, 5 μ m particle size): solvent gradient = 100% A at 0 min, 1% A at 5 min; solvent A = 0.035% TFA in Water; solvent B = 0.035% TFA in MeOH; flow rate : 2.5 mL/min. Purification of reaction products was carried out by flash chromatography using CombiFlash®Rf with Teledyne Isco Redi*Sep*®Rf High Performance Gold or Silicycle Silia*Sep*TM High Performance columns (4 g, 12 g, 24 g, 40 g or 80 g). The purity of all

compounds was over 95% and was analyzed with Waters LCMS system. ¹H NMR and ¹³C NMR spectra were obtained using a Varian Inova-400 (400 MHz for ¹H, and 75 MHz for ¹³C) spectrometer. Chemical shifts are reported relative to chloroform ($\delta = 7.24$) for ¹H NMR or dimethyl sulfoxide ($\delta = 2.50$) for ¹H NMR and dimethyl sulfoxide ($\delta = 39.51$) for ¹³C NMR. Data are reported as (br = broad, s = singlet, d = doublet, t = triplet, q = quartet, m = multiplet).

Synthesis of JH-II-127 (18)

7H-pyrrolo[2,3-d]pyrimidine-2,4-diol

To a suspended solution of 6-aminouracil (12.7 g, 100 mmol) and sodium acetate (8.2 g, 100 mmol) in H_2O (100 mL) at a temperature of 70-75 °C was added a solution of chloroacetaldehyde (50% in water, 23.6 g, 150 mmol). The resulting reaction mixture was stirred at 80 °C for 20min and then cooled to room temperature. The resulting solid was collected by filtration, washed with water and acetone, and dried *in vacuo* to give the title compound as a light-brown solid (14.74 g, 98% yield). ¹H NMR (DMSO-*d6*, 400 MHz) δ 11.38 (br, 1H), 11.10 (br, 1H), 10.46 (s, 1H), 6.53 (s, 1H), 6.18 (s, 1H), MS m/z 152.43 [M+1].

2,4-dichloro-7H-pyrrolo[2,3-d]pyrimidine (25)

POCl₃ (229 mmol) was slowly added at room temperature to a suspension of 7H-pyrrolo[2,3-d]pyrimidine-2,4-diol 11.5 g (76 mmol) in toluene (60 mL). The reaction mixture was heated to 70 °C and 26.5 mL of DIPEA (153 mmol) was added drop-wise over a period of two hours, at which time the reaction temperature was increased to 106 °C and the mixture stirred overnight. After cooling to room temperature, the reaction mixture was poured into a mixture of 200 mL EtOAc and 300 mL ice-water and then filtered through celite. The aqueous layer was extracted with EtOAc (3x200 mL) and the combined organic layers were washed with

brine, decolored with activated carbon, filtered through celite and concentrated to give the title compound. MS m/z 189.64 [M+1]

2,4,5-trichloro-7H-pyrrolo[2,3-d]pyrimidine (26)

To a solution of of 2,4-Dichloro-7H-pyrrolo[2,3-d]pyrimidine (2.0 g, 10.64 mmol) in DCM/THF (15mL/6mL) was added NCS (1.70g, 12.76 mmol). The mixture was heated to 90 °C under microwave irradiation for 2.5 hours. The solvent was removed in vacuo and the crude product was purified by flash column chromatography using a 9:1 v/v Hexane:Ethyl acetate as solvent to afford the title compound 2.2 g (93% yield) as a white crystalline solid. MS *m/z* 223.48 [M+1]

2,4,5-trichloro-7-((2-(trimethylsilyl)ethoxy)methyl)-7H-pyrrolo[2,3-d]pyrimidine (27)

To a solution of 2,4,5-trichloro-7H-pyrrolo[2,3-d]pyrimidine (2.0 g, 8.99 mmol) in *N,N*-dimethylformamide (30 mL) was added sodium hydride 60% (432 mg, 10.78 mmol) at 0 °C. The reaction mixture was allowed to stir for 30 min. after which time 2-(trimethylsilyl)ethoxymethyl chloride (1.91 mL, 10.78 mmol) was added. The reaction mixture was further stirred at room temperature for a period of 3 hours after which time water was added and stirred for 20 minutes. The resulting precipitate was filtered and dried to yield the title compound 2.85 g (90% yield) as a brown solid. MS *m/z* 353.84 [M+1]

2,5-dichloro-N-methyl-7-((2-(trimethylsilyl)ethoxy)methyl)-7H-pyrrolo[2,3-d]pyrimidin-4-amine (28)

To a solution 2,4,5-trichloro-7-((2-(trimethylsilyl)ethoxy)methyl)-7H-pyrrolo[2,3-d]pyrimidine (1 g, 2.8 mmol) in MeOH (15 mL) was added a solution of Methylamine 33%

in MeOH (0.29 mL, 3.11 mmol). The mixture was heated to 70 °C for 1 hour followed by cooling to room temperature at which time the solvent was removed *in vacuo* and the crude product was purified by flash column chromatography using a 9:1 v/v Hexane:Ethyl acetate as solvent to afford the title compound 916 mg (93% yield) as a white solid. MS m/z 348.08 [M+1]

(4-((5-chloro-4-(methylamino)-7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)-3-methoxyphenyl)(morpholino)methanone (18)

To a solution of 2,5-dichloro-N-methyl-7-((2-(trimethylsilyl)ethoxy)methyl)-7H-pyrrolo[2,3d]pyrimidin-4-amine (100 mg, 0.28 mmol) in sec-BuOH (5 mL) was added (4-amino-3methoxyphenyl)(morpholino)methanone (82 mg, 0.35 mmol) and K₂CO₃ (77.4 mg, 0.56 mmol). The reaction mixture was degassed for 5 min and then Pd₂(dba)₃ (15.4 mg, 0.017 mmol) and 2-dicyclohexylphosphino-2',4',6'-triisopropylbiphenyl (12 mg, 0.025 mmol) were added. The reaction flask was stirred at 90 °C for 6 hr. After cooling to room temperature, the reaction mixture was filtered through a pad of celite and partitioned between ethyl acetate and water. The organic layer was separated and the aqueous layer was extracted with ethyl acetate. The combined organic extracts were washed with brine, dried over MgSO₄, filtered and concentrated. The crude product was dissolved in DCM (10 mL) and TFA (2mL) was added and stirred at 50 °C for 3 hours at which time the solvent was removed in vacuo. The crude material was dissolved in THF (10 mL) and a saturated solution of NaHCO₃ (10 mL) was added and stirred for 6 hr. The mixture was extracted with ethyl acetate and the combined organic extracts were washed with brine, dried over MgSO₄, filtered and concentrated. Purification by HPLC gave the title compound 68 mg (58% yield) as a brown solid. ¹H NMR 400 MHz (DMSO- d_6) δ 11.43 (br, 1H), 8.62 (d, J = 4 Hz, 1H), 7.44 (s, 1H), 7.03 $(m, 1H), 6.6 (m, 1H), 3.95 (s, 3H), 3.62 - 3.5 (m, 8H), 2.48 (s, 3H); {}^{13}C NMR 100 MHz$ $(DMSO-d_6)$ δ 169.65, 156.91, 150.37, 146.98, 131.83, 127.29, 120.39, 116.70, 116.43, 110.00, 102.03, 95.63, 66.59, 56.42, 28.17; MS *m/z*: 417.34 [M+1].

Synthesis of (4-((5-chloro-4-methoxy-7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)-3-methoxyphenyl)(morpholino)methanone (9)

2-chloro-4-methoxy-7H-pyrrolo[2,3-d]pyrimidine

To a solution of 2-chloro-4-methoxy-7H-pyrrolo[2,3-d]pyrimidine (1 g, 5.4 mmol) in MeOH (15 mL) was added a solution of Methylamine 33% in MeOH (0.56 mL, 5.9 mmol). The mixture was heated to 70 °C for 1 hour followed by cooling to room temperature at which time the solvent was removed *in vacuo* and the crude product was purified by flash column chromatography using a 9:1 v/v Hexane:Ethyl acetate as solvent to afford the title compound 965 mg (95% yield) as a white solid. MS *m/z* 184.08 [M+1]

(3-methoxy-4-((4-methoxy-7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)phenyl)(morpholino)methanone (9)

To a solution of 2-chloro-4-methoxy-7H-pyrrolo[2,3-d]pyrimidine (100 mg, 0.54 mmol) in *sec*-BuOH (5 mL) was added (4-amino-3-methoxyphenyl)(morpholino)methanone (154 mg, 0.65 mmol) and K_2CO_3 (150 mg, 1.08 mmol). The reaction mixture was degassed for 5 min and then $Pd_2(dba)_3$ (30 mg, 0.032 mmol) and 2-dicyclohexylphosphino-2',4',6'-triisopropylbiphenyl (23 mg, 0.049 mmol) were added. The reaction flask was stirred at 110 °C for 8 hr. After cooling to room temperature, the reaction mixture was filtered through a pad of celite and partitioned between ethyl acetate and water. The organic layer was separated and the aqueous layer was extracted with ethyl acetate. The combined organic extracts were washed with brine, dried over MgSO₄, filtered and concentrated. Purification by HPLC gave the title compound 64 mg (31% yield) as a brown solid. ¹H NMR 400 MHz (DMSO- d_6) δ 11.58 (br, 1H), 8.58 (d, J = 7 Hz, 1H), 7.67 (m, 1H), 7.07 – 6.99 (m, 2H), 6.34 – 6.31 (m, 1H), 4.01 (s, 1H), 3.92 (s, 3H), 2.52 (s, 1H), 2.51 – 2.46 (m, 8H) MS m/z: 384.58

(M + 1).

(3-methoxy-4-((4-(methylamino)-7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)phenyl)(morpholino)methanone (8)

¹HNMR (400 MHz, DMSO) δ 11.96 (br, 1H), 8.48 (d, J = 4 Hz, 1H), 7.04 (s, 1H), 6.96 (m, 1H), 6.54 (s, 1H), 3.95 (s, 3H), 2.48 (s, 3H) 2.44 – 2.39 (m, 8H)

$(4-((7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)-3-methoxyphenyl) (morpholino) methanone \\ (10)$

¹H NMR 400 MHz (DMSO- d_6) δ 12.05 (br, 1H), 8.81 (s, 1H), 8.56 (br, 1H), 8.38 (d, J = 7 Hz, 1H), 7.37 (m, 2H), 7.22 (m, 1H), 7.01 (m, 1H), 3.90 (s, 3H), 2.51 – 2.46 (m, 8H) MS m/z: 354.58 (M + 1).

(3-methoxy-4-((4-((2-methoxyethyl)amino)-7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)phenyl) (morpholino)methanone (11)

¹HNMR (400 MHz, DMSO- d_6) δ 11.87 (br, 1H), 8.40 (d, J = 9 Hz, 1H), 7.23 (s, 1H), 7.10 (m, 1H), 7.03 – 6.98 (m, 1H), 3.92 (s, 3H), 3.27 (s, 3H), 2.48 (m, 12H) MS m/z: 427.19 (M + 1).

(4-((4-(ethylamino)-7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)-3-methoxyphenyl)(morpholino)methanone (12)

¹HNMR (400 MHz, DMSO- d_6) δ 11.98 (br, 1H), 9.01 (m, 1H), 8.4 (m, 1H), 7.13 (s, 1H), 7.02 (m, 1H), 6.84 (m, 1H), 6.64 (s, 1H), 3.95 (s, 3H), 2.45 – 2.41 (m, 8H), 1.68 (q, J = 7 Hz, 2H), 0.88 (t, J = 7 Hz, 3H) MS m/z : 397.09 (M + 1).

(3-methoxy-4-((4-(propylamino)-7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)phenyl)(morpholino)methanone (13)

¹HNMR (400 MHz, DMSO- d_6) δ 11.91 (br, 1H), 8.43 (d, J = 4 Hz, 1H), 7.81 (m, 1H), 7.10 (s, 1H), 7.0 (m, 1H), 6.61 (s, 1H), 6.48 (m, 1H), 3.98 (s, 3H), 2.47 (s, 1H), 2.45 – 2.41 (m, 8H), 1.38 – 1.01 (m, 4H), 0.88 (t, J = 7 Hz, 3H) MS m/z : 411.33 (M + 1).

(3-methoxy-4-((4-(((tetrahydrofuran-2-yl)methyl)amino)-7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)phenyl)(morpholino)methanone (14)

¹HNMR (400 MHz, DMSO- d_6) δ 7.22 (s, 1H), 7.16 (d, J = 9 Hz, 1H), 7.23 (s, 1H), 4.01 (s, 3H), 3.14 (d, J = 4 Hz 3H), 2.63 – 2.6 (m, 12H) MS m/z: 453.07 (M + 1).

(4-((4-(cyclopropylamino)-7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)-3-methoxyphenyl)(morpholino)methanone (15)

¹HNMR (400 MHz, DMSO- d_6) δ 12.06 (br, 1H), 8.56 (d, J = 4 Hz, 1H), 7.98 (m, 1H), 7.08 (s, 1H), 6.64 (s, 1H), 4.01 (s, 3H), 2.56 (s, 3H) 2.48 (m, 8H), 0.99 (m, 1H), 0.81 – 0.55 (m, 4H) MS m/z: 409.21 (M + 1).

(2-fluoro-5-methoxy-4-((4-(methylamino)-7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)phenyl)(morpholino)methanone (16)

¹HNMR (400 MHz, DMSO- d_6) δ 11.94 (br, 1H), 8.26 (s, 1H), 7.64 (s, 1H), 6.96 (m, 1H), 6.54 (s, 1H), 3.95 (s, 3H), 2.48 (s, 3H) 2.44 – 2.39 (m, 8H) MS m/z: 401.36 (M + 1).

(4-((5-chloro-7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)-3-methoxyphenyl)(morpholino)methanone (17)

¹H NMR 400 MHz (DMSO- d_6) δ 12.05 (br, 1H), 8.81 (s, 1H), 8.56 (br, 1H), 8.38 (d, J = 7 Hz, 1H), 7.37 (m, 2H), 7.22 (m, 1H), 6.88 (s, 1H), 3.92 (s, 3H), 2.50 – 2.48 (m, 8H) MS m/z: 388.27 (M + 1).

(4-((5-chloro-4-(ethylamino)-7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)-3-methoxyphenyl)(morpholino)methanone (19)

¹H NMR 400 MHz (DMSO- d_6) δ 12.01 (br, 1H), 8.62 (d, J = 4 Hz, 1H), 7.68 (br, 1H), 7.42 (s, 1H), 6.98 (m, 1H), 6.6 (m, 1H), 4.01 (q, J = 7 Hz, 2H), 3.91 (s, 3H), 2.50 – 2.46 (m, 8H), 1.2 (m, 3H); MS m/z: 431.17 [M+1].

(4-((5-chloro-4-(propylamino)-7H-pyrrolo[2,3-d]pyrimidin-2-yl)amino)-3-methoxyphenyl)(morpholino)methanone (20)

¹H NMR 400 MHz (DMSO- d_6) δ 11.48 (br, 1H), 8.59 (m, 1H), 7.0 (m, 2H), 3.91 (s, 3H), 2.50 – 2.46 (m, 8H), 1.65 (m, 5H), 0.95 (m, 2H); MS m/z: 445.25 [M+1].

Synthesis of (3-methoxy-4-((6-(methylamino)-9H-purin-2-yl)amino)phenyl)(morpholino)methanone (21)

$$CI \xrightarrow{N} X \xrightarrow{H} X \xrightarrow{N} X$$

2-chloro-N-methyl-9H-purin-6-amine

To a solution of 2,6-dichloro-9H-purine (1 g, 5.29 mmol) in MeOH (15 mL) was added a solution of Methylamine 33% in MeOH (0.55 mL, 5.89 mmol). The mixture was heated to 70 °C for 1 hour followed by cooling to room temperature. The resulting precipitate was filtered, washed with ice-cold MeOH and dried to afford the title compound 880 mg (92% yield) as a

(3-methoxy-4-((6-(methylamino)-9H-purin-2-yl)amino)phenyl)(morpholino)methanone (21)

To a solution of 2-chloro-N-methyl-9H-purin-6-amine (100 mg, 0.54 mmol) in sec-BuOH (5 mL) was added (4-amino-3-methoxyphenyl)(morpholino)methanone (154 mg, 0.65 mmol) and K₂CO₃ (150 mg, 1.08 mmol). The reaction mixture was degassed for 5 min and then Pd₂(dba)₃ (30)0.032 mmol) and 2-dicyclohexylphosphino-2',4',6'mg, triisopropylbiphenyl (23 mg, 0.049 mmol) were added. The reaction flask was stirred at 110 °C for 8 hr. After cooling to room temperature, the reaction mixture was filtered through a pad of celite and partitioned between ethyl acetate and water. The organic layer was separated and the aqueous layer was extracted with ethyl acetate. The combined organic extracts were washed with brine, dried over MgSO₄, filtered and concentrated. Purification by HPLC gave the title compound 70 mg (35% yield) as a light-brown solid. ¹HNMR (400 MHz, DMSO) δ 8.39 (d, J = 9 Hz, 1H), 8.06 (s, 1H), 7.06 – 7.02 (m, 1H), 6.98 (d, J = 3Hz, 1H), 3.85 (s, 1H), 2.47 (s, 3H), 2.45 - 2.51 (m, 8H), MS m/z 384.19 [M+1]

Synthesis of (3-methoxy-4-((6-methoxy-9H-purin-2-yl)amino)phenyl)(morpholino)methanone (22)

2-chloro-6-methoxy-9H-purine

2-chloro-6-methoxy-9H-purine

To a solution of 2,6-dichloro-9H-purine (1 g, 5.29 mmol) in MeOH (15 mL) was added a

solution of NaOMe 25% in MeOH (1.26 mL, 5.89 mmol). The mixture was heated to 80 $^{\circ}$ C for 1 hour followed by cooling to room temperature. The resulting precipitate was filtered, washed with ice-cold MeOH and dried to afford the title compound 920 mg (94% yield) as a white solid. MS m/z 185.36 [M+1]

(3-methoxy-4-((6-methoxy-9H-purin-2-yl)amino)phenyl)(morpholino)methanone (22)

To a solution of 2-chloro-6-methoxy-9H-purine (100 mg, 0.54 mmol) in sec-BuOH (5 mL) was added (4-amino-3-methoxyphenyl)(morpholino)methanone (154 mg, 0.65 mmol) and K₂CO₃ (150 mg, 1.08 mmol). The reaction mixture was degassed for 5 min and then Pd₂(dba)₃ (30 0.032 mmol) and 2-dicyclohexylphosphino-2',4',6'mg, triisopropylbiphenyl (23 mg, 0.049 mmol) were added. The reaction flask was stirred at 110 °C for 8 hr. After cooling to room temperature, the reaction mixture was filtered through a pad of celite and partitioned between ethyl acetate and water. The organic layer was separated and the aqueous layer was extracted with ethyl acetate. The combined organic extracts were washed with brine, dried over MgSO₄, filtered and concentrated. Purification by HPLC gave the title compound 50 mg (24% yield) as a light-brown solid. ¹HNMR (400 MHz, DMSO) δ 8.46 (d, J = 9 Hz, 1H), 8.01 (s, 1H), 7.89 (s, 1H), 7.04 (m, 1H), 4.03 (s, 3H), 3.96 (s, 3H), 2.48 - 2.40 (m, 8H), MS m/z 385.62 [M+1]

Synthesis of (3-methoxy-4-((4-(methylamino)-1H-pyrazolo[3,4-d]pyrimidin-6-yl)amino)phenyl)(morpholino)methanone (23)

2,4,6-trichloropyrimidine-5-carbaldehyde

DMF (12 mL) was added drop-wise to POCl₃ (75 mL, 800 mmol) at 0 °C, followed by barbituric acid (15 g, 118 mmol) was added portion-wise to the mixture. The mixture was stirred at 120 °C for 16 hr. Excess POCl₃ was removed *in vacuo* and the resulting residue was gradually poured into ice water. The mixture was extracted with DCM and the organic layer was washed with sat. NaHCO₃ solution, dried and concentrated to give the title compound as a yellow solid, yield 19.0 g, 77%. ¹HNMR (CDCl₃, 400MHz): δ 10.45 (s, 1H). MS *m/z* 212.81 [M+1]

4,6-dichloro-1H-pyrazolo[3,4-d]pyrimidine

To a solution of 2,4,6-trichloropyrimidine-5-carbaldehyde (19.0 g, 990 mmol) in methanol (300 mL) was added drop-wise a solution of hydrazine monohydrate (4.8 mL) in methanol (80 mL) at 0 °C followed by drop-wise addition of triethylamine (13 mL) in methanol (80 mL) at 0 °C. The mixture was stirred at the same temperature for 30 min. The solvent was removed and the residue was purified by flash chromatography to afford the title compound (10.3 g, 62% yield) as a yellow solid. ¹H NMR (CDCl₃, 400MHz): δ 11.56 (br, 1H), 8.43 (s, 1H). MS *m/z* 190.68 [M+1]

6-chloro-N-methyl-1H-pyrazolo[3,4-d]pyrimidin-4-amine

To a solution of 4,6-dichloro-1H-pyrazolo[3,4-d]pyrimidine (1 g, 5.29 mmol) in MeOH (15 mL) was added a solution of Methylamine 33% in MeOH (0.55 mL, 5.89 mmol). The mixture was heated to 70 °C for 1 hour followed by cooling to room temperature. The resulting precipitate was filtered, washed with ice-cold MeOH and dried to afford the title

3-methoxy-4-((4-(methylamino)-1H-pyrazolo[3,4-d]pyrimidin-6-yl)amino)phenyl)(morpholino)methanone (23)

To a solution of 6-chloro-N-methyl-1H-pyrazolo[3,4-d]pyrimidin-4-amine (100 mg, 0.54 mmol) in sec-BuOH (5 mL) was added (4-amino-3-methoxyphenyl)(morpholino)methanone (154 mg, 0.65 mmol) and K_2CO_3 (150 mg, 1.08 mmol). The reaction mixture was degassed for 5 min and then $Pd_2(dba)_3$ (30 mg, 0.032 mmol) and 2-dicyclohexylphosphino-2',4',6'-triisopropylbiphenyl (23 mg, 0.049 mmol) were added. The reaction flask was stirred at 110 $^{\circ}$ C for 8 hr. After cooling to room temperature, the reaction mixture was filtered through a pad of celite and partitioned between ethyl acetate and water. The organic layer was separated and the aqueous layer was extracted with ethyl acetate. The combined organic extracts were washed with brine, dried over MgSO₄, filtered and concentrated. Purification by HPLC gave the title compound 64 mg (31% yield) as a brown solid. 1 H NMR 400 MHz (CD₃OD) δ 8.18 (m, 1H), 8.10 (br, 1H), 8.02 (m, 1H), 7.83 (m, 1H), 7.58 (m, 1H), 7.20 (m, 1H), 6.86 (m, 1H), 4.05 (m, 1H), 3.97 (m, 3.91, 3.80 (m, 2H), 3.50 (m, 2H), 2.20 (m, 2H), 1.98 (m, 2H), 1.18 (m, 3.96 (m, 3.97), 3.84.63 (m) 1.18 (m, 3.97), 3.80 (m, 3.91).

Synthesis of (3-methoxy-4-((4-methoxy-1H-pyrazolo[3,4-d]pyrimidin-6-yl)amino)phenyl)(morpholino)methanone (24)

6-chloro-4-methoxy-1H-pyrazolo[3,4-d]pyrimidine

To a solution of 6-chloro-4-methoxy-1H-pyrazolo[3,4-d]pyrimidine (1 g, 5.29 mmol) in

MeOH (15 mL) was added a solution of NaOMe 25% in MeOH (1.26 mL, 5.89 mmol). The mixture was heated to 80 °C for 1 hour followed by cooling to room temperature. The resulting precipitate was filtered, washed with ice-cold MeOH and dried to afford the title compound 910 mg (92% yield) as a white solid. MS *m/z* 185.46 [M+1]

(3-methoxy-4-((4-methoxy-1H-pyrazolo[3,4-d]pyrimidin-6-yl)amino)phenyl) (morpholino)methanone (24)

To a solution of 6-chloro-4-methoxy-1H-pyrazolo[3,4-d]pyrimidine (100 mg, 0.54 mmol) in *sec*-BuOH (5 mL) was added (4-amino-3-methoxyphenyl)(morpholino)methanone (154 mg, 0.65 mmol) and K_2CO_3 (150 mg, 1.08 mmol). The reaction mixture was degassed for 5 min and then $Pd_2(dba)_3$ (30 mg, 0.032 mmol) and 2-dicyclohexylphosphino-2',4',6'-triisopropylbiphenyl (23 mg, 0.049 mmol) were added. The reaction flask was stirred at 110 °C for 8 hr. After cooling to room temperature, the reaction mixture was filtered through a pad of celite and partitioned between ethyl acetate and water. The organic layer was separated and the aqueous layer was extracted with ethyl acetate. The combined organic extracts were washed with brine, dried over MgSO₄, filtered and concentrated. Purification by HPLC gave the title compound 58 mg (28% yield) as a light-brown solid. ¹HNMR 400 MHz (DMSO) δ 13.31 (br, 1H), 8.39 (d, J = 4 Hz, 1H), 7.98 (d, J = 8 Hz, 1H), 7.01 (m, 2H), 4.05 (s, 3H), 3.98 (s, 3H), 2.46 (m, 8H), MS m/z : 385.23 (M + 1).

Biological materials and methods

Reagents and General methods. Tissue-culture reagents were from Life Technologies. P81 phosphocellulose paper was from Whatman and [γ -32P]-ATP was from Perkin Elmer. Nictide and LRRKtide were synthesised by Pepceuticals. Protein G sepharose was from Amersham. DNA constructs used for transfection were purified from *Escherichia coli* DH5 α

using Qiagen or Invitrogen plasmid Maxi kits according to the manufacturer's protocol. All DNA constructs were verified by DNA sequencing, which was performed by The Sequencing Service, School of Life Sciences, University of Dundee, Scotland, U.K., using DYEnamic ET terminator chemistry (Amersham Biosciences) on Applied Biosystems automated DNA sequencers.

Cell culture, treatments and cell lysis. HEK293 and Swiss 3T3 cells were cultured in DMEM (Dulbecco's Modified Eagle's medium) supplemented with 10% FBS (fetal bovine serum), 2mM glutamine and 1× penicillin/streptomycin solution. HEK-293 T-Rex cell lines were cultured in DMEM supplemented with 10% (v/v) FBS and 2mM glutamine, 1× penicillin/streptomycin solution, 15 µg/ml blastocidin and 100 µg/ml hygromycin. T-Rex cultures were induced to express the indicated protein by inclusion of 0.1 µg/ml doxycycline in the culture medium for 24-48 hours. Human lymphoblastoid cells were maintained in RPMI 1640 with 10% FBS, 2 mM glutamine, 1× penicillin/streptomycin solution and were maintained at cell density of $0.3 \times 10^6 - 2 \times 10^6$ cells per ml. Epstein-Barr virus immortalized primary human lymphoblastoid cells from one control subject and one Parkinson's disease patient homozygous for the LRRK2[G2019S] mutation were kindly provided by Alastair Reith (GSK) and have been described previously ¹. For inhibitor experiments, JH-II-127 and/or LRRK2-IN-1 was dissolved in DMSO and utilised at the indicated concentrations. The concentration of DMSO in the culture media did not exceed 1%. Following treatment, cells were washed once with PBS and lysed with buffer containing 50 mM Tris/HCl, pH 7.5, 1 mM EGTA, 1 mM EDTA, 1 mM sodium orthovanadate, 10 mM sodium β-glycerophosphate, 50 mM NaF, 5 mM sodium pyrophosphate, 0.27 M sucrose, 1 mM benzamidine, 2 mM phenylmethanesulphonylfluoride (PMSF) and 1% Triton X-100. When not used immediately, all lysate supernatants were snap-frozen in liquid nitrogen and stored at −80 °C until use. Protein concentrations were determined following centrifugation of the lysate at 16,000 x g at 4 °C for 20 minutes using the Bradford method with BSA as the standard. Transient transfection of HEK293 cells was performed using the PEI method ².

IC₅₀ **determination.** Active GST-LRRK2 (1326-2527), GST-LRRK2[G2019S] (1326-2527), GST-LRRK2[A2016T] (1326-2527) and GST-LRRK2[A2016T+G2019S] (1326-2527) enzyme was purified with glutathione sepharose from HEK293 cell lysate 36 h following transient transfection of the appropriate cDNA constructs as described previously 3 . Peptide kinase assays, performed in triplicate, were set up in a total volume of 40 μl containing 0.5 μg LRRK2 kinase (which at approximately 10% purity gives a final concentration of 8 nM) in

50 mM Tris/HCl, pH 7.5, 0.1 mM EGTA, 10 mM MgCl₂, 20 μ M Nictide, 0.1 μ M [γ -32P]ATP (\sim 500 cpm/pmol) and the indicated concentrations of inhibitor dissolved in DMSO. After incubation for 15 min at 30 °C, reactions were terminated by spotting 35 μ l of the reaction mix onto P81 phosphocellulose paper and immersion in 50 mM phosphoric acid. Samples were washed extensively and the incorporation of [γ -32P]ATP into Nictide was quantified by Cerenkov counting. IC₅₀ values were calculated with GraphPad Prism using non-linear regression analysis.

Immunoblot procedures. Cell lysates from Swiss 3T3 cells, human lymphoblastoid cells, GFP-LRRK2 expressing stable cell lines and mouse tissues were eluted in 65 μl 2x LDS sample buffer (Invitorgen) with final concentration of 1 μg/μl. Following heating at 70 °C for 10 min, 15 μl aliquots were resolved on 8 % SDS polyacrylamide gels and transferred to nitrocellulose membranes for detection of LRRK2 phosphorylated at Ser910, LRRK2 phosphorylated at Ser935 and total LRRK2, using purified rabbit monoclonal antibodies (LRRK2 phospho-serine 910 clone (LRRK2α pS910 UDD1), LRRK2 phospho-serine 935 clone (LRRK2α pS935 UDD2) and LRRK2 100-500 clone (LRRK2 (100 - 500) UDD3)) in PBS with 0.1% sodium azide (Epitomics) at 0.1 μg/ml concentrations. Immunoblot films were scanned on an Epson 4990 scanner, and images were managed with Adobe Photoshop. Secondary antibodies coupled to horseradish peroxidase used for immunoblotting were obtained from Pierce.

Pharmacodynamic (PD) study. JH-II-127 (18) was dissolved in 5% 1-methyl-2-pyrrolidinone (NMP) /95% PEG 300 (Sigma) solution and administered by intraperitoneal injection into wild type male C57BL/6 mice at doses of 0, 3, 10, 30, 50 and 100 mg/kg and at 100 mg/kg LRRK2-IN-1 (1) as a comparative control. Control mice were treated with an equal volume of NMP/PEG solution. One hour after administration, mice were sacrificed by cervical dislocation and, kidney, spleen and brain tissues were rapidly dissected and snap-frozen in liquid nitrogen. Animal experiments were approved by the University of Dundee Ethics Committee and performed under a U.K. Home office project license.

Dundee profiling⁴

JH-II-127 (18) was tested at 10 and 1 uM. Results show percent remaining enzymatic activity relative to the DMSO control.

K	I1	K	I10
TGFBR1	128	BTK	126
Aurora A	124	EPH-B1	126
MARK3	120	TGFBR1	119
EPH-B1	119	WNK1	115
PKBb	118	Aurora A	112
MELK	118	Lck	111
RSK2	117	РКСү	109
p38g MAPK	116	MPSK1	106
BTK	116	PKBa	105
IKKe	114	PAK6	105
ΡΚСγ	113	TLK1	105
TAO1	113	p38a MAPK	104
MPSK1	113	AMPK	103
ROCK 2	113	ABL	103
AMPK	112	PKBb	103
MAP4K5	110	NEK2a	103
ULK2	110	FGF-R1	102
Src	110	PAK2	102
PAK2	110	S6K1	102
CHK1	110	EF2K	102
BRSK2	110	p38d MAPK	102
SIK2	109	Src	102
MARK4	109	EPH-A4	102
MSK1	109	ROCK 2	101
IRAK1	109	EPH-A2	101
PDGFRA	109	PKA	101
WNK1	108	PKCa	100
ERK8	108	TAO1	100
p38b MAPK	108	VEG-FR	100
p38d MAPK	107	MSK1	99
EPH-A4	107	BRSK2	98
ERK1	107	BRSK1	98
SIK3	107	MARK4	98
TLK1	106	IKKe	98

ABL	106	TTBK1	98
S6K1	106	MST4	97
HIPK3	106	JAK2	97
CDK9-Cyclin T1	105	GSK3b	97
PINK	105	SIK3	96
EF2K	105	p38g MAPK	96
TTBK2	105	LKB1	96
VEG-FR	104	IKKb	95
TIE2	104	MST3	95
AMPK (hum)	104	TESK1	95
EPH-A2	104	ULK1	95
TTBK1	103	ULK2	95
PRK2	103	RSK2	94
ULK1	103	PKCz	94
LKB1	103	TIE2	94
MST4	103	PRK2	93
MST3	102	CHK1	93
PKCa	102	HIPK3	93
EIF2AK3	102	DDR2	93
MNK1	102	MAP4K5	92
PAK4	102	EPH-B3	92
RIPK2	102	AMPK (hum)	92
PAK6	102	CSK	92
CDK2-Cyclin A	102	PAK4	92
NEK6	101	IRAK1	92
PKBa	101	MARK3	92
ZAP70	101	SIK2	92
MKK2	101	CDK9-Cyclin T1	91
MARK2	101	Aurora B	90
CK1γ2	100	TTBK2	90
IKKb	100	PDGFRA	89
JAK2	99	PINK	89
PKA	99	PDK1	88
DDR2	98	PIM1	88
MAP4K3	98	EPH-B4	86
MAPKAP-K3	97	YES1	85
Aurora B	97	MEKK1	85
NEK2a	97	ZAP70	85
Lck	97	MAPKAP-K3	84
TESK1	97	TrkA	84
PIM3	96	RIPK2	83
RSK1	96	NEK6	83

TAK1	96	MST2	82
IRAK4	95	IRAK4	82
MINK1	95	PIM2	82
HIPK1	95	MARK2	82
PIM2	95	SYK	81
PIM1	95	p38b MAPK	79
MEKK1	94	EIF2AK3	79
ERK2	94	MELK	79
HER4	94	MNK1	78
BRSK1	94	MARK1	77
CK2	94	SGK1	76
FGF-R1	94	PIM3	75
EPH-B3	92	MKK2	72
PDK1	91	ERK2	71
MKK1	91	ERK8	71
SYK	91	HER4	71
MST2	90	CK2	70
p38a MAPK	90	PAK5	70
MARK1	89	ERK1	68
CSK	89	MINK1	68
PAK5	89	TBK1	67
SRPK1	89	CDK2-Cyclin A	67
TBK1	88	RSK1	60
GCK	86	EPH-B2	58
SGK1	86	MKK1	56
MAPKAP-K2	86	ERK5	52
YES1	85	CK1 ₂ 2	45
TrkA	85	HIPK1	44
HIPK2	84	HIPK2	43
GSK3b	82	TAK1	42
MLK1	81	MAP4K3	41
EPH-B4	80	GCK	41
CAMK1	80	MAPKAP-K2	40
MLK3	78	MLK1	38
ERK5	76	MLK3	35
PKCz	76	CAMK1	34
MNK2	75	CK1δ	33
NUAK1	75	SRPK1	31
CK18	74	DYRK3	30
DYRK3	74	MNK2	25
TTK	69	BRK	19
DYRK2	67	DYRK2	18

BRK	62	NUAK1	17
PLK1	62	CAMKKb	16
OSR1	61	STK33	15
EPH-B2	60	PLK1	13
ASK1	56	TTK	12
CAMKKb	50	OSR1	12
DYRK1A	49	MKK6	11
PKD1	47	IRR	10
PRAK	43	IR	10
JNK1	38	ASK1	10
MKK6	37	IGF-1R	9
IRR	31	JNK3	9
JNK3	29	DYRK1A	8
JNK2	27	PKD1	7
STK33	22	TSSK1	7
DAPK1	21	PRAK	6
IR	21	JNK1	6
TSSK1	14	DAPK1	5
CLK2	13	JNK2	4
PHK	13	CLK2	2
IGF-1R	13	PHK	1
SmMLCK	9	CHK2	1
CHK2	5	SmMLCK	1

Supplementary Fig.1 IC50s of JH-II-127 against SmMLCK and CHK2

KinomeScan analysis ⁵

Compound ID	JH-II-127
Compound Concentration (uM)	1

AAK1	88
ABL1(E255K)-phosphorylated	95
ABL1(F317I)-nonphosphorylated	96
ABL1(F317I)-phosphorylated	100
ABL1(F317L)-nonphosphorylated	93
ABL1(F317L)-phosphorylated	100
ABL1(H396P)-nonphosphorylated	100
ABL1(H396P)-phosphorylated	95
ABL1(M351T)-phosphorylated	80
ABL1(Q252H)-nonphosphorylated	88
ABL1(Q252H)-phosphorylated	94
ABL1(T315I)-nonphosphorylated	91
ABL1(T315I)-phosphorylated	96
ABL1(Y253F)-phosphorylated	97
ABL1-nonphosphorylated	100
ABL1-phosphorylated	100
ABL2	96
ACVR1	91
ACVR1B	82
ACVR2A	100
ACVR2B	100
ACVRL1	85
ADCK3	100
ADCK4	100
AKT1	97
AKT2	93
AKT3	93
ALK	70
ALK(C1156Y)	68
ALK(L1196M)	74
AMPK-alpha1	97
AMPK-alpha2	98
ANKK1	95
ARK5	89
ASK1	84

ACKS	100
ASK2	100
AURKA	100
AURKB	80
AURKC	100
AXL	97
BIKE	97
BLK	100
BMPR1A	91
BMPR1B	98
BMPR2	100
BMX	100
BRAF	100
BRAF(V600E)	100
BRK	97
BRSK1	100
BRSK2	100
ВТК	100
BUB1	100
CAMK1	100
CAMK1D	100
CAMK1G	85
CAMK2A	92
САМК2В	86
CAMK2D	99
CAMK2G	95
CAMK4	100
CAMKK1	96
CAMKK2	100
CASK	88
CDC2L1	85
CDC2L2	100
CDC2L5	100
CDK11	63
CDK2	95
CDK3	96
<u> </u>	5 0

CDK4-cyclinD1 95 CDK5 100 CDK7 95 CDK8 74 CDK9 92 CDKL1 87 CDKL2 86 CDKL3 86 CDKL5 100 CHEK1 100 CHEK2 31 CIT 92 CLK1 87 CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1B 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74		1
CDK5 100 CDK7 95 CDK8 74 CDK9 92 CDKL1 87 CDKL2 86 CDKL3 86 CDKL5 100 CHEK1 100 CHEK2 31 CIT 92 CLK1 87 CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1 80 CSNK1B 100 CSNK1G 100 CSNK1G 100 CSNK1G 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CDK4-cyclinD1	95
CDK7 95 CDK8 74 CDK9 92 CDKL1 87 CDKL2 86 CDKL3 86 CDKL5 100 CHEK1 100 CHEK2 31 CIT 92 CLK1 87 CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSN 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74		92
CDK8 74 CDK9 92 CDKL1 87 CDKL2 86 CDKL3 86 CDKL5 100 CHEK1 100 CHEK2 31 CIT 92 CLK1 87 CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1 97 CSNK1B 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CDK5	100
CDK9 92 CDKL1 87 CDKL2 86 CDKL3 86 CDKL5 100 CHEK1 100 CHEK2 31 CIT 92 CLK1 87 CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1G 97 CSNK1G 97 CSNK1G 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CDK7	95
CDKL1 87 CDKL2 86 CDKL3 86 CDKL5 100 CHEK1 100 CHEK2 31 CIT 92 CLK1 87 CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1G 97 CSNK1G 97 CSNK1G 97 CSNK1G 97 CSNK2A1 93 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CDK8	74
CDKL2 86 CDKL5 100 CHEK1 100 CHEK2 31 CIT 92 CLK1 87 CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1B 100 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CDK9	92
CDKL3 86 CDKL5 100 CHEK1 100 CHEK2 31 CIT 92 CLK1 87 CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1E 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CDKL1	87
CDKL5 100 CHEK1 100 CHEK2 31 CIT 92 CLK1 87 CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1B 100 CSNK1G 97 CSNK1G 97 CSNK1G 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CDKL2	86
CHEK1 100 CHEK2 31 CIT 92 CLK1 87 CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1G 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CDKL3	86
CHEK2 31 CIT 92 CLK1 87 CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1E 100 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CDKL5	100
CIT 92 CLK1 87 CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1E 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CHEK1	100
CLK1 87 CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1E 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CHEK2	31
CLK2 56 CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1E 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CIT	92
CLK3 99 CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1E 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CLK1	87
CLK4 63 CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1E 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CLK2	56
CSF1R 79 CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1E 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CLK3	99
CSF1R-autoinhibited 47 CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1E 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CLK4	63
CSK 100 CSNK1A1 80 CSNK1A1L 97 CSNK1D 82 CSNK1E 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CSF1R	79
CSNK1A1L 80 CSNK1D 82 CSNK1E 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CSF1R-autoinhibited	47
CSNK1A1L 97 CSNK1D 82 CSNK1E 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CSK	100
CSNK1D 82 CSNK1E 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CSNK1A1	80
CSNK1E 100 CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CSNK1A1L	97
CSNK1G1 97 CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CSNK1D	82
CSNK1G2 100 CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CSNK1E	100
CSNK1G3 100 CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CSNK1G1	97
CSNK2A1 93 CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CSNK1G2	100
CSNK2A2 70 CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CSNK1G3	100
CTK 96 DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CSNK2A1	93
DAPK1 54 DAPK2 40 DAPK3 52 DCAMKL1 74	CSNK2A2	70
DAPK2 40 DAPK3 52 DCAMKL1 74	СТК	96
DAPK3 52 DCAMKL1 74	DAPK1	54
DCAMKL1 74	DAPK2	40
	DAPK3	52
DCAMKL2 100	DCAMKL1	74
	DCAMKL2	100

DCAMKL3	77
DDR1	100
DDR2	93
DLK	89
DMPK	98
DMPK2	84
DRAK1	89
DRAK2	86
DYRK1A	95
DYRK1B	66
DYRK2	89
EGFR	93
EGFR(E746-A750del)	100
EGFR(G719C)	91
EGFR(G719S)	88
EGFR(L747-E749del, A750P)	81
EGFR(L747-S752del, P753S)	94
EGFR(L747-T751del,Sins)	84
EGFR(L858R)	83
EGFR(L858R,T790M)	100
EGFR(L861Q)	57
EGFR(S752-I759del)	97
EGFR(T790M)	97
EIF2AK1	100
EPHA1	100
EPHA2	88
EPHA3	88
EPHA4	94
EPHA5	98
EPHA6	88
EPHA7	98
EPHA8	99
EPHB1	100
EPHB2	98
ЕРНВ3	89

EPHB6 100 ERBB2 88 ERBB3 84 ERBB4 100 ERK1 100 ERK2 96 ERK3 62 ERK4 100 ERK5 85 ERK8 100 ERN1 92 FAK 93 FER 95 FES 99 FGFR1 99 FGFR2 87 FGFR3 83 FGFR3(G697C) 100 FGFR4 98 FGR 81 FLT1 97 FLT3 FLT3 100 FLT3(D835H) 98 FLT3(D835Y) 100 FLT3(N841I) FLT3(R834Q) 100 FLT3(R834Q) 100 FLT3-autoinhibited 100 FYN 90 GAK 655 GCN2(Kin.Dom.2,S808G) 91	EPHB4	83
ERBB2 88 ERBB3 84 ERBB4 100 ERK1 100 ERK2 96 ERK3 62 ERK4 100 ERK5 85 ERK8 100 ERN1 92 FAK 93 FER 95 FES 99 FGFR1 99 FGFR2 87 FGFR3 83 FGFR4 98 FGR 81 FLT1 97 FLT3 100 FLT3(D835H) 98 FLT3(D835Y) 100 FLT3(K663Q) 100 FLT3(R841I) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
ERBB3 84 ERBB4 100 ERK1 100 ERK2 96 ERK3 62 ERK4 100 ERK5 85 ERK8 100 ERN1 92 FAK 93 FER 95 FES 99 FGFR1 99 FGFR2 87 FGFR3 83 FGFR3(G697C) 100 FGFR4 98 FGR 81 FLT1 97 FLT3 100 FLT3(D835H) 98 FLT3(ITD) 94 FLT3(K663Q) 100 FLT3(R834Q) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
ERBB4 100 ERK1 100 ERK2 96 ERK3 62 ERK4 100 ERK5 85 ERK8 100 ERN1 92 FAK 93 FER 95 FES 99 FGFR1 99 FGFR2 87 FGFR3 83 FGFR3(G697C) 100 FGFR4 98 FGR 81 FLT1 97 FLT3 100 FLT3(D835H) 98 FLT3(ITD) 94 FLT3(ITD) 94 FLT3(N841I) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
ERK1 100 ERK2 96 ERK3 62 ERK4 100 ERK5 85 ERK8 100 ERN1 92 FAK 93 FER 95 FES 99 FGFR1 99 FGFR2 87 FGFR3 83 FGFR3(G697C) 100 FGFR4 98 FGR 81 FLT1 97 FLT3 (D835H) 98 FLT3(D835H) 98 FLT3(ITD) 94 FLT3(ITD) 94 FLT3(R841I) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FVN 90 GAK 65		
ERK2 96 ERK3 62 ERK4 100 ERK5 85 ERK8 100 ERN1 92 FAK 93 FER 95 FES 99 FGFR1 99 FGFR2 87 FGFR3 83 FGFR3(G697C) 100 FGFR4 98 FGR 81 FLT1 97 FLT3 100 FLT3(D835H) 98 FLT3(D835Y) 100 FLT3(ITD) 94 FLT3(K663Q) 100 FLT3(R834Q) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
ERK3 62 ERK4 100 ERK5 85 ERK8 100 ERN1 92 FAK 93 FER 95 FES 99 FGFR1 99 FGFR2 87 FGFR3 83 FGFR3(G697C) 100 FGFR4 98 FGR 81 FLT1 97 FLT3 100 FLT3(D835H) 98 FLT3(D835H) 98 FLT3(D835Y) 100 FLT3(K663Q) 100 FLT3(N841I) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
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ERK8 100 ERN1 92 FAK 93 FER 95 FES 99 FGFR1 99 FGFR2 87 FGFR3 83 FGFR3(G697C) 100 FGFR4 98 FGR 81 FLT1 97 FLT3 100 FLT3(D835H) 98 FLT3(D835Y) 100 FLT3(ITD) 94 FLT3(K663Q) 100 FLT3(N841I) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
ERN1 92 FAK 93 FER 95 FES 99 FGFR1 99 FGFR2 87 FGFR3 83 FGFR3(G697C) 100 FGR4 98 FGR 81 FLT1 97 FLT3 100 FLT3(D835H) 98 FLT3(D835Y) 100 FLT3(ITD) 94 FLT3(N841I) 100 FLT3(R834Q) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
FAK 93 FER 95 FES 99 FGFR1 99 FGFR2 87 FGFR3 83 FGFR3(G697C) 100 FGFR4 98 FGR 81 FLT1 97 FLT3 100 FLT3(D835H) 98 FLT3(D835H) 98 FLT3(D835Y) 100 FLT3(K663Q) 100 FLT3(N841I) 100 FLT3(R834Q) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
FER 95 FES 99 FGFR1 99 FGFR2 87 FGFR3 83 FGFR3(G697C) 100 FGFR4 98 FGR 81 FLT1 97 FLT3 100 FLT3(D835H) 98 FLT3(D835Y) 100 FLT3(ITD) 94 FLT3(K663Q) 100 FLT3(N841I) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
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FGFR1 99 FGFR2 87 FGFR3 83 FGFR3(G697C) 100 FGFR4 98 FGR 81 FLT1 97 FLT3 100 FLT3(D835H) 98 FLT3(D835Y) 100 FLT3(ITD) 94 FLT3(K663Q) 100 FLT3(N841I) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
FGFR2 87 FGFR3 83 FGFR3(G697C) 100 FGFR4 98 FGR 81 FLT1 97 FLT3 100 FLT3(D835H) 98 FLT3(D835Y) 100 FLT3(ITD) 94 FLT3(K663Q) 100 FLT3(N841I) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
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FGFR4 98 FGR 81 FLT1 97 FLT3 100 FLT3(D835H) 98 FLT3(D835Y) 100 FLT3(ITD) 94 FLT3(K663Q) 100 FLT3(N841I) 100 FLT3(R834Q) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
FGR 81 FLT1 97 FLT3 100 FLT3(D835H) 98 FLT3(D835Y) 100 FLT3(ITD) 94 FLT3(K663Q) 100 FLT3(N841I) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
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FLT3(D835Y) 100 FLT3(ITD) 94 FLT3(K663Q) 100 FLT3(N841I) 100 FLT3(R834Q) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
FLT3(ITD) 94 FLT3(K663Q) 100 FLT3(N841I) 100 FLT3(R834Q) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
FLT3(K663Q) 100 FLT3(N841I) 100 FLT3(R834Q) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
FLT3(N841I) 100 FLT3(R834Q) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65	, ,	
FLT3(R834Q) 100 FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65	<u> </u>	
FLT3-autoinhibited 100 FLT4 93 FRK 100 FYN 90 GAK 65		
FLT4 93 FRK 100 FYN 90 GAK 65	·	
FRK 100 FYN 90 GAK 65		
FYN 90 GAK 65		
GAK 65		

GRK1	85
GRK4	90
GRK7	99
GSK3A	100
GSK3B	84
HASPIN	76
НСК	99
HIPK1	90
HIPK2	100
HIPK3	91
HIPK4	85
HPK1	100
HUNK	91
ICK	81
IGF1R	95
IKK-alpha	100
IKK-beta	98
IKK-epsilon	100
INSR	83
INSRR	100
IRAK1	100
IRAK3	98
IRAK4	100
ITK	88
JAK1(JH1domain-catalytic)	77
JAK1(JH2domain-pseudokinase)	99
JAK2(JH1domain-catalytic)	100
JAK3(JH1domain-catalytic)	100
JNK1	60
JNK2	78
JNK3	67
KIT	79
KIT(A829P)	100
KIT(D816H)	99
KIT(D816V)	76

1,	
KIT(L576P)	51
KIT(V559D)	82
KIT(V559D,T670I)	92
KIT(V559D,V654A)	94
KIT-autoinhibited	87
LATS1	78
LATS2	93
LCK	99
ШМК1	100
LIMK2	82
LKB1	100
LOK	100
LRRK2	11
LRRK2(G2019S)	6.5
LTK	59
LYN	100
LZK	92
MAK	98
MAP3K1	90
MAP3K15	100
MAP3K2	98
MAP3K3	100
MAP3K4	100
MAP4K2	97
MAP4K3	100
MAP4K4	100
MAP4K5	100
МАРКАРК2	95
МАРКАРК5	99
MARK1	97
MARK2	100
MARK3	84
MARK4	82
MAST1	100
MEK1	98
	1

MEK2	100
MEK3	79
MEK4	72
MEK5	92
MEK6	99
MELK	91
MERTK	66
MET	97
MET(M1250T)	80
MET(Y1235D)	100
MINK	94
МКК7	100
MKNK1	85
MKNK2	91
MLCK	88
MLK1	100
MLK2	100
MLK3	100
MRCKA	99
MRCKB	100
MST1	100
MST1R	100
MST2	80
MST3	91
MST4	77
MTOR	100
MUSK	100
MYLK	61
MYLK2	82
MYLK4	81
MYO3A	100
мүозв	85
NDR1	92
NDR2	100
NEK1	96

NEK10	100
NEK11	100
NEK2	96
NEK3	100
NEK4	90
NEK5	83
NEK6	100
NEK7	100
NEK9	100
NIK	62
NIM1	100
NLK	100
OSR1	88
p38-alpha	96
p38-beta	82
p38-delta	100
p38-gamma	95
PAK1	100
PAK2	100
PAK3	100
PAK4	100
PAK6	100
PAK7	83
PCTK1	93
PCTK2	88
PCTK3	100
PDGFRA	80
PDGFRB	78
PDPK1	86
PFCDPK1(P.falciparum)	96
PFPK5(P.falciparum)	100
PFTAIRE2	72
PFTK1	82
PHKG1	88
PHKG2	66

PIK3C2B	98
PIK3C2G	100
PIK3CA	87
PIK3CA(C420R)	95
PIK3CA(E542K)	100
PIK3CA(E545A)	99
PIK3CA(E545K)	100
PIK3CA(H1047L)	90
PIK3CA(H1047Y)	100
PIK3CA(I800L)	97
PIK3CA(M1043I)	91
PIK3CA(Q546K)	82
PIK3CB	86
PIK3CD	83
PIK3CG	100
PIK4CB	90
PIM1	98
PIM2	98
PIM3	99
PIP5K1A	84
PIP5K1C	100
PIP5K2B	62
PIP5K2C	100
PKAC-alpha	98
PKAC-beta	100
PKMYT1	100
PKN1	100
PKN2	100
PKNB(M.tuberculosis)	100
PLK1	89
PLK2	100
PLK3	100
PLK4	97
PRKCD	100
PRKCE	100

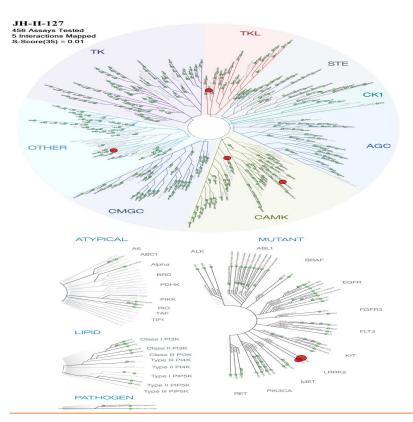
PRKCI 100 PRKCI 100 PRKCQ 93 PRKD1 85 PRKD2 98 PRKD3 100 PRKG1 72 PRKG2 99 PRKR 96 PRKX 100 PRP4 93 PYK2 100 QSK 86 RAF1 100 RET 96 RET(M918T) 96 RET(W804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.1-N-t		
PRKCQ 93 PRKD1 85 PRKD2 98 PRKD3 100 PRKG1 72 PRKG2 99 PRKR 96 PRKX 100 PRP4 93 PYK2 100 QSK 86 RAF1 100 RET 96 RET(M918T) 96 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 14 RPS6KA5(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.2-C-terminal) 88	PRKCH	
PRKD1 85 PRKD2 98 PRKD3 100 PRKG1 72 PRKG2 99 PRKR 96 PRKX 100 PRP4 93 PYK2 100 QSK 86 RAF1 100 RET (M918T) 96 RET(W904L) 92 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.2-C-terminal) 88	PRKCI	100
PRKD2 98 PRKD3 100 PRKG1 72 PRKG2 99 PRKR 96 PRKX 100 PRP4 93 PYK2 100 QSK 86 RAF1 100 RET 96 RET(M918T) 96 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 10 RPS6KA5(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.2-C-terminal) 88	PRKCQ	93
PRKD3 100 PRKG1 72 PRKG2 99 PRKR 96 PRKX 100 PRP4 93 PYK2 100 QSK 86 RAF1 100 RET 96 RET(M918T) 96 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.2-C-terminal) 88 RSK1(Kin.Dom.2-C-terminal) 88	PRKD1	85
PRKG1 72 PRKG2 99 PRKR 96 PRKX 100 PRP4 93 PYK2 100 QSK 86 RAF1 100 RET 96 RET(M918T) 96 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	PRKD2	98
PRKG2 99 PRKR 96 PRKX 100 PRP4 93 PYK2 100 QSK 86 RAF1 100 RET 96 RET(M918T) 96 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	PRKD3	100
PRKX 100 PRP4 93 PYK2 100 QSK 86 RAF1 100 RET 96 RET(M918T) 96 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	PRKG1	72
PRKX 100 PRP4 93 PYK2 100 QSK 86 RAF1 100 RET 96 RET(M918T) 96 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	PRKG2	99
PRP4 93 PYK2 100 QSK 86 RAF1 100 RET 96 RET(M918T) 96 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	PRKR	96
PYK2 100 QSK 86 RAF1 100 RET 96 RET(M918T) 96 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	PRKX	100
QSK 86 RAF1 100 RET 96 RET(M918T) 96 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	PRP4	93
RAF1 100 RET 96 RET(M918T) 96 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROCK1 88 ROCK2 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.1-N-terminal) 14 RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 69	PYK2	100
RET 96 RET(M918T) 96 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	QSK	86
RET(M918T) 96 RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RAF1	100
RET(V804L) 92 RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RET	96
RET(V804M) 100 RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RET(M918T)	96
RIOK1 91 RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RET(V804L)	92
RIOK2 100 RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RET(V804M)	100
RIOK3 67 RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RIOK1	91
RIPK1 76 RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.1-N-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RIOK2	100
RIPK2 96 RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RIOK3	67
RIPK4 82 RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RIPK1	76
RIPK5 100 ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RIPK2	96
ROCK1 88 ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RIPK4	82
ROCK2 93 ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RIPK5	100
ROS1 93 RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	ROCK1	88
RPS6KA4(Kin.Dom.1-N-terminal) 97 RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	ROCK2	93
RPS6KA4(Kin.Dom.2-C-terminal) 14 RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	ROS1	93
RPS6KA5(Kin.Dom.1-N-terminal) 100 RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RPS6KA4(Kin.Dom.1-N-terminal)	97
RPS6KA5(Kin.Dom.2-C-terminal) 78 RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RPS6KA4(Kin.Dom.2-C-terminal)	14
RSK1(Kin.Dom.1-N-terminal) 69 RSK1(Kin.Dom.2-C-terminal) 88	RPS6KA5(Kin.Dom.1-N-terminal)	100
RSK1(Kin.Dom.2-C-terminal) 88	RPS6KA5(Kin.Dom.2-C-terminal)	78
	RSK1(Kin.Dom.1-N-terminal)	69
RSK2(Kin.Dom.1-N-terminal) 85	RSK1(Kin.Dom.2-C-terminal)	88
	RSK2(Kin.Dom.1-N-terminal)	85

RSK2(Kin.Dom.2-C-terminal)	97		
	97		
RSK3(Kin.Dom.1-N-terminal)	100		
RSK3(Kin.Dom.2-C-terminal)	96		
RSK4(Kin.Dom.1-N-terminal)	93		
RSK4(Kin.Dom.2-C-terminal)	90		
S6K1	99		
SBK1	96		
SGK	100		
SgK110	74		
SGK2	100		
SGK3	100		
SIK	94		
SIK2	97		
SLK	96		
SNARK	100		
SNRK	99		
SRC	99		
SRMS	98		
SRPK1	76		
SRPK2	100		
SRPK3	100		
STK16	100		
STK33	42		
STK35	100		
STK36	94		
STK39	81		
SYK	80		
TAK1	91		
TAOK1	100		
TAOK2	85		
TAOK3	100		
TBK1	100		
TEC	100		
TESK1	100		
TGFBR1	99		

TGFBR2	80
TIE1	94
TIE2	97
TLK1	92
TLK2	98
TNIK	100
TNK1	100
TNK2	100
TNNI3K	97
TRKA	99
TRKB	100
TRKC	88
TRPM6	100
TSSK1B	100
TTK	13
TXK	100
TYK2(JH1domain-catalytic)	99
TYK2(JH2domain-pseudokinase)	100
TYRO3	68
ULK1	100
ULK2	100
ULK3	96
VEGFR2	91
VRK2	99
WEE1	100
WEE2	97
WNK1	95
WNK3	96
YANK1	92
YANK2	100
YANK3	100
YES	92
YSK1	89
YSK4	81
ZAK	100

ZAP70 97

S-Score Table JH-II-127



Compound Name	Selectivity Score Type	Number of Hits	Number of Non-Mutant Kinases	Screening Concentration (nM)	Selectivity Score
JH-II-127	S(35)	4	392	1000	0.01
JH-II-127	S(10)	0	392	1000	0
JH-II-127	S(1)	0	392	1000	0

HG-10-102-01 Vehicle 10 mg/Kg 50 mg/Kg 3 hrs 3hrs 6hrs 3hrs 6hrs 2 5 10 11 12 13 14 15 6 8 Brain αLRRK2 pS910 αLRRK2 pS935 αLRRK2 Spleen αLRRK2 pS910 αLRRK2 pS935 αLRRK2 **Kidney** αLRRK2 pS910

Formulation: 10% 1-methyl-2-pyrrolidinone (NMP) /90% PEG;

IP doses: 0, 10 (3 hrs), 10 (6hrs), 50 (3 hrs) and 50 (6hrs) mg/kg of HG-10-102-01;

αLRRK2 pS935

αLRRK2

IP injection volume: 200 µL/20g.

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