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Supporting Information

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Synthesis of compounds 9

General Methods for Chemistry

All reagents and solvents were purchased from suppliers without further purification. Analytical TLC was carried out on silica gel plates with fluorescence F254 and UV light visualization. 1 H NMR spectra were recorded on a Bruker AV-400 spectrometer at 400 MHz. 13 CNMR spectra were recorded on a Bruker AV-500 spectrometer at 125 MHz. Chemical shifts(δ) of NMR are reported in parts per million (ppm) units relative to residual undeuterated solvent, and coupling constants (J values) are given in hertz (Hz). Splitting patterns and apparent multiplicities are described as below: s (singlet), d (doublet), t (triplet), q (quartet), m (multiplet), br (broad signal), dd (doublet of doublets). High-resolution mass spectra (HRMS) were measured by an Applied Biosystems Q-STAR Elite ESI-LC-MS/MS mass spectrometer. Purity of the final compounds were determined with reverse-phase HPLC analysis to be over 95% (SI). HPLC instrument: Dionex Summit HPLC (Column: Diamonsil C18, 5.0μ m, 4.6×250 mm (Agilent Technologies); detector: PDA-100 photodiode array; injector: ASI-100 autoinjector; pump: p-680A). A flow rate of 1.0 mL/min was used with mobile phase of MeOH in H₂O with 0.1% modifier (ammonia, v/v).

Scheme S1. Synthesis of Compounds **9**. Reagents and conditions: a) K_2CO_3 , DMF, rt, 60-84.2%. b) i) Compounds **13**, $Pd(PhCN)_2CI_2$, $(o-MeC_6H_4)_3P$, DIPEA, THF, $70\,^{\circ}C$, Ar; ii) Ac_2O , $80\,^{\circ}C$, 20-30%; (two steps). c) i) Compounds **15**, TFA, 2-Butanol, $100\,^{\circ}C$; ii) TFA, DCM, rt, 60-80% (two steps). d) acryloyl chloride, DIPEA, DCM, $0\,^{\circ}C$, 55-70%.

tert-Butyl (3-((5-bromo-2-chloropyrimidin-4-yl)amino)phenyl)carbamate (**S3a**). To a solution of 5-bromo-2, 4-dichloropyrimidine (0.45 g, 2.0 mmol) and tert-butyl (3-aminophenyl)carbamate (0.42 g, 2.0 mmol) in DMF (3 mL), K_2CO_3 (0.55 g, 4.0 mmol) was added. The suspension was stirred overnight. 20 mL of water was added to the reaction mixture and the precipitate was collected by filtration. The solid was washed with ether and dried to yield compound **S3a** (0.67 g, 84.2%). ¹H NMR (400 MHz, DMSO- d_6) δ 8.29 (s, 1 H), 7.78 (s, 1 H), 7.45 (d, 1 H, J = 7.2 Hz), 7.32-7.28 (m, 2 H), 7.03 (dd, 1 H, J = 1.2, 8.0 Hz), 6.56 (s, 1 H), 1.53 (s, 9 H).

tert-Butyl (3-(2-chloro-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)carbamate (**10a**). Compound **S3a** (2.57 g, 6.43 mmol) and acrylic acid (2.22 mL, 64.3 mmol) were mixed in THF (40 mL) and DIPEA (11.2 mL) under argon. The slurry was stirred, evacuated and refilled with argon before bis(benzonitrile)palladium(II) dichloride (0.12 g, 5%) and tri-otolylphosphine (96 mg, 5%) were added. The mixture was then heated and stirred at 70 $^{\circ}$ C for 24 hrs and then 1.5 mL of Ac₂O was added. The reaction mixture was heated and stirred at 80 $^{\circ}$ C for an additional 24 hrs. The solvent was removed under reduced pressure and the residue was diluted with DCM. The organic layer was separated and washed with 1N HCl (100 mL) and brine, dried over anhydrous Na₂SO₄, concentrated and purified by silica gel chromatography to afford compound **10a** (0.71g, 30%). ¹H NMR (400 MHz, DMSO- d_6) δ 9.61 (s, 1 H), 9.07 (s, 1 H), 8.13 (d, 1 H, J = 9.6 Hz), 7.50 (s, 1 H), 7.45-7.39 (m, 2 H), 6.92-6.89 (m, 1 H), 6.85 (d, 1 H, J = 9.6 Hz), 1.46 (s, 9 H).

8-(3-Aminophenyl)-2-((2-methoxy-4-(4-methylpiperazin-1-

yl)phenyl)amino)pyrido[2,3-d]pyrimidin-7(8H)-one (12a). To a solution of compound 10a (260.9 mg, 0.7 mmol) in 2-butanol (8 mL), 2-methoxy-4-(4-methylpiperazin-1-yl)aniline (162.6 mg, 0.7 mmol) and trifluoroacetic acid (56.0 μ L, 0.7 mmol) were added in a sealed tube. The reaction was heated to 95 °C for 12 hrs. The reaction mixture was then allowed to cool to room temperature. The mixture was transferred to a round-bottom flask and then the solvent was removed under reduced pressure. The residue was dissolved in DCM (3.0 mL) and TFA (3.0 mL), and the resulting mixture was stirred at room temperature overnight. The solvent was removed under reduced pressure, and the residue was neutralized with saturated NaHCO₃ solution. The water layer was extracted with DCM. The organic layer was combined and washed with brine, dried over Na₂SO₄, filtered, concentrated, and purified by silica gel chromatography to afford 12a as a yellow solid (250 mg, 78% for two steps). ¹H NMR (400 MHz, DMSO- d_6) δ 8.71 (s, 1 H), 8.08 (s, 1 H), 7.87 (d, 1 H, J = 9.6 Hz), 7.44 (d, 1 H, J = 8.8 Hz), 7.18 (t, 1 H, J = 8.0 Hz),

6.70 (d, 1 H, *J* = 8.4 Hz), 6.55 (d, 1 H, *J* = 2.4 Hz), 6.41-6.36 (m, 3 H), 6.15 (br, 1 H), 5.25 (br, 2 H), 3.79 (s, 3 H), 3.06 (t, 4 H, *J* = 4.8 Hz), 2.45 (t, 4 H, *J* = 4.8 Hz), 2.23 (s, 3 H).

N-(3-(2-((2-methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (9a). Acryloyl chloride (45.4 μL, 0.55 mmol) was added dropwise to a mixture of 12a (250 mg, 0.55 mmol) and DIPEA (96.1 μL, 0.55 mmol) in dry DCM (10 mL) at 0 °C. The reaction mixture was stirred at 0 °C for 1.0 hr and concentrated under reduced pressure. The resulting crude product was purified by silica gel chromatography to afford 9a as a yellow solid (182.9 mg, 65%). ¹H NMR (400 MHz, DMSO- d_6) δ 10.36 (s, 1 H), 8.74 (s, 1 H), 8.15 (s, 1 H), 7.92-7.86 (m, 2 H), 7.60 (s, 1 H), 7.51 (t, 1 H, J = 8.0 Hz), 7.27 (d, 1 H, J = 8.8 Hz), 7.00 (d, 1 H, J = 8.0 Hz), 6.52 (s,1 H), 6.47-6.41 (m, 2 H), 6.27-6.23 (s,1 H), 6.03 (br, 1 H), 5.78-5.75 (m, 1 H), 3.77 (s, 3 H), 3.03 (m, 4 H), 2.43 (m, 4 H), 2.22 (s, 3 H). ¹³C NMR (125 MHz, DMSO- d_6) δ 163.2, 162.5, 158.7, 156.4, 139.9, 137.5, 136.9, 131.7, 129.4, 127.0, 123.9, 119.6, 118.8, 117.2, 106.2, 105.7, 99.6, 55.7, 54.5, 48.5, 45.7. HRMS (ESI) for C₂₈H₂₉N₇O₃ [M + H]+, calcd: 512.2405, found: 512.2400.

N-(3-(2-((2-Methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-6-methyl-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9b**). 1 H NMR (400 MHz, DMSO- d_6) δ 10.34 (s ,1 H), 8.68 (s, 1 H), 8.01 (s, 1 H), 7.87 (d, 1 H, J = 8.0 Hz), 7.78 (s, 1 H), 7.60 (s, 1 H), 7.51 (t, 1 H, J = 8.0 Hz), 7.30 (d, 1 H, J = 8.8 Hz), 7.00 (d, 1 H, J = 8.0 Hz), 6.52 (s, 1 H), 6.43 (dd, 1 H, J = 10.0, 16.8 Hz), 6.27-6.23 (m, 1 H), 6.01 (br, 1 H), 5.77-5.75 (m, 1 H), 3.77 (s, 3 H), 3.02 (m, 4 H), 2.44 (m, 4 H), 2.23 (s, 3 H), 2.11 (s, 3 H). 13 C NMR (125 MHz, Acetic- d_4) δ 166.2, 166.0, 157.7, 157.3, 157.0, 150.8, 146.8, 140.4, 138.2, 135.3, 132.0, 131.0, 129.1, 128.4, 125.8, 123.1, 121.7, 121.6, 109.5, 107.7, 101.8, 56.4, 54.0, 48.3, 43.7, 17.2. HRMS (ESI) for C₂₉H₃₂N₇O₃ [M + H]⁺, calcd: 526.2561, found: 526.2564.

N-(3-(2-((2-Methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-5-methyl-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9f**). 1 H NMR (400 MHz, DMSO- d_6) δ 10.34 (s, 1 H), 8.80 (s, 1 H), 8.10 (s, 1 H), 7.88 (d, 1 H, J = 8.0 Hz), 7.56 (s, 1 H), 7.50 (t, 1 H, J = 8.0 Hz), 7.26 (d, 1 H, J = 8.8 Hz), 6.97 (d, 1 H, J = 7.6 Hz), 6.52 (s, 1 H), 6.43 (dd,

1 H, J = 10.0, 16.8 Hz), 6.32 (s, 1 H), 6.27-6.23 (m, 1 H), 5.99 (br, 1 H), 5.78-5.75 (m, 1 H), 3.77 (s, 3 H), 3.02 (m, 4 H), 2.46-2.43 (m, 7 H), 2.22 (s, 3 H). ¹³C NMR (125 MHz, DMSO- d_6) δ 163.2, 162.1, 156.5, 156.3, 146.9, 139.9, 137.0, 131.7, 129.4, 127.1, 124.0, 119.7, 118.7, 116.5, 106.2, 106.1, 99.6, 55.7, 54.6, 48.5, 45.7, 17.0. HRMS (ESI) for $C_{29}H_{31}N_7O_3$ [M + H]⁺, calcd: 526.2561, found: 526.2558.

N-(3-(5-ethyl-2-((2-methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9g**). ¹H NMR (400 MHz, DMSO- d_6) δ 10.34 (s, 1 H), 8.85 (s, 1 H), 8.09 (s, 1 H), 7.87 (d, 1 H, J = 7.2Hz), 7.57 (s, 1 H), 7.50 (t, 1 H, J = 8.0Hz), 7.26 (d, 1 H, J = 8.8Hz), 6.98 (d, 1 H, J = 8.0Hz), 6.52 (s, 1 H), 6.43 (dd, 1 H, J = 10.0Hz, 16.8Hz), 6.30 (s, 1 H), 6.27-6.23 (m, 1 H), 6.00 (br, 1 H), 5.78-5.75 (m, 1 H), 3.77 (s, 3 H), 3.02 (m, 4 H), 2.88 (q, 2 H, J = 7.2 Hz), 2.43 (m, 4 H),2.22 (s, 3 H), 1.28 (t, 3 H, J = 7.2 Hz). ¹³C NMR (125 MHz, DMSO- d_6) δ 163.6, 162.7, 156.9, 165.5, 152.4, 140.3, 137.5, 132.1, 129.9, 127.5, 124.4, 120.1, 119.1, 115.0, 106.6, 105.7, 100.0, 56.1, 55.0, 49.0, 46.2, 23.4, 13.3. HRMS (ESI) for C₃₀H₃₃N₇O₃ [M + H]+, calcd: 540.2718, found: 540.2714.

N-(3-(2-((2-Methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-7-oxo-5-propylpyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9h**). ¹H NMR (400 MHz, DMSO- d_6) δ 10.34 (s, 1 H), 8.86(s, 1 H), 8.09 (s, 1 H), 7.87 (d, 1 H, J = 7.6 Hz), 7.57 (s, 1 H), 7.50 (t, 1 H, J = 8.0 Hz), 7.25 (d, 1 H, J = 8.8 Hz), 6.99-6.97 (m, 1 H), 6.51 (d, 1 H, J = 2.0 Hz), 6.43 (dd, 1 H, J = 10.0, 16.8 Hz), 6.29 (s, 1 H), 6.25 (dd, 1 H, J = 2.0, 17.2 Hz), 5.99 (br, 1 H), 5.76 (dd, 1 H, J = 1.6, 10.0 Hz), 3.77 (s, 3 H), 3.02 (m, 4 H), 2.81 (t, 2 H, J = 7.2 Hz), 2.42 (t, 4 H, J = 4.8 Hz), 2.21 (s, 3 H), 1.74-1.64 (m, 2 H), 1.01 (t, 3 H, J = 7.2 Hz). ¹³C NMR (125 MHz, DMSO- d_6) δ 168.4, 167.4, 161.7, 155.7, 145.1, 142.3, 136.9, 134.7, 132.3, 129.2, 124.9, 123.9, 120.7, 111.4, 110.5, 104.8, 60.9, 59.8, 53.8, 51.0, 37.0, 27.1, 18.9. HRMS (ESI) for $C_{31}H_{35}N_7O_3$ [M + H]⁺, calcd: 554.2874, found: 554.2875.

N-(3-(2-((2-methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-4,5-dimethyl-7-oxopyrido [2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9o**). ¹H NMR (400 MHz, DMSO-

 d_6) δ 10.33 (s, 1 H), 7.91 (d, 1 H, J = 8.4 Hz), 7.84 (s, 1 H), 7.52-7.48 (m, 2 H), 7.16 (d, 1 H, J = 7.2 Hz), 6.92 (d, 1 H, J = 7.6 Hz), 6.50 (s, 1 H), 6.42 (dd, 1 H, J = 10.0, 16.8 Hz), 6.31 (s, 1 H), 6.27-6.22 (m, 1 H), 5.95 (br, 1 H), 5.77-5.74 (m, 1 H), 3.78 (s, 3 H), 3.00 (m, 4 H), 2.77 (s, 3 H), 2.62 (s, 3 H), 2.42 (m, 4 H), 2.22 (s, 3 H). ¹³C NMR (125 MHz, Acetic- d_4) δ 169.0, 166.1, 165.1, 158.9, 156.9, 151.2, 150.7, 146.7, 140.5, 138.8, 132.0, 131.1, 129.1, 125.7, 123.0, 121.5, 121.4, 119.4, 109.5, 107.9, 101.6, 56.4, 54.0, 48.3, 43.7, 25.9, 25.0. HRMS (ESI) for $C_{30}H_{34}N_7O_3$ [M + H]⁺, calcd: 540.2718, found: 540.2720.

N-(3-(5-Methyl-2-((4-(4-methylpiperazin-1-yl)phenyl)amino)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9p**). 1 H NMR (400 MHz, DMSO- d_6) δ 10.37 (s, 1 H), 9.84 (s, 1 H), 8.82 (s, 1 H), 7.93-7.91 (m, 1 H), 7.57 (s, 1 H), 7.53 (t, 1 H, J = 8.0 Hz), 7.18 (d, 2 H, J = 6.0 Hz), 6.99 (d, 1 H, J = 8.4 Hz), 6.54 (d, 2 H, J = 6.8 Hz), 6.43 (dd, 1 H, J = 10.0, 16.8 Hz), 6.30(s, 1 H), 6.25 (dd, 1 H, J = 2.0, 16.8 Hz), 5.76 (dd, 1 H, J = 1.6, 10.0 Hz), 2.95-2.94 (m, 4 H), 2.46 (s, 3 H), 2.41 (t, 4 H, J = 4.8 Hz), 2.20 (s, 3 H). 13 C NMR (125 MHz, DMSO- d_6) δ 163.5, 162.4, 158.5, 156.6, 147.2, 146.3, 140.1, 137.4, 131.8, 131.7, 129.8, 127.4, 124.2, 119.8, 118.9, 116.4, 115.4, 106.0, 54.7, 48.7, 45.9, 17.1. HRMS (ESI) for $C_{28}H_{29}N_7O_2$ [M + H]+, calcd: 496.2456, found: 496.2456.

N-(3-(5-Methyl-2-((2-methyl-4-(4-methylpiperazin-1-yl)phenyl)amino)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9q**). 1 H NMR (400 MHz, DMSO- d_6) δ 10.29 (s, 1 H), 8.83 (s, 1 H), 8.75 (s, 1 H), 7.73 (d, 1 H, J = 8.4 Hz), 7.54 (s, 1 H), 7.41 (t, 1 H, J = 8.0 Hz), 7.07 (s, 1 H), 6.91 (d, J = 6.4 Hz, 1 H), 6.63 (s, 1 H), 6.45 (dd, 1 H, J = 10.0, 17.2 Hz), 6.28-6.23 (m, 2 H), 5.77 (dd, 1 H, J = 2.0, 10.0 Hz), 2.99 (m, 4 H), 2.43-2.40 (m, 7 H), 2.21 (s, 3 H), 2.08 (s, 3 H). 13 C NMR (125 MHz, DMSO- d_6) δ 163.4, 162.4, 156.7, 156.3, 147.0, 139.8, 136.9, 131.8, 129.2, 128.8, 127.2, 124.2, 119.8, 118.6, 117.1, 116.1, 112.7, 106.1, 54.63, 48.7, 45.8, 18.3, 17.0. HRMS (ESI) for $C_{29}H_{31}N_7O_2$ [M + H]+, calcd: 510.2612, found: 510.2603.

N-(3-(2-((2-Ethoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-5-methyl-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9r**). ¹H NMR (400 MHz, DMSO-

 d_6) δ 10.35 (s, 1 H), 8.80 (s, 1 H), 8.02 (s, 1 H), 7.89 (d, 1 H, J = 7.2 Hz), 7.57 (s, 1 H), 7.51 (t, 1 H, J = 8.0 Hz), 7.28 (d, 1 H, J = 8.8 Hz), 6.98 (d, 1 H, J = 8.4 Hz), 6.51 (d, 1 H, J = 2.4 Hz), 6.45 (dd, 1 H, J = 10.0, 16.8 Hz), 6.33 (s, 1 H), 6.25 (dd, 1 H, J = 2.0, 16.8 Hz), 5.99 (br, 1 H), 5.76 (dd, 1 H, J = 2.0, 10.0 Hz), 4.04 (q, 2 H, J = 6.8 Hz), 3.00 (m, 4 H), 2.46 (s, 3 H), 2.42 (t, 4 H, J = 4.8 Hz), 2.21 (s, 3 H), 1.32 (t, 3 H, J = 6.8 Hz). ¹³C NMR (125 MHz, DMSO- d_6) δ 163.2, 162.1, 156.5, 156.3, 146.8, 139.9, 137.0, 131.7, 129.5, 127.1, 124.0, 120.0, 119.6, 118.7, 116.6, 106.2, 106.1, 100.4, 63.9, 54.6, 48.6, 45.8, 17.0, 16.6. HRMS (ESI) for $C_{30}H_{33}N_7O_3$ [M + H]+, calcd: 540.27176, found: 540.27173.

N-(3-(5-Methyl-2-((4-(4-methylpiperazin-1-yl)-2-propoxyphenyl)amino)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9s**). ¹H NMR (400 MHz, DMSO- d_6) δ 10.35 (s, 1 H), 8.80 (s, 1 H), 8.00 (s, 1 H), 7.88 (d, 1 H, J = 8.0 Hz), 7.57 (s, 1 H), 7.51 (t, 1 H, J = 8.0 Hz), 7.27 (d, 1 H, J = 8.8 Hz), 6.98 (d, 1 H, J = 7.2 Hz), 6.51 (d, 1 H, J = 2.4 Hz), 6.43 (dd, 1 H, J = 10.0, 16.8 Hz), 6.33 (s, 1 H), 6.25 (dd, 1 H, J = 2.0, 17.2 Hz), 5.99 (br, 1 H), 5.76 (dd, 1 H, J = 2.0, 10.4 Hz), 3.94 (t, 1 H, J = 6.4 Hz), 3.00 (m, 4 H), 2.46 (s, 3 H), 2.42 (t, 4 H, J = 4.8 Hz), 2.22 (s, 3 H), 1.75-1.70 (m, 2 H), 0.95 (t, 3 H, J = 7.2 Hz). ¹³C NMR (125 MHz, DMSO- d_6) δ 163.2, 162.1, 156.5, 156.3, 146.8, 139.9, 137.0, 131.7, 129.4, 127.0, 124.0, 120.0, 119.6, 118.7, 116.6, 106.3, 106.1, 100.5, 69.7, 54.6, 48.6, 45.8, 22.0, 17.0, 10.4. HRMS (ESI) for C₃₁H₃₅N₇O₃ [M + H]+, calcd: 554.28741, found: 554.28711.

N-(3-(2-((2-Isopropoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-5-methyl-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9t**). 1 H NMR (400 MHz, DMSO- d_6) δ 10.36 (s, 1 H), 8.80 (s, 1 H), 7.93-7.90 (m, 2 H), 7.58 (s, 1 H), 7.52 (t, 1 H, J = 8.0 Hz), 7.31 (d, 1 H, J = 9.2 Hz), 6.99 (d, 1 H, J = 8.4 Hz), 6.54 (d, 1 H, J = 2.4 Hz), 6.43 (dd, 1 H, J = 10.0, 16.8 Hz), 6.34 (d, 1 H, J = 1.2 Hz), 6.25 (dd, 1 H, J = 2.0, 16.8 Hz), 5.99 (br, 1 H), 5.76 (dd, 1 H, J = 2.0, 10.0 Hz), 4.66-4.60 (m, 1 H), 2.99-2.98 (m, 4 H), 2.46 (s, 3 H), 2.42 (t, 4 H, J = 4.8 Hz), 2.21 (s, 3 H), 1.25 (d, 6 H, J = 6.0 Hz). 13 C NMR (125 MHz, DMSO- d_6) δ 163.2, 162.0, 156.5, 156.3, 146.8, 139.9, 137.1, 131.7, 129.5, 127.1, 124.0, 121.0, 119.7,

118.7, 116.6, 106.6, 106.1, 102.1, 70.7, 54.6, 48.6, 45.8, 21.8, 17.0. HRMS (ESI) for $C_{31}H_{35}N_7O_3$ [M + H]+, calcd: 554.2874, found: 554.2874.

S5 S6c-d 12c-d
$$\frac{\mathbf{9c-d}}{\mathbf{0}}$$

Scheme S2. Synthesis of Compounds **9c-d**. Reagents and conditions: a) R^2CH_2COOEt (**18**), LHDMS (1M in THF), THF, -78 °C to rt, 36-40%. b) 1) m-CPBA, DCM, 0 °C to rt; 2) 2-methoxy-4-(4-methylpiperazin-1-yl)aniline, TFA, 2-Butanol, 110 °C; 3) TFA, DCM, rt, 30-40%. c) acryloyl chloride, DIPEA, DCM, 0 °C, 58-75%.

N-(3-(2-((2-Methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-7-oxo-6-phenylpyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9c**). ¹H NMR (400 MHz, DMSO- d_6) δ 10.38 (s, 1 H), 8.82 (s, 1 H), 8.21 (s, 1 H), 8.13 (s, 1 H), 7.90 (d, 1 H, J = 7.2 Hz), 7.71 (d, 1 H, J = 7.6 Hz), 7.66 (s, 1 H), 7.53 (t, 1 H, J = 8.0 Hz), 7.43 (t, 1 H, J = 8.0 Hz), 7.36 (t, 1 H, J = 7.2Hz), 7.33 (d, 1 H, J = 8.8 Hz), 7.08 (d, 1 H, J = 7.6 Hz), 6.53 (d, 1 H, J = 2.0 Hz), 6.49 (dd, 1 H, J = 10.0, 16.8 Hz), 6.28 (dd, 1 H, J = 2.0, 16.8 Hz), 6.04 (br, 1 H), 5.78 (dd, 1 H, J = 2.0,10.0 Hz), 3.78 (s, 3 H), 3.06 (m, 4 H), 2.29 (m, 4 H). ¹³C NMR (125 MHz, DMSO- d_6) δ 163.3, 161.7, 158.9, 155.7, 140.0, 137.3, 136.1, 135.1, 131.8, 129.5, 128.6, 128.0, 127.7, 127.4, 127.1, 124.0, 120.0, 119.7, 118.8, 106.3, 106.1, 99.8, 55.7, 53.9, 47.8, 44.7. HRMS (ESI) for C₃₄H₃₄N₇O₃ [M + H]+, calcd: 588.2718, found: 588.2717.

tert-Butyl (3-(6-benzyl-2-(methylthio)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl) carbamate (**S6d**). LHMDS (1.0 M in THF, 6 mL, 6.0 mmol) was added into dry THF (5 mL) under argon protection, and then the mixture was stirred at -78 °C. Ethyl 3-phenylpropanoate (1.06 mL, 6.0 mmol) was then slowly added to the mixture and stirred for 20 min. Compound **S5** (720.0 mg, 2.0 mmol) dissolved in dry THF (5 mL) was

added to the mixture and stirred for another 1.0 hr at -78 °C. The reaction mixture slowly warmed to room temperature, stirred overnight. The reaction was quenched with saturated NH₄Cl solution and was extracted with DCM. The organic layer was washed with brine and dried over Na₂SO₄, filtered, concentrated to afford the crude product. The crude product was recrystallized from ethyl acetate to give a light yellow product **S6d** (347 mg, 36.6%). ¹H NMR (400 MHz, DMSO- d_6) δ 9.55 (s, 1 H), 8.87 (s, 1 H), 7.72 (s, 1 H), 7.46 (s, 1 H), 7.44-7.37 (m, 2 H), 7.35-7.32 (m, 4 H), 7.27-7.22 (m, 1 H), 6.89 (d, 1 H, J = 7.2 Hz), 3.86 (s, 2 H), 2.19 (s, 3 H), 1.45 (s, 9 H).

8-(3-Aminophenyl)-6-benzyl-2-((2-methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino) pyrido [2,3-d] pyrimidin-7(8H)-one (12d) To a solution of compound S6d (118.6 mg, 0.25 mmol) in DCM (5 mL), 75% m-CPBA (172.6 mg, 0.75 mmol) was added dropwise at 0 °C. The mixture was slowly warmed to room temperature and stirred for 4.5 hrs. Subsequently, the reaction mixture was quenched with a saturated solution of $Na_2S_2O_3$. The organic layer was separated and washed with saturated aqueous $NaHCO_3$ solution (three times), and brine. The combined organic layer was dried over Na_2SO_4 , filtered, and concentrated under reduced pressure to give tert-butyl (3-(6-benzyl-2-(methylsulfonyl)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)carbamate as a crude product which was used in the next step without further purification.

To a solution of the crude product in 2-butanol (5.0 mL), 2-methoxy-4-(4-methylpiperazin-1-yl)aniline (55.3 mg, 0.25 mmol) and trifluoroacetic acid (19.5 μ L, 0.25 mmol) were added in a sealed tube. The reaction was heated to 110 °C for 24 hrs. The reaction mixture was allowed to cool to room temperature. The mixture was transferred to a round-bottom flask and then the solvent was removed under reduced pressure to afford the crude product. The crude product was then dissolved in DCM (2 mL) and TFA (2 mL), and the resulting mixture was stirred at room temperature overnight. The solvent was removed under reduced pressure, and the residue was neutralized by saturated NaHCO₃ solution. The water layer was extracted with DCM. The

organic layer was washed with brine and dried over Na₂SO₄, filtered, concentrated, and purified by silica gel chromatography to afford **12d** as a solid (47.9 mg, total yield of three steps 35.0 %). ¹H NMR (400 MHz, DMSO) δ 8.65 (s, 1 H), 8.00 (s, 1 H), 7.61 (s, 1 H), 7.44 (d, 1 H, J = 8.0 Hz), 7.34-7.30 (m, 4 H), 7.24-7.21 (m, 1 H), 7.17 (t, 1 H, J = 8.0 Hz), 6.70 (d, 1H, J = 8.0 Hz), 6.54 (d, 1 H, J = 2.4 Hz), 6.39-6.36 (m, 2 H), 6,15 (br, 1 H), 5.24 (s, 2 H), 3.81 (s, 2 H), 3.78 (s, 3 H), 3.05 (t, 4 H, J = 4.8 Hz), 2.45 (t, 4 H, J = 4.8 Hz), 2.22(s, 3 H).

N-(3-(6-Benzyl-2-((2-methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9d**). The synthetic procedure of **9d** from **12d** was similar to that of **9a**. ¹H NMR (400 MHz, DMSO- d_6) δ 10.33 (s, 1 H), 8.69 (s, 1 H), 8.06 (s, 1 H), 7.86 (d, 1 H, J = 6.8 Hz), 7.65 (s, 1 H), 7.58 (s, 1 H), 7.51 (t, 1 H, J = 8.0 Hz), 7.33-7.23 (m, 6 H), 7.00 (d, 1 H, J = 7.2 Hz), 6.51 (s, 1 H), 6.43 (dd, 1 H, J = 9.6, 16.8 Hz), 6.26-6.22 (m, 1 H), 6.02 (brs, 1 H), 5.77-5.75 (m, 1 H), 3.83 (s, 2 H), 3.76 (s, 3 H), 3.02 (m, 4 H), 2.42 (m, 4 H), 2.22(s, 3 H). ¹³C NMR (125 MHz, DMSO- d_6) δ 163.2, 162.5, 158.1, 155.5, 139.9, 139.3, 137.2, 133.9, 131.7, 129.4, 128.8, 128.4, 127.1, 126.2, 124.0, 119.8, 119.5, 118.8, 106.2, 105.6, 99.6, 55.7, 54.6, 48.6, 45.7, 35.7. HRMS (ESI) for $C_{35}H_{36}N_7O_3$ [M + H]⁺, calcd: 602.2874, found: 602.2876.

Scheme S3. Synthesis of Compounds **9e.** .Reagents and conditions: a) LiOH·H₂O, THF:H₂O (1:1, v:v), 60 °C, 85%. b) *N, O*-dimethylhydroxylamine hydrochloride, DMTMM, CH₃CN, rt, 60.6%. c) Methylmagnesium bromide (1M in THF), THF, -78 °C to -10 °C, 49.3%. d) 3-nitroaniline, DIPEA, CH₃CN, 50 °C, 32.8%. e) Ph₃PCHCOOMe, toluene, 110 °C, Ar, 78%. f) Br₂, DCM, 0 °C to rt, 58.9%. g) Phenyboronic acid, Pd(dppf)₂DCM·Cl₂, Et₃N, DME:H₂O (10:1, v:v), rt, 83%. h) 1) m-CPBA, DCM, 0 °C to rt, ii) 2-methoxy-4-(4-methylpiperazin-1-yl)aniline, TFA, 2-Butanol, 110 °C, 63.2% (two steps). i) Pd/C, H₂ (1 atmo), AcOH, rt, 46.3%. j) acryloyl chloride, DIPEA, DCM, 0 °C, 74.5%.

4-Chloro-2-(methylthio)pyrimidine-5-carboxylic acid ($\mathbf{S8}$). LiOH·H₂O (2.52 g, 60 mmol) was slowly added into a mixed solution of ethyl 4-chloro-2-(methylthio)pyrimidine-5-carboxylate (11.64 g, 50.0 mmol) in THF (50 mL and H₂O (50 mL), then the mixture was heated and stirred at 60 °C overnight. The solution was acidified with 1M HCl after THF was partly removed under reduced pressure. The mixture was extracted with ethyl acetate. The organic layer was washed with brine and dried over Na₂SO₄, filtered,

concentrated to give **\$8** as a white solid (8.70 g, 85%). 1 H NMR (400 MHz, DMSO- d_{6}) δ 8.99 (s, 1 H), 2.58 (s, 3 H).

4-Chloro-*N*-methoxy-*N*-methyl-2-(methylthio)pyrimidine-5-carboxamide (**S9**). To a mixture of compound **S8** (204.6 mg, 1 mmol), DMTMM (276.7 mg, 1 mmol) and *N*, *O*-dimethylhydroxylamine hydrochloride (97.5 mg, 1 mmol) in acetonitrile (5 mL) was slowly added DIPEA (0.17mL, 1 mmol). The reaction mixture was stirred at room temperature for 1h and then the solvent was removed under reduced pressure. DCM was added to dissolve the crude product, and then the organic layer was washed with water, brine once and dried over Na_2SO_4 , filtered, concentrated, and purified by silica gel chromatography to afford **S9** as a white solid (150.0 mg, 60.6%). ¹H NMR (400 MHz, CDCl₃) δ 8.43 (s, 1 H), 3.55 (s, 3 H), 3.37 (s, 3 H), 2.59 (s, 3 H).

1-(4-Chloro-2-(methylthio)pyrimidin-5-yl)ethanone (**\$10**). Methylmagnesium bromide (1M in THF, 2.00 mL, 2.0 mmol) was slowly added to a solution of compound **\$9** (495.4 mg, 2.0 mmol) in dry THF (5 mL) under Ar condition at -78 °C. The resulting mixture was slowly warmed to -10 °C, and then stirred for 1h. The reaction was quenched with 1N HCl (2 mL) and then partitioned between water and DCM. The organic layer was washed with brine and dried over Na₂SO₄, filtered, concentrated, and purified by silica gel chromatography to afford **\$10** (200.0 mg, 49.3%). ¹H NMR (400 MHz, CDCl₃) δ 8.76 (s, 1 H), 2.69 (s, 3 H), 2.61 (s, 3 H).

1-(2-(Methylthio)-4-((3-nitrophenyl)amino)pyrimidin-5-yl)ethanone (**S11**). To a solution of 1-(4-chloro-2-(methylthio)pyrimidin-5-yl)ethanone (202.7 mg, 1.0 mmol) and 3-nitroaniline (138.1 mg,1.0 mmol) in 5 mL of acetonitrile was added DIPEA (0.17 mL, 1 mmol). The suspention was stirred at $50\,^{\circ}\text{C}$ overnight. The reaction mixture was allowed to cool to room temperature. The precipitate was filtered to give **S11** as a light yellow solid (100.0 mg, 32.8%). ¹H NMR (400 MHz, DMSO- d_6) δ 11.46 (s, 1 H), 9.00-8.99 (m, 2 H), 7.99 (d, 1 H, J = 8.4 Hz), 7.91 (d, 1 H, J = 8.0 Hz), 7.66 (t, 1 H, J = 8.0 Hz), 2.64 (s, 3 H), 2.58 (s, 3 H).

5-Methyl-2-(methylthio)-8-(3-nitrophenyl)pyrido[2,3-d]pyrimidin-7(8H)-one (**\$12**). A mixture of compound **S11** (918.0)mg, 3.0 mmol) and methyl (triphenylphosphoranylidene)acetate (1.21 g, 3.6 mmol) in dry toluene (10 mL) was heated to reflux under argon for 48 hrs and then the solvent was removed under reduced pressure. The resulting crude product was purified by silica gel chromatography to afford compound **S12** (776.0 mg, 78.0%). 1 H NMR (400 MHz, DMSO- d_{6}) δ 9.00 (s, 1 H), 8.37-8.34 (m, 2 H), 7.88-7.82 (m, 2 H), 6.62 (d, 1 H, J = 1.2 Hz), 2.52 (s, 3 H), 2.15 (s, 3 H).

6-Bromo-5-methyl-2-(methylthio)-8-(3-nitrophenyl)pyrido[2,3-d]pyrimidin-7(8H)-one (**S13**). Bromine (0.12 mL, 2.36 mmol) was slowly added dropwise to a solution of compound **S12** (776.0 mg, 2.36 mmol) at 0 °C under argon. The reaction mixture was stirred overnight at room temperature. Subsequently, the reaction mixture was quenched with a saturated solution of Na₂S₂O₃. The organic layer was separated and washed with water, then brine. The combined organic layer was dried over Na₂SO₄, filtered, concentrated and purified by silica gel chromatography to afford **S13** as a white solid (567.0 mg, 58.9%). ¹H NMR (400 MHz, DMSO- d_6) δ 9.15 (s, 1 H), 8.41-8.36 (m, 2 H), 7.91-7.84 (m, 2 H), 2.72 (s, 3 H), 2.14 (s, 3 H).

5-Methyl-2-(methylthio)-8-(3-nitrophenyl)-6-phenylpyrido[2,3-d]pyrimidin-7(8H)-one (**\$14**). Compound **\$13** (407.2mg, 1mmol), phenylboronic acid (182.9mg, 1.5mmol), Pd(dppf)₂DCM·Cl₂ (81.6mg, 0.1 mmol) and Et₃N (0.42mL, 3.0 mmol) were added to a mixture solution of DME (8 mL) and H₂O (0.8 mL). The mixture was evacuated and refilled with argon and then heated to 90 °C, stirred for 24 hrs. The mixture was diluted with DCM and partitioned between water and DCM. The organic layer was washed with brine and dried over Na₂SO₄, filtered, concentrated, and purified by silica gel chromatography to afford **\$14** as a white solid (337.0 mg, 83.0%). ¹H NMR (400 MHz, DMSO- d_6) δ 9.10 (s, 1 H), 8.40 (t, 1 H, J = 2.0 Hz), 8.37-8.34 (m, 1 H), 7.93-7.91 (m, 1H), 7.85 (t, 1 H, J = 8.0 Hz), 7.46 (t, 2 H, J = 7.2 Hz), 7.40 (t, 1 H, J = 7.6 Hz), 7.33-7.31 (m, 2 H), 2.36 (s, 3 H), 2.18 (s, 3 H).

2-((2-Methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-5-methyl-8-(3-nitrophenyl)-6-phenylpyrido[2,3-d]pyrimidin-7(8H)-one (S15). To a solution of compound S14 (337.0 mg, 0.83 mmol) in DCM (10mL), 75% m-CPBA(507.0 mg, 2.52 mmol) was added dropwise at 0 °C. The mixture was slowly warmed to room temperature and stirred for 4 hrs. Subsequently, the reaction mixture was quenched with a saturated solution of Na₂S₂O₃. The organic layer was separated and washed with saturated aqueous NaHCO₃ solution (three times), then brine. The combined organic layer was dried over Na₂SO₄, filtered, and concentrated under reduced pressure to give 5-methyl-2-(methylsulfonyl)-8-(3-nitrophenyl)-6-phenylpyrido[2,3-d]pyrimidin-7(8H)-one as a crude product which was used in the next step without further purification.

To a solution of the crude product in 2-butanol (8 mL), 2-methoxy-4-(4-methylpiperazin-1-yl)aniline (183.3 mg, 0.83 mmol) and TFA (66.5 μ L, 0.83 mmol) were added in a sealed tube (25 mL). The mixture was heated to 110 °C, and then stirred for 24 hrs. The mixture was allowed to cool to room temperature and was transferred to a round-bottom flask. The solvent was removed under reduced pressure and the resulting crude product was dissolved in DCM, and then washed with saturated aqueous NaHCO₃ solution and brine. The combined organic layer was dried over Na₂SO₄, filtered, concentrated and purified by silica gel chromatography to give **S15** as a yellow solid (303.0 mg, 63.2%.). ¹H NMR (400 MHz, DMSO- d_6) δ 8.90 (s, 1 H), 8.37-8.32 (m, 3 H), 7.87-7.81 (m, 2 H), 7.44 (t, 2 H, J = 7.2 Hz), 7.38-7.34 (m, 1 H), 7.31-7.29 (m, 2 H), 7.05 (br, 1H), 6.49 (s, 1 H), 5.99 (br, 1 H), 3.74 (s, 3 H), 3.04 (m, 4 H), 2.45 (t, 4 H, J = 4.4 Hz), 2.31 (s, 3 H), 2.23 (s, 3 H).

8-(3-Aminophenyl)-2-((2-methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-5-methyl-6-phenylpyrido[2,3-d]pyrimidin-7(8H)-one (12e). A suspension of compound S15 (303.0 mg, 0.52 mmol) and Pd/C (20 mg) in glacial acetic acid (10 mL) at room temperature was stirred under 1 atom H₂ for 6 hrs. The suspension was then filtered through Dicalite, and the filtrate was collected, concentrated under reduced pressure.

The resulting residue was neutralized by saturated NaHCO₃ solution. The water layer was extracted with DCM. The organic layer was washed with brine and dried over Na₂SO₄, filtered and concentrated to give compound **12e** as a yellow solid (241.0 mg, 46.3%). 1 H NMR (400 MHz, DMSO- d_6) δ 8.86 (s, 1 H), 8.04 (s, 1 H), 7.49-7.41 (m, 3 H), 7.35 (t, 1 H, J = 7.6 Hz), 7.37-7.34 (m, 2 H), 7.18 (t, 1 H, J = 8.0 Hz), 6.70 (d, 1 H, J = 8.0 Hz), 6.56 (d, 1 H, J = 2.4 Hz), 6.44-6.43 (m, 1 H), 6.41-6.39 (m, 1 H), 6.15 (br s, 1 H), 5.25 (s, 2 H), 3.81 (s, 3 H), 3.06-3.05 (m, 4 H), 2.45 (t, 4 H, J = 4.8 Hz), 2.27 (s, 3 H), 2.23 (s, 3 H).

N-(3-(2-((2-Methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-5-methyl-7-oxo-6-phenylpyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9e**). The synthetic procedure of **9e** is similar to that of **9a**. ¹H NMR (400 MHz, DMSO- d_6) δ 10.34 (s, 1 H), 8.89 (s, 1 H), 8.10 (s, 1 H), 7.86 (d, 1 H, J = 8.0 Hz), 7.64 (s, 1 H), 7.51 (t, 1 H, J = 8.0 Hz), 7.44 (t, 1 H, J = 7.2 Hz), 7.38-7.34 (m, 1 H), 7.31-7.29 (m, 3 H), 7.03 (d, 1 H, J = 8.0 Hz,), 6.53 (s, 1 H), 6.43 (dd, 1 H, J = 10.0, 17.2 Hz), 6.27-6.23 (m, 1 H), 6.02 (br, 1 H), 5.78-5.75 (m, 1 H), 3.79 (s, 3 H), 3.03 (m, 4 H), 2.45-2.44 (m, 4 H), 2.30 (s, 3 H), 2.23 (s, 3 H). ¹³C NMR (125 MHz, DMSO- d_6) δ 163.6, 162.2, 157.4, 155.8, 142.7, 140.3, 137.8, 135.9, 132.2, 130.9, 129.9, 127.8, 127.5, 124.5, 120.3, 120.1, 119.1, 106.7, 106.7, 100.0, 56.1, 55.0, 49.0, 46.2, 15.7. HRMS (ESI) for $C_{35}H_{35}N_7O_3$ [M + H]+, calcd: 602.2874, found: 602.2873.

Scheme S4. Synthesis of **9i-k**. Reagents and conditions: a) R¹MgBr (in THF), THF, -78 $^{\circ}$ C, Ar, 57.8-60% or CF₃TMS, THF, 0 $^{\circ}$ C to rt, Ar, 27%. b) MnO₂, DCM, reflux, 80-86.6%. c) Ph₃PCHCOOMe, toluene, 110 $^{\circ}$ C, 40-50%. d) 1) m-CPBA, DCM, 0 $^{\circ}$ C to rt; 2) 2-methoxy-4-(4-methylpiperazin-1-yl)aniline, TFA, 2-Butanol, 110 $^{\circ}$ C; 3) TFA, DCM, rt, 30-40%. e) acryloyl chloride, DIPEA, DCM, 0 $^{\circ}$ C, 70-75%.

tert-Butyl (3-((5-(cyclopropyl(hydroxy)methyl)-2-(methylthio)pyrimidin-4-yl)amino)phenyl)carbamate (**S16i**). Compound **S5** (1.44 g, 4 mmol) was dissolved in dry THF (10 mL) under Ar then cooled to -78°C , to which cyclopropylmagnesium bromide (20 mL, 10 mmol, 0.5M in THF) was slowly added and stirred for 4h. The reaction was quenched with 1N HCl (10 mL) and then partitioned between water and DCM. The organic layer was washed with brine and dried over Na₂SO₄, filtered, concentrated, and purified by silica gel chromatography to afford **S16i** (0.93 g, 57.8%). ¹H NMR (400 MHz, DMSO- d_6) δ 9.36 (s, 1H), 8.93 (s, 1 H), 8.11 (s, 1 H), 7.80 (s, 1 H), 7.35 (d, 1H, J = 8.0 Hz), 7.19 (t, 1 H, J = 8.0 Hz), 7.01 (d, 1 H, J = 8.0 Hz), 6.04 (d, 1 H, J = 3.2 Hz), 4.11-4.09 (m, 1 H), 2.45 (s, 3 H), 1.47 (s, 9 H), 0.60-0.55 (m, 1H), 0.47-0.42 (m, 1 H), 0.32-0.31 (m, 1 H).

tert-Butyl (3-((2-(methylthio)-5-(2,2,2-trifluoro-1-hydroxyethyl)pyrimidin-4-yl)amino)phenyl)carbamate ($\mathbf{S16k}$). To a mixture of compound $\mathbf{S5}$ (1.8 g, 5 mmol) and CsF (10 mg) in dry THF (25 mL) was added CF₃TMS (2.95 mL, 20 mmol) under Ar. The

reaction mixture was then stirred at room temperature for 48 h. The reaction was quenched with 0.5N HCl and partitioned between water and DCM. The organic layer was washed with brine and dried over Na₂SO₄, filtered, concentrated, and purified by silica gel chromatography to afford **S16k** (0.58 g, 27%). ¹H NMR (400 MHz, DMSO- d_6) δ 9.38 (s, 1 H), 8.93 (s, 1 H), 8.24 (s, 1 H), 7.81 (s, 1H), 7.39 (d, 1 H, J = 5.6 Hz), 7.24 (d, 1 H, J = 8.0 Hz), 7.20 (t, 1 H, J = 8.0 Hz), 7.06 (d, 1 H, J = 7.6 Hz), 5.66-5.63 (m, 1 H), 3.33 (s, 1 H), 2.41 (s, 3 H), 1.47 (s, 1 H).

tert-Butyl (3-((5-(cyclopropanecarbonyl)-2-(methylthio)pyrimidin-4-yl)amino)phenyl) carbamate (**\$17i**). To a solution of compound **\$16i** (0.93 g, 2.31 mmol) in DCM (20 mL), MnO₂ (2.01 g, 23.1 mmol) was added. The reaction mixture was then heated to reflux for 24 hrs. The solids was filtered off through a Celite pad and washed with DCM (3 × 10 mL). The filtrate was dried over Na₂SO₄, filtered, concentrated to afford **\$17i** (0.80g, 86.6.0%). 1 H NMR (400 MHz, DMSO- d_6) δ 11.39 (s, 1 H), 9.43 (s, 1 H), 9.22 (s, 1 H), 7.89 (s, 1 H), 7.36 (d, 1 H, J = 8.0 Hz), 7.24 (t, 1 H, J = 8.0 Hz), 7.11 (d, 1 H, J = 8.0 Hz), 3.01-2.96 (m, 1 H), 2.53 (s, 3 H), 1.48 (s, 9 H), 1.14-1.04 (m, 4 H).

tert-Butyl (3-(5-cyclopropyl-2-(methylthio)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl) carbamate (**10i**). A mixture of compound **S17i** (0.8 g, 2.0 mmol) and methyl (triphenylphosphoranylidene)acetate (1.0 g, 3.0 mmol) in toluene (10 mL) was heated to reflux under argon for 48 hrs and then the solvent was removed under reduced pressure. The resulting crude product was purified by silica gel chromatography to afford **10i** (0.35 g, 40.5%). ¹H NMR (400 MHz, DMSO- d_6) δ 9.54 (s, 1 H), 9.25 s, 1 H), 7.44-7.36 (m, 3 H), 7.87(d, 1 H, J = 7.6 Hz), 6.31 (s, 1 H), 2.41-2.38 (m, 1 H), 2.20 (s, 3 H), 1.46 (s, 9 H), 1.12-1.09 (m, 2 H), 0.87-0.84 (m, 2 H).

N-(3-(5-Cyclopropyl-2-((2-methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-7-oxopyrido [2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9i**). 1 H NMR (400 MHz, DMSO- d_6) δ 10.33 (s, 1H), 9.09 (s, 1 H), 8.11 (s, 1 H), 7.86 (d, 1 H, J = 7.6 Hz), 7.56 (s, 1 H), 7.49(t, 1 H, J = 8.0 Hz), 7.26 (d, 1 H, J = 9.2 Hz), 6.97 (d, 1 H, J = 8.4 Hz), 6.52 (d, 1 H, J = 2.4 Hz),

6.45 (dd, 1 H, J = 10.0, 16.8 Hz), 6.25 (dd, 1 H, J = 2.0, 16.8 Hz), 6.05-5.99 (m, 2 H), 5.76 (dd, 1 H, J = 2.0, 10.0 Hz), 3.77 (s, 3 H), 3.01 (m, 4 H), 2.43-2.37 (m, 5 H), 2.22 (s, 3 H), 1.09-1.07 (m, 2 H), 0.92-0.89 (m, 2 H). ¹³C NMR (125 MHz, DMSO- d_6) δ 163.7, 162.9, 156.6, 152.7, 140.1, 137.3, 131.8, 129.8, 127.6, 124.3, 120.0, 119.9, 119.0, 111.1, 106.7, 106.5, 99.8, 56.0, 54.7, 48.7, 45.9, 10.7, 8.6. HRMS (ESI) for $C_{31}H_{33}N_7O_3$ [M + H]⁺, calcd: 552.2718, found: 552.2712.

N-(3-(5-Isopropyl-2-((2-methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9j**). ¹H NMR (400 MHz, DMSO- d_6) δ 10.34 (s, 1 H), 8.92 (s, 1 H), 8.09 (s, 1 H), 7.86 (d, 1 H, J = 8.4 Hz), 7.58 (s, 1 H), 7.50 (t, 1 H, J = 8.0 Hz), 7.25 (d, 1 H, J = 8.8 Hz), 6.99-6.97 (m, 2 H), 6.51 (s, 1 H), 6.43 (dd ,1 H, J = 10.0, 16.8 Hz), 6.32 (s, 1 H), 6.27-6.23 (m, 1 H), 5.99 (br, 1 H), 5.77-5.75 (m, 1 H), 3.77 (s, 3 H), 3.50-3.47 (m, 1 H), 3.02 (m, 4 H), 2.43 (m, 4 H), 2.22 (s, 3 H), 1.29 (d, 6 H, J = 6.8 Hz). ¹³C NMR (125 MHz, DMSO- d_6) δ 163.5, 162.8, 156.8, 156.7, 156.1, 140.0, 137.3, 131.8, 129.7, 127.5, 124.2, 119.9, 119.8, 118.9, 112.6, 106.4, 104.9, 99.7, 55.9, 54.7, 48.6, 45.8, 27.1, 22.2. HRMS (ESI) for C₃₁H₃₅N₇O₃ [M + H]⁺, calcd: 554.2874, found: 554.2873.

N-(3-(2-((2-methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-7-oxo-5-(trifluoromethyl) pyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9k**). ¹H NMR (400 MHz, DMSO- d_6) δ 10.37 (s, 1 H), 8.71 (s, 1 H), 8.47 (s, 1 H), 7.88 (s, 1 H), 7.63 (s, 1 H), 7.51 (t, 1 H, J = 8.0 Hz), 7.16 (d, 1 H, J = 8.8 Hz), 7.04 (d, 1 H, J = 8.4 Hz), 6.89 (s, 1 H), 6.50(s, 1 H), 6.44 (dd, 1 H, J = 10.0, 16.8 Hz), 5.96 (br, 1 H), 5.77 (dd, 1 H, J = 2.0, 10.4 Hz), 3.76 (s, 3 H), 3.03 (m, 4 H), 2.42 (t, 4 H, J = 4.8 Hz), 2.22 (s, 3 H). ¹³C NMR (125 MHz, DMSO- d_6) δ 163.6, 161.3, 157.4, 155.5, 140.2, 136.7, 135.0, 134.8, 131.8, 129.9, 127.5, 125.6, 123.4, 121.2, 119.1 (q, 1C, J = 275 Hz), 123.9, 119.6, 119.3, 116.8, 106.2, 100.6, 99.6, 55.9, 54.7, 48.5, 45.9. HRMS (ESI) for $C_{29}H_{28}F_3N_7O_3$ [M + H]+, calcd: 580.2279, found: 580.2272.

Scheme S5. Synthesis of **9l-n**. Reagents and conditions: a) NaOH (1 M in H₂O), THF, 50 $^{\circ}$ C, 95%. b) 1H-benzo[d][1,2,3]triazole, EDCI, DCM, rt, 85%. c) EA, KHMDS (1 M in THF), THF, -78 $^{\circ}$ C to rt, Ar, 17.5%. d) Tf₂O, Et₃N, DCM, -78 $^{\circ}$ C, 82.9%. e) Me₂NH·HCl, DIPEA, THF, 40 $^{\circ}$ C, 98.3% or R₁B(OH)₂, Pd(PPh₃)₂Cl₂ (5%), K₂CO₃, THF/H₂O (5:1, v:v), r.t., Ar, 58-60%. f) 1) m-CPBA, DCM, 0 $^{\circ}$ C to rt; 2) 2-methoxy-4-(4-methylpiperazin-1-yl)aniline, TFA, 2-Butanol, 110 $^{\circ}$ C; 3) TFA, DCM, rt, 30-45%. g) acryloyl chloride, DIPEA, DCM, 0 $^{\circ}$ C, 60-77%.

4-((3-((Tert-butoxycarbonyl)amino)phenyl)amino)-2-(methylthio)pyrimidine-5-carboxylic acid (**S19**). 1 M NaOH (20 mL, 20 mmol) was slowly added into a solution of compound **S18** (4.0 g, 10 mmol) in 20 mL THF, then the mixture was heated at 50 °C for 4 h. The solution was acidified with 1M HCl after the solvent was partly removed under reduced pressure. The formed solid was collected by filtration and dried in a vacuum oven to give **S19** as a white solid (3.5 g, 95%). 1 H NMR (400 MHz, DMSO- d_6) δ 12.75 (br, 1 H), 9.34 (s, 1 H), 8.58 (s, 1 H), 7.78 (s, 1 H), 7.47 (d, 1 H, J = 8.0 Hz), 7.19 (t, 1 H, J = 8.0 Hz), 7.03 (d, 1 H, J = 8.0 Hz).

tert-Butyl (3-((5-(1H-benzo[d)[1,2,3]triazole-1-carbonyl)-2-(methylthio)pyrimidin-4-yl)amino)phenyl)carbamate (**\$20**). To a mixture of compound **\$19** (43.55 g, 115.7 mmol) and EDCI (19.1 g, 115.7 mmol) in dry DCM (400 mL) was added benzotriazole (11.91 g, 115.7 mmol). The reaction mixture was stirred at room temperature for 4h and then the mixture was purified by silica gel chromatography to afford **\$20** (46.96 g, 85%). ¹H NMR (400 MHz, DMSO- d_6) δ 9.76 (s, 1 H), 9.42 (s, 1 H), 8.77 (s, 1 H), 8.29 (t, 1H, J = 8.0 Hz), 7.84 (t, 1 H, J = 7.6 Hz), 7.65 (t, 1 H, J = 8.0 Hz), 7.27-7.20 (m, 2 H), 7.10 (d, 1 H, J = 8.0 Hz), 2.52 (s, 3 H), 1.47 (s, 9 H).

tert-Butyl (3-(5-hydroxy-2-(methylthio)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)carbamate (**S21**). A solution of potassium hexamethyldisilazide (KHMDS, 1M in THF, 87.8 mL, 87.8 mmol) was slowly added to a solution of EA (8.54 mL, 87.8 mmol) in dry THF (100 mL) under Ar at -78 °C. The reaction mixture was stirred at -78 °C for 0.5 h and then a solution of compound **S20** (16.76 g, 35.1 mmol) in dry THF (200 mL) was slowly added to the mixture during 0.5h. The resulting mixture was stirred at -78 °C for another hour and then slowly warmed to room temperature, stirred for 10h. The reaction was quenched with 1N HCl and then treated with 6N HCl until pH = 2. The resulting mixture was stirred for another hour, was then diluted with DCM and partitioned between water and DCM. The organic layer was washed with brine and dried over Na₂SO₄, filtered, concentrated, and purified by silica gel chromatography to afford **S21** (2.2 g, 15.7%). H NMR (400 MHz, DMSO- d_6) δ 12.15 (br, 1 H), 9.52 (s, 1 H), 8.89 (s, 1 H), 7.42-7.35 (s, 1 H), 6.85 (d, 1 H, J = 6.0 Hz), 5.84 (s, 1 H), 2.19 (s, 3 H), 1.46 (s, 9 H).

8-(3-((tert-Butoxycarbonyl)amino)phenyl)-2-(methylthio)-7-oxo-7,8-dihydropyrido[2,3-d] pyrimidin-5-yl trifluoromethanesulfonate (**S22**). To a solution of compound **S21** (2.2 g, 5.51 mmol) and Et₃N (2.30 mL, 11.02 mmol) in dry DCM (30 mL), Tf₂O (0.98 mL, 5.51 mmol) was added at -78 °C. The mixture was stirred at -78 °C for 2 hrs and then the reaction was treated with H₂O. The organic layer was separated and

washed with brine, dried over Na_2SO_4 , filtered, concentrated, and purified by silica gel chromatography to afford **S22** (2.43g , 82.9%). ¹H NMR (400 MHz, DMSO- d_6) δ 9.59 (s, 1 H), 8.88 (s, 1 H), 7.59 (s, 1 H), 7.42-7.40 (m, 2 H), 6.98-6.95 (m, 2 H), 1.46 (s, 9 H).

tert-butyl (3-(5-(dimethylamino)-2-(methylthio)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl) phenyl)carbamate (**10I**). To a solution of compound **S22** (668.0 mg, 1.25 mmol) in THF (15 mL) dimethylamine hydrochloride (153.5 mg, 1.88 mmol) and DIPEA (0.67 mL, 3.75 mmol) were added. The mixture was heated at $40\,^{\circ}\text{C}$ overnight. The solvent was then removed under reduced pressure. 15.0 mL of DCM was added to the resulting crude product. The organic layer was washed with water, brine, dried over Na₂SO₄, filtered, concentrated, and purified by silica gel chromatography to afford **10I** as a white solid (525.0 mg, 98.3%). ¹H NMR (400 MHz, CDCl₃) δ 8.80 (s, 1 H), 7.46 (s, 1 H), 7.39 (t, 1 H, J = 8.0 Hz), 7.29 (s, 1 H), 7.28 (s, 1 H), 6.91 (d, 1 H, J = 7.6 Hz), 6.64 (s, 1 H), 5.92 (s, 1 H), 2.15 (s, 3 H), 3.02 (s, 6 H), 1.49 (s, 9 H).

N-(3-(5-(dimethylamino)-2-((2-methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9l**). 1 H NMR (400 MHz, DMSO- d_6) δ 10.32 (s, 1 H), 8.78 (s, 1 H), 8.03 (s, 1 H), 7.85 (d, 1 H, J = 6.4 Hz), 7.54(s, 1 H), 7.48(t, 1 H, J = 8.0 Hz), 7.24(d, 1 H, J = 8.8 Hz), 6.93 (d, 1 H, J = 7.6 Hz), 6.51 (s, 1 H), 6.43 (dd, 1 H, J = 10.0, 16.8 Hz), 6.23-6.27 (m, 1 H), 5.99 (br, 1 H), 5.78-5.75 (m, 1 H), 5.59 (s, 1 H), 3.77 (s, 1 H), 3.00 (m, 10 H), 2.43 (m, 4 H), 2.22 (s, 3 H). 13 C NMR (125 MHz, Acetic- d_4) δ 166.5, 166.1, 158.7, 158.4, 157.5, 156.9, 150.7, 146.9, 140.4, 138.2, 132.0, 130.9, 129.0, 126.0, 122.7, 122.0, 121.7, 109.5, 103.6, 101.6, 56.4, 54.0, 48.2, 43.7, 43.5. HRMS (ESI) for C₃₀H₃₅N₈O₃ [M + H]⁺, calcd: 555.2827, found: 555.2824.

tert-Butyl (3-(2-(methylthio)-7-oxo-5-phenylpyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl) carbamate (**10m**). Compound **S22** (0.61 g, 1.14 mmol), phenylboronic acid (153.6 mg, 1.26 mmol), Bis(triphenylphosphine)palladium(II) dichloride (40.53 mg, 5%) and K₂CO₃ (473.6 mg, 3.43 mmol) were added to a mixture solution of THF (10 mL) and H₂O (2 mL). The mixture was stirred, evacuated, refilled with argon and then stirred at room

temperature for 24 hrs. The mixture was diluted with DCM. The organic layer was washed with brine and dried over Na_2SO_4 , filtered, concentrated, and purified by silica gel chromatography to afford **10m** (300 mg, 58%). ¹H NMR (400 MHz, DMSO- d_6) δ 9.59(s, 1 H), 8.57 (s, 1 H), 7.63-7.59 (m, 5 H), 7.54 (s, 1 H), 7.47-7.41 (m, 2 H), 6.97 (d, 1 H, J = 5.6 Hz), 6.64 (s, 1 H), 2.20 (s, 3 H), 1.47 (s, 9 H).

N-(3-(2-((2-Methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-7-oxo-5-phenylpyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide (**9m**). 1 H NMR (400 MHz, DMSO- d_{6}) δ 10.45 (s, 1 H), 8.42 (s, 1 H), 8.27 (s, 1 H), 7.90 (s, 1 H), 7.64-7.51 (m, 7 H), 7.28 (d, 1 H, J = 8.8 Hz), 7.06 (d, 1 H, J = 7.6 Hz), 6.57 (s, 1 H), 6.48 (dd, 1 H, J = 10.0, 16.8 Hz), 6.38 (s, 1 H), 6.29-6.25 (m, 1 H), 6.03 (br, 1 H), 5.79-5.76 (m, 1 H), 3.78 (s, 3 H), 3.12-2.97 (m, 4 H), 2.67 (m, 4 H). 13 C NMR (125 MHz, DMSO- d_{6}) δ 163.3, 161.9, 157.3, 156.9, 149.5, 140.0, 137.1, 134.8, 131.8, 129.5, 129.4, 129.0, 128.7, 127.2, 124.0, 119.6, 118.9, 116.5, 106.5, 105.0, 100.1, 55.8, 52.9, 46.7. HRMS (ESI) for $C_{34}H_{33}N_{7}O_{3}$ [M + H]⁺, calcd: 588.2718, found: 588.2717.

N-(3-(5-([1,1'-Biphenyl]-4-yl)-2-((2-methoxy-4-(4-methylpiperazin-1-yl)phenyl)amino)-7-oxopyrido[2,3-d]pyrimidin-8(7H)-yl)phenyl)acrylamide ($\mathbf{9n}$). 1 H NMR (400 MHz, Acetic- d_4) δ 8.70 (s, 1H), 8.15 (d, 1 H, J = 8.0 Hz), 7.88 (d, 2 H, J = 8.4 Hz), 7.75 (d, 2 H, J = 7.6 Hz), 7.71 (d, 2 H, J = 8.0 Hz), 7.64-7.59 (m, 2 H), 7.52-7.39 (m, 4 H), 7.20 (d, 1 H, J = 7.6 Hz), 6.77 (s, 1 H), 6.59 (d, 1 H, J = 2.0 Hz), 6.45 (d, 2 H, J = 5.6 Hz), 6.21 (d, 1 H, J = 8.8 Hz), 5.80 (t, 1 H, J = 5.6 Hz), 3.85 (s, 3 H), 3.71 (m, 4 H), 3.22 (m, 4 H), 2.95 (s, 3 H). 13 C NMR (125 MHz, DMSO- d_6) δ 163.3, 161.9, 157.3, 156.9, 149.1, 140.0, 139.3, 137.1, 133.8, 131.7, 129.6, 129.4, 129.1, 127.9, 127.2, 126.8, 123.9, 119.6, 118.8, 116.3, 106.1, 104.9, 99.6, 55.7, 54.6, 48.5, 45.8. HRMS (ESI) for $C_{40}H_{37}N_7O_3$ [M + H]+, calcd: 664.3031, found: 664.3028.

Table S1. Purity of the synthesized compounds 9.

Compd.	MeOH:H ₂ O (v:v)	Ret. time (min)	Purity (%)
9a	80:20	6.09	98.9
9b	85:15	5.22	99.3
9c	85:15	6.67	98.9
9d	85:15	8.68	96.9
9e	85:15	6.82	98.9
9f	85:15	5.50	98.8
9g	75:25	10.68	98.7
9h	85:15	6.96	95.0
9i	85:15	6.32	96.3
9 j	85:15	6.67	96.9
9k	85:15	7.37	98.5
91	85:15	5.29	99.4
9m	85:15	5.50	97.1
9n	85:15	13.35	96.7
90	75:25	10.66	97.8
9р	85:15	4.48	97.5
9q	85:15	4.67	96.8
9r	85:15	5.97	97.4
9s	85:15	6.88	97.9
9t	85:15	6.30	98.2

Method:Dionex Summit, Column: Diamonsil C18, $5.0\mu m$, 4.6×250 mm (Agilent Technologies); detector: PDA-100 photodiode array; pump: p-680A). A flow rate of 1.0 mL/min was used with mobile phase of MeOH in H2O with 0.1% modifier (ammonia, v/v)

Enzyme-linked immunosorbent assay (ELISA) kinase assay

Poly (Glu, Tyr)_{4:1} (Sigma, St. Louis, MO) (20 μg/mL) was precoated in 96-well ELISA plates as the substrate. The active kinases were incubated with indicated compounds in 1×reaction buffer (50 mmol/L HEPES pH 7.4, 20 mmol/L MgCl₂, 0.1 mmol/L MnCl₂, 0.2 mmol/L Na₃VO₄, 1 mmol/L DTT) containing 5 μmol/L ATP at 37 °C for 1 h. After incubation, the wells was washed with PBS, and then incubated with antiphosphotyrosine (PY99) antibody (Santa Cruz, CA) and horseradish peroxidase (HRP) – conjugated secondary antibody in sequence. The wells were visualized using ophenylenediamine (OPD) and read using a multiwell spectrophotometer (VERSAmaxTM, Molecular Devices, Sunnyvale, CA, USA) at 492 nm.

Kinase profiling study and K_d determination.

The kinase profiling study and K_d determination were conducted using the DiscoveRx Kinome screening platform (http://www..com).

For most assays, kinase-tagged T7 phage strains were prepared in an E. coli host derived from the BL21 strain. E. coli were grown to log-phase and infected with T7 phage and incubated with shaking at 32°C until lysis. The lysates were centrifuged and filtered to remove cell debris. The remaining kinases were produced in HEK-293 cells and subsequently tagged with DNA for qPCR detection. Streptavidin-coated magnetic beads were treated with biotinylated small molecule ligands for 30 minutes at room temperature to generate affinity resins for kinase assays. The liganded beads were blocked with excess biotin and washed with blocking buffer (SeaBlock (Pierce), 1% BSA, 0.05% Tween 20, 1 mM DTT) to remove unbound ligand and to reduce non-specific binding. Binding reactions were assembled by combining kinases, liganded affinity beads, and test compounds in 1x binding buffer (20% SeaBlock, 0.17x PBS, 0.05% Tween 20, 6 mM DTT). All reactions were performed in polystyrene 96-well plates in a final volume of 0.135 ml. The assay plates were incubated at room temperature with shaking for 1 hour and the affinity beads were washed with wash buffer (1x PBS, 0.05% Tween 20). The beads were then re-suspended in elution buffer (1x PBS, 0.05% Tween 20, 0.5 μM non-biotinylated affinity ligand) and incubated at room temperature with shaking for 30 minutes. The kinase concentration in the eluates was measured by qPCR.

An 11-point 3-fold serial dilution of each test compound was prepared in 100% DMSO at 100x final test concentration and subsequently diluted to 1x in the assay (final DMSO concentration = 1%). Most Kds were determined using a compound top concentration = 30,000 nM. If the initial Kd determined was < 0.5 nM (the lowest concentration tested), the measurement was repeated with a serial dilution starting at a lower top concentration. A Kd value reported as 40,000 nM indicates that the Kd was determined to be >30,000 nM.

Binding constants (Kds) were calculated with a standard dose-response curve using the Hill equation:

Response = Background +
$$\frac{\text{Signal - Background}}{1 + (Kd^{Hill Slope} / Dose^{Hill Slope})}$$

The Hill Slope was set to -1.

Curves were fitted using a non-linear least square fit with the Levenberg-Marquardt algorithm.

Western blot analysis

Western blotting was conducted as previously reported. In brief, cells were seeded to six-well plates and incubated overnight, and then starved with serum-free medium for 24 h. Starved cells were treated with or without different concentrations of compounds for 2 h and then stimulated with or without EGF (50 ng/ml) for 15 min. Cell samples were then lysed in 1×SDS lysis buffer. Proteins were resolved by SDS-PAGE and transferred onto polyvinylidene difluoride membranes (Millipore), which were blocked with nonfat milk and hybridized with specific primary antibodies. The bands were visualized using an enhanced chemiluminescence reagent (GE Healthcare) after hybridization with a HRP-conjugated secondary antibody. For the washing-out experiments, cells were seeded in 96-well plates and incubated overnight. After the treatment of compounds at indicated concentration for 2 h, cells were collected at indicated time points and processed for Western blot analysis.

Irreversibility assessments for compounds

The mobility shift assay were used to examine irreversible binding mode of compound **9f** with EGFR L858R/T790M. Compound **4** as used as positive control in this assay. The EGFR L858R/T790M protein and compound **9f** or **4** are incubated together for 30 min, and then diluted into standard enzyme assay mixtures containing ATP and substrate peptide (5-FAM-EEPLYWSFPAKKK-CONH₂, 1.5 M). Meanwhile, control reactions with inhibitor added to the reaction mix without being incubated with enzyme were also conducted. The microplate was then placed in the EZ Reader (Caliper Life Sciences, MA) and wells were continuously sampled for 120 min.

In general, preincubation of a reversible inhibitor with the protein results in similar kinase activity with that of none preinbubation control, whereas, preincubation of an irreversible inhibitor will dramatically decreased the kinase activity compared with control. The results in Figure S1 showed that preincubation of the EGFR^{L858R/T790M} protein with **9f** caused an obvious decrease of kinase activity relative to the non-incubation sample, which is quite similar to that of compound **4**, supporting the irreversible binding of **9f** with the protein.

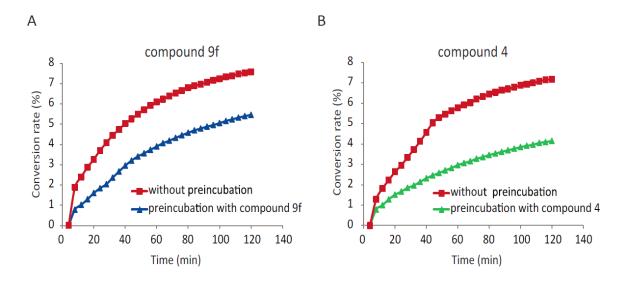


Fig. S1. 9f irreversibly binds with EGFR^{L858R/T790M}.

Cell lines and proliferation assay

NCI-H1975, A431 cancer cells and, GES-1, LO2, WI-38 and MCF-10A normal cell lines were obtained from American Type Culture Collection (ATCC, Rockville, MD) and maintained in strict accordance with the instruction and established procedures. The cell proliferation assay was evaluated using SRB (Sulforhodamine B) assay treated for 72 hrs with different concentrations of compounds.

Table S2. Anti-proliferative activities of the new EGFR T790M mutant inhibitors. ^a

-				
Cods	Anti-proliferation (IC ₅₀ , nM)			
- Sp. 11	H1975	A431	Selectivity Ratio ^b	
9a	3.7 ± 2.7	<16	<5.0	
9e	28.9 ± 11.6	236.8 ± 165.2	8.2	
9f	2.8 ± 2.0	865.6 ± 426.1	311.3	
9i	24.8 ± 17.2	4250.8 ± 2602.0	171.5	
9n	>10000	>50000	NA	
9p	6.8 ± 0.4	1104.0 ± 589.8	162.8	
9r	5.3 ± 1.2	2186.5 ± 575.4	415.1	
9t	11.6 ± 10.2	7468.7 ± 948.4	645.2	

9u	16.1 ± 10.0	562.9 ± 124.6	35.0
9v	21.0 ± 9.4	687.4 ± 35.3	32.7
3	69.8 ± 43.9	16.7 ± 4.0	0.2
4	50.1 ± 23.2	5550.4 ± 1680.3	110.7
5	43.1 ± 12.1	10117.1 ± 2919.5	274.0

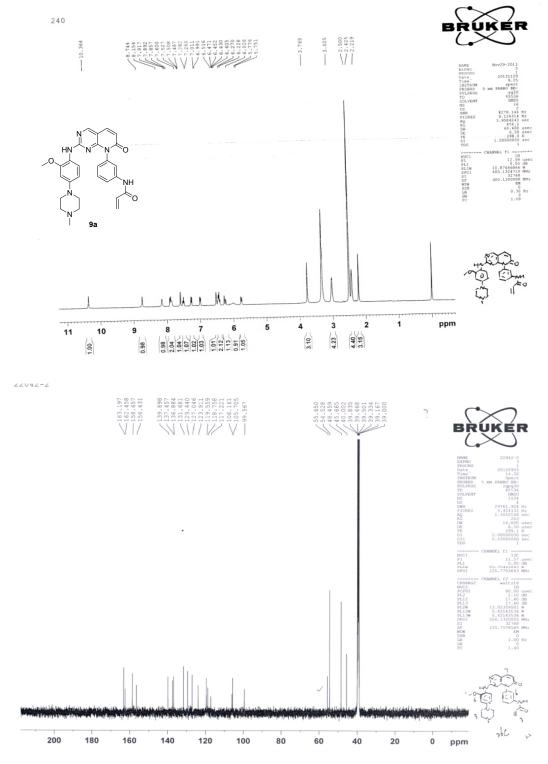
^a The anti-proliferative activities of the compounds were evaluated using SRB assay. The data were means from at least four independent experiments. ^b Ratio = IC_{50} (A431) / IC_{50} (H1975). NA means not available.

Table S3. Anti-proliferative activities of **9f** against a panel of normal cell lines.

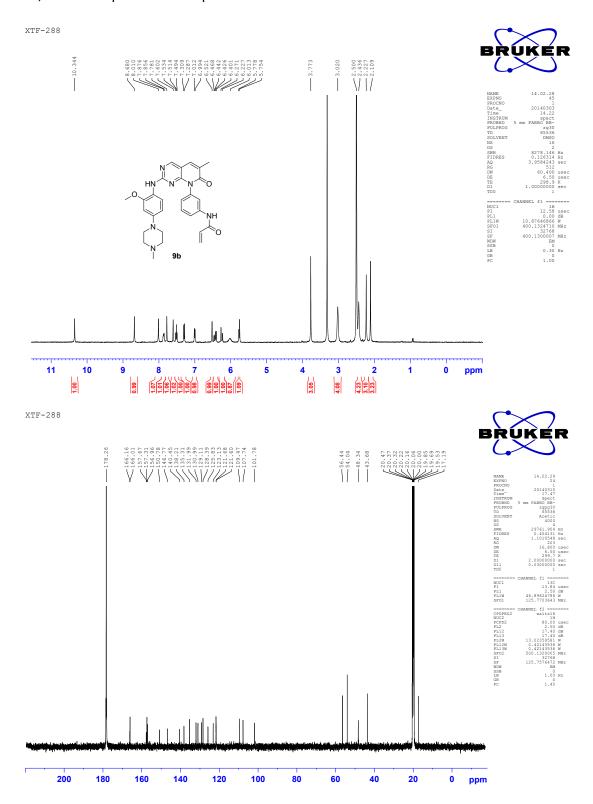
Cpds	Anti-proliferation (IC ₅₀ , μM)			
Op 0.0	GES-1	LO2	WI-38	MCF-10A
9f	4.591±0.248	2.493±0.683	7.231±0.489	2.016±0.316

¹H NMR and ¹³C NMR spectra of compounds 9

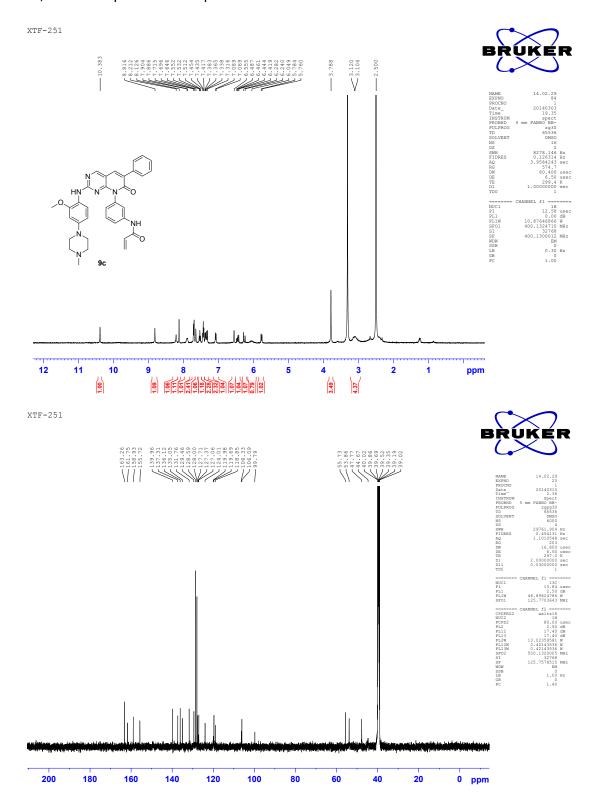
¹H, ¹³C NMR spectra of compound **9a**



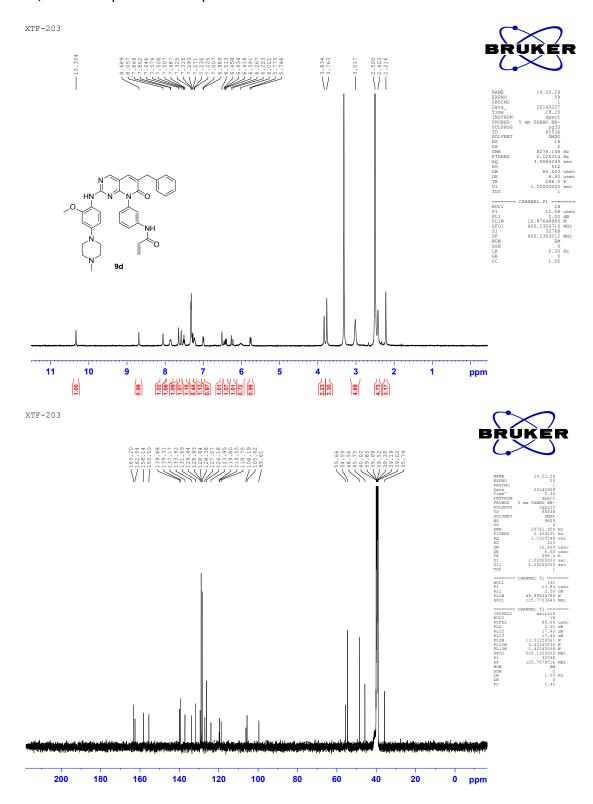
¹H, ¹³C NMR spectra of compound **9b**



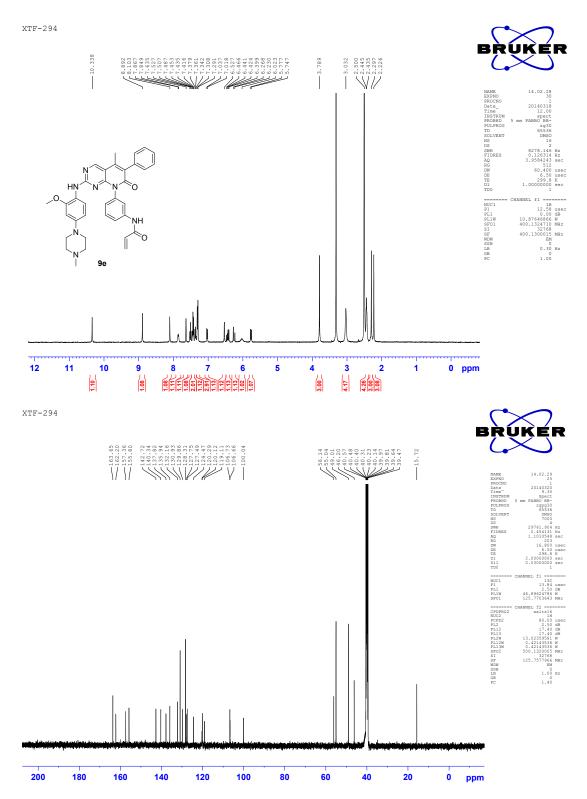
¹H, ¹³C NMR spectra of compound **9c**



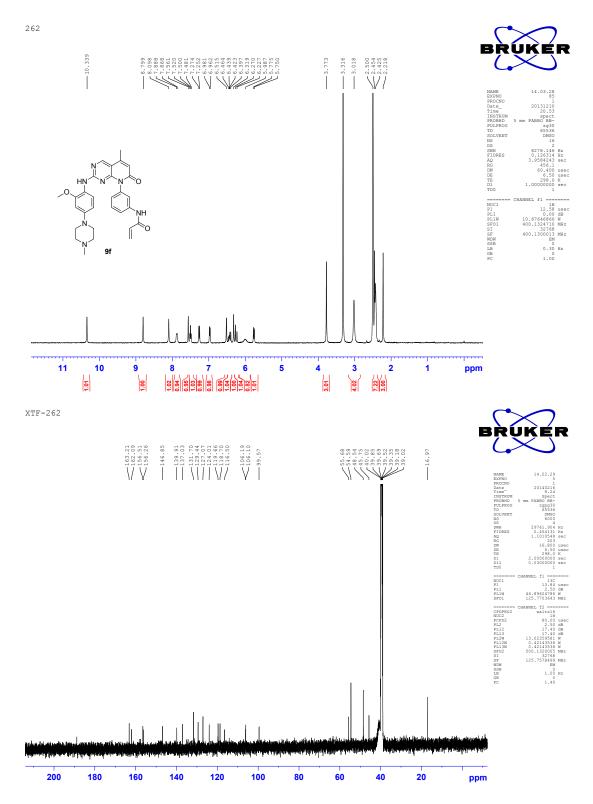
¹H, ¹³C NMR spectra of compound **9d**



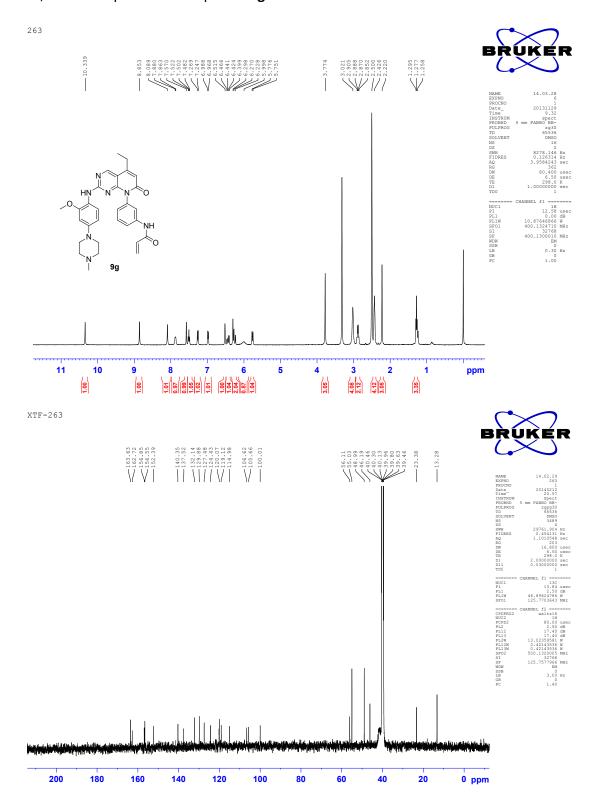
¹H, ¹³C NMR spectra of compound **9e**



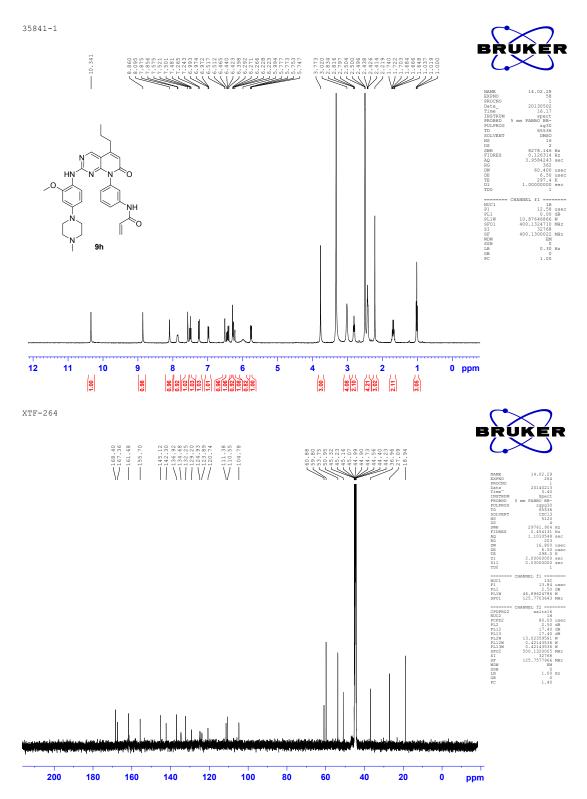
¹H, ¹³C NMR spectra of compound **9f**



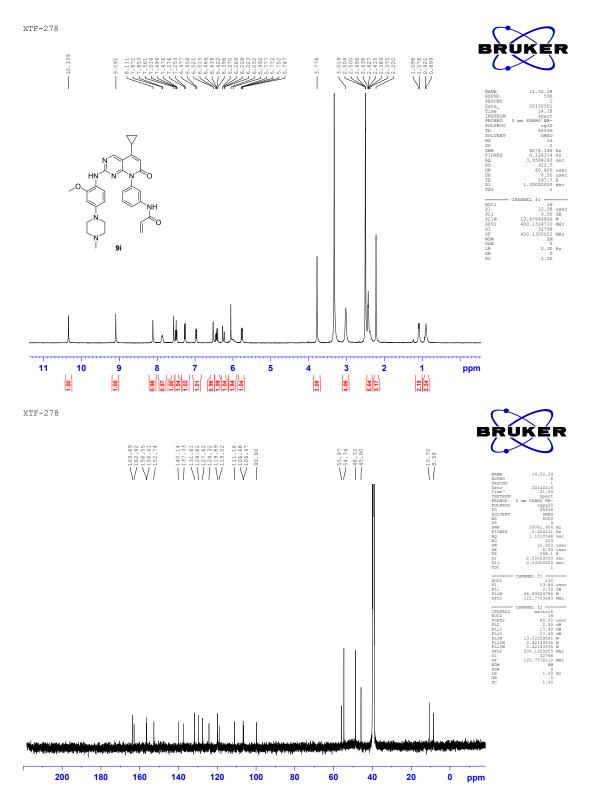
¹H, ¹³C NMR spectra of compound **9g**



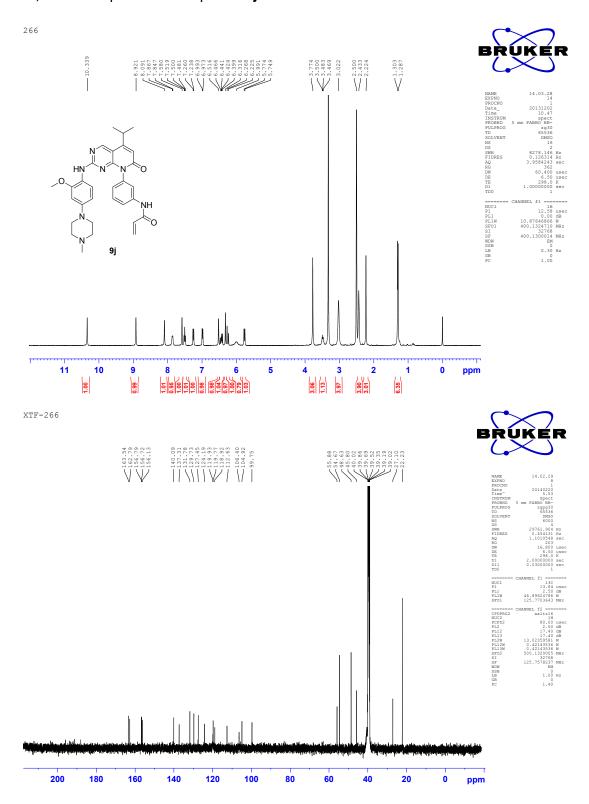
¹H, ¹³C NMR spectra of compound **9h**



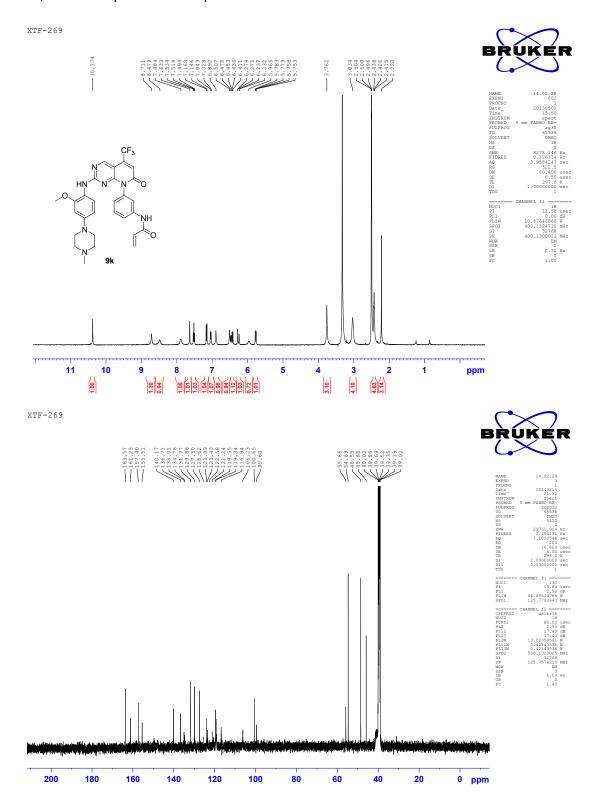
¹H, ¹³C NMR spectra of compound **9i**



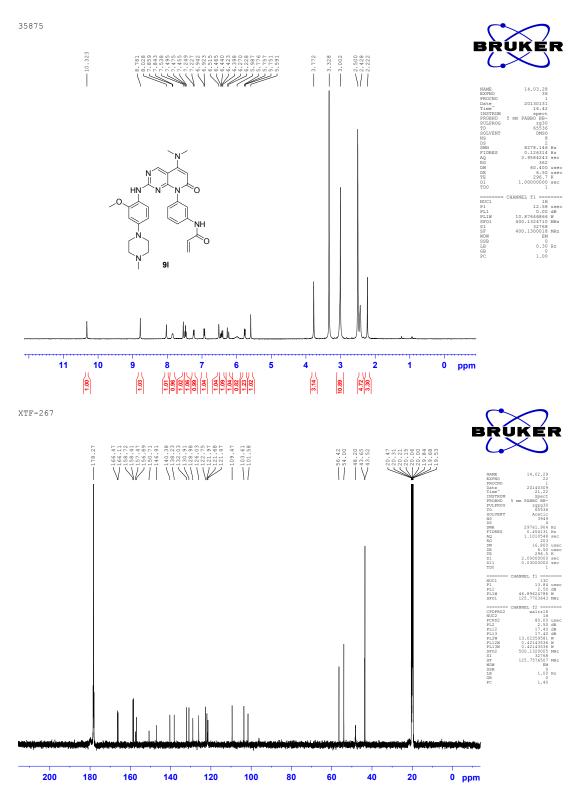
¹H, ¹³C NMR spectra of compound **9j**



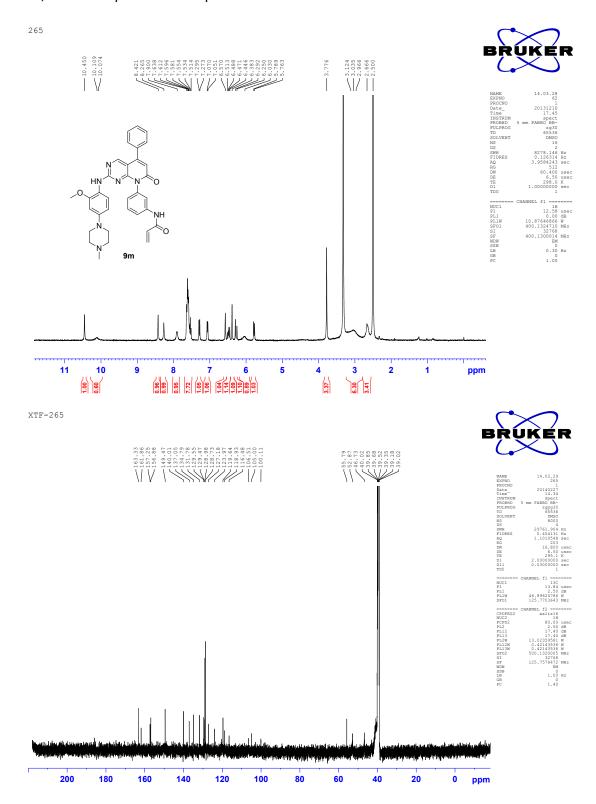
¹H, ¹³C NMR spectra of compound **9k**



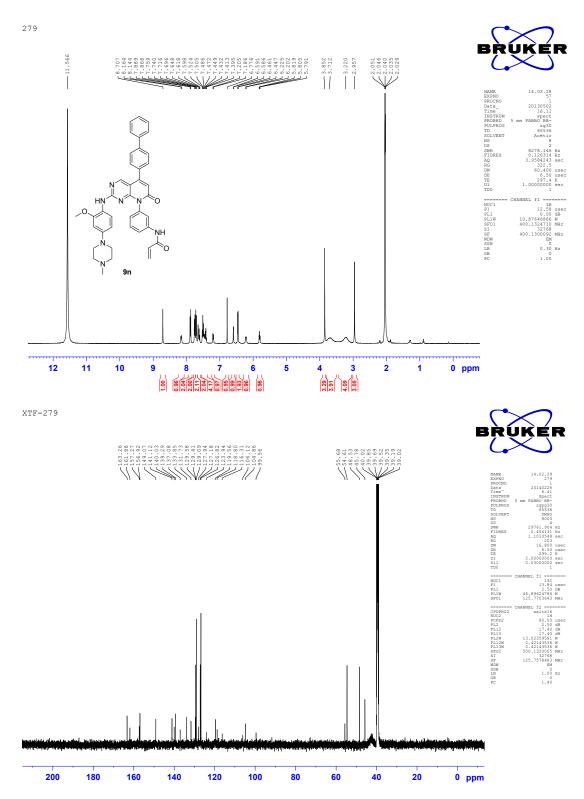
¹H, ¹³C NMR spectra of compound **9I**



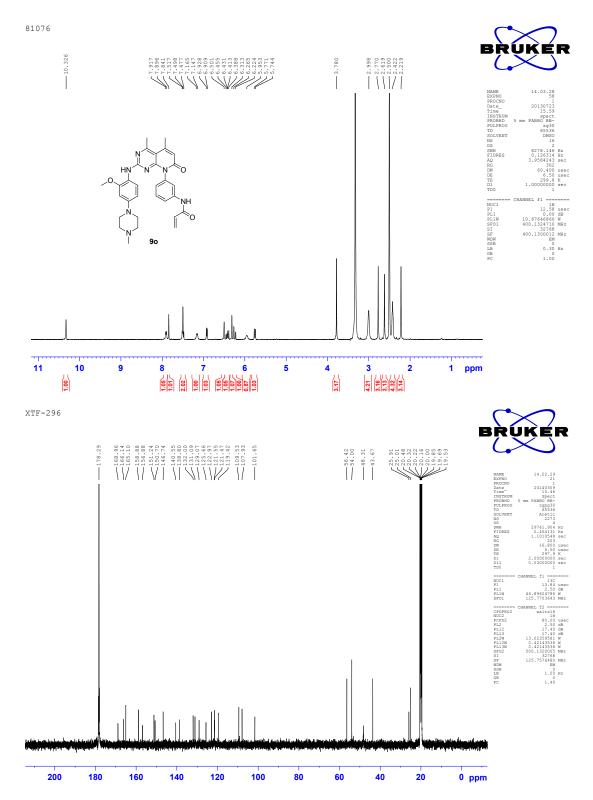
¹H, ¹³C NMR spectra of compound **9m**



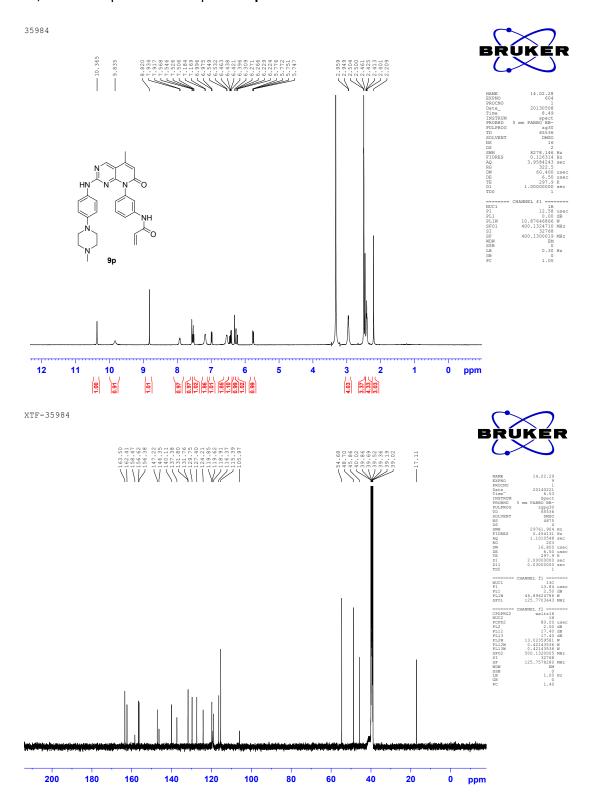
¹H, ¹³C NMR spectra of compound **9n**



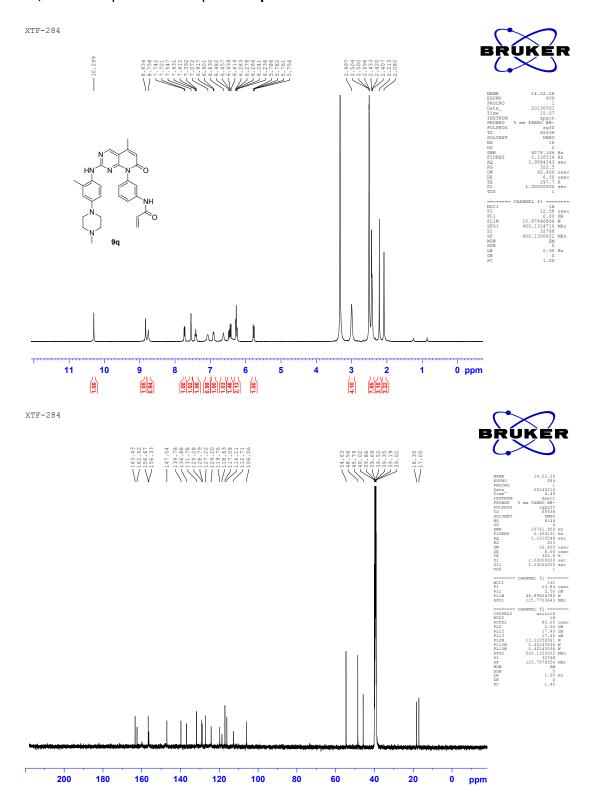
¹H, ¹³C NMR spectra of compound **90**



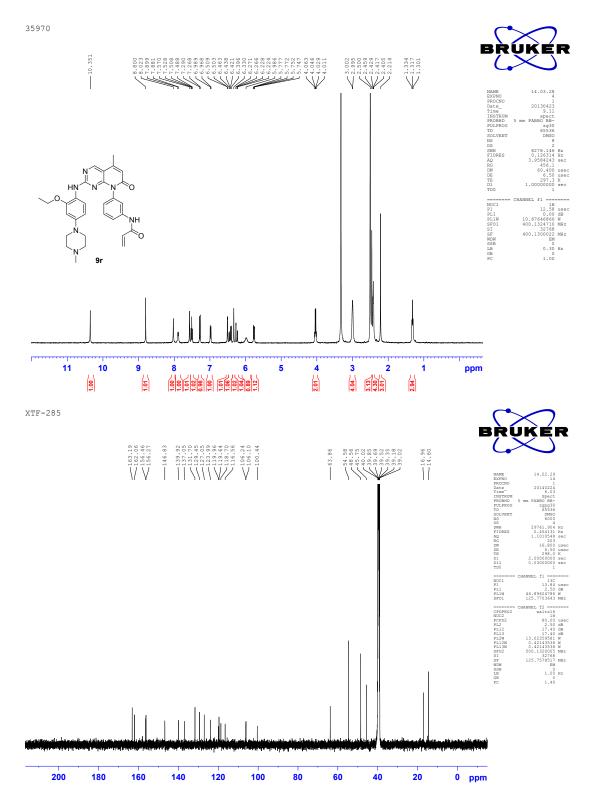
¹H, ¹³C NMR spectra of compound **9p**



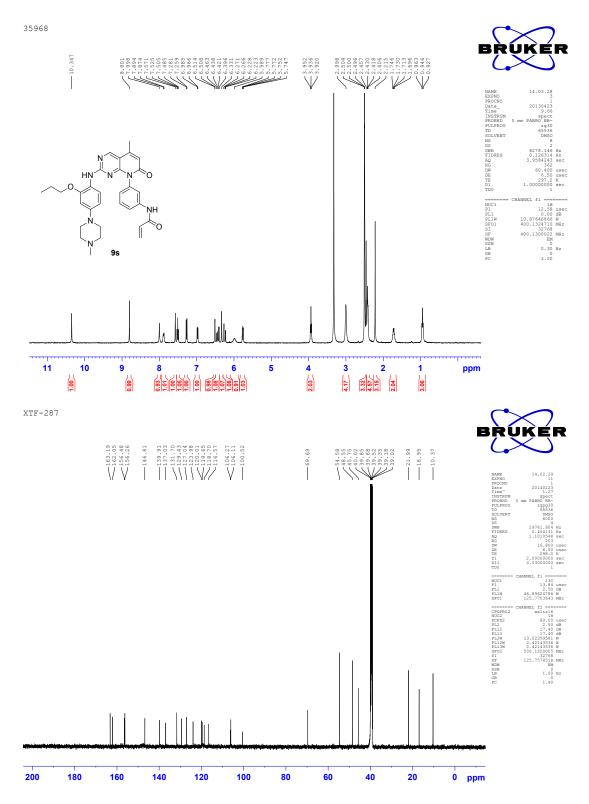
¹H, ¹³C NMR spectra of compound **9q**



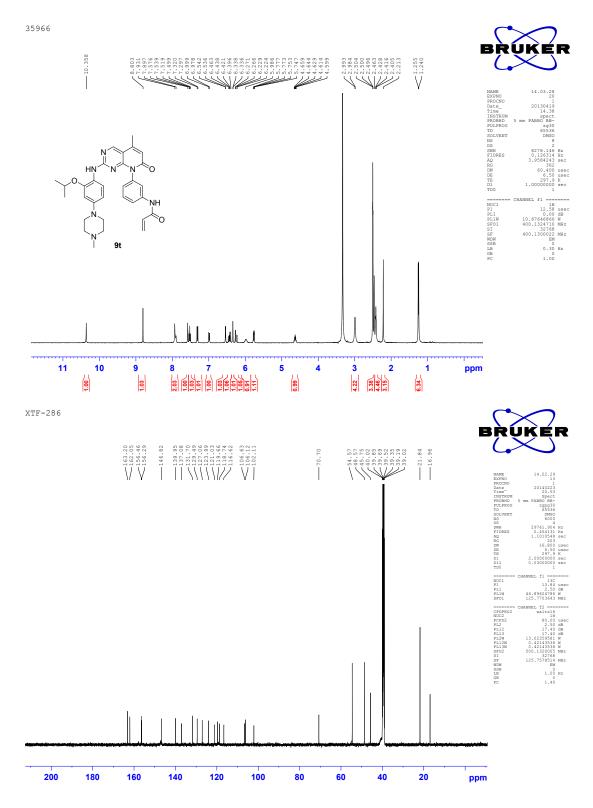
¹H, ¹³C NMR spectra of compound **9r**



¹H, ¹³C NMR spectra of compound **9s**



¹H, ¹³C NMR spectra of compound **9t**



The Kinase Profiling Results of Compound 9f

Target	XTF-262
Gene Symbol	%Ctrl @ 100nM
AAK1	100
ABL1(E255K)-phosphorylated	56
ABL1(F317I)-nonphosphorylated	87
ABL1(F317I)-phosphorylated	100
ABL1(F317L)-nonphosphorylated	74
ABL1(F317L)-phosphorylated	100
ABL1(H396P)-nonphosphorylated	100
ABL1(H396P)-phosphorylated	78
ABL1(M351T)-phosphorylated	84
ABL1(Q252H)-nonphosphorylated	75
ABL1(Q252H)-phosphorylated	84
ABL1(T315I)-nonphosphorylated	82
ABL1(T315I)-phosphorylated	82
ABL1(Y253F)-phosphorylated	100
ABL1-nonphosphorylated	65
ABL1-phosphorylated	88
ABL2	92
ACVR1	92
ACVR1B	96
ACVR2A	100
ACVR2B	88
ACVRL1	95
ADCK3	86
ADCK4	95
AKT1	76
AKT2	96
AKT3	100
ALK	92
ALK(C1156Y)	75
ALK(L1196M)	100
AMPK-alpha1	100
AMPK-alpha2	85
ANKK1	98
ARK5	100
ASK1	94
ASK2	84
AURKA	98
AURKB	82
AURKC	98
AXL	100
BIKE	94
BLK	68
BMPR1A	100
BMPR1B	91
BMPR2	99
BMX	83
BRAF	
	92
BRAF(V600E)	100

Target	XTF-262
Gene Symbol	%Ctrl @ 100nM
BRK	93
BRSK1	100
BRSK2	100
BTK	100
BUB1	93
CAMK1	100
CAMK1D	100
CAMK1G	100
CAMK2A	94
CAMK2B	100
CAMK2D	100
CAMK2G	100
CAMK4	100
CAMKK1	100
CAMKK2	100
CASK	68
CDC2L1	100
CDC2L2	100
CDC2L5	99
CDK11	98
CDK2	100
CDK3	76
CDK4-cyclinD1	100
CDK4-cyclinD3	89
CDK5	100
CDK7	83
CDK8	100
CDK9	
	99
CDKL1	97
CDKL2	100
CDKL3	92
CDKL5	78
CHEK1	96
CHEK2	98
CIT	92
CLK1	78
CLK2	100
CLK3	100
CLK4	81
CSF1R	100
CSF1R-autoinhibited	93
CSK	100
CSNK1A1	78
CSNK1A1L	100
CSNK1D	99
CSNK1E	100
CSNK1G1	93
CSNK1G2	100
	86

Target	XTF-262
Gene Symbol	%Ctrl @ 100nM
CSNK2A1	80
CSNK2A2	100
СТК	100
DAPK1	88
DAPK2	97
DAPK3	98
DCAMKL1	65
DCAMKL2	100
DCAMKL3	100
DDR1	84
DDR2	89
DLK	89
DMPK	75
DMPK2	91
DRAK1	90
DRAK2	100
DYRK1A	91
DYRK1B	100
DYRK2	90
EGFR	40
EGFR(E746-A750del)	70
EGFR(G719C)	77
	56
EGFR(G719S) EGFR(L747-E749del, A750P)	19
EGFR(L747-S752del, P753S)	33 27
EGFR(L747-T751del,Sins)	
EGFR(L858R)	28
EGFR(L858R,T790M)	0.3
EGFR(L861Q)	26
EGFR(S752-I759del)	47
EGFR(T790M)	0.2
EIF2AK1	100
EPHA1	100
EPHA2	80
EPHA3	100
EPHA4	99
EPHA5	94
EPHA6	100
EPHA7	100
EPHA8	100
EPHB1	100
EPHB2	98
EPHB3	100
EPHB4	96
EPHB6	79
ERBB2	75
ERBB3	100
ERBB4	33
ERK1	100

Gene Symbol %Ctrl @ 100nM ERK2 98 ERK3 100 ERK4 100 ERK8 100 ERN1 78 FAK 100 FER 93 FES 100 FGFR1 89 FGFR2 85 FGFR3 86 FGFR3(G697C) 100 FGFR4 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(K663Q) 100 FLT3(R844I) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3B 98 HASPIN 88 <th>Target</th> <th>XTF-262</th>	Target	XTF-262
ERK2 98 ERK3 100 ERK4 100 ERK5 100 ERK8 100 ERK8 100 ERN1 78 FAK 100 FER 93 FES 100 FGFR1 89 FGFR2 85 FGFR3 86 FGFR3(G697C) 100 FGFR 100 FGR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(K663Q) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HIPK1 96 HUNK 95 ICK 100 INSR 98 IKK-epsilon 100 INSR 98 IKK-epsilon 100 INSR 90		
ERK3 100 ERK4 100 ERK5 100 ERK8 100 ERN1 78 FAK 100 FER 93 FES 100 FGFR1 89 FGFR2 85 FGFR3 86 FGFR3(G697C) 100 FGFR 100 FGFR 100 FGFR 100 FGTR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(K663Q) 100 FLT3(N841I) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK1 96 HUNK 95 ICK 100 INSR 98 IKK-epsilon 100 INSR 90		
ERK4 100 ERK5 100 ERK8 100 ERN1 78 FAK 100 FER 93 FES 100 FGFR1 89 FGFR2 85 FGFR3 86 FGFR3(G697C) 100 FGR 100 FGR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 INSR 98 IKK-alpha 92 IKK-alpha 92 IKK-beta 100 INSR 80		
ERK5 100 ERK8 100 ERN1 78 FAK 100 FER 93 FES 100 FGFR1 89 FGFR2 85 FGFR3 86 FGFR3(G697C) 100 FGFR4 100 FGR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(ITD) 100 FLT3(ITD) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-apta 98 IKK-apta 98 IKK-apta 98 IKK-apta 98 IKK-apta 98 IKK-apta 99 IKK-apt		
ERK8 100 ERN1 78 FAK 100 FER 93 FES 100 FGFR1 89 FGFR2 85 FGFR3 86 FGFR3(G697C) 100 FGFR4 100 FGR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(K663Q) 100 FLT3(K8663Q) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HIPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80		
ERN1 78 FAK 100 FER 93 FES 100 FGFR1 89 FGFR2 85 FGFR3 86 FGFR3(G697C) 100 FGFR4 100 FGR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(K663Q) 100 FLT3(K663Q) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HIPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80		
FAK 100 FER 93 FES 100 FGFR1 89 FGFR2 85 FGFR3 86 FGFR3(G697C) 100 FGFR4 100 FGR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(K663Q) 100 FLT3(K663Q) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HIPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80		
FER 93 FES 100 FGFR1 89 FGFR2 85 FGFR3 86 FGFR3(G697C) 100 FGFR4 100 FGR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835H) 82 FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(K663Q) 100 FLT3(R834U) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HIPK1 96 HUNK 95 ICK 100 IGF1R 100 INSR 80		
FES 100 FGFR1 89 FGFR2 85 FGFR3 86 FGFR3(G697C) 100 FGFR4 100 FGR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R IKK-epsilon 100 INSR 80		
FGFR1 89 FGFR2 85 FGFR3 86 FGFR3 86 FGFR3(G697C) 100 FGFR4 100 FGR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 IKK-alpha 92 IKK-beta 98 IKK-epsilon 100 INSR 80		
FGFR2 85 FGFR3 86 FGFR3(G697C) 100 FGFR4 100 FGR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 IKK-epsilon 100 INSR 80		
FGFR3 FGFR3(G697C) FGFR4 FGFR4 FGR FGR FGR FLT3 FLT3 FLT3 FLT3 FLT3(D835H) FLT3(D835H) FLT3(ITD) FLT3(K663Q) FLT3(K663Q) FLT3(R834Q) FLT3(R834Q) FLT3-autoinhibited FLT4 FRK FYN FR FY FR FR FY FR FY FR		
FGFR3(G697C) 100 FGR4 100 FGR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80		
FGFR4 100 FGR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80		
FGR 100 FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80		
FLT1 98 FLT3 96 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(R841I) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80		
FLT3 96 FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(N841I) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80		100
FLT3(D835H) 82 FLT3(D835Y) 100 FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80	FLT1	98
FLT3(D835Y) 100 FLT3(K663Q) 100 FLT3(K663Q) 100 FLT3(R841I) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80		96
FLT3(ITD) 100 FLT3(K663Q) 100 FLT3(N841I) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80	FLT3(D835H)	82
FLT3(K663Q) 100 FLT3(N841I) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80	FLT3(D835Y)	100
FLT3(N841I) 100 FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80	FLT3(ITD)	100
FLT3(R834Q) 95 FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80	FLT3(K663Q)	100
FLT3-autoinhibited 94 FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-beta 98 IKK-epsilon 100 INSR 80	FLT3(N841I)	100
FLT4 79 FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80	FLT3(R834Q)	95
FRK 98 FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80	FLT3-autoinhibited	94
FYN 98 GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80	FLT4	79
GAK 100 GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-epsilon 100 INSR 80	FRK	98
GCN2(Kin.Dom.2,S808G) 79 GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-epsilon 100 INSR 80	FYN	98
GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-epsilon 100 INSR 80	GAK	100
GRK1 77 GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-epsilon 100 INSR 80	GCN2(Kin.Dom.2.S808G)	79
GRK4 86 GRK7 85 GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-epsilon 100 INSR 80		77
GRK7 GSK3A GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-epsilon INSR 80		
GSK3A 100 GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-epsilon 100 INSR 80		
GSK3B 98 HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-epsilon 100 INSR 80		
HASPIN 88 HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-epsilon 100 INSR 80		
HCK 100 HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-beta 98 IKK-epsilon 100 INSR 80		
HIPK1 75 HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-beta 98 IKK-epsilon 100 INSR 80		
HIPK2 80 HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-beta 98 IKK-epsilon 100 INSR 80		
HIPK3 87 HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-beta 98 IKK-epsilon 100 INSR 80		
HIPK4 100 HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-beta 98 IKK-epsilon 100 INSR 80		
HPK1 96 HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-beta 98 IKK-epsilon 100 INSR 80		
HUNK 95 ICK 100 IGF1R 95 IKK-alpha 92 IKK-beta 98 IKK-epsilon 100 INSR 80		
ICK 100 IGF1R 95 IKK-alpha 92 IKK-beta 98 IKK-epsilon 100 INSR 80		
IGF1R 95 IKK-alpha 92 IKK-beta 98 IKK-epsilon 100 INSR 80		
IKK-alpha 92 IKK-beta 98 IKK-epsilon 100 INSR 80		
IKK-beta 98 IKK-epsilon 100 INSR 80		95
IKK-epsilon 100 INSR 80	-	
INSR 80		98
	IKK-epsilon	100
INSRR 100	INSR	80
	INSRR	100

Target	XTF-262
Gene Symbol	%Ctrl @ 100nM
IRAK1	100
IRAK3	100
IRAK4	87
ITK	70
JAK1(JH1domain-catalytic)	100
JAK1(JH2domain-pseudokinase)	68
JAK2(JH1domain-catalytic)	74
JAK3(JH1domain-catalytic)	11
JNK1	73
JNK2	86
JNK3	99
KIT	97
KIT(A829P)	82
KIT(D816H)	83
KIT(D816V)	83
KIT(L576P)	87
KIT(V559D)	79
KIT(V559D,T670I)	95
KIT(V559D,V654A)	98
KIT-autoinhibited	80
LATS1	100
LATS2	91
LCK	96
LIMK1	96
LIMK2	100
LKB1	95
LOK	100
LRRK2	89
LRRK2(G2019S)	94
LTK	100
LYN	100
LZK	100
MAK	100
MAP3K1	79
MAP3K15	95
MAP3K2	73
MAP3K3	100
MAP3K4	100
MAP4K2	100
MAP4K3	100
MAP4K4	85
MAP4K5	92
MAPKAPK2	90
MAPKAPK5	100
MARK1	97
MARK2	79
MARK3	100
MARK4	92
MAST1	100

Gene Symbol %Ctrl @ 100nM MEK1 100 MEK2 76 MEK3 73 MEK4 100 MEK5 76 MEK6 99 MELK 87 MERTK 77 MET 100 MET(M1250T) 73 MET(Y1235D) 100 MINK 100 MKK7 76 MKNK1 79 MKNK2 88 MLCK 100 MLK3 100 MLK2 100 MLK3 100 MRCKA 100 MRCKA 100 MRCKB 98 MST1 99 MST1R 100 MST2 100 MST3 100 MST4 74 MTOR 100 MUSK 85 MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK3 74 NEK1 189 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK6 100 NEK7 100 NEK8 98 NEK5 90 NEK6 100 NEK7 100 NEK6 100 NEK7 100 NEK7 100 NEK8 98 NEK5 90 NEK6 100 NEK7 100 NEK9 100 NIK 100	Torgot	XTF-262
MEK1 100 MEK2 76 MEK3 73 MEK4 100 MEK5 76 MEK6 99 MELK 87 MERTK 77 MET 100 MET(M1250T) 73 MET(Y1235D) 100 MINK 100 MKK7 76 MKNK1 79 MKNK2 88 MLCK 100 MLK1 100 MLK2 100 MLK3 100 MRCKA 100 MRCKB 98 MST1 99 MST1R 100 MST2 100 MST3 100 MST4 74 MTOR 100 MUSK 85 MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK3 74 NEK6 198 NEK1 89 NEK1 89 NEK1 190 NEK1 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK3 74 NEK6 100 NEK7 100 NEK7 100 NEK7 100 NEK7 100 NEK7 100 NEK6 100 NEK7 100 NEK7 100 NEK7 100 NEK7 100 NEK7 100 NEK7 100 NEK9 100 NIK 100	Care Cumbal	
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MLK2 100 MLK3 100 MRCKA 100 MRCKB 98 MST1 99 MST1R 100 MST2 100 MST3 100 MST4 74 MTOR 100 MUSK 85 MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK6 100 NEK7 100 NIK 100	MLCK	100
MLK3 100 MRCKA 100 MRCKB 98 MST1 99 MST1R 100 MST2 100 MST3 100 MST4 74 MTOR 100 MUSK 85 MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK6 100 NEK7 100 NIK 100	MLK1	100
MRCKA 100 MRCKB 98 MST1 99 MST1R 100 MST2 100 MST3 100 MST4 74 MTOR 100 MUSK 85 MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK1 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK7 100 NEK9 100 NIK 96	MLK2	100
MRCKB 98 MST1 99 MST1R 100 MST2 100 MST3 100 MST3 100 MST4 74 MTOR 100 MUSK 85 MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK1 100 NEK1 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK7 100 NEK9 100 NIK 96	MLK3	100
MST1 99 MST1R 100 MST2 100 MST3 100 MST4 74 MTOR 100 MUSK 85 MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK9 100 NIK 96	MRCKA	100
MST1R 100 MST2 100 MST3 100 MST3 100 MST4 74 MTOR 100 MUSK 85 MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK6 100 NEK7 100 NIK 96	MRCKB	98
MST2 100 MST3 100 MST4 74 MTOR 100 MUSK 85 MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK6 100 NEK7 100 NIK 96	MST1	99
MST3 100 MST4 74 MTOR 100 MUSK 85 MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK7 100 NEK7 100 NEK9 100 NIK 100 NIK 100 NIK 96	MST1R	100
MST4 74 MTOR 100 MUSK 85 MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK7 100 NEK9 100 NIK 100 NIK 100 NIK 100 NIK 96	MST2	100
MTOR 100 MUSK 85 MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK7 100 NIK 100 NIK 100 NIK 100 NIK 96		100
MUSK 85 MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK7 100 NEK9 100 NIK 100 NIK 100 NIK 100 NIK 96	MST4	74
MYLK 72 MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK6 100 NEK7 100 NIK 100 NIK 100 NIK 100 NIK 100 NIK 100 NIK 96	MTOR	100
MYLK2 96 MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK7 100 NIK 100 NIK 100 NIK 100 NIK 100 NIK 96	MUSK	85
MYLK4 100 MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK7 100 NEK9 100 NIK 100 NIM1 100 NLK 96	MYLK	72
MYO3A 100 MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK9 100 NIK 100 NIK 100 NIK 96	MYLK2	96
MYO3B 99 NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK9 100 NIK 100 NIM 100 NILK 96	MYLK4	100
NDR1 92 NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK9 100 NIK 100 NIM 100 NIM 96	MYO3A	100
NDR2 97 NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK9 100 NIK 100 NIM1 100 NLK 96	MYO3B	99
NEK1 100 NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK9 100 NIK 100 NIM1 100 NLK 96	NDR1	92
NEK10 88 NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK9 100 NIK 100 NIM1 100 NLK 96	NDR2	97
NEK11 89 NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK9 100 NIK 100 NIM1 100 NLK 96	NEK1	100
NEK2 100 NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK9 100 NIK 100 NIM1 100 NLK 96	NEK10	88
NEK3 74 NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK9 100 NIK 100 NIM1 100 NLK 96	NEK11	89
NEK4 98 NEK5 90 NEK6 100 NEK7 100 NEK9 100 NIK 100 NIM1 100 NLK 96	NEK2	100
NEK5 90 NEK6 100 NEK7 100 NEK9 100 NIK 100 NIM1 100 NLK 96	NEK3	74
NEK6 100 NEK7 100 NEK9 100 NIK 100 NIM1 100 NLK 96	NEK4	98
NEK7 100 NEK9 100 NIK 100 NIM1 100 NLK 96	NEK5	90
NEK9 100 NIK 100 NIM1 100 NLK 96	NEK6	100
NEK9 100 NIK 100 NIM1 100 NLK 96	NEK7	100
NIK 100 NIM1 100 NLK 96	NEK9	100
NIM1 100 NLK 96		
NLK 96		
	OSR1	100

Target	XTF-262
Gene Symbol	%Ctrl @ 100nM
p38-alpha	100
p38-beta	94
p38-delta	98
p38-gamma	76
PAK1	91
PAK2	94
PAK3	100
PAK4	90
PAK6	100
PAK7	100
PCTK1	90
PCTK2	100
PCTK3	100
PDGFRA	100
PDGFRB	95
PDPK1	100
PFCDPK1(P.falciparum)	86
PFPK5(P.falciparum)	88
PFTAIRE2	100
PFTK1	91
PHKG1	100
PHKG2	94
PIK3C2B	97
PIK3C2G	88
PIK3CA	81
PIK3CA(C420R)	71
PIK3CA(E542K)	80
PIK3CA(E545A)	49
PIK3CA(E545K)	66
PIK3CA(H1047L)	97
PIK3CA(H1047Y)	54
PIK3CA(I800L)	61
PIK3CA(M1043I)	100
PIK3CA(Q546K)	81
PIK3CB	96
PIK3CD	100
PIK3CG	70
PIK4CB	100
PIM1	88
PIM2	100
PIM3	98
PIP5K1A	80
PIP5K1C	100
PIP5K2B	100
PIP5K2C	100
PKAC-alpha	99
PKAC-beta	99
PKMYT1	83
PKN1	100
TANT	100

Gene Symbol %Ctrl @ 100nM PKN2 93 PKNB(M.tuberculosis) 100 PLK1 87 PLK2 55 PLK3 82 PLK4 82 PRKCD 88 PRKCE 100 PRKCH 95 PRKCI 100 PRKCQ 100 PRKD1 100 PRKD2 99 PRKD3 96 PRKG1 93 PRKG2 97 PRKR 81 PRKX 100 PRP4 92 PYK2 100 QSK 63 RAF1 85 RET 97 RET(M918T) 78 RET(W804L) 76 RET(W804M) 100 RIOK2 100 RIOK3 100 RIPK1 100 RIPK2 97 RIPK4 97	Target	XTF-262
PKN2 93 PKNB(M.tuberculosis) 100 PLK1 87 PLK2 55 PLK3 82 PLK4 82 PRKCD 88 PRKCE 100 PRKCH 95 PRKCI 100 PRKCQ 100 PRKD1 100 PRKD2 99 PRKD3 96 PRKG1 93 PRKG2 97 PRKX 100 PRPKX 100 PRYK2 100 QSK 63 RAF1 85 RET 97 RET(M918T) 78 RET(W804L) 76 RET(W804M) 100 RIOK2 100 RIOK3 100 RIOK4 97 RIPK4 97 RIPK5 100 ROCK1 97 ROCK2 82 ROS1 <th></th> <th></th>		
PLK1 87 PLK2 55 PLK3 82 PLK4 82 PRKCD 88 PRKCE 100 PRKCH 95 PRKCI 100 PRKCQ 100 PRKD1 100 PRKD2 99 PRKD3 96 PRKG1 93 PRKG2 97 PRKR 81 PRKX 100 PRP4 92 PYK2 100 QSK 63 RAF1 85 RET 97 RET(M918T) 78 RET(W804L) 76 RET(W804M) 100 RIOK2 100 RIOK3 100 RIPK4 97 RIPK4 97 RIPK5 100 ROCK1 97 ROCK2 82 ROS1 100 RPS6KA4(Kin.Dom.1-N-termina		
PLK1 87 PLK2 55 PLK3 82 PLK4 82 PRKCD 88 PRKCE 100 PRKCH 95 PRKCI 100 PRKCQ 100 PRKD1 100 PRKD2 99 PRKD3 96 PRKG1 93 PRKG2 97 PRKR 81 PRKX 100 PRP4 92 PYK2 100 QSK 63 RAF1 85 RET 97 RET(M918T) 78 RET(W804L) 76 RET(W804M) 100 RIOK2 100 RIOK3 100 RIPK4 97 RIPK4 97 RIPK5 100 ROCK1 97 ROCK2 82 ROS1 100 RPS6KA4(Kin.Dom.1-N-termina	PKNB(M.tuberculosis)	100
PLK3 82 PLK4 82 PRKCD 88 PRKCE 100 PRKCH 95 PRKCI 100 PRKCQ 100 PRKD1 100 PRKD2 99 PRKD3 96 PRKG1 93 PRKG2 97 PRKR 81 PRKX 100 PRP4 92 PYK2 100 QSK 63 RAF1 85 RET (M918T) 78 RET(W804L) 76 RET(W804M) 100 RIOK2 100 RIOK3 100 RIPK1 100 RIPK2 97 RIPK4 97 RIPK5 100 ROCK1 97 ROCK2 82 ROS1 100 RPS6KA4(Kin.Dom.2-C-terminal) 88 RPS6KA5(Kin.Dom.2-C-terminal) 100		87
PLK3 82 PLK4 82 PRKCD 88 PRKCE 100 PRKCH 95 PRKCI 100 PRKCQ 100 PRKD1 100 PRKD2 99 PRKD3 96 PRKG1 93 PRKG2 97 PRKR 81 PRKX 100 PRP4 92 PYK2 100 QSK 63 RAF1 85 RET (M918T) 78 RET(W804L) 76 RET(W804M) 100 RIOK2 100 RIOK3 100 RIPK1 100 RIPK2 97 RIPK4 97 RIPK5 100 ROCK1 97 ROCK2 82 ROS1 100 RPS6KA4(Kin.Dom.2-C-terminal) 88 RPS6KA5(Kin.Dom.2-C-terminal) 100	PLK2	55
PLK4 82 PRKCD 88 PRKCE 100 PRKCH 95 PRKCI 100 PRKCQ 100 PRKD1 100 PRKD2 99 PRKD3 96 PRKG1 93 PRKG2 97 PRKR 81 PRKX 100 PRP4 92 PYK2 100 QSK 63 RAF1 85 RET 97 RET(M918T) 78 RET(V804L) 76 RET(V804M) 100 RIOK1 90 RIOK2 100 RIOK3 100 RIPK4 97 RIPK5 100 ROCK1 97 ROCK2 82 ROS1 100 RPS6KA4(Kin.Dom.2-C-terminal) 88 RPS6KA5(Kin.Dom.1-N-terminal) 98 RPS6KA5(Kin.Dom.2-C-terminal)		
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ROCK1 97 ROCK2 82 ROS1 100 RPS6KA4(Kin.Dom.1-N-terminal) 88 RPS6KA4(Kin.Dom.2-C-terminal) 98 RPS6KA5(Kin.Dom.1-N-terminal) 80 RPS6KA5(Kin.Dom.2-C-terminal) 100 RSK1(Kin.Dom.1-N-terminal) 100 RSK2(Kin.Dom.2-C-terminal) 78 RSK2(Kin.Dom.2-C-terminal) 99 RSK3(Kin.Dom.1-N-terminal) 100 RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98	RIPK4	97
ROCK2 82 ROS1 100 RPS6KA4(Kin.Dom.1-N-terminal) 88 RPS6KA4(Kin.Dom.2-C-terminal) 98 RPS6KA5(Kin.Dom.1-N-terminal) 100 RSK1(Kin.Dom.1-N-terminal) 100 RSK1(Kin.Dom.2-C-terminal) 100 RSK2(Kin.Dom.1-N-terminal) 78 RSK2(Kin.Dom.2-C-terminal) 99 RSK3(Kin.Dom.1-N-terminal) 100 RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98	RIPK5	100
ROS1 100 RPS6KA4(Kin.Dom.1-N-terminal) 88 RPS6KA4(Kin.Dom.2-C-terminal) 98 RPS6KA5(Kin.Dom.1-N-terminal) 80 RPS6KA5(Kin.Dom.2-C-terminal) 100 RSK1(Kin.Dom.1-N-terminal) 100 RSK2(Kin.Dom.2-C-terminal) 78 RSK2(Kin.Dom.2-C-terminal) 99 RSK3(Kin.Dom.1-N-terminal) 100 RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98	ROCK1	97
RPS6KA4(Kin.Dom.1-N-terminal) 88 RPS6KA4(Kin.Dom.2-C-terminal) 98 RPS6KA5(Kin.Dom.1-N-terminal) 80 RPS6KA5(Kin.Dom.2-C-terminal) 100 RSK1(Kin.Dom.1-N-terminal) 100 RSK2(Kin.Dom.2-C-terminal) 78 RSK2(Kin.Dom.2-C-terminal) 99 RSK3(Kin.Dom.1-N-terminal) 100 RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98	ROCK2	82
RPS6KA4(Kin.Dom.2-C-terminal) 98 RPS6KA5(Kin.Dom.1-N-terminal) 80 RPS6KA5(Kin.Dom.2-C-terminal) 100 RSK1(Kin.Dom.1-N-terminal) 100 RSK2(Kin.Dom.2-C-terminal) 78 RSK2(Kin.Dom.2-C-terminal) 99 RSK3(Kin.Dom.1-N-terminal) 100 RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98	ROS1	100
RPS6KA5(Kin.Dom.1-N-terminal) 80 RPS6KA5(Kin.Dom.2-C-terminal) 100 RSK1(Kin.Dom.1-N-terminal) 100 RSK2(Kin.Dom.2-C-terminal) 78 RSK2(Kin.Dom.2-C-terminal) 99 RSK3(Kin.Dom.1-N-terminal) 100 RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98	RPS6KA4(Kin.Dom.1-N-terminal)	88
RPS6KA5(Kin.Dom.2-C-terminal) 100 RSK1(Kin.Dom.1-N-terminal) 100 RSK1(Kin.Dom.2-C-terminal) 100 RSK2(Kin.Dom.1-N-terminal) 78 RSK2(Kin.Dom.2-C-terminal) 99 RSK3(Kin.Dom.1-N-terminal) 100 RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98	RPS6KA4(Kin.Dom.2-C-terminal)	98
RSK1(Kin.Dom.1-N-terminal) 100 RSK1(Kin.Dom.2-C-terminal) 100 RSK2(Kin.Dom.1-N-terminal) 78 RSK2(Kin.Dom.2-C-terminal) 99 RSK3(Kin.Dom.1-N-terminal) 100 RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98	RPS6KA5(Kin.Dom.1-N-terminal)	80
RSK1(Kin.Dom.2-C-terminal) 100 RSK2(Kin.Dom.1-N-terminal) 78 RSK2(Kin.Dom.2-C-terminal) 99 RSK3(Kin.Dom.1-N-terminal) 100 RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98	RPS6KA5(Kin.Dom.2-C-terminal)	100
RSK2(Kin.Dom.1-N-terminal) 78 RSK2(Kin.Dom.2-C-terminal) 99 RSK3(Kin.Dom.1-N-terminal) 100 RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98	RSK1(Kin.Dom.1-N-terminal)	100
RSK2(Kin.Dom.1-N-terminal) 78 RSK2(Kin.Dom.2-C-terminal) 99 RSK3(Kin.Dom.1-N-terminal) 100 RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98	RSK1(Kin.Dom.2-C-terminal)	100
RSK3(Kin.Dom.1-N-terminal) 100 RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98	RSK2(Kin.Dom.1-N-terminal)	78
RSK3(Kin.Dom.1-N-terminal) 100 RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98		99
RSK3(Kin.Dom.2-C-terminal) 100 RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98		100
RSK4(Kin.Dom.1-N-terminal) 67 RSK4(Kin.Dom.2-C-terminal) 98	, ,	100
RSK4(Kin.Dom.2-C-terminal) 98	-	67
		98
		81

Target	XTF-262
Gene Symbol	%Ctrl @ 100nM
SBK1	86
SGK	100
SgK110	86
SGK2	99
SGK3	92
SIK	100
SIK2	100
SLK	99
SNARK	100
SNRK	96
SRC	84
SRMS	79
SRPK1	73
SRPK2	100
SRPK3	99
STK16	100
STK33	100
STK35	100
STK36	78
STK39	100
SYK	100
TAK1	88
TAOK1	100
TAOK2	49
TAOK3	83
TBK1	66
TEC	100
TESK1	100
TGFBR1	86
TGFBR2	100
TIE1	87
TIE2	97
TLK1	100
TLK2	93
TNIK	100
TNK1	100
TNK2	95
TNNI3K	100
TRKA	96
TRKB	86
TRKC	96
TRPM6	73
TSSK1B	100
TTK	94
TXK	74
TYK2(JH1domain-catalytic)	76
TYK2(JH2domain-pseudokinase)	100
TYRO3	97
ULK1	78

Target	XTF-262
Gene Symbol	%Ctrl @ 100nM
ULK2	83
ULK3	87
VEGFR2	84
VRK2	85
WEE1	98
WEE2	99
WNK1	84
WNK3	85
YANK1	100
YANK2	96
YANK3	100
YES	95
YSK1	95
YSK4	100
ZAK	100
ZAP70	89

Fig. S4. the kinase profiling results of compound 9f (XTF-262) at 100nM.