## Supporting Information:

## Fragment-Based Design of a Potent MAT2a

## Inhibitor and in Vivo Evaluation in an MTAP Null

## Xenograft Model

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## Additional figures:

MAT2b C-terminus


Fragment 2


Figure S1: Ligand-induced changes to the allosteric site of MAT2a, which is displayed viewed from solvent (top), and in cross section (bottom). Left: View of the allosteric site bound to MAT2b, with Phe333 (highlighted) constricting the width of the site. Right: The crystal structure of 2 (PDB 7bhs, $1.1 \AA$ ) shows two molecules of the ligand occupying the symmetric site. Ligand binding re-orients the side chain of Phe333, which re-shapes the site into a long and narrow cleft and partially occludes solvent access.


Figure S2: WaterMap analysis (see Computational Details for references) of $\mathbf{5}$ bound to MAT2a. The calculated excess free energies $\Delta G$ of water sites 14 are $+6.8,+4.1,+3.0$, and $+5.4 \mathrm{kcal} / \mathrm{mol}$, respectively.



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Figure S3: QM optimised structures of 4-aminoquinazolinones 5 (nonplanar), 19 (planar), and 20 (nonplanar). The calculated dihedral angles $\tau_{1}$ and $\tau_{2}$, defined to the left, are -169.7 and 16.1, 179.9 and 0.2 , and -173.1 and -13.0 degrees for $\mathbf{5}, \mathbf{1 9}$, and 20, respectively. For 5 bound to MAT2a, the corresponding experimental values of $\tau_{1}$ and $\tau_{2}$ are -169.8 and -11.4 degrees, as determined by X-ray crystallography (Figure 3).


Figure S4: Mouse plasma concentrations of $\mathbf{2 8}$ after a single SC dose at $100 \mathrm{mg} / \mathrm{kg}$ (blue circles), 14 once daily SC doses of $100 \mathrm{mg} / \mathrm{kg}$ (red triangles), and 14 once daily SC doses of $50 \mathrm{mg} / \mathrm{kg}$ (green diamonds).

## Equation S1:

$$
\text { Methionine }+ \text { ATP }+\mathrm{H}_{2} \mathrm{O} \rightarrow \mathrm{SAM}+\text { pyrophosphate }+ \text { phosphate }
$$

Table S1: Crystallographic data collection and refinement statistics. Values in parentheses are for the highest resolution shell.

| Protein Compound | $\begin{gathered} \hline \text { MAT2a } \\ 1 \\ \hline \end{gathered}$ | $\begin{gathered} \text { MAT2a } \\ 2 \end{gathered}$ | $\begin{gathered} \text { MAT2a } \\ 5 \end{gathered}$ | $\begin{gathered} \hline \text { MAT2a } \\ 26 \end{gathered}$ |
| :---: | :---: | :---: | :---: | :---: |
| PDB ID | 7BHR | 7BHS | 7BHT | 7BHU |
| Data collection |  |  |  |  |
| DLS beamline | 104 | 104 | 103 | 103 |
| Date | 2018-07-13 | 2018-11-30 | 2020-09-22 | 2019-05-10 |
| Detector | Pilatus 6M | Pilatus 6M | Eiger X 16M | Eiger X 16M |
| Wavelength ( A ) | 0.9750 | 0.9763 | 0.8969 | 0.9763 |
| Data processing |  |  |  |  |
| Software | DIALS/Aimless | XDS/Staraniso | XDS/Staraniso | XDS/Staraniso |
| Space group | 1222 | 1222 | 1222 | 1222 |
| Cell dimensions |  |  |  |  |
| $a, b, c(\AA)$ | 67.43 | 67.92 | 67.87 | 68.24 |
|  | 94.21 | 94.02 | 93.93 | 94.17 |
|  | 117.28 | 117.09 | 116.97 | 117.30 |
| $\alpha, \beta, Y\left({ }^{\circ}\right)$ | 90 | 90 | 90 | 90 |
|  | 90 | 90 | 90 | 90 |
|  | 90 | 90 | 90 | 90 |
| Resolution ( $\AA$ ) | 58.64-1.08 | 40.11-1.05 | 19.82-1.05 | 58.98-1.15 |
|  | (1.11-1.08) | (1.15-1.05) | (1.14-1.05) | (1.19-1.15) |
| $R_{\text {pim }}$ | 0.027 (0.529) | 0.058 (0.597) | 0.042 (0.570) | 0.025 (0.319) |
| Reflections, total | 892550 | 798752 | 1618327 | 677640 |
| Reflections, unique | 144857 (6068) | 132040 (6603) | 120837 (6042) | 119589 (5980) |
| // $\sigma$ / | 11.6 (1.1) | 9.4 (2.2) | 12.1 (1.8) | 19.1 (2.3) |
| Completeness, spherical (\%) | 91.2 (52.4) | 76.4 (16.9) | 70.3 (16.3) | 90.6 (47.2) |
| Completeness, ellipsoidal (\%) |  | 93.8 (49.1) | 94.9 (72.3) | 91.4 (49.1) |
| Multiplicity | 6.2 (3.9) | 6.0 (4.0) | 13.4 (14.4) | 5.7 (2.5) |
| CC 1/2 | 0.997 (0.626) | 0.998 (0.535) | 0.999 (0.736) | 0.999(0.741) |
| Refinement (Buster) |  |  |  |  |
| Resolution ( $\AA$ ) | 58.64-1.08 | 40.11-1.05 | 19.82-1.05 | 16.90-1.15 |
| $R_{\text {work }} / R_{\text {free }}$ | 0.167 / 0.183 | 0.170 / 0.185 | $0.156 / 0.175$ | $0.153 / 0.169$ |
| No. atoms |  |  |  |  |
| Protein | 2943 | 2939 | 2983 | 2945 |
| Inhibitor | 13 | 19 | 15 | 20 |
| SAM | 27 | 27 | 27 | 27 |
| Water | 322 | 286 | 400 | 357 |
| $B$-factors ( $\AA^{2}$ ) |  |  |  |  |
| Protein | 14.3 | 11.4 | 11.8 | 12.6 |
| Inhibitor | 13.9 | 16.7 | 11.3 | 10.8 |
| SAM | 14.3 | 10.8 | 9.3 | 12.7 |
| Water | 25.7 | 23.6 | 23.5 | 24.4 |
| R.m.s. deviations |  |  |  |  |
| Bond lengths ( $\AA$ ) | 0.010 | 0.010 | 0.009 | 0.010 |
| Bond angles ( ${ }^{\circ}$ ) | 1.100 | 1.110 | 1.102 | 1.093 |
| Model Validation (MolProbity) |  |  |  |  |
| Ramachandran favoured (\%) | 97.88 | 97.62 | 97.62 | 97.62 |
| Ramachandran allowed (\%) | 2.12 | 2.38 | 2.38 | 2.38 |
| Ramachandran outliers (\%) | 0.00 | 0.00 | 0.00 | 0.00 |
| Ramachandran Z-score | 0.34 | 0.10 | -0.15 | 0.12 |


| Protein Compound | $\begin{gathered} \hline \text { MAT2a } \\ 28 \\ \hline \end{gathered}$ | $\begin{gathered} \hline \text { MAT2a } \\ 29 \\ \hline \end{gathered}$ | $\begin{gathered} \hline \text { MAT2a } \\ 31 \\ \hline \end{gathered}$ |
| :---: | :---: | :---: | :---: |
| PDB ID | 7BHV | 7BHW | 7BHX |
| Data collection |  |  |  |
| DLS beamline | 103 | 103 | 103 |
| Date | 2019-04-26 | 2019-05-10 | 2020-09-22 |
| Detector | Eiger X 16M | Eiger X 16M | Eiger X 16M |
| Wavelength ( A ) | 0.9763 | 0.9763 | 0.8153 |
| Data processing |  |  |  |
| Software | XDS/Staraniso | XDS/Staraniso | XDS/Staraniso |
| Space group | 1222 | 1222 | 1222 |
| Cell dimensions |  |  |  |
| $a, b, c(\AA)$ | 68.18 | 68.58 | 68.04 |
|  | 93.85 | 94.20 | 94.03 |
|  | 117.07 | 117.40 | 117.05 |
| $\alpha, \beta, y\left({ }^{\circ}\right)$ | 90 | 90 | 90 |
|  | 90 | 90 | 90 |
|  | 90 | 90 | 90 |
| Resolution ( $\AA$ ) | 73.23-1.16 | 59.22-1.16 | 58.83-1.08 |
|  | (1.20-1.16) | (1.21-1.16) | (1.17-1.08) |
| $R_{\text {pim }}$ | 0.019 (0.110) | 0.021 (0.271) | 0.037 (0.537) |
| Reflections, total | 666288 | 681000 | 1654788 |
| Reflections, unique | 118844 (5943) | 113536 (5678) | 121124 (6056) |
| / / $\sigma$ / | 27.4 (4.6) | 21.5 (2.3) | 13.4 (1.8) |
| Completeness, spherical (\%) | 91.9 (49.8) | 88.1 (40.8) | 75.6 (17.4) |
| Completeness, ellipsoidal (\%) | 92.2 (50.4) | 90.4 (46.9) | 94.0 (68.5) |
| Multiplicity | 5.6 (2.1) | 6.0 (3.4) | 13.7 (13.7) |
| CC 1/2 | 0.999 (0.980) | 1.000 (0.854) | 1.000 (0.743) |
| Refinement (Buster) |  |  |  |
| Resolution ( $\AA$ ) | 16.98-1.08 | 16.90-1.16 | 15.80-1.08 |
| $R_{\text {work }} / R_{\text {free }}$ | 0.149 / 0.157 | 0.156 / 0.167 | $0.158 / 0.173$ |
| No. atoms |  |  |  |
| Protein | 2945 | 2950 | 2977 |
| Inhibitor | 20 | 22 | 21 |
| SAM | 27 | 27 | 27 |
| Water | 357 | 342 | 375 |
| $B$-factors ( $\AA^{2}$ ) |  |  |  |
| Protein | 12.6 | 14.8 | 13.1 |
| Inhibitor | 10.8 | 11.5 | 11.1 |
| SAM | 12.7 | 14.4 | 10.7 |
| Water | 24.4 | 26.2 | 24.0 |
| R.m.s. deviations |  |  |  |
| Bond lengths ( A ) | 0.010 | 0.010 | 0.010 |
| Bond angles ( ${ }^{\circ}$ ) | 1.078 | 1.103 | 1.080 |
| Model Validation (MolProbity) |  |  |  |
| Ramachandran favoured (\%) | 97.62 | 98.15 | 97.88 |
| Ramachandran allowed (\%) | 2.38 | 1.85 | 2.12 |
| Ramachandran outliers (\%) | 0.00 | 0.00 | 0.00 |
| Ramachandran Z-score | 0.04 | 0.12 | -0.11 |

## Compound Synthesis

Flash column chromatography was carried out using self-packed silica cartridges and C18flash chromatography was performed with a combine flash (Agela MP 200) using C18 cartridges from Agela.
Preparative reverse phase HPLC was performed on a Waters instrument (2545, 2767 and 2489) fitted with a QDa or SQ Detector 2 ESCi mass spectrometers and a Waters X-Bridge or Waters XSelect or Waters SunFire reverse-phase column (C-18, $5 \mu \mathrm{~m}, 30 \mathrm{~mm}$ diameter and 150 mm length with a flow rate of $60 \mathrm{ml} / \mathrm{min}$ or C-18, $5 \mu \mathrm{~m}, 19 \mathrm{~mm}$ diameter and 250 mm length with a flow rate of $25 \mathrm{ml} / \mathrm{min}$ ) or an Agilent 1290 Infinity II Preparative system equipped with a SQ MS detector (Multimode ESI/APCI source), with a Waters CSH C18 OBD column ( 5 microns silica, 30 mm diameter, 100 mm length, flow rate of $50 \mathrm{ml} / \mathrm{min}$ ) using decreasingly polar mixtures of water (containing 0.1-0.3\% aqueous ammonium) or water (containing $0.1 \%$ formic acid) and acetonitrile as eluents.
NMR: NMR spectra were recorded on a Bruker AVANCE III HD $400(400 \mathrm{MHz})$ or Bruker AVANCE NEO $400(400 \mathrm{MHz})$ or Bruker AVANCE III $400(400 \mathrm{MHz})$ or Bruker AVANCE II 300 ( 300 MHz ) or Bruker AVANCE III 300 ( 300 MHz ) or Bruker AVANCE III HD 300 $(300 \mathrm{MHz})$. Characteristic chemical shifts ( $\delta$ ) are given in parts-per-million downfield from tetramethylsilane ( 0.00 ppm ) or solvent peaks as the internal reference using conventional abbreviations for designation of major peaks: e.g. s, singlet; d, doublet; $t$, triplet; q, quarter; m, multiplet; br, broad.
Analytical LCMS: LC/MS experiments were performed using a Shimadzu LCMS-2020 with electrospray ionization in positive ion detection mode with 20ADXR pump, SIL-20ACXR autosampler, CTO-20AC column oven, M20A PDA Detector and LCMS 2020 MS detector. The MS detector was configured with electrospray ionization as ionizable source.

All animal studies were conducted in accordance with the guidelines established by the internal IACUC (Institutional Animal Care and Use Committee) and reported following the ARRIVE (Animal Research: Reporting In Vivo experiments) guidelines

## General methods:

2,4,7-Trichloroquinazoline is commercially available, but it can also be synthesised according to the following methods:
Method 1: 7-Chloroquinazoline-2,4-diol ( $600 \mathrm{mg}, 3.05 \mathrm{mmol}$ ), and DIEA ( $2.7 \mathrm{~mL}, 15.3 \mathrm{mmol}$ ) were stirred in $\mathrm{POCl}_{3}(6.0 \mathrm{~mL}, 64.4 \mathrm{mmol})$ at $0{ }^{\circ} \mathrm{C}$ for 1 h . The solvent was removed under reduced pressure and the crude product ( 700 mg ) was used in the following step directly without further purification. ${ }^{1}$
Method 2: To a stirred suspension of 7-chloroquinazoline-2,4( $1 \mathrm{H}, 3 \mathrm{H}$ )-dione ( $5.00 \mathrm{~g}, 25.4$ mmol ) and DIPEA ( $8.86 \mathrm{~mL}, 50.9 \mathrm{mmol}$ ) in toluene ( 100 mL ) at rt was added $\mathrm{POCl}_{3}(3.56$ $\mathrm{mL}, 38.1 \mathrm{mmol}$ ) dropwise. The resultant suspension was heated at $90^{\circ} \mathrm{C}$ for 3.5 h . The reaction mixture was allowed to cool to rt and was carefully added dropwise to an ice cold solution of saturated $\mathrm{NaHCO}_{3}(700 \mathrm{~mL})$ with vigorous stirring. EtOAc was added and the layers were separated. The aqueous was re-extracted with EtOAc and the combined organic phases were
dried by passing through a phase separating cartridge. The filtrate was concentrated to dryness to afford 2,4,7-trichloroquinazoline ( $4.29 \mathrm{~g}, 72.2 \%$ ) as an orange solid. ${ }^{1} \mathrm{H}$ NMR ( 400 MHz , DMSO) $\delta: 7.94(\mathrm{dd}, J=8.9,2.1 \mathrm{~Hz}, 1 \mathrm{H}), 8.22(\mathrm{dd}, J=2.1,0.5 \mathrm{~Hz}, 1 \mathrm{H}), 8.33(\mathrm{dd}, J=8.9,0.5$ $\mathrm{Hz}, 1 \mathrm{H}) . \mathrm{m} / \mathrm{z}$ : no mass ion observed.

## 4-(Dimethylamino)-6-ethoxy-1,3,5-triazin-2(1H)-one (1)



2,4,6-Trichloro-1,3,5-triazine ( $5.00 \mathrm{~g}, 27.1 \mathrm{mmol}$ ), ethanol ( $1.27 \mathrm{~mL}, 27.1 \mathrm{mmol}$ ), and $\mathrm{K}_{2} \mathrm{CO}_{3}$ $(3.75 \mathrm{~g}, 27.1 \mathrm{mmol})$ were stirred in $\mathrm{MeCN}(30 \mathrm{~mL})$ at rt for 4 h . Water $(489 \mu \mathrm{~L}, 27.1 \mathrm{mmol})$ was then added and the mixture was stirred at rt for 30 min . The solids were filtered out, the solvent was evaporated and the residue ( 4.78 g ) was used directly in the following step. Crude 4-chloro-6-ethoxy-1,3,5-triazin-2 $(1 H)$-one ( 120 mg ) and dimethylamine ( $47 \mu \mathrm{~L}, 2 \mathrm{M}$ in THF, 0.68 mmol ) were stirred in water $(5.0 \mathrm{~mL})$ at $50^{\circ} \mathrm{C}$ for 3 h . The solvent was evaporated and the crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $0.05 \% \mathrm{TFA}$ ) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 4 -(dimethylamino)-6-ethoxy-1,3,5-triazin-2(1H)-one $(30.0 \mathrm{mg}, 24 \%$ over 2 steps) as a white solid; $\mathbf{1 H}$ NMR ( 300 MHz , Methanol-d4) $\delta 1.46(\mathrm{t}$, $3 \mathrm{H}), 3.26(\mathrm{~s}, 3 \mathrm{H}), 3.37(\mathrm{~s}, 3 \mathrm{H}), 4.62(\mathrm{q}, 2 \mathrm{H}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+185$.

## 4-(Dimethylamino)-6-ethoxy-1-methyl-1,3,5-triazin-2(1H)-one (3)



A mixture of 4-(dimethylamino)-6-ethoxy-1,3,5-triazin-2(1H)-one $1(1.00 \mathrm{~g}, 5.43 \mathrm{mmol})$, iodomethane ( $0.771 \mathrm{~g}, 5.43 \mathrm{mmol}$ ), and $\mathrm{K}_{2} \mathrm{CO}_{3}(0.750 \mathrm{~g}, 5.43 \mathrm{mmol})$ in $\mathrm{MeCN}(10 \mathrm{~mL})$ was stirred at $50{ }^{\circ} \mathrm{C}$ for 4 h . The mixture was cooled to rt and the crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $0.05 \% \mathrm{NH}_{3} \cdot \mathrm{H}_{2} \mathrm{O}$ ) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 4-(dimethylamino)-6-ethoxy-1-methyl-1,3,5-triazin-2(1H)-one ( $0.300 \mathrm{~g}, 28 \%$ ) as a white solid. 1H NMR ( 300 MHz , DMSO-d6) $\delta 1.35(\mathrm{t}, 3 \mathrm{H}$ ), $3.03(\mathrm{~s}, 3 \mathrm{H}), 3.11(\mathrm{~s}, 3 \mathrm{H}), 3.16(\mathrm{~s}$, $3 \mathrm{H}), 4.43(\mathrm{q}, 2 \mathrm{H}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+199$.

## 4-(Dimethylamino)-6-ethoxy-1-phenyl-1,3,5-triazin-2(1H)-one (4)



Commercially available 4-(dimethylamino)-6-hydroxy-1-phenyl-1,3,5-triazin-2(1H)-one (200 $\mathrm{mg}, 0.860 \mathrm{mmol}$ ), bromoethane ( $103 \mathrm{mg}, 0.950 \mathrm{mmol}$ ), and silver carbonate ( $950 \mathrm{mg}, 3.44$ $\mathrm{mmol})$ were suspended in toluene $(20 \mathrm{~mL})$ and the mixture was stirred at $100^{\circ} \mathrm{C}$ for 15 h . The reaction mixture was cooled to rt and filtered through celite and the solvent was removed under
reduced pressure. The crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $0.05 \% \mathrm{NH}_{4} \mathrm{HCO}_{3}$ ) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 4-(dimethylamino)-6-ethoxy-1-phenyl-1,3,5-triazin-2( $1 H$ )-one ( $60.0 \mathrm{mg}, 27 \%$ ) as a white solid. $\mathbf{1 H}$ NMR ( 400 MHz, DMSO-d6) $\delta 1.17(3 \mathrm{H}, \mathrm{t}), 3.10(3 \mathrm{H}, \mathrm{s}), 3.19(3 \mathrm{H}, \mathrm{s}), 4.38(2 \mathrm{H}, \mathrm{q}), 7.26(2 \mathrm{H}, \mathrm{m}), 7.40$ $(1 \mathrm{H}, \mathrm{m}), 7.45(2 \mathrm{H}, \mathrm{dd}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+261$.

## 7-Chloro-4-(dimethylamino)quinazolin-2(1H)-one (5)



Dimethylamine ( $0.47 \mathrm{~mL}, 2.0 \mathrm{M}$ in THF, 0.940 mmol ) was added to 2,4,7-trichloroquinazoline ( $200 \mathrm{mg}, 0.860 \mathrm{mmol}$ ) in THF ( 2.0 mL ) and the solution was stirred at rt for 2 h . The reaction mixture was diluted with $\operatorname{EtOAc}(50 \mathrm{~mL})$, and washed sequentially with saturated $\mathrm{NaHCO}_{3}(20$ $\mathrm{mL} \times 2$ ), water ( $20 \mathrm{~mL} \times 2$ ), and saturated brine ( $20 \mathrm{~mL} \times 2$ ). The organic layer was dried over $\mathrm{Na}_{2} \mathrm{SO}_{4}$, filtered, evaporated and the crude product ( 160 mg ) was used in the following step without further purification. Acetic acid ( 3.0 mL ) was added to the crude 2,7 -dichloro- $N, N$ -dimethylquinazolin- 4 -amine ( $150 \mathrm{mg}, 0.620 \mathrm{mmol}$ ) and the resulting mixture was stirred at $100^{\circ} \mathrm{C}$ for 12 h . After cooling the reaction mixture to rt , the solvent was removed under reduced pressure. The crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $0.05 \% \mathrm{NH}_{4} \mathrm{OH}$ ) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 7-chloro-4-(dimethylamino)quinazolin- $2(1 \mathrm{H}$ )-one ( $40.0 \mathrm{mg}, 22 \%$ over two steps) as a white solid. $\mathbf{1 H}$ NMR ( 300 MHz, DMSO-d6) $\delta 3.25(6 \mathrm{H}, \mathrm{s}), 7.10(1 \mathrm{H}, \mathrm{dd}), 7.20(1 \mathrm{H}, \mathrm{d}), 7.94(1 \mathrm{H}, \mathrm{d}), 10.82$ $(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+224$.

## 4-(Dimethylamino)quinazolin-2(1H)-one (6)



A solution of 2,4-dichloroquinazoline ( $150 \mathrm{mg}, 0.750 \mathrm{mmol}$ ), dimethylamine ( $450 \mu \mathrm{~L}, 2 \mathrm{M}$ in THF, 0.900 mmol$)$, DIPEA $(0.131 \mathrm{~mL}, 0.750 \mathrm{mmol})$ in $\mathrm{MeCN}(2.1 \mathrm{~mL})$ was heated at $70{ }^{\circ} \mathrm{C}$ for 3 h . The reaction mixture was cooled to rt, diluted with EtOAc ( 10 mL ), and washed sequentially with water ( $10 \mathrm{~mL} \times 2$ ), and saturated brine ( $10 \mathrm{~mL} \times 2$ ). The organic layer was dried with $\mathrm{MgSO}_{4}$, filtered and evaporated. The crude product was purified by flash silica chromatography, elution gradient 0 to $50 \% \mathrm{EtOAc}$ in heptane. Pure fractions were evaporated to dryness to afford 2 -chloro- $N, N$-dimethylquinazolin- 4 -amine ( $125 \mathrm{mg}, 80 \%$ ) as a white solid. 1H NMR ( $500 \mathrm{MHz}, \mathrm{CDCl}_{3}$ ) $\delta 3.42(\mathrm{~s}, 6 \mathrm{H}), 7.39$ (ddd, 1 H ), 7.69 (ddd, 1H), $7.75-7.8$ (m, $1 \mathrm{H}), 7.99-8.05(\mathrm{~m}, 1 \mathrm{H}) ; \mathbf{m} / \mathbf{z} \mathrm{ES}+[\mathrm{M}+\mathrm{H}]+208$.
A solution of 2-chloro- $N, N$-dimethylquinazolin- 4 -amine ( $118 \mathrm{mg}, 0.570 \mathrm{mmol}$ ) in acetic acid $(5.7 \mathrm{~mL})$ was stirred at $90^{\circ} \mathrm{C}$ overnight. The reaction mixture was cooled to rt and the solvent was evaporated (as an azeotrope with toluene). The crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $1 \% \mathrm{NH}_{3} \cdot \mathrm{H}_{2} \mathrm{O}$ ) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 4-(dimethylamino)quinazolin- $2(1 \mathrm{H}$ )-one ( $69.8 \mathrm{mg}, 65 \%$ ) as a white solid. 1H NMR (DMSO-d6,
$500 \mathrm{MHz}) \delta 3.24(6 \mathrm{H}, \mathrm{s}), 7.08(1 \mathrm{H}, \mathrm{ddd}), 7.18(1 \mathrm{H}, \mathrm{dd}), 7.53(1 \mathrm{H}, \mathrm{ddd}), 7.91(1 \mathrm{H}, \mathrm{dd}), 10.72$ $(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+190$.

## 7-Chloroquinazolin-2(1H)-one (7)



2,2,2-Trichloroacetyl chloride ( $386 \mathrm{mg}, 2.12 \mathrm{mmol}$ ) was added to 2-amino-4chlorobenzaldehyde ( $300 \mathrm{mg}, 1.93 \mathrm{mmol}$ ) and DMAP ( $471 \mathrm{mg}, 3.86 \mathrm{mmol}$ ) in DCM ( 5.0 mL ) under $\mathrm{N}_{2}$. The resulting mixture was stirred at rt for 12 h . The reaction mixture was quenched with water $(2.0 \mathrm{~mL})$, extracted with $\mathrm{DCM}(2.0 \mathrm{~mL} x 2)$ and the organic layer was dried over $\mathrm{Na}_{2} \mathrm{SO}_{4}$, filtered and evaporated. The residue was purified by preparative TLC (EtOAc: petroleum ether = 1: 10), to afford 2,2,2-trichloro- $N$-(5-chloro-2-formylphenyl) acetamide (400 $\mathrm{mg}, 69 \%$ ) as a white solid; $\mathbf{1 H}$ NMR (DMSO-d6, 300 MHz ) $\delta 7.61(1 \mathrm{H}, \mathrm{dd}), 8.08(1 \mathrm{H}, \mathrm{d}), 8.36$ $(1 \mathrm{H}, \mathrm{d}), 10.03(1 \mathrm{H}, \mathrm{d}), 12.34(1 \mathrm{H}, \mathrm{s})$; no MS signal detected.
Ammonium acetate ( $77.0 \mathrm{mg}, 1.00 \mathrm{mmol}$ ) was added to $2,2,2$-trichloro- N -( 5 -chloro-2formylphenyl)acetamide ( $150 \mathrm{mg}, 0.500 \mathrm{mmol}$ ) in DMSO ( 3.0 mL ) under $\mathrm{N}_{2}$. The resulting mixture was stirred at $80^{\circ} \mathrm{C}$ for 12 h . After cooling to rt , the crude product was purified by preparative HPLC using decreasingly polar mixtures of water and MeCN as eluents. Fractions containing the desired compound were freeze-dried to afford 7 -chloroquinazolin- $2(1 \mathrm{H})$-one $(13.0 \mathrm{mg}, 14 \%)$ as a white solid. 1H NMR (DMSO-d6, 300 MHz ) $\delta 7.25-7.37(2 \mathrm{H}, \mathrm{m}), 7.93$ $(1 \mathrm{H}, \mathrm{d}), 9.29(1 \mathrm{H}, \mathrm{s}), 11.95(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+181$.

## 7-Chloro-4-(methylamino)quinazolin-2(1H)-one (8)



2,4,7-Trichloroquinazoline ( $400 \mathrm{mg}, 1.72 \mathrm{mmol}$ ) and methylamine in ethanol ( $30 \mathrm{~mL}, 33 \%$ $\mathrm{w} / \mathrm{w}$ ) were stirred in $\mathrm{MeCN}(30 \mathrm{~mL})$ at rt for 30 min . The solvent was removed under reduced pressure and the crude mixture was taken up in formic acid $(10 \mathrm{~mL})$ to give a yellow solution. The resulting mixture was stirred at rt for 24 h . The solvent was removed under reduced pressure and the reaction mixture was diluted with MeCN . The precipitate was collected by filtration, washed with MeCN and DMSO and dried under vacuum to afford 7-chloro-4-(methylamino)quinazolin- $2(1 \mathrm{H}$ )-one ( $250 \mathrm{mg}, 70 \%$ over two steps) as a white solid. 1H NMR ( 300 MHz, DMSO-d6) $\delta 3.09(3 \mathrm{H}, \mathrm{d}), 7.30(1 \mathrm{H}, \mathrm{d}), 7.39(1 \mathrm{H}, \mathrm{dd}), 8.44(1 \mathrm{H}, \mathrm{d}), 10.79(1 \mathrm{H}, \mathrm{s})$, $11.93(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+210$.

4-Amino-7-chloroquinazolin-2(1H)-one (9)


2,4,7-Trichloroquinazoline ( $200 \mathrm{mg}, 0.862 \mathrm{mmol}$ ) and ammonia ( $10 \mathrm{~mL}, 1 \mathrm{M}$ in THF) were stirred in $\mathrm{MeCN}(30 \mathrm{~mL})$ at rt for 30 min . The solvent was removed under reduced pressure and the crude product was purified by C18-flash chromatography, elution gradient 10 to $50 \%$ MeCN in water. Fractions containing the product were evaporated to dryness providing 100 mg of impure compound. 50 mg of the crude product were purified by preparative HPLC using decreasingly polar mixtures of water (containing $10 \mathrm{mM} \mathrm{NH} 4 \mathrm{HCO}_{3}+0.1 \% \mathrm{NH}_{3} \cdot \mathrm{H}_{2} \mathrm{O}$ ) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 4-amino-7-chloroquinazolin-2( 1 H )-one ( $20.0 \mathrm{mg}, 24 \%$ ) as a white solid. 1H NMR (300 MHz, DMSO-d6) $\delta 7.13(2 \mathrm{H}, \mathrm{m}), 7.88(2 \mathrm{H}, \mathrm{br} \mathrm{s}), 7.99(1 \mathrm{H}, \mathrm{d}), 10.72(1 \mathrm{H}, \mathrm{br} \mathrm{s}) ; \mathbf{m} / \mathbf{z}$ (ES+), [M+H]+ 196.

## $N$-(7-chloro-2-oxo-1,2-dihydroquinazolin-4-yl)acetamide (10)



2,4,7-Trichloroquinazoline ( $200 \mathrm{mg}, 0.862 \mathrm{mmol}$ ) and ammonia ( $10 \mathrm{~mL}, 1 \mathrm{M}$ in THF) were stirred in $\mathrm{MeCN}(30 \mathrm{~mL})$ at rt for 30 min . The solvent was removed under reduced pressure and the crude product was purified by C18-flash chromatography, elution gradient 10 to $50 \%$ MeCN in water. Fractions containing the intermediate were evaporated to dryness providing 100 mg of impure compound. 4-Amino-7-chloroquinazolin-2( 1 H )-one ( 100 mg , crude) was suspended in acetic anhydride ( 5.0 mL ) and the resulting mixture was stirred at $80^{\circ} \mathrm{C}$ for 2 h . The mixture was cooled to rt and a precipitate was observed. The precipitate was collected by filtration, washed with $\mathrm{MeCN}(75 \mathrm{~mL})$ and dried under vacuum to afford $N$-(7-chloro-2-oxo-1,2-dihydroquinazolin-4-yl)acetamide ( $36.9 \mathrm{mg}, 18 \%$ over two steps) as a white solid. $\mathbf{1 H}$ NMR (400 MHz, DMSO-d6) $\delta 2.27(3 \mathrm{H}, \mathrm{s}), 7.24(2 \mathrm{H}, \mathrm{m}), 8.09(1 \mathrm{H}, \mathrm{d}), 11.61(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}$ (ES+), $[\mathrm{M}+\mathrm{H}]+238$.

## 7-Chloro-4-methoxyquinazolin-2(1H)-one (11)



Methanol ( $0.188 \mathrm{~mL}, 4.65 \mathrm{mmol}$ ) was added to a solution of 2,4,7-trichloroquinazoline ( 200 $\mathrm{mg}, 0.862$ ) and DIEA ( $0.650 \mathrm{~mL}, 3.72 \mathrm{mmol}$ ) in MeCN ( 10 mL ). The resulting mixture was stirred at rt for 30 min . The solvent was removed under reduced pressure, NaOH ( $186 \mathrm{mg}, 4.65$ $\mathrm{mmol})$ in water $(10 \mathrm{~mL})$ was added and the resulting mixture was stirred at rt for 2 days. The crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $0.05 \% \mathrm{NH}_{3} \cdot \mathrm{H}_{2} \mathrm{O}$ ) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 7-chloro-4-methoxyquinazolin-2(1H)-one ( $40.0 \mathrm{mg}, 22 \%$ over two steps) as a white solid. 1H NMR ( 300 MHz , DMSO-d6) $\delta 4.01(3 \mathrm{H}$, s), $7.24(2 \mathrm{H}, \mathrm{m}), 7.84(1 \mathrm{H}, \mathrm{dd}), 11.44(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+211$.

## 7-Chloro-4-phenoxyquinazolin-2(1H)-one (12)



2,4,7-Trichloroquinazoline ( $150 \mathrm{mg}, 0.646 \mathrm{mmol}$ ), DIEA ( $0.487 \mathrm{~mL}, 2.79 \mathrm{mmol}$ ), and phenol ( $263 \mathrm{mg}, 2.79 \mathrm{mmol}$ ) were stirred in $\mathrm{MeCN}(20 \mathrm{~mL})$ at rt for 30 min . The solvent was removed under reduced pressure. The reaction mixture was diluted with MeCN . The precipitate was collected by filtration, washed with MeCN and dried under vacuum to afford 7-chloro-4-phenoxyquinazolin- $2(1 \mathrm{H}$ )-one ( $140 \mathrm{mg}, 80 \%$ ) as a white solid. $\mathbf{1 H}$ NMR ( 300 MHz , DMSOd6) $\delta 7.14(1 \mathrm{H}, \mathrm{dd}), 7.27(4 \mathrm{H}, \mathrm{m}), 7.48(2 \mathrm{H}, \mathrm{t}), 7.98(1 \mathrm{H}, \mathrm{d}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+273$.

## 7-Chloro-4-phenylquinazolin-2(1H)-one (13)



Phenylmagnesium bromide ( $4.92 \mathrm{~mL}, 1.0 \mathrm{M}$ in THF, 4.92 mmol ) was added dropwise to 2-amino-4-chlorobenzonitrile ( $250 \mathrm{mg}, 1.64 \mathrm{mmol}$ ) in THF ( 20 mL ) at $25^{\circ} \mathrm{C}$ under $\mathrm{N}_{2}$. The resulting mixture was stirred at $50^{\circ} \mathrm{C}$ for 10 h . After cooling the reaction mixture to $0^{\circ} \mathrm{C}$, methyl chloroformate ( $155 \mathrm{mg}, 1.64 \mathrm{mmol}$ ) was added and the resulting mixture was stirred at rt for 16 h . The solvent was removed under reduced pressure, water was added and the mixture was extracted with DCM ( $3 \times 75 \mathrm{~mL}$ ). The organic layers were dried over $\mathrm{Na}_{2} \mathrm{SO}_{4}$, filtered and evaporated to afford a yellow solid. The residue was suspended in MeCN and the precipitate was collected by filtration, washed with DMSO and dried under vacuum to afford 7-chloro-4-phenylquinazolin- $2(1 \mathrm{H}$ )-one ( $173 \mathrm{mg}, 41 \%$ ) as a yellow solid. $\mathbf{1 H}$ NMR ( 300 MHz , DMSO$\left.\mathrm{d} 6+\mathrm{CF}_{3} \mathrm{COOD}\right) \delta 7.28(1 \mathrm{H}, \mathrm{m}), 7.44(1 \mathrm{H}, \mathrm{d}), 7.63(6 \mathrm{H}, \mathrm{m}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+257$.

## 7-Chloro-4-(ethyl(methyl)amino)quinazolin-2(1H)-one (14)



2,4,7-Trichloroquinazoline ( 150 mg , crude, synthesised according to method 1) and N methylethanamine $(0.100 \mathrm{~mL}, 1.16 \mathrm{mmol})$ were stirred in $\mathrm{MeCN}(20 \mathrm{~mL})$ for 30 min at rt . The crude product was purified by flash C18-flash chromatography, elution gradient 20 to $50 \%$ MeCN in water. Pure fractions were evaporated to dryness to afford 2( $\left\{\right.$ bis $[$ ethyl(methyl)amino]phosphoryl $\}$ oxy)-7-chloro- $N$-ethyl- $N$-methylquinazolin-4-amine. ${ }^{2}$ ( $110 \mathrm{mg}, 42 \%$ over two steps from 7 -chloroquinazoline-2,4-diol) as a colourless oil. 1H NMR ( 300 MHz, DMSO-d6) $\delta 1.07(6 \mathrm{H}, \mathrm{t}), 1.28(3 \mathrm{H}, \mathrm{t}), 2.64(6 \mathrm{H}, \mathrm{m}), 3.05(4 \mathrm{H}, \mathrm{m}), 3.36(3 \mathrm{H}, \mathrm{s})$, $3.74(2 \mathrm{H}, \mathrm{q}), 7.35(1 \mathrm{H}, \mathrm{dd}), 7.58(1 \mathrm{H}, \mathrm{d}), 8.12(1 \mathrm{H}, \mathrm{d}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+400$.
2-(\{bis[ethyl(methyl)amino]phosphoryl\}oxy)-7-chloro- $N$-ethyl- $N$-methylquinazolin-4-amine $(100 \mathrm{mg}, 0.250 \mathrm{mmol})$ and formic acid $(5.0 \mathrm{~mL}, 130 \mathrm{mmol})$ were stirred at rt for 3 days and 45 ${ }^{\circ} \mathrm{C}$ for 3 h . The solvent was removed under reduced pressure. The crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $0.05 \% \mathrm{NH}_{3} \cdot \mathrm{H}_{2} \mathrm{O}$ )
and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 7 -chloro-4-(ethyl(methyl)amino)quinazolin-2( 1 H )-one ( $50.0 \mathrm{mg}, 84 \%$ ) as a white solid. 1H NMR ( 300 MHz , DMSO-d6) $\delta 1.25(3 \mathrm{H}, \mathrm{t}), 3.22(3 \mathrm{H}, \mathrm{s}), 3.64(2 \mathrm{H}, \mathrm{q}), 7.10(1 \mathrm{H}$, dd), $7.20(1 \mathrm{H}, \mathrm{d}), 7.86(1 \mathrm{H}, \mathrm{d}), 10.73(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+238$.

## 4-((2-Aminoethyl)(methyl)amino)-7-chloroquinazolin-2(1H)-one (15)



2,4,7-Trichloroquinazoline ( 180 mg , 0.776 mmol ), DIEA ( $0.585 \mathrm{~mL}, 3.35 \mathrm{mmol}$ ), and benzyl (2-(methylamino)ethyl)carbamate ( $697 \mathrm{mg}, 3.35 \mathrm{mmol}$ ) were stirred in $\mathrm{MeCN}(15 \mathrm{~mL})$ at rt for 30 min . The solvent was removed under reduced pressure and the crude product was purified by flash C18-flash chromatography, elution gradient 30 to $70 \% \mathrm{MeCN}$ in water. Pure fractions were evaporated to dryness to afford benzyl (2-((7-chloro-2-hydroxyquinazolin-4yl)(methyl)amino)ethyl)carbamate ( $50.0 \mathrm{mg}, 17 \%$ ) as a white solid. 1H NMR ( 300 MHz , DMSO-d6) $\delta 3.40$ (t, 2H) , 3.42(s, 3H), 3.86 (t, 2H); 4.98 (s, 2H), $7.40-7.20$ (m, 4H), 7.52 $7.41(\mathrm{~m}, 2 \mathrm{H}), 7.72(\mathrm{~d}, 1 \mathrm{H}), 7.95(\mathrm{dd}, 1 \mathrm{H}), 8.26-8.17(\mathrm{~m}, 1 \mathrm{H}), 8.34(\mathrm{~d}, 1 \mathrm{H}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+)$, $[\mathrm{M}+\mathrm{H}]+387$
Benzyl (2-((7-chloro-2-hydroxyquinazolin-4-yl)(methyl)amino)ethyl)carbamate ( 50.0 mg , $0.130 \mathrm{mmol})$ and acetic acid ( $5.0 \mathrm{~mL}, 87.3 \mathrm{mmol}$ ) were stirred at $100^{\circ} \mathrm{C}$ for 20 h . After cooling the mixture to rt, the solvent was removed under reduced pressure. The crude product was purified by flash C18-flash chromatography, elution gradient 30 to $70 \% \mathrm{MeCN}$ in water. Pure fractions were evaporated to dryness to afford 4-((2-aminoethyl)(methyl)amino)-7-chloroquinazolin- $2(1 \mathrm{H}$ )-one ( $25.0 \mathrm{mg}, 76 \%$ ) as a white solid. 1H NMR ( 400 MHz , DMSOd6) $\delta 2.60(3 \mathrm{H}, \mathrm{s}), 3.17(2 \mathrm{H}, \mathrm{t}), 3.73(2 \mathrm{H}, \mathrm{d}), 7.20(2 \mathrm{H}, \mathrm{m}), 8.10(1 \mathrm{H}, \mathrm{d}), 8.70(2 \mathrm{H}, \mathrm{s}), 10.87$ ( $1 \mathrm{H}, \mathrm{s}$ ); $\mathbf{~ m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+253$.

7-Chloro-4-(methyl(phenyl)amino)quinazolin-2(1H)-one (16)


2,4,7-Trichloroquinazoline ( $200 \mathrm{mg}, 0.862$ ), DIEA ( $0.487 \mathrm{~mL}, 2.79 \mathrm{mmol}$ ) and N methylaniline ( $299 \mathrm{mg}, 2.79 \mathrm{mmol}$ ) were stirred in MeCN $(20 \mathrm{~mL})$ at rt for 30 min . The solvent was removed under reduced pressure and subsequently formic acid ( 20 mL ) was added. The resulting solution was stirred at rt for 20 h . The solvent was removed under reduced pressure and the crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $10 \mathrm{mM} \mathrm{NH} \mathrm{NCO}_{3}+0.1 \% \mathrm{NH}_{3} \cdot \mathrm{H}_{2} \mathrm{O}$ ) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 7-chloro-4-(methyl(phenyl)amino)quinazolin-2( 1 H )-one ( $120 \mathrm{mg}, 49 \%$ over two steps) as a white solid. 1H NMR ( 300 MHz, DMSO-d6) $\delta 3.44(3 \mathrm{H}, \mathrm{s}), 6.60(1 \mathrm{H}, \mathrm{d}), 6.72(1 \mathrm{H}, \mathrm{dd}), 7.18(1 \mathrm{H}, \mathrm{d}), 7.33$ (3H, m), $7.45(2 \mathrm{H}, \mathrm{dd}), 11.14(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+286$.

## 4-(Benzyl(methyl)amino)-7-chloroquinazolin-2(1H)-one (17)



2,4,7-Trichloroquinazoline ( $150 \mathrm{mg}, 0.646 \mathrm{mmol}$ ) and $N$-methyl-1-phenylmethanamine ( 100 $\mu \mathrm{L}, 0.780 \mathrm{mmol})$ were stirred in $\mathrm{MeCN}(20 \mathrm{~mL})$ at rt for 30 min . The solvent was removed under reduced pressure and the crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $10 \mathrm{mM} \mathrm{NH}_{4} \mathrm{HCO}_{3}+0.1 \% \mathrm{NH}_{3} \cdot \mathrm{H}_{2} \mathrm{O}$ ) and MeCN as eluent. Fractions containing the desired compound were evaporated to dryness to afford 4 -(benzyl(methyl)amino)-7-chloroquinazolin- $2(1 \mathrm{H}$ )-one $(60.0 \mathrm{mg}, 31 \%)$ as a white solid. 1H NMR ( 300 MHz , DMSO-d6) $\delta 3.21(3 \mathrm{H}, \mathrm{s}), 4.92(2 \mathrm{H}, \mathrm{s}), 7.07(1 \mathrm{H}, \mathrm{dd}), 7.23(1 \mathrm{H}$, d), $7.35(5 \mathrm{H}, \mathrm{m}), 7.84(1 \mathrm{H}, \mathrm{d}), 10.92(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+300$.

## 7-Chloro-4-(methyl(phenethyl)amino)quinazolin-2(1H)-one (18)



2,4,7-Trichloroquinazoline ( $160 \mathrm{mg}, 0.690$ ) and $N$-methyl-2-phenylethan-1-amine ( $100 \mu \mathrm{~L}$, $1.04 \mathrm{mmol})$ were stirred in $\mathrm{MeCN}(20 \mathrm{~mL})$ at rt for 30 min . The solvent was removed under reduced pressure and formic acid ( 5.7 mL ) was added. The resulting solution was stirred at rt for 20 h and the solvent was removed under reduced pressure. The crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $10 \mathrm{mM} \mathrm{NH}_{4} \mathrm{HCO}_{3}$ $+0.1 \% \mathrm{NH}_{3} \cdot \mathrm{H}_{2} \mathrm{O}$ ) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 7-chloro-4-(methyl(phenethyl)amino)quinazolin-2(1H)-one ( $138 \mathrm{mg}, 64 \%$ over two steps) as a white solid. 1H NMR ( 300 MHz, DMSO-d6) $\delta 3.00(2 \mathrm{H}$, $\mathrm{m}), 3.27(3 \mathrm{H}, \mathrm{s}), 3.84(2 \mathrm{H}, \mathrm{m}), 7.09(1 \mathrm{H}, \mathrm{dd}), 7.20(2 \mathrm{H}, \mathrm{m}), 7.28(4 \mathrm{H}, \mathrm{d}), 7.85(1 \mathrm{H}, \mathrm{d}), 10.83$ $(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+314$.

## 4-(Azetidin-1-yl)-7-chloroquinazolin-2(1H)-one (19)



2,4,7-Trichloroquinazoline ( $150 \mathrm{mg}, 0.646 \mathrm{mmol}$ ), DIEA ( $0.487 \mathrm{~mL}, 2.79 \mathrm{mmol}$ ), and azetidine ( $80.0 \mathrm{mg}, 1.40 \mathrm{mmol}$ ) were stirred in $\mathrm{MeCN}(15 \mathrm{~mL})$ at rt for 30 min . A precipitate formed during the reaction and this was collected by filtration, washed with MeCN and dried under vacuum to afford 4-(azetidin-1-yl)-2,7-dichloroquinazoline ( 60.0 mg ) as a white solid, which was used without further purification.
4-(Azetidin-1-yl)-2,7-dichloroquinazoline ( 55.0 mg ) was stirred in acetic acid ( 5.0 mL ) at 110 ${ }^{\circ} \mathrm{C}$ for 2 days. A precipitate formed during the reaction and was collected by filtration, washed with MeCN and dried under vacuum to afford 4-(azetidin-1-yl)-7-chloroquinazolin-2(1H)-one ( 40.0 mg , $29 \%$ over two steps) as a white solid. 1H NMR ( 300 MHz , DMSO-d6) $\delta 2.08$ ( 2 H , m), $3.55(2 \mathrm{H}, \mathrm{t}), 3.94(2 \mathrm{H}, \mathrm{t}), 7.32(1 \mathrm{H}, \mathrm{d}), 7.40(1 \mathrm{H}, \mathrm{dd}), 8.11(1 \mathrm{H}, \mathrm{dd}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+$ 236.

7-Chloro-4-(pyrrolidin-1-yl)quinazolin-2(1H)-one (20)


2,4,7-Trichloroquinazoline ( 150 mg , crude, synthesised according to method 1 and pyrrolidine $(0.100 \mathrm{~mL}, 1.20 \mathrm{mmol})$ were stirred in $\mathrm{MeCN}(20 \mathrm{~mL})$ at rt for 30 min . The solvent was removed under reduced pressure to afford 7-chloro-4-(pyrrolidin-1-yl)quinazolin-2-yl di(pyrrolidin-1-yl)phosphinate ${ }^{2}$ ( 200 mg , crude) as a yellow oil. Formic acid ( 15 mL ) was added to 7 -chloro-4-(pyrrolidin-1-yl)quinazolin-2-yl di(pyrrolidin-1-yl)phosphinate ( 195 mg ) and the resulting mixture was stirred at rt for 1 day. The solvent was removed under reduced pressure and the crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $0.1 \%$ formic acid) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 7-chloro-4-(pyrrolidin-1-yl)quinazolin- $2(1 \mathrm{H}$ )-one ( $60.0 \mathrm{mg}, 40 \%$ over three steps from 7 -chloroquinazoline-2,4-diol) as a yellow solid. 1H NMR ( 300 MHz , DMSO-d6) $\delta 1.94(4 \mathrm{H}, \mathrm{m}), 3.78(4 \mathrm{H}, \mathrm{m}), 7.10(1 \mathrm{H}, \mathrm{dd})$, $7.19(1 \mathrm{H}, \mathrm{d}), 8.05(1 \mathrm{H}, \mathrm{d}), 10.74(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+250$.

## 7-Chloro-4-(piperidin-1-yl)quinazolin-2(1H)-one (21)



2,4,7-Trichloroquinazoline ( 150 mg , crude, synthesised according to method 1 ) and piperidine $(0.100 \mathrm{~mL}, 1.01 \mathrm{mmol})$ in $\mathrm{MeCN}(20 \mathrm{~mL})$ were stirred at rt for 30 min . The crude product was purified by flash C18-flash chromatography, elution gradient 50 to $100 \% \mathrm{MeCN}$ in water. Fractions containing the product were evaporated to dryness to afford 7-chloro-4-(piperidin-1-yl)quinazolin-2-yl di(piperidin-1-yl)phosphinate ${ }^{2}$ ( 110 mg , crude) as a yellow oil. Formic acid $(10 \mathrm{~mL})$ was added to 7-chloro-4-(piperidin-1-yl)quinazolin-2-yl di(piperidin-1yl)phosphinate ( 100 mg , crude) and the resulting mixture was stirred at $45^{\circ} \mathrm{C}$ for 1 day. The solvent was removed under reduced pressure. The crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $0.1 \%$ formic acid) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 7-chloro-4-(piperidin-1-yl)quinazolin-2( 1 H )-one ( $50.0 \mathrm{mg}, 32 \%$ over three steps from 7-chloroquinazoline-2,4-diol) as a white solid. 1H NMR ( 300 MHz , DMSO-d6) $\delta$ 1.62-1.70 (6 $\mathrm{H}, \mathrm{m}), 3.60-3.67(4 \mathrm{H}, \mathrm{m}), 7.12(1 \mathrm{H}, \mathrm{dd}), 7.21(1 \mathrm{H}, \mathrm{d}), 7.69(1 \mathrm{H}, \mathrm{d}), 10.92(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+)$, $[\mathrm{M}+\mathrm{H}]+264$.
$N$-(1-(7-chloro-2-oxo-1,2-dihydroquinazolin-4-yl)pyrrolidin-3-yl)acetamide (23)


2,4,7-Trichloroquinazoline ( 200 mg , crude, synthesised according to method 1), $\mathrm{K}_{2} \mathrm{CO}_{3}$ (643 $\mathrm{mg}, 4.65 \mathrm{mmol}$ ) and $N$-(pyrrolidin-3-yl)acetamide ( $596 \mathrm{mg}, 4.65 \mathrm{mmol}$ ) were stirred in MeCN $(50 \mathrm{~mL})$ at rt for 1 h . The solvent was removed under reduced pressure and the crude product was purified by flash C18-flash chromatography, elution gradient 20 to $80 \% \mathrm{MeCN}$ in water. Fractions containing the product were evaporated to dryness to afford $N$-(1-(7-chloro-2-hydroxyquinazolin-4-yl)pyrrolidin-3-yl)acetamide (crude) as a white solid. Half of the crude was used in the following reaction. The other half of the crude product was further purified by preparative HPLC using decreasingly polar mixtures of water (containing $0.1 \%$ formic acid) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford $N$-(1-(7-chloro-2-oxo-1,2-dihydroquinazolin-4-yl)pyrrolidin-3-yl)acetamide (30.0 $\mathrm{mg}, 22 \%$ over two steps from 7 -chloroquinazoline-2,4-diol) as a yellow solid. 1H NMR (300 MHz, DMSO-d6) $\delta 1.81(3 \mathrm{H}, \mathrm{s}), 1.90(1 \mathrm{H}, \mathrm{m}), 2.11(1 \mathrm{H}, \mathrm{dd}), 3.62(1 \mathrm{H}, \mathrm{m}), 3.92(3 \mathrm{H}, \mathrm{m}), 4.29$ $(1 \mathrm{H}, \mathrm{m}), 7.11(1 \mathrm{H}, \mathrm{dd}), 7.19(1 \mathrm{H}, \mathrm{d}), 8.00(1 \mathrm{H}, \mathrm{d}), 8.16(1 \mathrm{H}, \mathrm{d}), 10.79(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+)$, $[\mathrm{M}+\mathrm{H}]+307$.

## 4-(3-Aminopyrrolidin-1-yl)-7-chloroquinazolin-2(1H)-one (22)


$N$-(1-(7-chloro-2-hydroxyquinazolin-4-yl)pyrrolidin-3-yl)acetamide 23 ( $100 \mathrm{mg}, 0.33 \mathrm{mmol}$ ) and $\mathrm{HCl}\left(4.0 \mathrm{~mL}, 4 \mathrm{M}\right.$ in dioxane, 132 mmol ) in water ( 6.0 mL ) were stirred at $50^{\circ} \mathrm{C}$ for 3 days. After cooling the reaction to rt, the solvent was removed under reduced pressure and the crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $0.1 \%$ formic acid) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 4-(3-aminopyrrolidin-1-yl)-7-chloroquinazolin- $2(1 \mathrm{H}$ )-one ( $30.0 \mathrm{mg}, 31 \%$ ) as a white solid. $\mathbf{1 H}$ NMR ( 300 MHz , DMSOd6) $\delta 1.94(1 \mathrm{H}, \mathrm{s}), 2.15(1 \mathrm{H}, \mathrm{m}), 3.83(5 \mathrm{H}, \mathrm{dt}), 7.12(1 \mathrm{H}, \mathrm{dd}), 7.20(1 \mathrm{H}, \mathrm{d}), 8.00(1 \mathrm{H}, \mathrm{d})$; $\mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+265$.

## 7-Chloro-4-(indolin-1-yl)quinazolin-2(1H)-one (24)



2,4,7-Trichloroquinazoline ( $200 \mathrm{mg}, 0.930 \mathrm{mmol}$ ), DIEA ( $0.487 \mathrm{~mL}, 2.79 \mathrm{mmol}$ ) and indoline $(443 \mathrm{mg}, 3.72 \mathrm{mmol})$ were stirred in $\mathrm{MeCN}(20 \mathrm{~mL})$ at rt for 30 min . The solvent was removed under reduced pressure and formic acid ( 20 mL ) was added. The resulting solution was stirred at rt for 20 h . The solvent was removed under reduced pressure and $\mathrm{MeCN}(20 \mathrm{~mL})$ was added. The precipitate was collected by filtration, washed with MeCN and dried under vacuum to afford 7-chloro-4-(indolin-1-yl)quinazolin- $2(1 \mathrm{H}$ )-one ( $50.0 \mathrm{mg}, 18 \%$ over two steps) as a pale yellow solid. 1H NMR ( 300 MHz , DMSO-d6) $\delta 3.14(2 \mathrm{H}, \mathrm{t}), 4.38(2 \mathrm{H}, \mathrm{t}), 7.02(1 \mathrm{H}, \mathrm{t}), 7.16$ ( $2 \mathrm{H}, \mathrm{ddd}$ ), $7.30(2 \mathrm{H}, \mathrm{m}), 7.58(1 \mathrm{H}, \mathrm{d}), 7.96(1 \mathrm{H}, \mathrm{d}), 11.23(1 \mathrm{H}, \mathrm{s}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+298$.

## 7-Chloro-4-(dimethylamino)-1-methylquinazolin-2(1H)-one (25)


$\mathrm{NaH}(91.2 \mathrm{mg}, \sim 60 \%$ dispersion in mineral oil, 2.28 mmol ) was added in small portions to 7-chloro-4-(dimethylamino)quinazolin-2( 1 H )-one $5(170 \mathrm{mg}, 0.760 \mathrm{mmol}$ ) in DMF ( 10 mL ) at $0^{\circ} \mathrm{C}$ under $\mathrm{N}_{2}$. The resulting mixture was stirred at $15^{\circ} \mathrm{C}$ for 30 min . Iodomethane ( 324 mg , 2.28 mmol ) was added and the resulting mixture was stirred at $15{ }^{\circ} \mathrm{C}$ for 5 h . The reaction mixture was quenched with water ( 20 mL ) and extracted with $\mathrm{CHCl}_{3}(2 \times 25 \mathrm{~mL})$. The combined organic layers were evaporated to afford a yellow oil and the crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $0.1 \%$ formic acid) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 7-chloro-4-(dimethylamino)-1-methylquinazolin-2( 1 H )-one ( 63.0 mg , $35 \%$ ) as a white solid. 1H NMR ( 300 MHz , DMSO-d6) $\delta 3.22(6 \mathrm{H}, \mathrm{s}), 3.41(3 \mathrm{H}, \mathrm{s}), 7.19(1 \mathrm{H}$, dd), $7.44(1 \mathrm{H}, \mathrm{d}), 7.95(1 \mathrm{H}, \mathrm{d}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+238$.

## 7-Chloro-4-(dimethylamino)-1-(2-hydroxyethyl)quinazolin-2(1H)-one (26)


$\mathrm{Cs}_{2} \mathrm{CO}_{3}(1.16 \mathrm{~g}, 3.58 \mathrm{mmol})$ was added to 7-chloro-4-(dimethylamino)quinazolin-2(1H)-one 5 ( $200 \mathrm{mg}, 0.890 \mathrm{mmol}$ ) and 2-((tert-butyldimethylsilyl)oxy)ethan-1-ol ( $473 \mathrm{mg}, 2.68 \mathrm{mmol}$ ) in 1,4-dioxane ( 10 mL ) under $\mathrm{N}_{2}$. The resulting mixture was stirred at $100^{\circ} \mathrm{C}$ for 30 h . The reaction mixture was allowed to cool to rt and the crude product was purified by flash C18flash chromatography, elution gradient 25 to $80 \% \mathrm{MeCN}$ in water. Fractions containing the product were evaporated to dryness to afford 1-(2-((tert-butyldimethylsilyl)oxy)ethyl)-7-chloro-4-(dimethylamino)quinazolin-2( 1 H )-one ( 200 mg , crude) as a yellow oil. $\mathrm{HCl}(10 \mathrm{~mL}$, 4 M in dioxane, 40.0 mmol$)$ was added to 1-(2-((tert-butyldimethylsilyl)oxy)ethyl)-7-chloro-4-(dimethylamino)quinazolin-2( 1 H )-one ( 150 mg , crude) and the resulting mixture was stirred at rt for 1 h . The solvent was removed under reduced pressure and the crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing 10 mM $\mathrm{NH}_{4} \mathrm{HCO}_{3}+0.1 \% \mathrm{NH}_{3} \cdot \mathrm{H}_{2} \mathrm{O}$ ) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 7-chloro-4-(dimethylamino)-1-(2-hydroxyethyl)quinazolin- $2(1 \mathrm{H}$ )-one ( $60.0 \mathrm{mg}, 34 \%$ over two steps) as a white solid. 1H NMR ( 400 MHz, DMSO-d6) $\delta 3.21(6 \mathrm{H}, \mathrm{s}), 3.61(2 \mathrm{H}, \mathrm{t}), 4.09(2 \mathrm{H}, \mathrm{t}), 4.83(1 \mathrm{H}, \mathrm{s}), 7.17(1 \mathrm{H}, \mathrm{dd})$, $7.60(1 \mathrm{H}, \mathrm{d}), 7.93(1 \mathrm{H}, \mathrm{d}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+268$.

## 7-Chloro-1-cyclopropyl-4-(dimethylamino)quinazolin-2(1H)-one (27)



7-Chloro-4-(dimethylamino)quinazolin-2(1H)-one 5 ( $200 \mathrm{mg}, 0.900 \mathrm{mmol}$ ), potassium cyclopropyltrifluoroborate ( $662 \mathrm{mg}, 4.48 \mathrm{mmol}$ ), $\mathrm{K}_{2} \mathrm{CO}_{3}(370 \mathrm{mg}, 2.68 \mathrm{mmol}$ ) and copper (II) acetate $(81.2 \mathrm{mg}, 0.440 \mathrm{mmol})$ were stirred in water $(6.0 \mathrm{~mL})$ and toluene $(20 \mathrm{~mL})$ under an atmosphere of $\mathrm{O}_{2}$ at 1 atm and $70{ }^{\circ} \mathrm{C}$ for 2 days. The mixture was filtered, evaporated, and purified by flash C18-flash chromatography, elution gradient 30 to $60 \% \mathrm{MeCN}$ in water. Pure fractions were evaporated to dryness to afford 7-chloro-1-cyclopropyl-4-(dimethylamino)quinazolin-2(1H)-one ( $8.90 \mathrm{mg}, 4 \%$ ) as a white solid. 1H NMR ( 300 MHz , DMSO-d6) $\delta 0.62(2 \mathrm{H}, \mathrm{m}), 1.14(2 \mathrm{H}, \mathrm{td}), 2.76(1 \mathrm{H}, \mathrm{tt}), 3.19(6 \mathrm{H}, \mathrm{s}), 7.18(1 \mathrm{H}, \mathrm{dd}), 7.63(1 \mathrm{H}$, d), $7.88(1 \mathrm{H}, \mathrm{d}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+265$.

7-Chloro-4-(dimethylamino)-1-(m-tolyl)quinazolin-2(1H)-one (29)

$\mathrm{K}_{2} \mathrm{CO}_{3}(204 \mathrm{mg}, 1.48 \mathrm{mmol})$ was added to 7-chloro-4-(dimethylamino)quinazolin-2( 1 H )-one $5(110 \mathrm{mg}, 0.490 \mathrm{mmol})$, 1-iodo-3-methylbenzene ( $214 \mathrm{mg}, 0.980 \mathrm{mmol}$ ), copper(I) iodide ( $94.0 \mathrm{mg}, 0.490 \mathrm{mmol}$ ) and copper ( $31.3 \mathrm{mg}, 0.490 \mathrm{mmol}$ ) in DMA ( 2.0 mL ) under $\mathrm{N}_{2}$. The resulting mixture was stirred at $140{ }^{\circ} \mathrm{C}$ for 20 h . The reaction mixture was filtered through celite and the crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $10 \mathrm{mM} \mathrm{NH} \mathrm{H}_{4} \mathrm{HCO}_{3}+0.1 \% \mathrm{NH}_{3} \cdot \mathrm{H}_{2} \mathrm{O}$ ) and MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 7-chloro-4-(dimethylamino)-1-(m-tolyl)quinazolin-2(1H)-one ( $25.0 \mathrm{mg}, 16 \%$ ) as a white solid. 1H NMR ( 400 MHz , DMSO-d6) $\delta 2.38(3 \mathrm{H}, \mathrm{s}), 3.3(6 \mathrm{H}, \mathrm{s}), 6.35(1 \mathrm{H}, \mathrm{d}), 7.09(2 \mathrm{H}, \mathrm{m}), 7.19(1 \mathrm{H}, \mathrm{dd})$, $7.33(1 \mathrm{H}, \mathrm{d}), 7.48(1 \mathrm{H}, \mathrm{t}), 8.03(1 \mathrm{H}, \mathrm{d}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+314$.

7-Chloro-4-(dimethylamino)-1-(pyridin-2-yl)quinazolin-2(1H)-one (30)

$\mathrm{K}_{2} \mathrm{CO}_{3}$ (226 mg, 1.64 mmol ) was added to 7-chloro-4-(dimethylamino)quinazolin-2( 1 H )-one 5 ( $122 \mathrm{mg}, 0.550 \mathrm{mmol}$ ), 2-iodopyridine ( $224 \mathrm{mg}, 1.09 \mathrm{mmol}$ ), copper(I) iodide ( $104 \mathrm{mg}, 0.550$ mmol ) and copper ( $34.7 \mathrm{mg}, 0.550 \mathrm{mmol}$ ) in DMA ( 8.0 mL ) under $\mathrm{N}_{2}$. The resulting mixture was stirred at $140^{\circ} \mathrm{C}$ for 20 h . After cooling the mixture to rt , the crude product was purified by preparative HPLC using decreasingly polar mixtures of water (containing $0.05 \% \mathrm{NH}_{3} \cdot \mathrm{H}_{2} \mathrm{O}$ ) as MeCN as eluents. Fractions containing the desired compound were evaporated to dryness to afford 7-chloro-4-(dimethylamino)-1-(pyridin-2-yl)quinazolin-2( 1 H )-one ( $65.0 \mathrm{mg}, 40 \%$ ) as a white solid. 1H NMR ( 300 MHz, DMSO-d6) $\delta 3.30(6 \mathrm{H}, \mathrm{s}), 6.29(1 \mathrm{H}, \mathrm{d}), 7.21(1 \mathrm{H}, \mathrm{dd}), 7.54$ $(2 \mathrm{H}, \mathrm{m}), 8.05(2 \mathrm{H}, \mathrm{m}), 8.67(1 \mathrm{H}, \mathrm{dd}) ; \mathbf{m} / \mathbf{z}(\mathrm{ES}+),[\mathrm{M}+\mathrm{H}]+301$.

## HPLC/UPLC traces for key compounds

7-Chloro-4-(dimethylamino)quinazolin-2(1H)-one (5)


Retention time: 0.725
MS Spectrum
Spectrum Mode:Averaged 0.716-0.733(92-94) Base Peak:224.00(74605)
BG Mode:Calc Segment l-Event 1


7-Chloro-4-(dimethylamino)-1-phenylquinazolin-2(1H)-one (28)



7-Chloro-4-(dimethylamino)-1-(pyridin-3-yl)quinazolin-2(1H)-one (31)


7-Chloro-4-(ethyl(methyl)amino)-1-phenylquinazolin-2(1H)-one (32)



7-Chloro-4-(ethyl(methyl)amino)-1-(pyridin-3-yl)quinazolin-2(1H)-one -(33)

3: UV Detector: TAC: Wavelength Range: (220 - 320) Smooth (Mn, 2x3)
1.252e+1 Range: $1.264 e+1$



## Biological data and errors

| Compound | Activity | Mean <br> $\mathbf{p K}_{\mathbf{d}}$ | Std <br> $\mathbf{d e v}$ | Top conc. <br> $(\boldsymbol{\mu M})$ | Replicates |
| :---: | :---: | :---: | :---: | :---: | :---: |
| $\mathbf{1}$ | A | 3.6 | 0.19 | 1000 | 9 |
| $\mathbf{2}$ | A | 5.2 | 0.06 | 100 | 4 |
| $\mathbf{3}$ | N | - | - | 200 | 1 |
| $\mathbf{4}$ | A | 4.4 | 0.00 | 500 | 3 |
| $\mathbf{5}$ | A | 5.3 | 0.11 | 50 | 5 |


| Compound | pIC50 | Std <br> Dev | Mean Max conc inh | Replicates |
| :---: | :---: | :---: | :---: | :---: |
| 1 | 3.7 | 0.039 | -96 | 4 |
| 2 | <3 | - | - | 1 |
| 3 | <3 | - | - | 1 |
| 4 | 3.5 | 0.29 | -84 | 6 |
| 5 | 5.1 | 0.075 | -99 | 3 |
| 6 | <4 | - | - | 4 |
| 7 | 3.2 | 0.033 | -85 | 3 |
| 8 | 4.3 | 0.079 | -72 | 5 |
| 9 | <4 | - | - | 1 |
| 10 | <4 | - | - | 2 |
| 11 | <4 | - | - | 1 |
| 12 | <3 | - | - | 1 |
| 13 | <4 | - | - | 1 |
| 14 | 6.1 | 0.010 | -110 | 3 |
| 15 | <4 | - | - | 1 |
| 16 | <3 | - | - | 1 |
| 17 | 4.2 | 0.05 | -86 | 3 |
| 18 | <4 | - | - | 4 |
| 19 | <4 | - | - | 2 |
| 20 | 5.3 | 0.13 | -110 | 3 |
| 21 | 4.1 | 0.032 | -86 | 3 |
| 22 | 4.3 | 0.15 | -70 | 3 |
| 23 | <4 | - | - | 3 |
| 24 | <4 | - | - | 2 |


| $\mathbf{2 5}$ | 3.7 | 0.35 | -65 | 5 |
| :---: | :---: | :---: | :---: | :---: |
| $\mathbf{2 6}$ | 5.1 | 0.029 | -100 | 3 |
| $\mathbf{2 7}$ | 5.1 | 0.082 | -99 | 3 |
| $\mathbf{2 8}$ | 7.6 | 0.087 | -110 | 5 |
| $\mathbf{2 9}$ | 7.8 | 0.14 | -110 | 3 |
| $\mathbf{3 0}$ | 4.7 | 0.21 | -78 | 4 |
| $\mathbf{3 1}$ | 7.7 | 0.16 | -110 | 5 |
| $\mathbf{3 2}$ | 7.9 | 0.18 | -110 | 3 |
| $\mathbf{3 3}$ | 7.7 | 0.083 | -110 | 5 |


| Compound | Mean <br> Cell SDMA <br> pIC $_{50}$ | Std <br> dev | Replicates | Mean <br> Prolif $^{\text {pGl }_{50}}$ | Std <br> dev | Replicates |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| $\mathbf{2 8}$ | 7.6 | 0.22 | 7 | 6.5 | 0.18 | 10 |
| $\mathbf{3 1}$ | 7.8 | 0.26 | 4 | 6.5 | 0.23 | 5 |
| $\mathbf{3 2}$ | 8.2 | 0.25 | 3 | 6.8 | 0.13 | 3 |
| $\mathbf{3 3}$ | 8.3 | 0.18 | 3 | 6.8 | 0.07 | 2 |

Representative dose-response curves for cellular SDMA and proliferation assays:

Cmpd
Cell IC50
(SDMA levels)

29


Log Concentration [M]

31


32



Log Concentration [M]

Proliferation IC50
(HCT110 MTAP KO)



Log Concentration [M]


Log Concentration [M]


## In vivo Methods

## Animals

Female Ncr Nude mice were purchased from Taconic Biosciences. Mice were housed under pathogen-free conditions in individual ventilated cages (IVC) at our AAALAC (Association for the Assessment and Accreditation of Laboratory Animal Care) accredited facility in Waltham, MA. All animal manipulations were conducted in a biosafety cabinet maintained under positive pressure. Mice used were 5-6 weeks old at the time of tumour implantation.

## Xenograft Efficacy Studies

Ten million HCT116 MTAP KO tumour cells were injected subcutaneously in the right flank of Ncr female mice in a volume of 0.1 mL . Tumour volumes (measured by caliper), animal body weight, and tumour condition were recorded twice weekly for the duration of the study. The tumour volume was calculated (taking length to be the longest diameter across the tumour and width to be the corresponding perpendicular diameter) using the formula: length (mm) x width $(\mathrm{mm})^{2} / 0.52$. For efficacy studies, growth inhibition from the start of treatment was assessed by comparison of the differences in tumour volume between control and treated groups. Because the variance in mean tumour volume data increases proportionally with volume (and is therefore disproportionate between groups), data were log transformed to remove any size dependency before statistical evaluation. Statistical significance was evaluated using a one-tailed, 2-sample $t$ test. For efficacy studies, mice were randomized based on tumour volumes using stratified sampling, and enrolled into control and treatment groups. Dosing began when mean tumour size reached approximately $200 \mathrm{~mm}^{3}$. Compound $\mathbf{2 8}$ was formulated in $0.5 \%$ hydroxypropyl methylcellulose (HPMC)/ $0.1 \%$ Tween 80 in water and dosed subcutaneously at $5 \mathrm{~mL} / \mathrm{kg}$ using a 25 gauge syringe.

Quantitative LCMSMS to evaluate SAM in tumour: LC/MS/MS quantitation of tumour lysates was performed using an Agilent 6490 triple-quadrupole mass spectrometer with electrospray ionization in positive ion detection mode with an Agilent 1290 UHPLC system consisting of a binary pump, HTS autosampler, and a Waters Acquity UPLC BEH Amide 2.1 x $100 \mathrm{~mm}, 1.7 \mu \mathrm{~m}$ column. Chromatography was performed under HILIC conditions with a gradient of $97 \%$ B to $30 \%$ B over 3.5 min , using water and MeCN with $0.1 \%$ formic acid. Internal standard was d-citrulline (CDN Isotopes \# D-6396) at $1 \mathrm{mg} / \mathrm{L}$ in $\mathrm{MeCN} /$ methanol/water/formic acid 40/40/20/0.05 $\mathrm{v} / \mathrm{v} / \mathrm{v} / \mathrm{v}$. External standard was $S$ adenosylmethionine (Cayman Chemical \# 16376) prepared at concentrations from 5 -10,000
nM in methanol. MRM transitions for detection were $399.15>250$, collision energy 10 V (SAM) and $180>74$, collision energy 28 V (d-citrulline.)
Tumour lysates were prepared by taking $50-100 \mathrm{mg}$ of frozen tissue and adding $\mathrm{MeCN} /$ methanol $/$ water $40 / 40 / 20 \mathrm{v} / \mathrm{v} / \mathrm{v}$ to the samples at a concentration of $1 \mathrm{~mL} / 100 \mathrm{mg}$ tissue. Samples were gently agitated for 2 min and then spun at $\sim 21000 \mathrm{~g}$ at $0^{\circ} \mathrm{C}$ for 10 min . Clear supernatant was transferred, then the solvent addition, shaking, and centrifugation were repeated with the second supernatant added to the original. Prior to LC-MS-MS analysis, 15 $\mu \mathrm{L}$ of each lysate was combined with $135 \mu \mathrm{~L}$ of internal standard then vortexed thoroughly.

## MAT2a protein production

Full-length human MAT2a protein was recombinantly expressed in E.coli using a plasmid obtained from the Structural Genomics Consortium ${ }^{1}$. $N$-terminally His 6 -tagged MAT2a was purified by $\mathrm{Ni}^{2+}$-immobilised metal affinity chromatography (IMAC), proteolytic cleavage of the hexahistidine tag and size exclusion chromatography. Final protein concentration was 20 $\mathrm{mg} / \mathrm{mL}(0.47 \mathrm{mM})$ in a buffer containing 10 mM Hepes $\mathrm{pH} 7.5,500 \mathrm{mM} \mathrm{NaCl}, 5 \%$ glycerol and 0.5 mM TCEP.

To allow immobilisation, an expression plasmid containing His6-TEV-Avi-MAT2a (2-395) was generated and Avi-tagged protein was purified as above. In vitro biotinylation was carried out using the BirA biotinylation kit from Avidity according to the manufacturer's instructions. Excess reagents were removed by passing over a desalting column.

Protein was snap-frozen and stored in aliquots until use.

## X-ray crystallography of MAT2a

Full-length human MAT2a protein was recombinantly expressed in E.coli using a plasmid obtained from the Structural Genomics Consortium ${ }^{3}$. Final protein concentration was 20 $\mathrm{mg} / \mathrm{mL}(0.47 \mathrm{mM})$ in a buffer containing 10 mM Hepes $\mathrm{pH} 7.5,500 \mathrm{mM} \mathrm{NaCl}, 5 \%$ glycerol and 0.5 mM TCEP. For crystallisation, a 1.5 -fold molar excess of S-adenosylmethionine was added before dispensing sitting drop vapour diffusion experiments with 150 nL protein and 150 nL of a reservoir solution containing 8-12 \% PEG 8000, 8— $12 \%$ ethylene glycol, 0.1 M Hepes pH 8.0. Bar shaped crystals appeared within minutes, reaching their maximum dimensions within 12 h . Crystallisation reliability was enhanced by micro-seeding. Crystals remained stable for approximately 4 weeks without loss of diffraction quality, although the $S$ adenosylmethionine appears partially hydrolysed in crystals harvested after more than a few
days. Inhibitors were introduced by soaking the crystals in reservoir solution supplemented with $10 \%$ of compound solution ( 100 mM in DMSO). Soaked crystals were frozen after 224 h without further cryoprotection. X-ray diffraction data were collected at Diamond Light Source beamlines I03 and I04, and images were processed with autoproc ${ }^{4}$, STARANISO ${ }^{5}$ and additional programmes from the CCP4 suite ${ }^{6}$. The structures were solved by molecular replacement using $\mathrm{AMoRe}^{7}$ and refined with Buster ${ }^{8}$ and Coot $^{9}$. Initial ligand restraint dictionaries were generated with Grade ${ }^{10}$. Atomic coordinates and structure factors are available from the Protein Data Bank.

## Computational details

Protein structures were prepared before analysis using the Protein Preparation Wizard ${ }^{11}$ in Maestro ${ }^{12}$. Ligand tautomers and protonation states were assigned using Epik ${ }^{13,14}$, and protonation states of the protein and water orientations were determined and optimised at pH 7.0. Hydrogen atoms of ligand, protein, and water molecules were minimised.

WaterMap ${ }^{15,16}$ calculations involved analysing simulated explicit-solvent waters in the binding sites near bound ligands. Water sites within $8 \AA$ of the ligand of interest were analysed. X-ray waters were included and treated as solvent. Simulation time was 2.0 ns .
Quantum-mechanical geometry optimisations were done at DFT level using Jaguar ${ }^{17}$ in Maestro ${ }^{12}$. All optimisations were performed using B3LYP-D3 functionals, $6-31 \mathrm{G}^{* *}$ basis sets, and a PBF solvent (water) model.

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(1) In some cases after evaporation the residual $\mathrm{POCl}_{3}$ caused the formation of a phosphorylated species that was hydrolysed in the last step with formic acid or acetic acid to give the final product. The yields account for the by-product.
(2) This phosphorous by-product originates from residual $\mathrm{POCl}_{3}$ (synthesis 2,4,7trichloroquinazoline, method 1) and the yield is calculated based on the starting material in method 1 .
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