Synthesis and profiling of a 3-aminopyridin-2-one-based kinase targeted fragment library: Identification of 3-amino-5-(pyridin-4-yl)pyridin-2(*1H*)-one scaffold for monopolar spindle 1 (MPS1) and Aurora kinases inhibition

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Synthesis

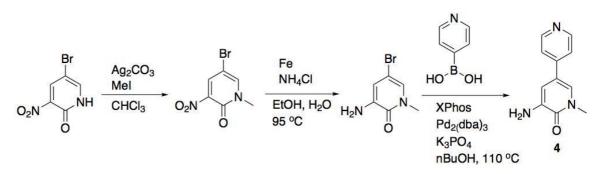


Figure 1: Synthesis of 1-methyl-3-amino-5-(pyridin-5-yl)pyridin-2-one 4

1-Methyl-3-nitro-5-bromopyridin-2-one

Silver carbonate (4.09 g, 14.84 mmol, 1.3 equivalents) and methyl iodide (7.11 mL, 114 mmol, 10 equivalents) were added to a covered flask containing 5-bromo-3nitropyridin-2-one (2.5 g, 11.42 mmol, 1 equivalent) and CHCl₃ (40 mL) and the reaction allowed to stir at ambient temperature for 2 days. The solution was then filtered through a pad of celite (5 g) and concentrated, giving a yellow solid which was purified by column chromatography (20% EtOAc in hexane, followed by DCM:EtOAc 1:1) to give 1.11 g (42%) of 1-methyl-3-nitro-5-bromopyridin-2-one as an orange solid. Rf = 0.33 (DCM:EtOAc 1:1). mp: 124 °C. ¹H NMR (500 MHz, CDCl₃) 3.70 (3H, s), 7.84 (1H, d, J = 2.8 Hz), 8.39 (1H, d, J = 2.8 Hz). ¹³C (126 MHz, CDCl₃) 153.4 (C), 144.6 (CH), 141.2 (CH), 99.9 (C), 94.1 (C), 39.0 (CH₃). HRMS: Found 232.9558 and 234.9539, calculated for C₆H₆⁷⁹BrN₂O₃ (M+H)⁺: 234.9536. v_{max} cm⁻¹ 1344 (NO₂), 1594 (C=O).

1-Methyl-3-amino-5-bromopyridin-2-one

Iron powder (1 g, 18 mmol, 4.2 equivalents) and ammonium chloride (0.964 g, 18 mmol, 4.2 equivalents) were added to a solution of 1-methyl-3-nitro-5bromo-pyridin-2-one (1 g, 4.3 mmol, 1 equivalent) in ethanol (8 mL) and water (6 mL) and stirred at 95 °C for 3 hours. The solution was then filtered through a pad of celite and concentrated. The residue was then taken up in water (20 mL) and extracted with EtOAc (3 x 30 mL). The organic layers were dried over magnesium sulfate and concentrated to give a brown oil that solidified upon standing. The residue was purified by SCX to give the desired product as a dark red solid (300 mg, 34%). R_f = 0.5 (DCM:EtOAc 1:1). mp: 200 °C. ¹H NMR (500 MHz, CDCl₃) 3.55 (3H, s), 4.37 (2H, s), 6.57 (1H, d, J = 2.4 Hz), 6.83 (1H, d, J = 2.4 Hz).¹³C NMR (126 MHz, DMSO-d₆) 156.5 (C), 139.8 (C), 124.2 (CH), 111.9 (CH), 98.7 (C), 37.3 (CH₃). HRMS: Found 202.9815 and 204.9794, calculated for $C_6H_8^{79}BrN_2O$ (M+H)⁺: 202.9815, calculated for $C_6H_8^{81}BrN_2O$ (M+H)⁺: 204.9794. ν_{max} cm⁻¹ 3314 (NH₂), 1568 (C=O).

1-Methyl-3-amino-5-(pyridin-5-yl)pyridin-2-one (4)

4-Pyridyl boronic acid (73 mg, 0.591 mmol, 1.2 equivalents), 2-dicyclohexylphosphino-2,4,6-triisopropylbiphenyl (47 mg, 0.99 mmol, 0.2 equivalents), potassium phosphate (315 mg, 1.478 mmol, 3 equivalents) and tris(dibenzylideneacetone) dipalladium(0) (23 mg, 0.025 mmol, 0.05 equivalents) were added to 1-methyl-3-amino-5-bromopyridin-2-one (100 mg, 0.493 mmol, 1 equivalent) in butan-1-ol (5 mL) and the reaction mixture was heated to 110 °C for 3 hours. The solution was then cooled and filtered through a pad of celite, which was washed with MeOH. The solution was concentrated and the residue taken up in MeOH and filtered through a SCX column and then purified by column chromatography (5% MeOH in CHCl₃) to give a brown solid (45 mg, 46%). R_f = 0.67 (DCM:EtOAc 1:1). mp: 236 °C. ¹H NMR (500 MHz, DMSO-d₆) 3.53 (3H, s), 5.31 (2H, s), 6.87 (1H, d, J = 2.5 Hz), 7.50 (2H, dd, J = 1.5 Hz, 4.7 Hz), 7.62 (1H, d, J = 2.5 Hz), 8.54 (2H, dd, J = 1.5 Hz, 4.7 Hz).¹³C NMR (126 MHz, DMSO-d₆) 157.5 (C), 150.5 (CH), 144.9 (C), 138.8 (C), 124.9 (CH), 119.9 (CH), 116.0 (C), 107.6 (CH), 37.6 (CH₃). HRMS: Found 202.0975, calculated for C₁₁H₁₂N₃O (M+H)⁺: 202.0975.

3-Amino-5-(quinolin-4-yl)pyridin-2(1H)-one (6)

5-Bromo-2-methoxypyridin-3-amine (100 mg, 0.49 mmol, 1 equivalent), quinoline-4boronic acid (102 mg, 0.59 mmol, 1.2 equivalents), potassium phosphate (314 mg, 1.48 mmol, 3 equivalents), 2-dicyclohexylphosphino-2,4,6-triisopropylbiphenyl (47 mg, 0.10 mmol, 0.2 equivalents) and tris(dibenzylideneacetone)dipalladium(0) (22.55 mg, 0.025 mmol, 0.05 equivalents) were dissolved in *n*-butanol (4 mL) and stirred at 110 °C for 3 hours. The solution was then cooled and filtered through a pad of celite, which was then washed with methanol. The solution was concentrated and purified by SCX column (eluting at room temperature with 2 M ammonia in methanol) and concentrated. The residue was then dissolved in acetonitrile and sodium iodide (222 mg, 1.484 mmol, 3 equivalents) was added followed by dropwise addition of trimethylsilyl chloride (0.190 mL, 1.484 mmol, 3 equivalents) and the reaction mixture stirred for 16 hours. The solution was concentrated, taken up in methanol and purified by SCX column (eluting at room temperature with 2 M ammonia in methanol) followed by column chromatography (5% MeOH in EtOAc) to give the product as a grey solid (33 mg, 24%). $R_f = 0.31$ (5% MeOH in EtOAc). mp: 191 °C. ¹H NMR (500 MHz, DMSO-d₆) 5.30 (2H, s), 6.66 (1H, d, J = 2.3 Hz), 6.86 (1H, d, J = 2.3 Hz), 7.40 (1H, d, J = 4.4 Hz), 7.62 (1H, ddd, J = 1.4 Hz, 6.8 Hz, 8.4 Hz), 7.78 (1H, ddd, J = 1.4 Hz, 6.8 Hz, 8.4 Hz), 8.00 - 8.15 (2H, m), 8.87 (1H, d, J = 4.4 Hz), 11.72 (1H, s).¹³C NMR (126 MHz, DMSO-d₆) 157.6 (C), 150.7 (CH), 148.7 (C), 145.1 (C), 139.1 (C), 130.0 (CH), 129.8 (CH), 127.2 (CH), 126.6 (C), 125.9 (CH), 121.4 (CH), 120.7 (CH), 116.7 (C), 112.1 (CH). HRMS: Found 238.0980, calculated for C₁₄H₁₂N₃O (M+H)⁺: 238.0975.

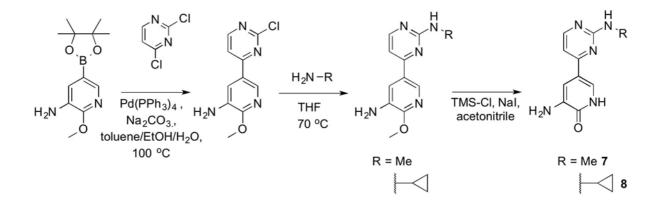


Figure 2: Synthesis of 3-amino-5-(2-(methylamino)pyrimidin-4-yl)pyridin-2(1H)-one **7** and 3-amino-5-(2-(cyclopropylamino)pyrimidin-4-yl)pyridin-2(1H)-one **8**

2-Methoxy-3-amino-5-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)pyridine

5-Bromo-2-methoxypyridin-3-amine (2.0 g, 9.85 mmol, 1 equivalent) was added to dioxane followed by bis(pinacolato)diboron (5.0 g, 19.7 mmol, 2 equivalents), potassium acetate (2.9 g, 29.6 mmol, 3 equivalents) and (1,1'-bis(diphenyl-phosphosphinol)ferrocene)dichloropalladium(II) dichloromethane adduct (0.804 g, 0.985 mmol, 0.1 equivalents) and stirred at 80 °C for 16 hours under argon. After cooling the solution was diluted with water and the product extracted with EtOAc. The combined organic layers were washed with brine, dried over magnesium sulfate and concentrated. The product was then purified by column chromotography (DCM:EtOAc 1:1) and concentrated. The residue was triturated with diethyl ether to give the product as a yellow solid (950 mg, 39%). $R_f = 0.73$ (DCM:EtOAc 1:1).

mp: 137 °C. ¹H NMR (500 MHz, DMSO-d₆) 1.27 (12H, s), 3.87 (3H, s), 4.92 (2H, s), 7.12 (1H, d, J = 1.7 Hz), 7.64 (1H, d, J = 1.7 Hz).¹³C NMR (126 MHz, DMSO-d₆) 154.6 (C), 139.9 (CH), 132.4 (C), 124.2 (CH), 83.9 (C), 53.4 (CH₃), 25.1 (CH₃). HRMS: Found 251.1566, calculated for $C_{12}H_{20}BN_2O_3$ (M+H)⁺: 251.1564.

2-Methoxy-3-amino-5-((2-chloro)pyrimidin-4-yl)pyridine

2-Methoxy-3-amino-5-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)pyridine (500 mg, 2.0 mmol, 1 equivalent) was dissolved in toluene (20 mL) and ethanol (5 mL) and then 2,4-dichloropyrimidine (357 mg, 2.40 mmol, 1.2 equivalents) was added followed by tetrakis(triphenylphosphine)palladium(0) (116 mg, 0.1 mmol, 0.05 equivalents). Sodium carbonate (1.46 g, 13.79 mmol, 6.9 equivalents) and water (7.5 mL) were then added and the reaction mixture heated to reflux for 2 hours under argon. After cooling the solution was diluted with 15 mL water and extracted with EtOAc (3x50 mL) and the organic layers collected and washed with brine. The organic layers were dried over magnesium sulfate and concentrated. The residue was purified using an SCX column, eluting at room temperature with 2 M ammo-nia in methanol, and triturated with diethyl ether to give the desired product as a grey solid (413 mg, 73%). R_f = 0.47 (5% MeOH in EtOAc). ¹H NMR (500 MHz, DMSO-d₆) 3.95 (3H, s), 5.32 (2H, s), 7.65 (1H, d, J = 1.8 Hz), 8.02 (1H, d, J = 5.3 Hz), 8.24 (1H, d, J = 1.8 Hz), 8.72 (1H, d, J = 5.3 Hz). HRMS: Found 238.0563, calculated for C₁₀H₁₀CIN₄O (M+H)⁺: 238.0564.

2-Methoxy-3-amino-5-((2-methylamino)pyrimidin-4-yl)pyridine

2-Methoxy-3-amino-5-((2-chloro)pyrimidin-4-yl)pyridine (80 mg, 0.338 mmol, 1 equivalent) was added to 2 M methylamine in THF (3.4 mL) and the reaction mixture stirred in a sealed tube at 70 °C for 24 hours. The solvent was then removed under reduced pressure and the product purified by SCX column (eluting at room temperature with 2 M ammonia in methanol) to give the desired product as a pale yellow solid (50 mg, 64%). $R_f = 0.47$ (1:1 DCM: EtOAc). ¹H NMR (500 MHz, CDCl₃) 3.07 (3H, d, J = 5.2 Hz), 3.91 (2H, brs), 4.06 (3H, s), 5.15 (1H, d, J = 5.2 Hz), 6.90 (1H, d, J = 5.2 Hz), 7.61 (1H, s), 8.23 (1H, s), 8.31 (1H, d, J = 5.2 Hz). ¹³C NMR (126 MHz, CDCl₃) 162.9 (C), 158.2 (CH), 149.8 (C), 134.6 (CH), 123.9 (C), 121.6 (C), 118.1 (CH), 105.7 (CH), 100.0 (C), 53.3 (CH₃), 28.3 (CH₃). HRMS: Found 232.1190, calculated for C₁₁H₁₄N₅O (M+H)⁺: 232.1198.

3-Amino-5-(2-(methylamino)pyrimidin-4-yl)pyridin-2(1H)-one (7)

Methylamine in THF (2 M, 5 mL) was added to 2-methoxy-3-amino-5-((2-chloro) pyrimidin-4-yl)pyridine (80 mg, 0.34 mmol, 1 equivalent) and stirred overnight at 70 °C. The solvent was then removed and the residue taken up in methanol and purified by SCX (eluted with 2 M ammonia in methanol). The residue was then taken up in acetonitrile (2 mL) and sodium iodide (162 mg, 1.08 mmol, 3.2 equivalents) was added followed by dropwise addition of trimethylsilyl chloride (0.138 mL, 1.08 mmol, 3.2 equivalents). After stirring for 2 hours the solvent was removed and the residue purified by semiprep HPLC (Method A) to give the product as a pale yellow solid (12 mg, 16%). R_f = 0.80 (10% MeOH in CHCl₃). mp: 278 °C. ¹H NMR (500 MHz, DMSO-d₆) 2.83 (3H, d, J = 4.4 Hz), 5.19 (2H, s), 6.84 (1H, d, J = 4.6 Hz), 6.93 (1H, d, J = 4.5 Hz), 7.16 (1H, s), 7.52 (1H, s), 8.19 (1H, d, J = 4.6 Hz), 11.53 (1H, brs).¹³C NMR (126 MHz, DMSO-d₆) 186.6 (C), 163.1 (C), 158.6 (CH), 158.4 (C), 138.7 (C), 121.6 (CH), 117.1 (C), 108.3 (CH), 100.6 (CH), 28.4 (CH₃). HRMS: Found 218.1037, calculated for C₁₀H₁₂N₅O (M+H)⁺: 218.1036.

2-Methoxy-3-amino-5-((2-cyclopropylamino)pyrimidin-4-yl)pyridine

Cyclopropylamine (0.7 mL, 10.1 mmol, 20 equivalents) was added to 2-methoxy-3amino-5-((2-chloro)pyrimidin-4-yl)pyridine (120 mg, 0.51 mmol, 1 equivalent) in THF (3.5 mL) and stirred at 70 °C for 16 hours. The solvent was then removed under reduced pressure to give an orange oil that was purified by SCX column (eluted with 2 M ammonia in methanol) to give a pale yellow solid (60 mg, 46%) R_f =0.47 (DCM:EtOAc 1:1). mp: 230 °C. ¹H NMR (500 MHz, DMSO-d₆) 0.45 - 0.52 (2H, m), 0.63 - 0.75 (2H, m), 2.77 (1H, m), 5.13 (2H, s), 7.02 (1H, d, J = 5.2 Hz), 7.31 (1H, d, J = 3.5 Hz), 7.55 (1H, brs), 8.11 (1H, brs), 8.29 (1H, d, J = 5.0 Hz).¹³C NMR (126 MHz, DMSO-d₆) 163.9 (C), 162.5 (C), 158.8 (CH), 154.0 (C), 132.8 (C), 132.4 (CH), 127.4 (C), 116.8 (CH), 106.0 (CH), 53.7 (CH₃), 24.4 (CH), 6.9 (CH₂). HRMS: Found 259.1381, calculated for $C_{13}H_{16}N_5O$ (M+H)⁺: 259.1376.

3-Amino-5-(2-(cyclopropylamino)pyrimidin-4-yl)pyridin-2(1H)-one (8)

Trimethylsilyl chloride (0.15 mL, 1.17 mmol, 5 equivalents) was added dropwise to a solution of 2-methoxy-3-amino-5-((2-cyclopropylamino)pyrimidin-4-yl)pyridine (60 mg, 0.23 mmol, 1 equivalent) and sodium iodide (175 mg, 1.17 mmol, 5 equiv-alents) in acetonitrile (2.5 mL) and stirred for 2 hours at room temperature. The solvent was

then removed under reduced pressure and the product purified by semiprep HPLC (Method A) to give a pale yellow solid (28 mg, 49%). $R_f = 0.51$ (10% MeOH in CHCl₃). mp: 218 °C. ¹H NMR (500 MHz, DMSO-d₆) 0.48 (2H, m), 0.67 (2H, m), 2.70 - 2.78 (1H, m), 5.19 (2H, s), 6.89 (1H, d, J = 5.3 Hz), 7.16 (1H, s), 7.22 (1H, d, J = 3.5 Hz), 7.53 (1H, s), 8.22 (1H, d, J = 5.3 Hz), 11.73 (1H, s).¹³C NMR (126 MHz, DMSO-d₆) 163.7 (C), 162.0 (C), 158.6 (CH), 158.4 (C), 138.7 (C), 121.7 (CH), 117.0 (C), 108.3 (CH), 104.9 (CH), 24.3 (CH), 6.9 (CH₂). HRMS: Found 244.1192, calculated for C₁₂H₁₄N₅O (M+H)⁺: 244.1193.

3-Amino-5-(pyridin-3-yl)pyridin-2(1H)-one (11)

Prepared using general procedure B to give a grey solid (24 mg, 65%): 6-Methoxy-3,3'-bipyridin-5-amine (40 mg, 0.199 mmol, 1 equivalent), trimethylsilyl chloride (0.994 mL, 0.994 mmol, 5 equivalents) and sodium iodide (149 mg, 0.994 mmol, 5 equivalents) in acetonitrile (1 mL). $R_f = 0.2$ (5% MeoOH in EtOAc). mp: 223 °C. ¹H NMR (500 MHz, DMSO-d₆) 5.22 (2H, s), 6.81 (1H, d, J = 2.4 Hz), 7.06 (1H, d, J = 2.4 Hz), 7.40 (1H, ddd, J = 0.8 Hz, 4.8 Hz, 8.0 Hz), 7.87 (1H, ddd, J = 1.6 Hz, 2.4 Hz, 8.0 Hz), 8.46 (1H, dd, J = 1.6 Hz, 4.7 Hz), 8.71 (1H, dd, J = 0.8 Hz, 2.4 Hz).¹³C NMR (126 MHz, DMSO-d₆) 157.7 (C), 147.9 (CH), 146.9 (CH), 139.5 (C), 133.6 (C), 133.1 (CH), 124.2 (C), 118.7 (CH), 116.5 (C), 109.5 (CH). HRMS: Found 188.0816, calculated for $C_{10}H_{10}N_3O$ (M+H)⁺: 188.0824.

5-(4-((Dimethylamino)methyl)phenyl)-2-methoxypyridin-3-amine

2-Methoxy-3-amino-5-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)pyridine (200 mg, 0.80 mmol, 1 equivalent) was dissolved in toluene (8 mL) and ethanol (2 mL) and then 1-(4-bromophenyl)-N,N-dimethylmethylamine (205 mg, 0.96 mmol, 1.2 equiv-alents) was added followed by tetrakis(triphenylphosphine)palladium(0) (46.2 mg, 0.04 mmol, 0.05 equivalents). Sodium carbonate (585 mg, 5.52 mmol, 6.9 equivalents) and water (3.5 mL) were then added and the reaction mixture heated to reflux for 2 hours under argon. After cooling the solution was diluted with water (15 mL) and extracted with EtOAc (3x50 mL) and the organic layers collected and washed with brine. The organic layers were then dried over magnesium sulfate and concentrated. The residue was purified using an SCX column (eluted with 2 M ammonia in methanol) to give a grey solid (147 mg, 71.4%). $R_f = 0.24$ (5%

MeOH in EtOAc). ¹H NMR (500 MHz, CD₃OD) 2.86 (6H, s), 4.02 (3H, s), 4.31 (2H, s), 7.27 (1H, d, J = 2.2 Hz), 7.56 (2H, d, J = 8.3 Hz), 7.69 (2H, d, J = 8.3 Hz), 7.72 (1H, d, J = 2.2 Hz).¹³C NMR (126 MHz, CD₃OD) 152.8 (C), 140.0, 132.2 (C), 131.4 (CH), 131.0 (CH), 129.8 (C), 127.0 (CH), 118.1 (CH), 113.1 (C), 60.7 (CH₂), 52.6 (CH₃), 41.7 (CH₃). HRMS: Found 258.1598, calculated for $C_{15}H_{20}N_{3}O$ (M+H)⁺: 258.1601.

3-Amino-5-(4((dimethylamino)methyl)phenyl)aminopyridin-2-one (12)

Prepared using general procedure B to give a grey solid (18 mg, 64%): 5-(4-((dimethylamino)methyl)phenyl)-2-methoxypyridin-3-amine (30 mg, 0.12 mmol, 1 equivalent), trimethylsilyl chloride (0.075 mL, 0.58 mmol, 5 equivalents) and sodium iodide (87 mg, 0.58 mmol, 5 equivalents) in acetonitrile (1 mL) and THF (1 mL). R_f = 0.22 (10% MeOH in CHCl₃). mp: 251 °C. ¹H NMR (500 MHz, DMSO-d₆) 2.15 (6H, s), 3.37 (2H, s), 5.16 (2H, s), 6.81 (1H, d, J = 2.4 Hz), 6.93 (1H, d, J = 2.4 Hz), 7.29 (2H, d, J = 8.3 Hz), 7.41 (2H, d, J = 8.2 Hz), 11.53 (1H, s).¹³C NMR (126 MHz, DMSO-d₆) 157.5 (C), 139.2 (C), 137.7 (C), 136.6 (C), 129.7 (CH), 125.4 (CH), 119.4 (C), 117.7 (CH), 110.2 (CH), 63.5 (CH₂), 45.47 (CH₃). HRMS: Found 244.1443, calculated for C₁₄H₁₈N₃O (M+H)⁺: 244.1444.

6-Methyl-2-oxo-5-(pyridin-4-yl)-1,2-dihydropyridine-3-carboxamide (14)

Concentrated sulfuric acid (350 μ L, 6.3 mmol, 14 equivalents) was added to 6methyl-2-oxo-5-(pyridin-4-yl)-1,2-dihydropyridine-3-carbonitrile (95 mg, 0.450 mmol, 1 equivalent), and the solution was heated to 120 °C for 15 minutes. The solution was then cooled to 0 °C and diluted with water, whereupon concentrated ammonium hydroxide was added to bring the pH to 10. The resulting precipitate was filtered and dried to give the desired product as a white solid (74 mg, 72%). R_f = 0.53 (DCM:EtOAc 1:1). mp: 286 °C. ¹H NMR (500 MHz, DMSO-d₆) 2.34 (3H, s), 7.42 (2H, dd, J = 1.5 Hz, 4.5 Hz), 7.60 (1H, d, J = 3.8 Hz), 8.21 (1H, s), 8.62 (2H, dd, J = 1.5 Hz, 4.5 Hz), 8.97 (1H, d, J = 3.8 Hz), 11.09 (1H, s). ¹³C NMR (126 MHz, DMSO-d₆) 164.9 (C), 162.6 (C), 150.3 (CH), 149.3 (C), 145.3 (CH), 145.1 (C), 124.4 (CH), 118.6 (C), 116.6 (C), 18.2 (CH₃). HRMS: Found 230.0923, calculated for C₁₂H₁₂N₃O₂ (M+H)⁺: 230.0924.

3-Amino-5-(pyridin-4-yl)-6-methylpyridin-2-one (15)

6-Methyl-2-oxo-5-(pyridin-4-yl)-1,2-dihydropyridine-3-carboxamide (60 mg, 0.262 mmol, 1 equivalent) was dissolved in 1 M NaOH (1.5 mL) with heating and then cooled to 0 °C. Bromine (16 μL, 0.314 mmol, 1.2 equivalents) was then added dropwise and the reaction mixture heated to 120 °C for 3 hours then cooled. The solution was then adjusted to pH = 7 with HCl (1 M) and cooled to 0 °C for 2 hours. The precipitate was then filtered and washed with water to give the desired product as an off-white solid (12 mg, 23%). R_f = 0.62 (5% MeOH in EtOAc). mp: 262 °C. ¹H NMR (500 MHz, DMSO-d₆) 2.12 (3H, s), 4.97 (2H, s), 6.47 (1H, s), 7.30 (2H, d, J = 5.2 Hz), 8.55 (2H, brs), 11.56 (1H, brs).¹³C NMR (126 MHz, DMSO-d₆) 158.3 (C), 150.0 (NCHCH), 147.2 (C), 137.1 (C), 127.8 (C), 124.3 (CH), 115.4 (C), 113.1 (CH), 16.8 (CH₃). HRMS: Found 202.0985, calculated for C₁₁H₁₂N₃O (M+H)⁺: 202.0975.

3,5-Difluoro-N-(2-oxo-5-(pyridin-4-yl)-1,2-dihydropyridin-3-yl)benzamide (20)

Prepared using general procedure C to give a yellow solid (48 mg, 54%): 3,5difluorobenzoylchloride (0.031 mL, 0.27 mmol, 1 equivalent), 3-amino-5-(pyridin-4yl)pyridin-2-one (50 mg, 0.27 mmol, 1 equivalent). $R_f = 0.51$ (DCM:EtOAc 1:1). mp: 293 °C. ¹H NMR (500 MHz, DMSO-d₆) 7.56 (1H, tt, J = 2.1 Hz, 9.0 Hz), 7.65 - 7.72 (2H, m), 8.21 (2H, d, J = 6.7 Hz), 8.27 (1H, s), 8.76 (1H, d, J = 2.6 Hz), 8.83 (2H, d, J = 6.6 Hz), 9.77 (1H, s), 13.00 (1H, s). ¹³C NMR (126 MHz, DMSO-d₆) 163.8 (C), 162.7 (dd, J = 247.8 Hz, 12.8 Hz, CF), 158.1 (C), 152.0 (C), 142.9 (CH), 137.8 (t, J = 8.6 Hz, C), 133.4 (CH), 129.2 (C), 124.6 (CH), 121.9 (CH), 112.9 (C), 111.6 (dd, J = 20.4 Hz, 6.5 Hz, CH), 108.0 (t, J = 25.9 Hz, C). HRMS: Found 328.0893, calculated for C₁₇H₁₂N₃O₂F₂ (M+H)⁺: 328.0892.

N-(2-Oxo-5-(pyridin-4-yl)-1,2-dihydropyridin-3-yl)benzenesulfonamide (**21**)

Prepared using general procedure C to give a yellow solid (18 mg, 21%): benzenesulfonylchloride (0.034 mL, 0.27 mmol, 1 equivalent), 3-amino-5-(pyridin-4yl)pyridin-2-one (50 mg, 0.27 mmol, 1 equivalent). R_f = 0.31 (5% MeOH in EtOAc). mp: 289 °C. ¹H NMR (500 MHz, DMSO-d₆) 7.52 (2H, dd, J = 1.6 Hz, 4.6 Hz), 7.56 (2H, t, J = 7.6 Hz), 7.61 - 7.66 (1H, m), 7.74 (1H, d, J = 2.5 Hz), 7.76 (1H, d, J = 2.5 Hz), 7.87 - 7.93 (2H, m), 8.57 (2H, dd, J = 1.6 Hz, 4.6 Hz), 9.76 (1H, s), 12.34 (1H, s).¹³C NMR (126 MHz, DMSO-d₆) 157.8 (C), 150.6, 150.0 (CH), 143.8 (C), 140.3 (C), 133.6 (CH), 129.7 (CH), 128.6 (C), 127.3 (CH), 124.5 (CH), 124.4 (CH), 120.0 (CH). HRMS: Found: 328.0750, calculated for $C_{16}H_{14}N_3O_3S (M+H)^+$: 328.0750.

4-Bromo-N-(2-oxo-5-(pyridin-4-yl)-1,2-dihydropyridin-3-yl)benzamide

Prepared using general procedure C to give a white solid (255 mg, 52%): 4-bromobenzoyl chloride (293 mg, 1.34 mmol, 1 equivalent), 3-amino-5-(pyridin-4-yl)pyridin-2-one (250 mg, 1.34 mmol, 1 equivalent). $R_f = 0.58$ (DCM:EtOAc 1:1). mp: 294 °C. ¹H NMR (500 MHz, DMSO-d₆) 7.78 (2H, d, J = 8.6 Hz), 7.91 (2H, d, J = 8.6 Hz), 8.13 (2H, d, J = 6.8 Hz), 8.19 (1H, s), 8.79 (2H, d, J = 6.8 Hz), 8.82 (1H, d, J = 2.6 Hz), 9.59 (1H, s), 12.94 (1H, d, J = 2.7 Hz).¹³C NMR (126 MHz, DMSO-d₆) 165.1 (C), 158.1 (C), 151.3 (C), 143.8 (CH), 133.3 (C), 132.3 (CH), 132.0 (CH), 130.0 (CH), 129.5 (C), 126.5 (C), 123.1 (CH), 121.7 (CH), 113.4 (C).

4-(4-Methylpiperazin-1-yl)-N-(2-oxo-5-(pyridin-4-yl)-1,2-dihydropyridin-3-yl)benzamide (22)

General Procedure D: 4-Bromo-*N*-(2-oxo-5-(pyridin-4-yl)-1,2-dihydropyridin-3-yl) benzamide (100 mg, 0.27 mmol, 1 equivalent) and 1-methylpiperazine (3 mL, 27 mmol, 100 equivalents) were stirred in *N*-methyl-2-pyrrolidinone (1.5 mL) at 200 °C in a microwave reactor for 6 hours. The excess 1-methylpiperazine was then removed under reduced pressure and the product filtered and washed with methanol to give a white solid (38 mg, 36%). R_f = 0.70 (DCM:EtOAc 1:1). mp: 280 °C. ¹H NMR (500 MHz, DMSO-d₆) 2.23 (3H, s), 2.42 - 2.47 (4H, m), 3.26 - 3.32 (4H, m), 7.05 (2H, d, J = 9.0 Hz), 7.60 (2H, dd, J = 1.6 Hz, 4.6 Hz), 7.77 (1H, d, J = 2.5 Hz), 7.80 (2H, d, J = 9.0 Hz), 8.58 (2H, dd, J = 1.6 Hz, 4.5 Hz), 8.77 (1H, d, J = 2.5 Hz), 9.15 (1H, s), 12.54 (1H, s). ¹³C NMR (126 MHz, DMSO-d₆) 165.2 (C), 157.9 (C), 153.9 (C), 150.7 (CH), 144.3 (C), 129.9 (C), 129.1 (CH), 127.1 (CH) , 122.6 (C), 121.3 (CH), 120.3 (CH), 115.8 (C), 114.2 (CH), 54.7 (CH₂), 47.2 (CH₂), 46.2 (CH₃). HRMS: Found: 390.1925, calculated for C₂₂H₂₄N₅O₂ (M+H)⁺: 390.1917.

N-(2-Oxo-5-(pyridin-4-yl)-1,2-dihydropyridin-3-yl)-4-piperidin-1-yl)benzamide (23)

Prepared using general procedure D to give a white solid (26 mg, 26%): 4-bromo-*N*-(2-oxo-5-(pyridin-4-yl)-1,2-dihydropyridin-3-yl)benzamide (100 mg, 0.27 mmol, 1 equivalent), piperidine (2.67 mL, 27 mmol, 100 equivalents). $R_f = 0.71$ (DCM: EtOAc 1:1). mp: 284 °C. ¹H NMR (500 MHz, DMSO-d₆) 1.59 (6H, brs), 3.34 (4H, brs), 7.01 (2H, d, J = 8.9 Hz), 7.60 (2H, d, J = 6.1 Hz), 7.76 (1H, d, J = 2.5 Hz), 7.77 (2H, d, J = 9.0 Hz), 8.58 (2H, d, J = 6.0 Hz), 8.77 (1H, d, J = 2.5 Hz), 9.12 (1H, s), 12.54 (1H, s).¹³C NMR (126 MHz, DMSO-d₆) 165.1 (C), 157.8 (C), 153.9 (C), 150.7 (CH), 144.2 (C), 130.0 (C), 129.1 (CH), 126.9 (CH), 121.7 (C), 120.9 (CH), 120.2

(CH), 115.8 (C), 114.1 (CH), 48.4 (CH₂), 25.3 (CH₂), 24.4 (CH₂). HRMS: Found: 375.1799, calculated for $C_{22}H_{23}N_4O_2$ (M+H)⁺: 375.1816.

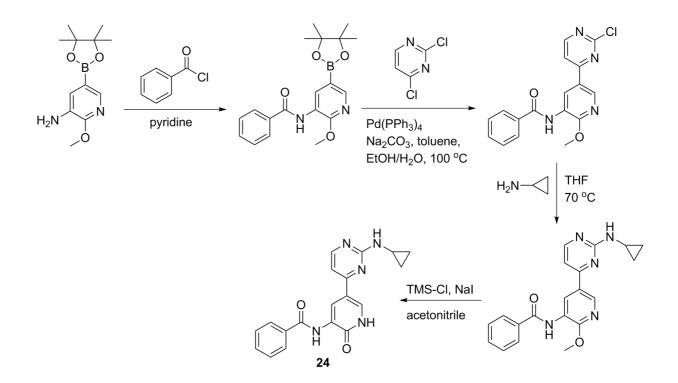


Figure 3: Synthesis of *N*-(5-(2-(cyclopropylamino)pyrimidin-4-yl)-2-oxo-1,2-dihydropyridin-3-yl)benzamide (**24**).

N-(2-Methoxy-5-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)pyridin-3-yl)benzamide

Benzoyl chloride (0.058 mL, 0.5 mmol, 1 equivalent) was added to a solution of 2methoxy-3-amino-5-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)pyridine (125 mg, 0.5 mmol, 1 equivalent) in pyridine (3.5 mL) and stirred for 16 hours. The solvent was then removed under reduced pressure and the residue was taken up methanol and filtered to give the product as a white solid (87 mg, 49%). $R_f = 0.54$ (DCM:EtOAc 1:1). mp: 251 °C. ¹H NMR (500 MHz, CDCl₃) 1.35 (12H, s), 4.10 (3H, s), 7.49 - 7.62 (3H, m), 7.91 (2H, d, J = 8.0 Hz), 8.28 (1H, s), 8.37 (1H, s), 9.06 (1H, s).¹³C (126 MHz, CDCl₃) 165.5 (C), 155.2 (C), 147.5 (CH), 134.7 (C), 132.3 (CH), 132.0 (CH), 128.9 (CH), 127.0 (CH), 122.4 (C), 83.9 (C), 54.2 (CH₃), 24.9 (CH₃). HRMS: Found 377.1640, calculated for C₁₉H₂₃N₂O₄BNa (M+Na)⁺: 377.1647.

N-(5-(2-Chloropyrimidin-4-yl)-2-methoxypyridin-3-yl)benzamide

N-(2-Methoxy-5-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)pyridin-3-yl)benzamide (175 mg, 0.494 mmol, 1 equivalent), 2,4-dichloropyrimidine (88 mg, 0.593 mmol,

1.2 equivalents) and tetrakis(triphenylphosphine)palladium(0) (28.5 mg, 0.025 mmol,

0.05 equivalents) were added to toluene (8 mL) and ethanol (2 mL) followed by addition of sodium carbonate (361 mg, 3.41 mmol, 6.9 equivalents) and water (3 mL). The reaction mixture was then heated to reflux for 2 hours and the solvent removed under reduced pressure and the residue purified by column chromatography (20% EtOAc in DCM) to give a pale yellow solid (80 mg, 48%). $R_f = 0.71$ (20% EtOAc in DCM). mp: >300 °C. ¹H NMR (500 MHz, CDCl₃) 4.18 (3H, s), 7.53 - 7.60 (2H, m), 7.60 - 7.65 (1H, m), 7.68 (1H, d, J = 5.3 Hz), 7.91 - 7.96 (2H, m), 8.48 (1H, s), 8.65 (1H, d, J = 5.3 Hz), 8.75 (1H, d, J = 2.3 Hz), 9.44 (1H, d, J = 2.3 Hz).¹³C NMR (126 MHz, CDCl₃) 168.6 (C), 159.9 (CH), 140.7 (CH), 132.6 (CH), 129.0 (CH), 127.2 (CH), 125.3 (C), 124.4 (CH), 123.2 (C), 114.7 (CH), 54.7 (CH₃). HRMS: Found 341.0805, calculated for $C_{17}H_{14}N_4O_2CI (M+H)^+$: 341.0800.

N-(5-(2-(Cyclopropylamino)pyrimidin-4-yl)-2-methoxypyridin-3-yl)benzamide

Prepared by general procedure D to give a yellow solid (19 mg, 51%): *N*-(5-(2-chloropyrimidin-4-yl)-2-methoxypyridin-3-yl)benzamide (35 mg, 0.10 mmol, 1 equivalent), cyclopropylamine (0.142 mL, 2.05 mmol, 20 equivalents) and THF (1 mL). The reaction mixture was stirred in a sealed vial at 70 °C for 24 hours. $R_f = 0.52$ (DCM:EtOAc 1:1). ¹H NMR (500 MHz, CDCl₃) 0.59 - 0.66 (2H, m), 0.84 - 0.94 (2H, m), 2.72 - 3.02 (1H, m), 4.15 (3H, s), 5.43 (1H, s), 7.06 (1H, d, J = 5.2 Hz), 7.55 (2H, t, J = 7.4 Hz), 7.61 (1H, t, J = 7.4 Hz), 7.91 - 7.97 (2H, m), 8.41 (1H, d, J = 5.2 Hz), 8.45 (1H, s), 8.65 (1H, s), 9.45 (1H, d, J = 2.1 Hz). HRMS: Found 362.1607, calculated for $C_{20}H_{20}N_5O_2$ (M+H)⁺: 362.4051.

N-(5-(2-(Cyclopropylamino)pyrimidin-4-yl)-2-oxo-1,2-dihydropyridin-3-yl)benzamide (**24**)

Prepared using general procedure B to give a yellow solid (12 mg, 65%): *N*-(5-(2-(cyclopropylamino)pyrimidin-4-yl)-2-methoxypyridin-3-yl)benzamide (19 mg, 0.053 mmol, 1 equivalent), sodium iodide (23.64 mg, 0.158 mmol, 3 equivalents), trimethylsilyl chloride (0.02 mL, 0.158 mmol, 3 equivalents) and acetonitrile 0.5 mL). R_f = 0.52 (5% MeOH in EtOAc). mp: 292 °C. ¹H NMR (500 MHz, DMSO-d₆) 0.38 - 0.59 (2H, m), 0.59 - 0.74 (2H, m), 2.63 - 2.87 (1H, m), 6.99 (1H, d, J = 5.2 Hz), 7.29 (1H, d, J = 3.2 Hz), 7.56 (2H, t, J = 7.3 Hz), 7.62 (1H, t, J = 7.3 Hz), 7.93 (2H, m), 8.14 (1H, brs), 8.25 (1H, d, J = 5.2), 8.98 (1H, brs), 9.41 (1H, brs).¹³C NMR (126 MHz, DMSO-d₆) 165.0 (C), 163.7 (C), 158.6 (CH), 134.6 (C), 132.3 (CH), 129.1 (CH), 127.5 (CH), 121.3 (CH), 115.8 (C), 104.5 (CH), 24.5 (CH), 7.1 (CH₂). HRMS: Found 348.1453, calculated for C₁₉H₁₈N₅O₂ (M+H)⁺: 348.1455.

Mobility Shift Assay Protocol

Screening of 26 kinases using a mobility shift assay

A mobility shift assay to determine kinase inhibition was carried out using a modified version of the ProfilerPro assay kit 1 (Perkin Elmer, Coventry, UK) including MPS1 and Aurora B. Each kit contained an enzyme plate, a substrate plate, reconstitution buffer, 1 M DTT, protease inhibitor cocktail and termination buffer. Test compounds were added as a solution in DMSO. The 3-aminopyridin-2-one compounds **1-3, 5-8** and **15** were tested at 100 μ M final concentration. The 3-benzamido-pyridin-2-one compounds **16, 22,** and **24** were tested at 30 μ M final concentration. Compound **17** was tested at 100 μ M final concentration. The peptide phosphorylation assay was carried out as per the manufacturer's instructions. The electrophoretic separation and analysis of the phosphorylated and non-phosphorylated peptides was carried out on a 4-sipper chip using a LabChip EZReader II (Perkin Elmer). Results were expressed as % inhibition of peptide phosphorylation relative to control wells for 0 % (no test compound) and 100 % (no ATP) inhibition. The percentage inhibition was calculated using the following equation:

% Inhibition = [1 - (% Conversion _{Sample} / % Conversion _{0% Inhibition Control})] x 100

Kinase	[ATP] (µM)	[Enzyme] (nM)	[Peptide Substrate] (µM)	MgCl ₂ (mM)
MPS1	10.0	6.0	5.0	10
Aurora A	20.0	5.0	1.5	5
Aurora B	15.0	1.0	1.5	5
SRC	38.0	0.219	1.5	10
RSK1	23.3	5.582	1.5	10
MAPKAPK5	5.0	3.143	1.5	10
PKD2	32.1	0.445	1.5	10
ΡΚϹζ	3.8	3.34	1.5	10
PKA	1.7	0.437	1.5	10
p38 α	396.5	3.733	1.5	10
MSK1	21.2	2.507	1.5	10
MET	79.5	2.819	1.5	10
MAPKAPK2	4.6	0.081	1.5	10
LYN	17.0	0.976	1.5	10
LCK	28.5	0.187	1.5	10
INSR	871.8	21.367	1.5	10
GSK3β	7.3	3.211	1.5	10
FYN	36.0	0.197	1.5	10
Erk2	62.1	1.298	1.5	10
Erk1	33.4	1.051	1.5	10
CK1d	16.3	2.416	1.5	10
CHK2	57.8	8.389	1.5	10
CHK1	33.0	2.805	1.5	10
AKT2	186.1	0.293	1.5	10
AKT1	48.0	0.174	1.5	10
ABL	14.0	0.401	1.5	10

 Table 1: Assay Conditions for Mobility Shift Assay

	% Inhibition							
Kinase	1	2	3	5	6	7	8	15
MPS1	0	41	50	41	69	71	87	20
AurA	58	100	0	46	62	98	87	49
AurB	20	96	43	35	61	94	79	77
SRC	31	86	12	8	24	89	46	21
RSK1	19	89	27	23	10	91	89	17
PRAK	19	99	0	1	15	88	24	24
PKD2	10	67	0	26	36	70	81	8
ΡΚϹζ	27	83	50	5	17	37	0	13
PKA	0	66	68	52	48	81	8	0
p38 α	9	39	0	0	0	25	12	8
MSK1	26	95	27	61	22	94	27	14
MET	33	81	0	18	19	76	26	14
MAPKAPK2	25	97	1	19	32	79	28	33
LYN	15	49	18	15	27	63	42	10
LCK	4	11	6	7	40	55	38	4
INSR	2	25	1	6	13	80	27	23
GSK3β	17	68	9	26	20	89	87	39
FYN	11	34	10	9	40	66	51	6
Erk2	3	68	0	12	15	53	45	11
Erk1	9	62	0	20	22	52	46	11
CK1d	45	72	40	38	37	79	63	33
CHK2	49	99	0	36	57	98	61	42
CHK1	37	77	68	1	26	79	26	4
AKT2	77	97	0	4	9	93	18	17
AKT1	42	97	12	22	15	89	18	10
ABL	5	58	11	26	66	67	43	5
S(30%)	0.31	0.92	0.23	0.27	0.42	0.96	0.58	0.23
S(50%)	0.08	0.77	0.08	0.08	0.19	0.92	0.35	0.04
S(80%)	0.00	0.46	0.00	0.00	0.00	0.42	0.19	0.00

Table 2: % Inhibition for 3-aminopyridin-2-one fragment library at 100 μM measured using a microfluidics-based assay

		% Inhibition			
Kinase	16	17 *	22	24	
MPS1	62	42	88	76	
AurA	59	89	92	99	
AurB	24	27	71	91	
SRC	60	59	98	99	
RSK1	19	38	41	98	
PRAK	1	10	10	71	
PKD2	30	22	46	69	
ΡΚϹζ	0	0	24	0	
PKA	0	6	0	75	
p38 α	4	3	6	36	
MSK1	1	15	8	83	
MET	9	23	24	65	
MAPKAPK2	29	9	12	96	
LYN	55	56	93	86	
LCK	45	50	97	75	
INSR	13	21	30	27	
GSK3β	92	93	94	99	
FYN	59	59	95	93	
Erk2	5	14	1	37	
Erk1	12	2	0	33	
CK1d	79	78	31	78	
CHK2	29	58	33	71	
CHK1	24	21	84	0	
AKT2	4	50	13	91	
AKT1	12	24	13	73	
ABL	55	75	96	88	
S(30%)	0.35	0.46	0.54	0.88	
S(50%)	0.31	0.31	0.38	0.77	
S(80%)	0.04	0.08	0.35	0.38	

Table 3: % Inhibition for 3-benzamido-pyridin-2-one based fragment library at 30 μ M (*100 μ M) measured using a microfluidics-based assay

		Library
MPS1	0.38	0.75
AurA	0.63	1.00
AurB	0.63	0.50
SRC	0.25	1.00
RSK1	0.38	0.25
PRAK	0.25	0.25
PKD2	0.38	0.25
ΡΚϹζ	0.13	0.00
PKA	0.50	0.25
p38 α	0.00	0.00
MSK1	0.38	0.25
MET	0.25	0.25
MAPKAPK2	0.25	0.25
LYN	0.13	1.00
LCK	0.13	0.50
INSR	0.13	0.00
GSK3β	0.38	1.00
FYN	0.25	1.00
Erk2	0.25	0.00
Erk1	0.25	0.00
CK1d	0.38	0.75
CHK2	0.50	0.50
CHK1	0.38	0.25
AKT2	0.38	0.25
AKT1	0.25	0.25
ABL	0.38	1.00

S_{kinase}(50%) Kinase 3-Aminopyridinone 3-Benzamidopyridinone Fragment Library Library

Table 4: Skinase(50%) of 3-aminopyridin-2-one fragment library and 3-benzamidopyridin-2-one library

IC₅₀ Determination

For all compounds, IC_{50} values for MPS1, Aurora A, and Aurora B were determined using MPS1 and Aurora A/B mobility shift assays as described previously.¹ IC_{50} values were calculated from a four-parameter logistic fit of percentage inhibition versus concentration (at 8 compound concentrations) using GraphPad Prism 5 (GraphPad Software, Inc., CA, USA). Mean IC_{50} values were determined from two independent measurements each carried out in duplicate. K_i values were estimated from the mean IC_{50} values using the Cheng-Prusoff equation.

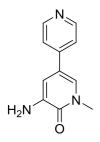
Crystallisation Structure Solution and Refinement

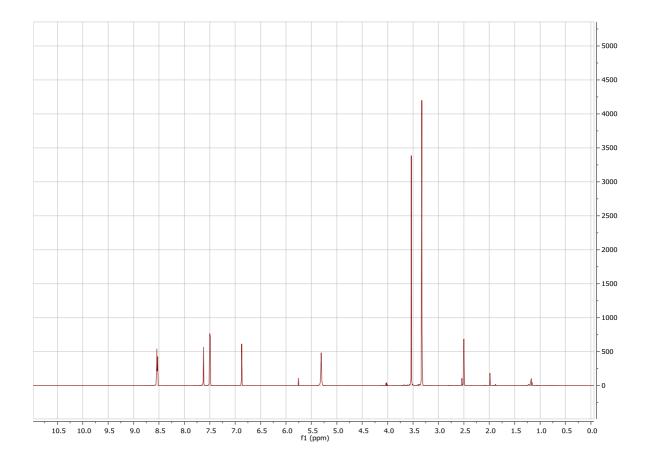
X-ray diffraction data were collected at the Diamond Light Source, Oxford, UK on beamline I04 or I24. Data were integrated and merged using MOSFLM^{2, 3} and AIMLESS.⁴ The structures were solved by molecular replacement using PHASER^{3, 5} and 2ZMC as the molecular replacement model with ligands and water removed. The protein-ligand structures were manually rebuilt in COOT⁶ and refined with BUSTER⁷ in iterative cycles. Ligand restraints were generated with Grade⁸ and mogul.⁹ The quality of the structures was assessed with MOLPROBITY.¹⁰

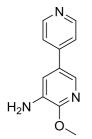
Compound 23 22 2 PDB Code 4CVA 4CV9 4CV8 Space Group /222 /222 /222 Lattice Constants - - 103 - b (Å) 107.16 108.72 110.31 - c (Å) 112.75 113.56 114.24 Data Collection - - - - X-ray Source Diamond Diamond Diamond Barmond Beamline 104 104 124 Resolution Range (Å) 40.72 - 2.50 2.60 - 2.50 3.18 - 3.00 Unique Reflections 15117 (1679) 15436 (1726) 9187 (1454) Completeness (%) 99.9 (100.0) 99.4 (99.9) 99.2 (0.354) Multiplicity 4.4 (4.5) 4.8 (5.0) 4.4 (4.	Compound	23	22	2	
Space Group /222 /222 /222 Lattice Constants a (Å) 70.46 70.38 70.81 b (Å) 107.16 108.72 110.31 c (Å) 112.75 113.56 114.24 Data Collection 104 104 124 Resolution Range (Å) 40.72 - 2.50 40.94 - 2.50 41.23 - 3.00 Highest Resolution Shell (Å) 2.60 - 2.50 2.60 - 2.50 3.18 - 3.00 Unique Reflections 15117 (1679) 15436 (1726) 9187 (1454) Completeness (%) 99.9 (100.0) 99.4 (99.9) 99.5 (99.7) Multiplicity 4.4 (4.5) 4.8 (5.0) 4.4 (4.4) Rmerge 0.09 (1.8) 0.06 (1.4) 0.14 (1.9) CC1/2 ¹¹ 0.998 (0.34) 0.999 (0.482) 0.992 (0.354) Mosaicity 0.18 0.19 0.14 Mean(I/ σ (I)) 10.7 (0.8) 13.8 (0.9) 9.2 (1.0) Refinement 2.55 No. waters 30 32 6 No. amino acids	•			2	
Lattice Constants a (Å) 70.46 70.38 70.81 b (Å) 107.16 108.72 110.31 c (Å) 112.75 113.56 114.24 Data Collection 104 104 124 Resolution Range (Å) 40.72 - 2.50 40.94 - 2.50 41.23 - 3.00 Highest Resolution Shell (Å) 2.60 - 2.50 2.60 - 2.50 3.18 - 3.00 Unique Reflections 15117 (1679) 15436 (1726) 9187 (1454) Completeness (%) 99.9 (100.0) 99.4 (99.9) 99.5 (99.7) Multiplicity 4.4 (4.5) 4.8 (5.0) 4.4 (4.4) Rmerge 0.09 (1.8) 0.06 (1.4) 0.14 (1.9) CC1/2 ¹¹ 0.998 (0.34) 0.999 (0.482) 0.992 (0.354) Mosaicity 0.18 0.19 0.14 Mean(l/ σ (l)) 10.7 (0.8) 13.8 (0.9) 9.2 (1.0) Refinement 255 No. waters 30 32 6 No. amino acids 254 255 25 No. waters					
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Data CollectionX-ray SourceDiamondDiamondDiamondBeamline104104124Resolution Range (Å)40.72 - 2.5040.94 - 2.5041.23 - 3.00Highest Resolution Shell (Å)2.60 - 2.502.60 - 2.503.18 - 3.00Unique Reflections15117 (1679)15436 (1726)9187 (1454)Completeness (%)99.9 (100.0)99.4 (99.9)99.5 (99.7)Multiplicity4.4 (4.5)4.8 (5.0)4.4 (4.4)Rmerge0.09 (1.8)0.06 (1.4)0.14 (1.9)CC1/2 ¹¹ 0.998 (0.34)0.999 (0.482)0.992 (0.354)Mosaicity0.180.190.14Mean(I/ σ (I))10.7 (0.8)13.8 (0.9)9.2 (1.0)Refinement R_{free} (%)21.5023.2322.56No. amino acids254254255No. waters30326No. ligands111No. Ethylene glycol264No. DMSO210 r.m.s. deviation 0.010.010.01Bond angles (⁵)1.081.121.15Ramachandran PlotTamachandran PlotTamachandran PlotFavoured (%)95.9895.1893.98	b (Å)	107.16	108.72	110.31	
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BeamlineI04I04I24Resolution Range (Å)40.72 - 2.5040.94 - 2.5041.23 - 3.00Highest Resolution Shell (Å)2.60 - 2.502.60 - 2.503.18 - 3.00Unique Reflections15117 (1679)15436 (1726)9187 (1454)Completeness (%)99.9 (100.0)99.4 (99.9)99.5 (99.7)Multiplicity4.4 (4.5)4.8 (5.0)4.4 (4.4)Rmerge0.09 (1.8)0.06 (1.4)0.14 (1.9)CC1/2 ¹¹ 0.998 (0.34)0.999 (0.482)0.992 (0.354)Mosaicity0.180.190.14Mean(I/ σ (I))10.7 (0.8)13.8 (0.9)9.2 (1.0)Refinement Z 255255No. amino acids254254255No. waters30326No. ligands111No. Ethylene glycol264No. DMSO210Bond lengths (Å)0.010.010.01Bond lengths (Å)0.010.010.01Bond lengths (Å)95.9895.1893.98	Data Collection				
Resolution Range (Å)40.72 - 2.5040.94 - 2.5041.23 - 3.00Highest Resolution Shell (Å)2.60 - 2.502.60 - 2.503.18 - 3.00Unique Reflections15117 (1679)15436 (1726)9187 (1454)Completeness (%)99.9 (100.0)99.4 (99.9)99.5 (99.7)Multiplicity4.4 (4.5)4.8 (5.0)4.4 (4.4)Rmerge0.09 (1.8)0.06 (1.4)0.14 (1.9)CC1/2 ¹¹ 0.998 (0.34)0.999 (0.482)0.992 (0.354)Mosaicity0.180.190.14Mean(I/\sigma(I))10.7 (0.8)13.8 (0.9)9.2 (1.0)RefinementImage: Signal Signa	X-ray Source	Diamond	Diamond	Diamond	
Highest Resolution Shell (Å) $2.60 - 2.50$ $2.60 - 2.50$ $3.18 - 3.00$ Unique Reflections15117 (1679)15436 (1726)9187 (1454)Completeness (%)99.9 (100.0)99.4 (99.9)99.5 (99.7)Multiplicity 4.4 (4.5) 4.8 (5.0) 4.4 (4.4)Rmerge 0.09 (1.8) 0.06 (1.4) 0.14 (1.9)CC1/2 ¹¹ 0.998 (0.34) 0.999 (0.482) 0.992 (0.354)Mosaicity 0.18 0.19 0.14 Mean(I/ σ (I)) 10.7 (0.8) 13.8 (0.9) 9.2 (1.0)Refinement $IS.54$ 18.03 Rfree (%) 21.50 23.23 22.56 No. amino acids 254 254 255 No. waters 30 32 6 No. ligands111No. Ethylene glycol 2 6 4 No. DMSO 2 1 0 Bond lengths (Å) 0.01 0.01 0.01 Bond angles (°) 1.08 1.12 1.15 Ramachandran Plot $Favoured$ (%) 95.98 95.18 93.98	Beamline	104	104	124	
Unique Reflections15117 (1679)15436 (1726)9187 (1454)Completeness (%)99.9 (100.0)99.4 (99.9)99.5 (99.7)Multiplicity4.4 (4.5)4.8 (5.0)4.4 (4.4)Rmerge0.09 (1.8)0.06 (1.4)0.14 (1.9)CC1/2 ¹¹ 0.998 (0.34)0.999 (0.482)0.992 (0.354)Mosaicity0.180.190.14Mean(I/ σ (I))10.7 (0.8)13.8 (0.9)9.2 (1.0)RefinementRtactor (%)18.9718.5418.03Rtree (%)21.5023.2322.56No. amino acids2542550No. ligands111No. Ethylene glycol264No. DMSO210 r.m.s. deviation 0.010.010.01Bond lengths (Å)0.010.010.01Bond angles (°)1.081.121.15Ramachandran PlotFavoured (%)95.9895.1893.98	Resolution Range (Å)	40.72 - 2.50	40.94 - 2.50	41.23 - 3.00	
Completeness (%)99.9 (100.0)99.4 (99.9)99.5 (99.7)Multiplicity $4.4 (4.5)$ $4.8 (5.0)$ $4.4 (4.4)$ Rmerge $0.09 (1.8)$ $0.06 (1.4)$ $0.14 (1.9)$ CC1/2 ¹¹ $0.998 (0.34)$ $0.999 (0.482)$ $0.992 (0.354)$ Mosaicity 0.18 0.19 0.14 Mean(I/ σ (I)) $10.7 (0.8)$ $13.8 (0.9)$ $9.2 (1.0)$ RefinementRfactor (%) 18.97 18.54 18.03 Rfree (%) 21.50 23.23 22.56 No. amino acids 254 254 255 No. waters 30 32 6 No. ligands 1 1 1 No. Ethylene glycol 2 6 4 No. DMSO 2 1 0 r.m.s. deviation 1.02 1.02 1.12 Bond lengths (Å) 0.01 0.01 0.01 Bond angles (°) 1.08 1.12 1.15 Ramachandran Plot $Favoured (\%)$ 95.98 95.18 93.98	Highest Resolution Shell (Å)	2.60 - 2.50	2.60 - 2.50	3.18 - 3.00	
Multiplicity4.4 (4.5)4.8 (5.0)4.4 (4.4)Rmerge $0.09 (1.8)$ $0.06 (1.4)$ $0.14 (1.9)$ CC1/2 ¹¹ $0.998 (0.34)$ $0.999 (0.482)$ $0.992 (0.354)$ Mosaicity 0.18 0.19 0.14 Mean(I/ σ (I)) $10.7 (0.8)$ $13.8 (0.9)$ $9.2 (1.0)$ RefinementRfactor (%) 18.97 18.54 18.03 Rfree (%) 21.50 23.23 22.56 No. amino acids 254 254 255 No. waters 30 32 6 No. ligands 1 1 1 No. Ethylene glycol 2 6 4 No. DMSO 2 1 0 r.m.s. deviation 1.08 1.12 1.15 Ramachandran Plot 1.08 93.98 93.98	Unique Reflections	15117 (1679)	15436 (1726)	9187 (1454)	
Rmerge $0.09 (1.8)$ $0.06 (1.4)$ $0.14 (1.9)$ CC1/2 ¹¹ $0.998 (0.34)$ $0.999 (0.482)$ $0.992 (0.354)$ Mosaicity 0.18 0.19 0.14 Mean(I/ σ (I)) $10.7 (0.8)$ $13.8 (0.9)$ $9.2 (1.0)$ RefinementRfactor (%) 18.97 18.54 18.03 Rfree (%) 21.50 23.23 22.56 No. amino acids 254 255 255 No. waters 30 32 6 No. ligands 1 1 1 No. Ethylene glycol 2 6 4 No. DMSO 2 1 0 r.m.s. deviationBond lengths (Å) 0.01 0.01 0.01 Bond angles (°) 1.08 1.12 1.15 Ramachandran PlotFavoured (%) 95.98 95.18 93.98	Completeness (%)	99.9 (100.0)	99.4 (99.9)	99.5 (99.7)	
CC1/2 ¹¹ 0.998 (0.34)0.999 (0.482)0.992 (0.354)Mosaicity0.180.190.14Mean(I/ σ (I))10.7 (0.8)13.8 (0.9)9.2 (1.0)RefinementRfactor (%)18.9718.5418.03Rtree (%)21.5023.2322.56No. amino acids254254255No. waters30326No. ligands111No. Ethylene glycol264No. DMSO210r.m.s. deviationBond lengths (Å)0.010.010.01Bond angles (°)1.081.121.15Ramachandran PlotFavoured (%)95.9895.1893.98	Multiplicity	4.4 (4.5)	4.8 (5.0)	4.4 (4.4)	
Mosaicity0.180.190.14Mean(I/ σ (I))10.7 (0.8)13.8 (0.9)9.2 (1.0)RefinementR_factor (%)18.9718.5418.03R_free (%)21.5023.2322.56No. amino acids254255No. amino acids254255No. waters30326No. ligands111No. Ethylene glycol264No. DMSO210r.m.s. deviationBond lengths (Å)0.010.01Bond angles (°)1.081.121.15Ramachandran Plot95.9895.1893.98	Rmerge	0.09 (1.8)	0.06 (1.4)	0.14 (1.9)	
Mean(I/ σ (I))10.7 (0.8)13.8 (0.9)9.2 (1.0)Refinement R_{factor} (%)18.9718.5418.03 R_{free} (%)21.5023.2322.56No. amino acids254254255No. waters30326No. ligands111No. Ethylene glycol264No. DMSO210r.m.s. deviationBond lengths (Å)0.010.010.01Bond angles (°)1.081.121.15Ramachandran PlotFavoured (%)95.9895.1893.98	CC1/2 ¹¹	0.998 (0.34)	0.999 (0.482)	0.992 (0.354)	
Refinement R _{factor} (%) 18.97 18.54 18.03 R _{free} (%) 21.50 23.23 22.56 No. amino acids 254 254 255 No. waters 30 32 6 No. ligands 1 1 1 No. Ethylene glycol 2 6 4 No. DMSO 2 1 0 F.m.s. deviation Bond lengths (Å) 0.01 0.01 0.01 Bond angles (°) 1.08 1.12 1.15 Ramachandran Plot Favoured (%) 95.98 95.18 93.98	Mosaicity	0.18	0.19	0.14	
$\begin{tabular}{ c c c c c c } \hline R_{factor}(\%) & 18.97 & 18.54 & 18.03 \\ \hline R_{free}(\%) & 21.50 & 23.23 & 22.56 \\ \hline No. amino acids & 254 & 254 & 255 \\ \hline No. waters & 30 & 32 & 6 \\ \hline No. ligands & 1 & 1 & 1 \\ \hline No. Ethylene glycol & 2 & 6 & 4 \\ \hline No. DMSO & 2 & 1 & 0 \\ \hline r.m.s. deviation & & & \\ \hline Bond lengths(Å) & 0.01 & 0.01 & 0.01 \\ \hline Bond angles(°) & 1.08 & 1.12 & 1.15 \\ \hline Ramachandran Plot & & \\ \hline Favoured(\%) & 95.98 & 95.18 & 93.98 \\ \hline \end tabular $	Mean(I/ σ (I))	10.7 (0.8)	13.8 (0.9)	9.2 (1.0)	
Rtree (%) 21.50 23.23 22.56 No. amino acids 254 254 255 No. waters 30 32 6 No. ligands 1 1 1 No. Ethylene glycol 2 6 4 No. DMSO 2 1 0 r.m.s. deviation Bond lengths (Å) 0.01 0.01 Bond angles (°) 1.08 1.12 1.15 Ramachandran Plot Favoured (%) 95.98 95.18 93.98	Refinement				
No. amino acids 254 254 255 No. waters 30 32 6 No. ligands 1 1 1 No. Ethylene glycol 2 6 4 No. DMSO 2 1 0 r.m.s. deviation Volume Volume Volume Bond lengths (Å) 0.01 0.01 0.01 Bond angles (°) 1.08 1.12 1.15 Ramachandran Plot Volume Volume Volume Volume Favoured (%) 95.98 95.18 93.98 93.98	R _{factor} (%)	18.97	18.54	18.03	
No. waters 30 32 6 No. ligands 1 1 1 No. Ethylene glycol 2 6 4 No. DMSO 2 1 0 r.m.s. deviation 0.01 0.01 0.01 Bond lengths (Å) 0.01 0.01 0.01 Bond angles (°) 1.08 1.12 1.15 Ramachandran Plot Favoured (%) 95.98 95.18 93.98	R _{free} (%)	21.50	23.23	22.56	
No. ligands 1 1 1 No. Ethylene glycol 2 6 4 No. DMSO 2 1 0 r.m.s. deviation V V V Bond lengths (Å) 0.01 0.01 0.01 Bond angles (°) 1.08 1.12 1.15 Ramachandran Plot V V V V Favoured (%) 95.98 95.18 93.98	No. amino acids	254	254	255	
No. Ethylene glycol 2 6 4 No. DMSO 2 1 0 r.m.s. deviation 0.01 0.01 0.01 Bond lengths (Å) 0.01 0.01 1.12 Bond angles (°) 1.08 1.12 1.15 Ramachandran Plot 95.98 95.18 93.98	No. waters	30	32	6	
No. DMSO 2 1 0 r.m.s. deviation 0.01 0.01 0.01 Bond lengths (Å) 0.01 0.01 0.01 Bond angles (°) 1.08 1.12 1.15 Ramachandran Plot 95.98 95.18 93.98	No. ligands	1	1	1	
r.m.s. deviation Bond lengths (Å) 0.01 0.01 0.01 Bond angles (°) 1.08 1.12 1.15 Ramachandran Plot Favoured (%) 95.98 95.18 93.98	No. Ethylene glycol	2	6	4	
Bond lengths (Å) 0.01 0.01 0.01 Bond angles (°) 1.08 1.12 1.15 Ramachandran Plot Favoured (%) 95.98 95.18 93.98	No. DMSO	2	1	0	
Bond angles (°) 1.08 1.12 1.15 Ramachandran Plot Favoured (%) 95.98 95.18 93.98	r.m.s. deviation				
Ramachandran Plot Favoured (%) 95.98 95.18 93.98	Bond lengths (Å)	0.01	0.01	0.01	
Favoured (%) 95.98 95.18 93.98	Bond angles (°)	1.08	1.12	1.15	
	Ramachandran Plot				
Forbidden (%) 0.00 0.00 0.00	Favoured (%)	95.98	95.18	93.98	
	Forbidden (%)	0.00	0.00	0.00	

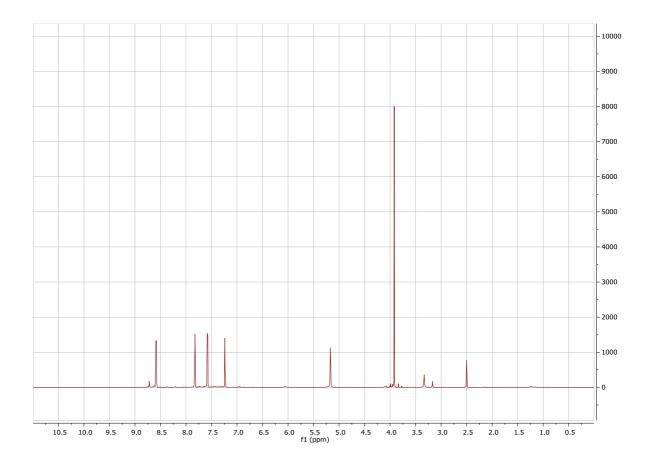
Table 5: Data collection and refinement statistics

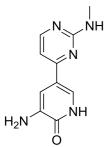
¹H-NMR spectra

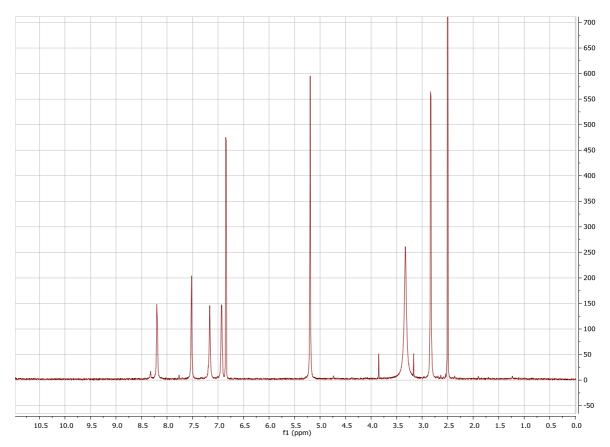


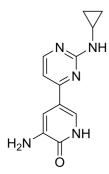


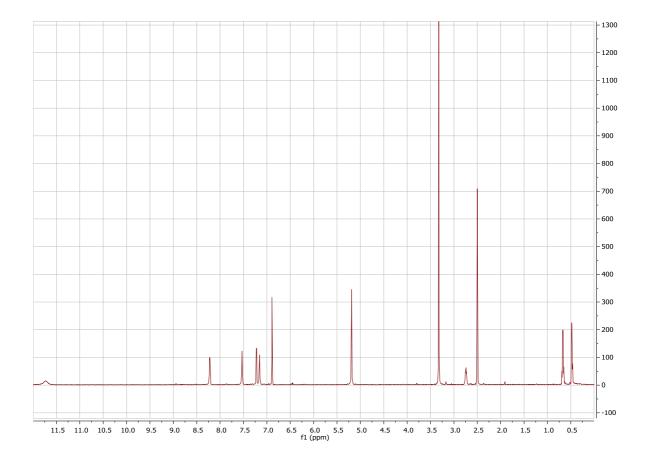


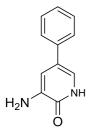


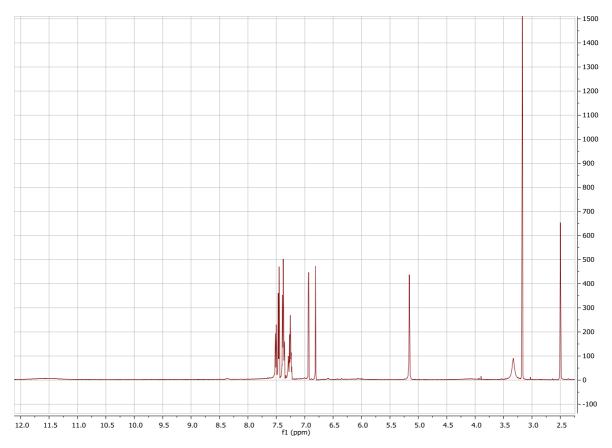


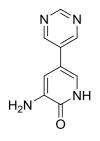


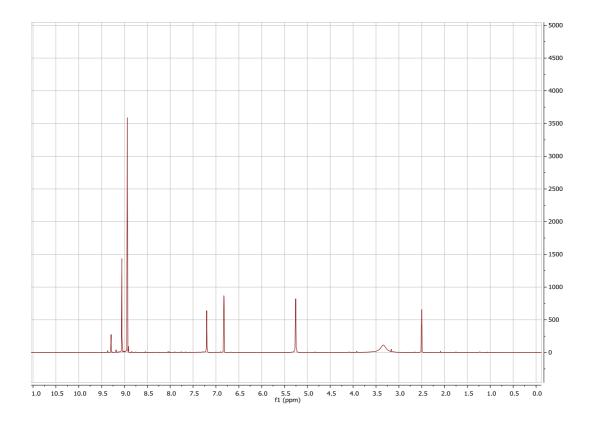


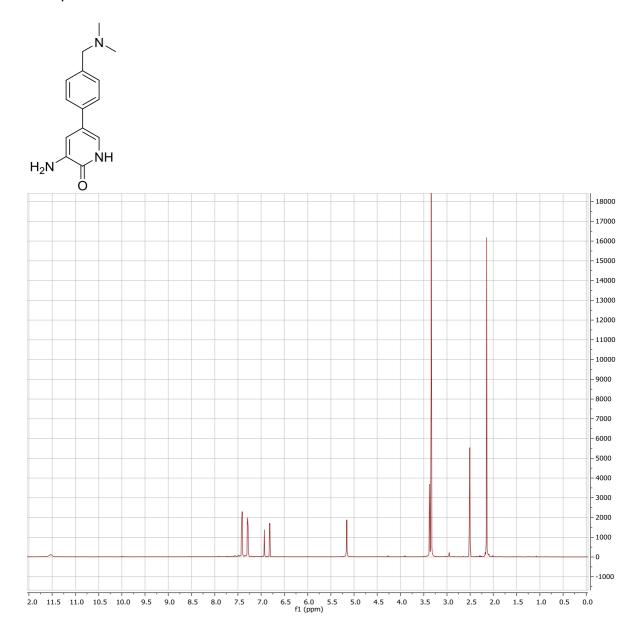


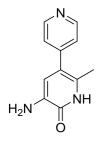


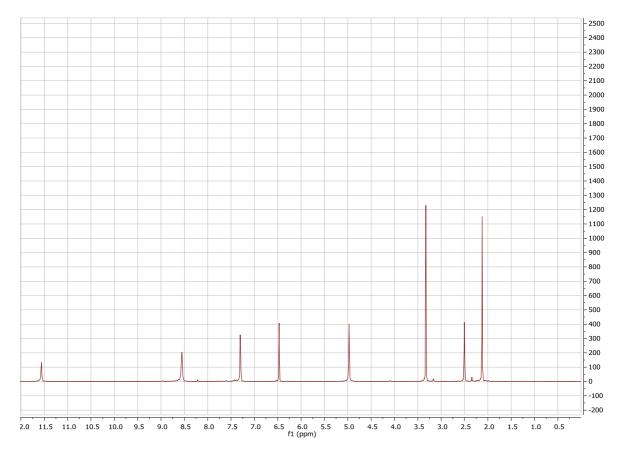


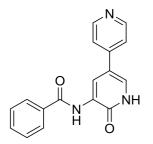


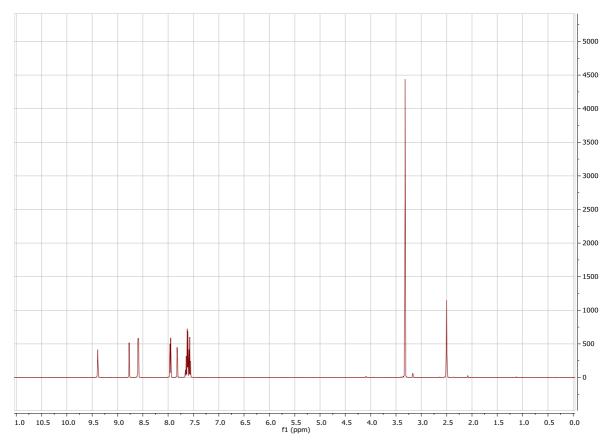


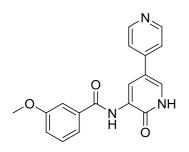


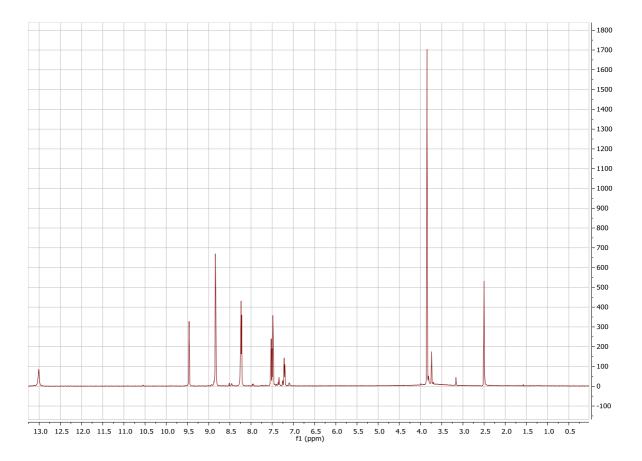


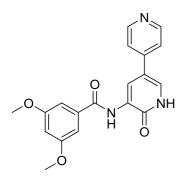


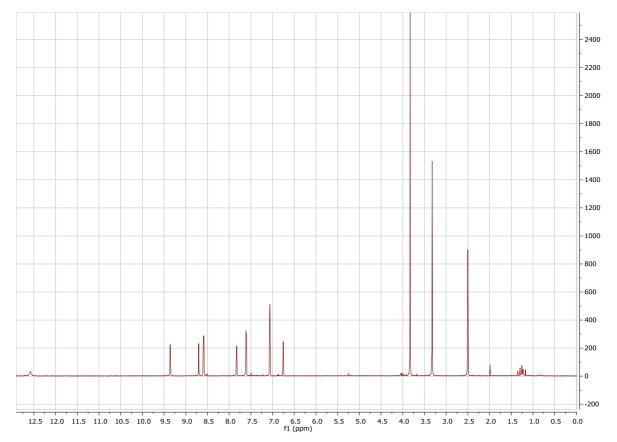


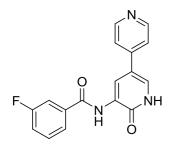


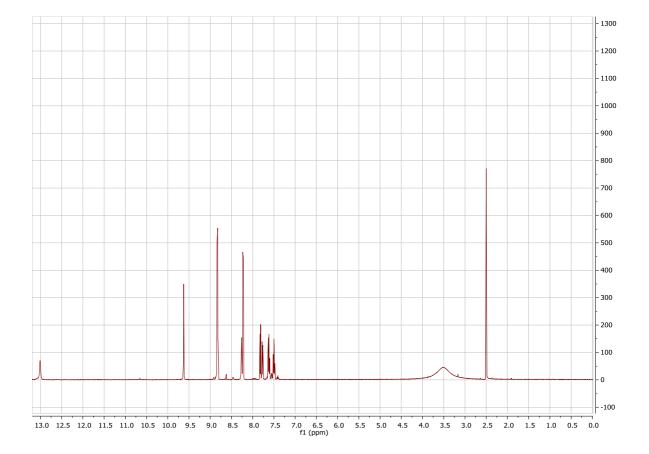


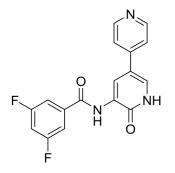


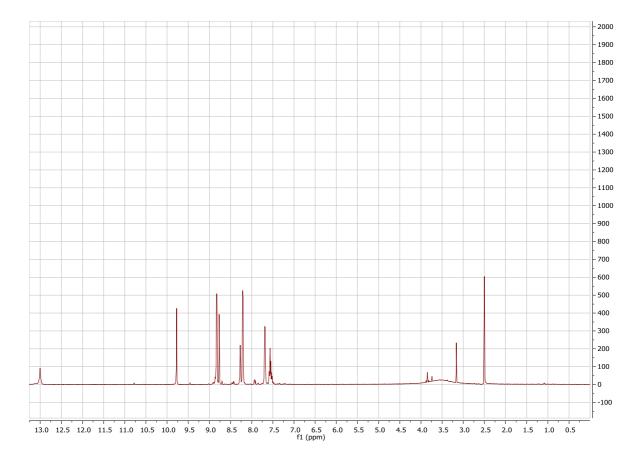


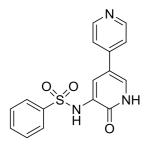


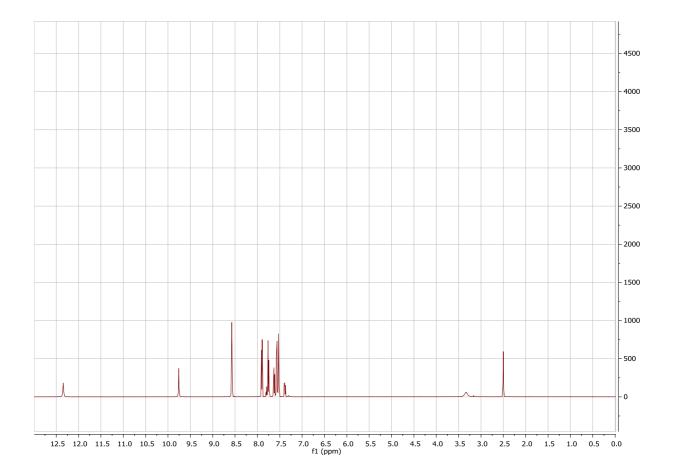


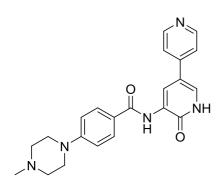


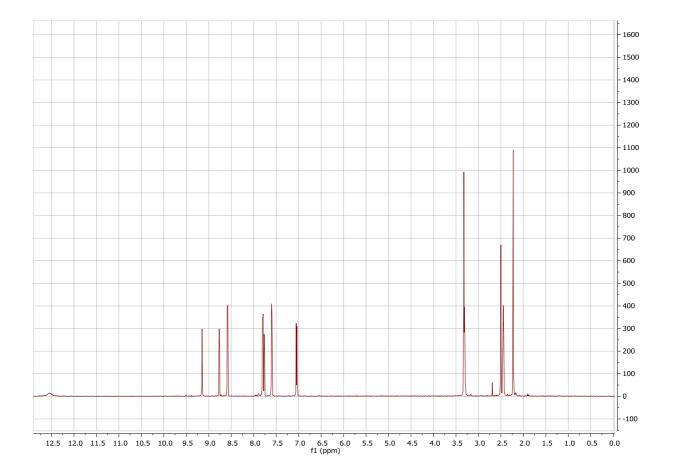


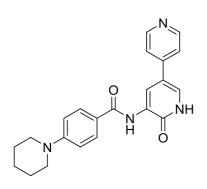


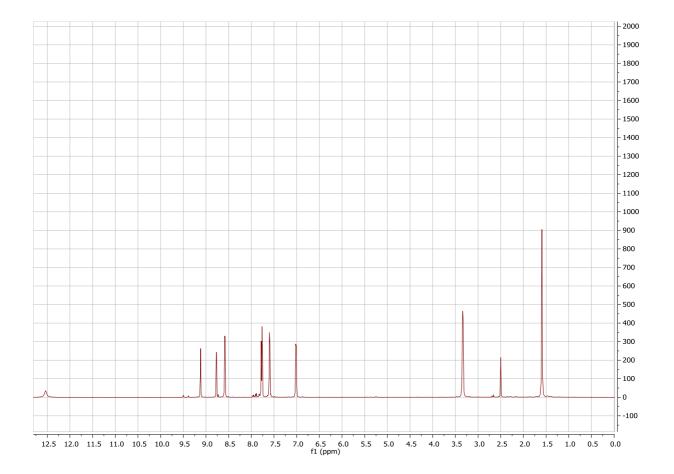


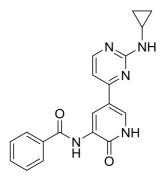


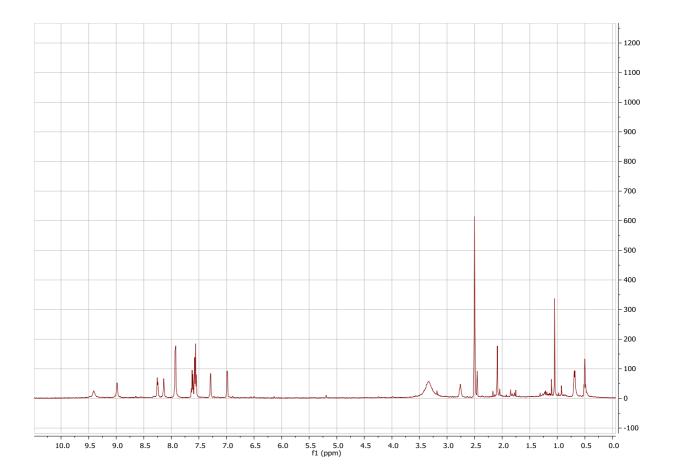












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