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Bisubstrate Inhibitors of Nicotinamide *N*-Methyltransferase (NNMT) with Enhanced Activity

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ABSTRACT: Nicotinamide *N*-methyltransferase (NNMT) catalyzes the methylation of nicotinamide to form *N*-methylnicotinamide. Overexpression of NNMT is associated with a variety of diseases, including a number of cancers and metabolic disorders, suggesting a role for NNMT as a potential therapeutic target. By structural modification of a lead NNMT inhibitor previously developed in our group, we prepared a diverse library of inhibitors to probe the different regions of the enzyme's active site. This investigation revealed that incorporation of a

naphthalene moiety intended to bind the hydrophobic nicotinamide binding pocket via π - π stacking interactions, significantly increases the activity of bisubstrate-like NNMT inhibitors (IC₅₀ 1.41 μ M). These findings are further supported by ITC binding assays as well as modeling studies. The most active NNMT inhibitor identified in the present study demonstrated a dose-dependent inhibitory effect on the cell proliferation of an HSC-2 oral cancer cell line.

INTRODUCTION

Nicotinamide *N*-methyltransferase (NNMT) is an important metabolic enzyme that catalyzes the transfer of a methyl group from the co-factor *S*-adenosyl-L-methionine (SAM) onto its various substrates, most notably nicotinamide (NA) and other pyridines, to form 1-methyl-nicotinamide (MNA) or the corresponding pyridinium ions.^{1–3} The past decade has seen a renewed interest in the biological function of NNMT in a range of human diseases. While it was previously assumed that NNMT's primary roles were limited to nicotinamide metabolism and xenobiotic detoxification of endogenous metabolites, broader roles for NNMT in human health and disease are becoming clearer.⁴ NNMT has been found to be overexpressed in a variety of diseases, including metabolic disorders^{5–7}, cardiovascular disease^{8,9}, cancer^{10–14} and Parkinson's disease^{15,16}. In general, overexpression of NNMT has been linked to disease progression in the aforementioned afflictions, with the exception of its role in Parkinson's disease where NNMT seems to be neuroprotective.^{17,18} Collectively, NNMT appears to play a unique role in the regulation of post-translational modifications and signal transduction, making it an attractive and viable therapeutic target.

Despite growing interest, few small molecule NNMT inhibitors have been described to date. Among these structures, the product of the enzymatic reaction, MNA, is a known inhibitor of NNMT and has generally been used in biochemical activity assays.¹⁹ Recently, Cravatt and

coworkers reported chloroacetamide-based covalent NNMT inhibitors that react with cysteine C165 in the SAM-binding pocket of the enzyme.²⁰ Notably, Sanofi researchers also recently reported a series of nicotinamide analogues that inhibit NNMT activity, leading to decreased MNA production, stabilization of insulin levels, glucose regulation, and weight loss in mouse models of metabolic disorders.^{21,22} In another approach, the group of Watowich focused on the development of inhibitors based on NNMT's alternative substrate quinoline. Their compounds showed improvement of symptoms in diet-induced obese mice.²³ Previous work in our group has focused on bisubstrate inhibitors designed to mimic the transition state of the methylation reaction catalyzed by NNMT with compound **1** (Fig. 1) showing activity on par with the known general methyltransferase inhibitor sinefungin.²⁴

Designing bisubstrate analogues as inhibitors is an established and effective strategy that has been applied to a range of methyltransferase enzymes including catechol *O*-methyltransferase (COMT),²⁵⁻²⁶ histone lysine methyltransferases,²⁷ arginine methyltransferases,^{28–30} and more recently nicotinamide *N*-methyltransferase.^{24,31} A recently published co-crystal structure of a bisubstrate inhibitor bound to NNMT (PDB ID: 6CHH) clearly delineates key interactions with residues in the enzyme active site, providing valuable information for further optimization of improved bisubstrate-like inhibitors.³¹ The work here described builds on our previous findings for "trivalent" inhibitor **1**, which is assumed to simultaneously bind in the adenosine, amino acid, and nicotinamide binding pockets of the NNMT active site. Based upon insights provided by recent NNMT crystal structures, we have designed new inhibitors wherein the nicotinamide moiety is replaced by other aromatic substituents accompanied by variation in the length of the linker connecting the amino acid moiety. Based on the high conservation of the residues in the

adenosine binding pocket, no changes were made to the adenosine group. A schematic overview of the design strategy is been presented in Figure 1.



Figure 1. Schematic overview of the design strategy of the second generation of inhibitors based on trivalent bisubstrate compounds 1^{24} and 2^{31}

RESULTS AND DISCUSSION

Design: The ternary crystal structure of NNMT (PDB ID: 3ROD) reveals the interactions of nicotinamide and *S*-Adenosyl-L-homocysteine (SAH) with the active site residues.³² The active site can be roughly divided into three binding regions for the adenosine group, the amino acid moiety, and the nicotinamide unit. The starting point was trivalent bisubstrate compound **1**, which was designed to bind all three binding regions. In order to find the optimal substitutions, a systematic approach was applied, where variations were made to the nicotinamide mimic on the one hand and the amino acid moiety on the other. The benzamide group, representing nicotinamide, was also replaced by methyl benzoate or benzoic acid moieties. Notably, the crystal structure of the NNMT–nicotinamide–SAH ternary complex reveals π - π stacking between tyrosine residue Y204 and the nicotinamide substrate.³² We therefore also prepared an analogue bearing a naphthalene unit in the presumed nicotinamide position with the aim of introducing stronger π - π stacking with the tyrosine residues of the NNMT active site. We also

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explored variation of the amino acid moiety as part of our design strategy: in some analogues the amine of the amino acid unit was omitted to reduce charge and in others the carboxylic acid replaced by the corresponding primary amide. In addition, variation in the length of the carbon chain linking the amino acid moiety was examined. Furthermore, inspired by the structure of histone methyltransferase DOTL1 inhibitor pinometostat,³³ we also investigated the incorporation of an isopropyl group to replace the amino acid moiety entirely.

Synthesis: Key aldehyde intermediates (compounds 6, 8, 9, 16, 17, 22, 23, 27, 28) required for the synthesis of the various bisubstrate analogues pursued were prepared from commercially available materials, in good overall yields, as summarized in Schemes 1-3. The trivalent inhibitors were then prepared via a convenient double reductive amination strategy starting from the commercially available 2'-3'-*O*-isopropylidene-6-aminomethyl-adenosine starting material and the corresponding aldehydes (Schemes 4 and 5).

The preparation of aromatic aldehydes 6, 8, and 9 began with the selective mono-deprotection of dimethyl isophthalate using sodium hydroxide (Scheme 1).³⁴ Monomethyl isophthalate (3) was subsequently transformed into trityl-protected amide 4 using tritylamine via its acid chloride intermediate and reduced by diisobutylaluminum hydride (DIBAL-H) to give alcohol 5. The alcohol was oxidized to aldehyde 6 using pyridinium dichromate (PDC). For aldehydes 8 and 9, the carboxylic acid of 3 was selectively reduced using a mixture of sodium borohydride and boron trifluoride diethyl etherate.³⁵ The resulting alcohol (7) was oxidized using PDC to yield the corresponding aldehyde (8). Following hydrolysis of the methyl ester in 8 and subsequent conversion to the *tert*-butyl ester, aldehyde 9 was obtained.³⁶



Scheme 1. Synthetic route for aldehydes 6, 8 and 9. *Reagents and conditions:* (a) NaOH, MeOH, rt, 16 h (95%); (b) i) SOCl₂, reflux, 2h, ii) tritylamine, CH_2Cl_2 , 0°C-rt, 2 h (72%); (c) DIBAL-H, -78°C-rt, 2 h (85%); (d) PDC, CH_2Cl_2 , rt, 2 h (53-64%); (e) NaBH₄, BF₃ Et₂O, THF, 0°C-rt, 2 h (89%); (f) LiOH, THF/H₂O (2:1); (g) 2-*tert*-butyl-1,3-diisopropylisourea, CH_2Cl_2 , *tert*-butanol (39% over 2 steps).

Aliphatic aldehydes **16** and **17** containing trityl-protected amide functionalities were prepared from succinimide and glutarimide respectively (Scheme 2). The cyclic amides were first trityl-protected and subsequently ring-opened using potassium hydroxide. Reduction to the corresponding alcohols and oxidization using PDC gave aldehydes **16** and **17**.^{37,38} In analogous fashion, aldehydes **22** and **23**, both containing *tert*-buytl ester moieties, were prepared by ring opening of succinic or glutaric anhydride with *tert*-buytl alcohol to obtain mono-esters **18** and **19**.^{39,40} The carboxylic acid functionalities were reduced to alcohols **20** and **21** and then oxidized using PDC to yield aldehydes **22** and **23**.



Scheme 2. Synthetic route for aldehydes 16, 17, 22 and 23. *Reagents and conditions:* (a) TrtCl, CH₃CN, K₂CO₃, rt, 48 h (20-28%); (b) KOH, EtOH, reflux, overnight (37-93%) (c) NaBH₄, BF₃Et₂O, THF, 0°C-rt, 2 h (64-81%); (d) PDC, CH₂Cl₂, rt, 2 h (65-78%); (e) *tert*-butanol, DMAP, *N*-Hydroxysuccinimide, Et₃N, toluene, overnight (25-93%).

Aldehydes **27** and **28**, both containing protected amino acid functionalities, were prepared starting from the appropriately protected aspartic acid and glutamic acid building blocks (Scheme 3). Conversion of the side chain carboxylates to their corresponding Weinreb amides yielded intermediates **24** and **25**. Reduction of aspartate-derived **24** with DIBAL-H gave amino acid aldehyde **27** in high yield. For the preparation of aldehyde **28**, a similar route was followed with the addition of a second Boc-protection of intermediate **25** to avoid an intramolecular cyclization side reaction.^{24,41}



Scheme 3. Synthetic route for aldehydes **27** and **28**. *Reagents and conditions:* (a) CH₃NHOCH₃·HCl, BOP, Et₃N, CH₂Cl₂, rt, 2 h (85-88%); (b) (Boc)₂O, Et₃N, DMAP, CH₂Cl₂ (94%); (c) DIBAL-H in hexanes (1 M), THF, -78°C, assumed quant.

With the necessary aldehyde building blocks in hand, assembly of the bisubstrate inhibitors was performed in each case starting from commercially available 2'-3'-O-isopropylidene-6-aminomethyl-adenosine (Scheme 4). Using a reliable reductive amination approach, aromatic aldehydes 6, 8, 9, and commercially available 2-naphthaldehyde were each coupled to the protected adenosine species to yield intermediates 29-32. These intermediates were next connected with aliphatic aldehydes 16, 17, 22, 23, 27, 28 or acetone via a second reductive amination step to give the corresponding protected tertiary amine intermediates 33-56 (Scheme 5). Global deprotection of the acid-labile protecting groups was carried out in CH₂Cl₂/TFA (1:1) with isopropylidene group cleavage facilitated by subsequent addition of water. The crude products were purified by preparative HPLC to yield bisubstrate analogues 1, 57-60.



Scheme 4. Synthesis of intermediate compounds **29-32**. *Reagents and conditions:* (a) NaBH(OAc)₃, AcOH, DCE, rt, overnight (50-74%).



Scheme 5. Representative scheme for the synthesis of the final compounds, shown for compounds 1 and 57-61. The same procedure was used starting from aldehydes 30-32 to form intermediate compounds 39-56 and 80 and final compounds 62-79 and 81 as detailed in the experimental section. *Reagents and conditions*: (a) aldehyde, NaBH(OAc)₃, AcOH, DCE, rt, overnight (49-77%); (b) i) TFA, CH₂Cl₂, rt, 2h, ii) H₂O, rt, 30 min (47-73%).

Inhibition Studies: The bisubstrate analogues were next tested for their NNMT inhibitory activity using a method recently developed in our group.² This assay employs Ultra High Performance (UHP) Hydrophilic Liquid Interaction Chromatography (HILIC) coupled to Quadrupole Time-Of-Flight Mass Spectrometry (QTOF-MS) to rapidly and efficiently assess NNMT inhibition by analysis of the formation of MNA. The NNMT inhibition of all compounds was initially screened at a fixed concentration of 250 μ M for all of the compounds. In cases where at least 50% inhibition was detected at this concentration full inhibition curves were measured in triplicate to determine the corresponding IC₅₀ values. As reference compounds, we included the well-established and general methyltransferase inhibitors sinefungin and SAH. In addition, we also synthesized two recently described NNMT inhibitors, compound **2** and 6-

(methylamino)-nicotinamide, following the procedures described in the corresponding publications.^{21,31} The structures of these reference compounds are provided in Figure 2.



Figure 2. Chemical structures of the reference compounds used in NNMT inhibition studies

The results of the NNMT inhibition studies are summarized in Table 1 and clearly show that only minor adjustments to the functional groups found in the enzyme's natural substrates are tolerated. Among the compounds studied, the most potent inhibition was observed when the aliphatic moiety corresponded to the same length in the amino acid side chain as present in the methyl donor SAM. Notably, the preferred aromatic moiety was found to be the naphthalene group, an apparent confirmation of our hypothesis that increased π - π stacking can lead to enhanced binding in the nicotinamide pocket. The bisubstrate analogue containing both of these elements (compound **78**), displayed the highest inhibitory activity against NNMT with an IC₅₀ of 1.41 µM. Interestingly, the amino acid and naphthyl moieties were also found to independently enhance the activity of the other inhibitors prepared. In this way a suboptimal moiety at one

position can be compensated – to an extent – by including either the SAM amino acid motif or the naphthalene unit at the other position. For example, bisubstrate analogues containing the benzamide, benzoic acid, or methyl benzoate groups only show inhibitory activity if they also contain the amino acid motif (compounds 1, 2, 66, 72) with IC₅₀ values of 4.36-23.4 μ M respectively. On the other hand, among the bisubstrate analogues lacking the amino acid motif, inclusion of the naphthalene moiety (compounds 74-79) enhances NNMT inhibition albeit with moderate IC₅₀ values in the range of 52.6-129.9 μ M.

Other notable findings were the results obtained with the reference compounds. The general methyltransferase inhibitors sinefungin and SAH showed inhibitory activities in line with those previously reported.²⁴ Interestingly, the 6-methylamino-NA compound recently described by Sanofi to be a submicromolar inhibitor,²¹ gave an IC₅₀ of 19.8 μ M in our assay. The recently published bisubstrate analogue **2** exhibited good activity (IC₅₀ 4.4 μ M) on par with published values.³¹ Given the potent inhibition measured for both compound **2** and **78**, we also prepared and tested compound **81**, an analogue of **78** bearing the same naphthyl moiety but with the amino acid motif containing an additional methylene unit as in **2**. Somewhat surprisingly, this linker elongation resulted in a complete loss of inhibitory activity (IC₅₀>250 μ M).

To gain insight into the selectivity of compound **78**, we also tested its activity against representative members of both the arginine and lysine families of methyltransferases, PRMT1 and NSD2 respectively. In both cases compound **78** was tested at a concentration of 50 μ M and showed no significant inhibition (>50% of the enzyme's activity remained), see Table S1.

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Table 1. Tabulated overview of the chemical structures and inhibition results of the final compounds and reference compounds

			IC ₅₀ values $(\mu M)^a$				
Reference compounds		Sinefungin		SAH		6-methylamino- nicotinamide	
IC ₅₀ (µM)		12.51 ± 2.11		35.30 ± 5.48		19.81 ± 2.50	
		R ₁					
		H ₂ N ² z _z () _n	MeO		HO		
R ₂	NH ₂	57 (n=1): >250	62 : >	>250	68 : >250		74 : 111.50 ± 28.79
	NH2	58 (n=1): >250	63: >	>250	69 : >250		75 : 52.62 ± 9.08
	O DH	59 (n=1): >250	64: >	>250	70 : >250		76 : >250
	O V	60 (n=1): >250	65: >	>250	71 : >250		77: >250
	O S ^{S²} (M NH ₂ OH	1 (m=1, n=1): 14.90 ± 2.07 2 (m=2, n=2): 4.36 ± 0.27	66 (1 17.4	m=1, n=1): 5 ± 2.65	72 (m=1, n=1): 23.41 ± 4.86		78 (m=1, n=1): 1.41 ± 0.16 81 (m=2, n=2): >250
		61 (n=1): >250	67: >	>250	73 : >250		79 : 129.90 ± 14.80

^aAssays performed in triplicate on at least six different inhibitor concentrations. Standard errors of the mean reported.

ITC binding studies: To further evaluate the binding interactions of the most active bisubstrate analogues with NNMT, isothermal titration calorimetry (ITC) studies were performed, Compounds, **1**, **66**, **72** and **78**, all containing the amino acid moiety but with varying aromatic substituents, were investigated. As illustrated in Figure 3, the dissociation constants (K_d) measured for these compounds track very well with the IC₅₀ values measured in the *in vitro* assay. Compounds **1** and **66** display similar binding to NNMT with K_d values of 36 µM and 25 µM respectively while compound **72** binds less tightly with a K_d of 124 µM. In good agreement with the results of the inhibition assay, the most active inhibitor, compound **78**, also displayed the highest binding affinity for NNMT with a K_d of 5.6 µM. As expected, the inhibitors were each found to bind the enzyme with a 1:1 stoichiometry.



Figure 3. ITC isotherms and thermograms including thermodynamic binding parameters measured for compounds 1, 66, 72, and 78 with hNNMT.

Modeling studies:

To further investigate the way in which the inhibitors bind within the NNMT active site modeling studies were performed. Working from the available crystal structure of NNMT protein bounded to nicotinamide and SAH (PDB ID: 3ROD)³², compounds 1, 2, 78, and 81 were modeled in the binding pocket. In an attempt to explain the significant difference in activity of 78 and 81 additional molecular dynamic simulations were also performed for compounds 1, 2, 78 and 81. While these simulations suggest differences in the binding interaction of the compounds (supplemental figure S1), the calculated binding energies for each are all very similar (supplemental table S2). In terms of their active site orientations, compounds 1, 2, 78 and 81 are all predicted to position their three branches roughly in the same regions of the active site, however their orientations and interactions are quite different.

From the modeling data, two distinct features are apparent. First, when the chain linking the amino acid moiety is shorter (as in compounds 1 and 78), the formation of an intramolecular hydrogen bond interaction was observed between the carboxylate of the amino acid moiety and the protonated tertiary amine (see Figure 4). This intramolecular interaction is highly stable for compound 78 and less stable for compound 1. This additional interaction reduces the entropic energy of the ligand, thereby potentially stabilizing its binding, and re-orients the amino acid part in the pocket, preventing the polar interactions with neighboring residues (e.g. Y25, D61, Y69, and T163) observed when the chain is longer (as present in compounds 2 and 81). This intramolecular hydrogen bond may explain the difference in activity observed between compounds 78 and 81. The second distinct feature is the tyrosine rich environment around the naphthalene moiety of 78 compared to the nicotinamide unit of 1. The orientation of the tyrosine residues surrounding this part of the molecule leads to π - π stacking interactions with the

naphthalene and hint at an explanation for the strong inhibition and high affinity of compound **78** with NNMT protein (Fig. 4).



Figure 4. Modeling results for compound **78** in the NNMT active site (PDB ID: 3ROD). Molecular dynamics simulation indicates the presence of an intramolecular hydrogen bond (2.7Å, shown in cyan) specific to compound **78** (in green) that would be expected to reduce the entropic energy of the ligand and potentially stabilize binding to NNMT (in white). Proposed intermolecular hydrogen bond network (in yellow) and π - π stacking interactions with Tyr residues (in purple) stabilize compound **78** in the NNMT active site (hydrogens omitted for clarity).

Cell based assays

To evaluate the cellular activity of the bisubstrate inhibitors, the compounds were tested for their effect on cell proliferation in the human oral cancer cell line HSC-2. We have shown recently that knockdown of NNMT in this particular cell line has a marked effect on cell proliferation and apoptosis.⁴² As shown in Figure 5 there were no significant differences in cell proliferation rate between HSC-2 cells treated with DMSO at 0.1% concentration and cells grown with culture medium only, at any time of each performed assay. Upon treatment with the NNMT inhibitors, cell proliferation was not significantly inhibited by compounds **1**, **2** and **81**

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(Figure 5). On the contrary, relative to the DMSO control, treatment with compound **78** led to a notable decrease in cell proliferation. In particular, cell proliferation was significantly (p < 0.05) inhibited by compound **78** at 10µM (20% reduction), 50 µM (21% reduction) and 100 µM (27% reduction) concentrations, 48 hours after treatment. Interestingly, at the longest 72 hours time point taken, treatment with compound **78** lead to an even greater and significant (p < 0.01) decrease in cell proliferation (44% reduction), at the highest concentration (100 µM) (Figure 5).

We next investigated the effect of compound **78** on cellular NNMT activity by assessing its impact on MNA production in the same HSC-2 cell line. Cells were treated with 100 μ M of **78** and MNA levels determined after 0, 1, 2, and 3 days. Cells treated with compound **78** show a significant (p < 0.01) decrease in the levels of MNA (50% reduction) compared to controls after 48 hours. Interestingly, at 72 hours an increase in cellular MNA production was detected, however, the same effect was also observed in the DMSO control (but not in the untreated control) suggesting an effect attributable to longer-term DMSO exposure. The results of the cellular MNA analysis are presented in supplemental Figure S2.



Figure 5. Results of the MTT cell viability assay on HSC-2 human oral cancer cells. Only compound **78** showed a significant effect on cell proliferation after 48 and 72 hours

CONCLUSION

Building from our earlier findings with first reported ternary bisubstrate NNMT inhibitor 1,²⁴ we designed and prepared a focused library of novel inhibitors to provide new structure-activity insights. In doing so, various structural motifs were investigated for the ability to enhance inhibitor activity and binding within the NNMT active site. By probing the SAM and NA binding pockets with different spacers and functional groups, we found that the optimal ligands are the endogenous amino acid side-chain and the naphthalene moiety. Among the naphthalene-

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containing bisubstrate analogues prepared, compound 78 showed the most potent NNMT inhibition. In this way the activity of our initial NNMT inhibitor 1 (IC₅₀ 14.9 μ M) was improved 10-fold with compound 78 displaying an IC_{50} value of 1.41 μ M. Notably, using an assay designed to directly measure NNMT product formation, compound 78 was shown to be more potent than most other NNMT inhibitor reported to date. ITC-based binding studies provided additional insights in the affinity of the inhibitors for the enzyme with measured K_d value following a trend similar to that observed for the IC_{50} data obtained in the *in vitro* inhibition assays. From modeling studies, the improved activity of compound 78 can be rationalized by the apparent presence of an intramolecular hydrogen bonding interaction predisposing the compound to an active conformation with lower entropic cost. In addition, the modeling indicates that the naphthalene group in 78 is properly oriented so as to benefit from additional π - π stacking interactions with several tyrosine residues in the nicotinamide binding pocket of the enzyme. The cellular data obtained for compound 78 shows a significant inhibitory effect on cell proliferation in HSC-2 oral cancer cells. These promising results provide important new insights for the design and further optimization of potent NNMT inhibitors.

EXPERIMENTAL PROCEDURES

General Procedures: All reagents employed were of American Chemical Society (ACS) grade or finer and were used without further purification unless otherwise stated. For compound characterization, ¹H NMR spectra were recorded at 400 MHz with chemical shifts reported in parts per million (ppm) downfield relative to tetramethylsilane (TMS), H₂O (δ 4.79), CHCl₃ (7.26) or DMSO (δ 2.50). ¹H NMR data are reported in the following order: multiplicity (s, singlet; d, doublet; t, triplet; q, quartet and m, multiplet), coupling constant (*J*) in hertz (Hz) and

the number of protons. Where appropriate, the multiplicity is preceded by br, indicating that the signal was broad. ¹³C NMR spectra were recorded at 101 MHz with chemical shifts reported relative to CDCl₃ (δ 77.16), methanol (δ 49.00) or DMSO (δ 39.52). The ¹³C NMR spectra of the compounds recorded in D₂O could not be referenced. High-resolution mass spectrometry (HRMS) analysis was performed using a Q-TOF instrument. Compounds 1,²⁴ 2,³¹ 3,³⁴ 7,³⁴ 8,³⁶ 9,³⁷ 10,⁴³ 12,³⁸ 14,³⁸ 16,⁴⁴ 18,⁴⁰ 19,⁴⁰ 20,⁴⁵ 21,⁴⁶ 22,⁴⁷ 23,⁴⁰ 24,⁴¹ 25,⁴¹ 26,⁴⁸ 27,⁴¹ 28⁴⁸ were prepared as previously described and had NMR spectra and mass spectra consistent with the assigned structures. Purity was confirmed to be \geq 95% by analytical RP-HPLC using a Phenomenex Kinetex C18 column (5 µm, 250 × 4.6 mm) eluted with a water–acetonitrile gradient moving from 0% to 100% CH₃CN (0.1% TFA) in 30 minutes. The compounds were purified via preparative HPLC using a Reprosil-Pur C18-AQ column (10 µm, 250 × 22 mm) eluted with a water–acetonitrile gradient moving from 0% to 100% CH₃CN (0.1% TFA) in 30 minutes. The compounds were purified via preparative HPLC using a Reprosil-Pur C18-AQ column (10 µm, 250 × 22 mm) eluted with a water–acetonitrile gradient moving from 0% to 50% CH₃CN (0.1% TFA) over 60 minutes at a flow rate of 12.0 ml min⁻¹ with UV detection at 214 nm and 254 nm.

Methyl 3-(tritylcarbamoyl)benzoate (4). Mono-methylisophthalate **3** (0.98 g, 5.4 mmol) was refluxed in 10 mL SOCl₂ at 90 °C for about one hour (until the reaction mixture is a clear solution). The SOCl₂ was removed under reduced pressure and the acid chloride intermediate was redissolved in 15 mL dry CH₂Cl₂ and transferred to a cooled (ice-bath) solution of tritylamine (1.41 g, 5.4 mmol) and 2 mL triethylamine in 30 mL CH₂Cl₂. The reaction was stirred overnight under N₂ atmosphere, allowing the mixture to warm to room temperature. After the reaction was completed (monitored by TLC (petroleum ether / CH₂Cl₂= 1:1)), the reaction mixture was washed with water and brine and the organic phase dried over Na₂SO₄ and concentrated. The crude product was purified by column chromatography (petroleum ether / CH₂Cl₂= 2:1) to give compound **4** as a white powder (1.64 g, 72% yield). ¹H NMR (400 MHz,

CDCl₃) δ 8.45 (t, J = 1.6 Hz, 1H), 8.18 (m, 1H), 8.03 (m, 1H), 7.53 (t, J = 7.8 Hz, 1H), 7.41 – 7.26 (m, 15H), 3.94 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 166.3, 165.4, 144.5, 135.6, 132.5, 131.7, 130.6, 128.9, 128.7, 128.1, 128.1, 127.6, 127.2, 71.0, 52.4. HRMS (ESI): calculated for C₂₈H₂₃NO₃ [M+Na]⁺ 444.1576, found 444.1581.

3-(Hydroxymethyl)-*N***-tritylbenzamide (5).** Methyl 3-(tritylcarbamoyl)benzoate **4** (0.56 g, 1.33 mmol) was dissolved in dry CH₂Cl₂ (20 mL) under a N₂ atmosphere, the reaction solution was cooled down to -78 °C, and then diisobutylaluminum hydride (DIBAL-H) (5.5 mL, 1.0 M hexane solution) was added slowly. The reaction mixture was stirred at -78°C for 2 hours. Saturated aq. NH₄Cl (50 mL) was added slowly to quench the reaction under -78°C, followed by the addition of a saturated Rochelle salt solution (100 mL). The mixture was stirred at room temperature overnight, extracted with CH₂Cl₂ and the organic layers dried over Na₂SO₄ and concentrated under reduced pressure. The crude product was purified by column chromatography (CH₂Cl₂/EtOAc = 9:1) to obtain **5** as a white powder (0.44 g, 85% yield). ¹H NMR (400 MHz, DMSO-*d*₆) δ 8.92 (s, 1H), 7.78 (s, 1H), 7.75 – 7.71 (m, 1H), 7.47 (d, *J* = 7.8 Hz, 1H), 7.40 (t, *J* = 7.6 Hz, 1H), 7.36 – 7.18 (m, 15H), 5.26 (br, 1H), 4.54 (s, 2H). ¹³C NMR (101 MHz, DMSO-*d*₆) δ 167.0, 145.3, 143.0, 135.5, 129.6, 128.9, 128.3, 127.9, 126.7, 126.5, 126.2, 79.6, 69.9, 69.9, 63.0. HRMS (ESI): calculated for C₂₇H₂₃NO₂ [2M+Na]⁺ 809.3355, found 809.3359.

3-Formyl-*N***-tritylbenzamide (6).** 3-(hydroxymethyl)-*N*-tritylbenzamide **5** (0.20 g, 0.51 mmol) and pyridinium dichromate (PDC) 0.23 g, 0.61 mmol) were placed in a 50 mL round bottom flask and 10 mL of dry CH_2Cl_2 was added under N_2 atmosphere at room temperature. The reaction was stirred till completion, as monitored by TLC (petroleum ether / $CH_2Cl_2 = 5:1$). The mixture was filtered and the organic layer was washed with brine, dried over anhydrous Na_2SO_4 and concentrated under reduced pressure. The resulting crude product was purified by

column chromatography (petroleum ether / $CH_2Cl_2 = 9:1$) to obtain **6** as a white powder (0.13 g, yield 64%). ¹H NMR (400 MHz, DMSO-*d*₆) δ 10.09 (s, 1H), 9.31 (s, 1H), 8.39 (s, 1H), 8.17 (d, *J* = 7.7 Hz, 1H), 8.06 (d, *J* = 7.7 Hz, 1H), 7.68 (t, *J* = 7.7 Hz, 1H), 7.41 – 7.17 (m, 15H). ¹³C NMR (101 MHz, CDCl₃) δ 191.5, 165.1, 144.4, 136.5, 136.2, 133.0, 132.5, 129.5, 128.6, 128.5, 128.1, 127.7, 127.3, 77.2, 71.1. HRMS (ESI): calculated for C₂₇H₂₁NO₂ [2M+Na]⁺ 805.3042, found 805.3047.

N-(triphenylmethyl)glutarimide (11). Glutarimide (2.8 g, 25 mmol), triphenylchloromethane (7.4 g, 25 mmol), and potassium carbonate (3.7 g, 25 mmol) were added to 100 mL acetonitrile and the mixture was stirred at room temperature overnight. Saturated aqueous NaHCO₃ (50 mL) was added and the mixture was extracted with EtOAc. The combined organic layers were dried with anhydrous Na₂SO₄ and the solvent was removed under reduced pressure. The crude product was purified by column chromatography (petroleum ether / EtOAc = 4:1) to obtain **11** as a white powder (1.8 g, yield 20%). ¹H NMR (400 MHz, DMSO-*d*₆) δ 7.45 – 7.35 (m, 6H), 7.20 (t, *J* = 7.8 Hz, 6H), 7.08 (t, *J* = 7.3 Hz, 3H), 2.66 (t, *J* = 6.4 Hz, 4H), 2.01 (p, *J* = 6.5 Hz, 2H). ¹³C NMR (101 MHz, CDCl₃) δ 172.4, 143.4, 128.5, 127.3, 125.9, 35.5, 16.7. HRMS (ESI): calculated for C₂₄H₂₁NO₂ [M+Na]⁺ 378.1470, found 378.1493.

5-Oxo-5-(tritylamino)pentanoic acid (13). To 2.80 g of KOH dissolved in 50 ml of ethanol was added *N*-tritylglutarimide **11** (1.00 g, 2.8 mmol) and the mixture was refluxed for 48 hours. The mixture was then concentrated to dryness and redissolved in H₂O. Acidification of the basic solution with conc. HCl to pH=2 and filtration of the product gave compound **13** as a white powder (0.96 g, yield 91%). ¹H NMR (400 MHz, CD₃OD) δ 7.30 – 7.17 (m, 15H), 2.37 (t, *J* = 7.4 Hz, 2H), 2.25 (t, *J* = 7.4 Hz, 2H), 1.79-1.87 (m 2H). ¹³C NMR (101 MHz, CD₃OD) δ 175.5,

173.3, 144.6, 128.6, 127.3, 127.2, 126.7, 126.3, 35.2, 32.6, 20.7. HRMS (ESI): calculated for $C_{24}H_{23}NO_3 [M+Na]^+ 396.1576$, found 396.1573.

5-Hydroxy-*N***-tritylpentanamide (15).** To a solution of **13** (2.60 g, 6.96 mmol) in dry THF (60 mL) cooled to 0°C was added NaBH(OAc)₃ (0.28 g, 7.3 mmol). The solution was stirred until evolution of H₂ stopped, and BF₃.OEt₂ (1.1 mL, 8.8 mmol) was added dropwise. The reaction was stirred at room temperature for 4 hours. The reaction was quenched by adding 50 mL H₂O at 0°C. The mixture was extracted with EtOAc and the combined organic layers were washed with sat. aq. Na₂CO₃, brine and dried over Na₂SO₄. The crude product was purified by column chromatography (100% EtOAc) to give compound **15** as a white powder (1.60 g, 64% yield). ¹H NMR (400 MHz, CDCl₃) δ 7.22 – 6.74 (m, 15H), 6.36 (br, 1H), 3.29 – 3.19 (br, 2H), 2.01 (t, *J* = 7.2 Hz, 2H), 1.46 – 1.36 (m, 2H), 1.24 (m, 2H). ¹³C NMR (101 MHz, CDCl₃) δ 171.9, 144.7, 128.6, 127.9, 127.0, 62.0, 37.0, 32.0, 21.4. HRMS (ESI): calculated for C₂₄H₂₅NO₂ [M+Na]⁺ 382.1783, found 382.1783.

5-Oxo-*N***-tritylpentanamide (17).** 5-hydroxy-*N*-tritylpentanamide **15** (1.30 g, 3.6 mmol) and PDC (2.00 g, 5.4 mmol) were dissolved in 50 mL of dry CH₂Cl₂ and stirred for 2 hours under N₂ atmosphere at room temperature. The mixture was filtered and the organic layer was washed with brine, dried over anhydrous Na₂SO₄ and concentrated under reduced pressure. The crude product was purified by column chromatography (100% CH₂Cl₂) to give compound **17** as an off-white powder (0.84 g, 65% yield). ¹H NMR (400 MHz, CDCl₃) δ 9.71 (s, 1H), 7.36 – 7.10 (m, 15H), 6.59 (s, 1H), 2.44 (t, *J* = 7.0 Hz, 2H), 2.32 (t, *J* = 7.2 Hz, 2H), 1.97 – 1.88 (m, 2H). ¹³C NMR (101 MHz, CDCl₃) δ 202.0, 170.8, 144.6, 128.6, 127.9, 127.0, 70.5, 42.9, 36.1, 17.9. HRMS (ESI): calculated for C₂₄H₂₃NO₂ [M+Na]⁺ 380.1626, found 380.1629.

3-(((((3aR.4R,6R.6aR)-6-(6-amino-9H-purin-9-vl)-2,2-dimethyltetrahydrofuro[3,4-d][1,3]dioxol-4-yl)methyl)amino)methyl)-N-tritylbenzamide (29). 3-Formyl-N-tritylbenzamide 6 (1.22 g, 3.12 mmol), 2'-3'-O-isopropylidene-6-aminomethyl-adenosine (1.00 g, 3.43 mmol) and acetic acid (0.45 mL, 8 mmol) were dissolved in 1,2-dichloroethane (DCE, 50 mL) and stirred at room temperature under a N₂ atmosphere. After 3 hours, NaBH(OAc)₃ (1.09 g, 5.15 mmol) was added and the reaction mixture was stirred overnight at room temperature. The reaction was quenched by adding 1 N NaOH solution (50 mL), and the product was extracted with CH₂Cl₂. The combined organic layers were washed with brine and dried over Na₂SO₄. The solvent was evaporated and the crude product was purified by column chromatography (10% MeOH in CH₂Cl₂) to give compound **29** as a white powder (1.25 g, 59% yield). ¹H NMR (400 MHz, DMSO- d_6) δ 8.89 (s, 1H), 8.34 (s, 1H), 8.06 (s, 1H), 7.79 (s, 1H), 7.71 (d, J = 7.7 Hz, 1H), 7.43 (d, J = 7.7 Hz, 1H) 1H), 7.39 - 7.24 (m, 15H), 7.20 (m, 3H), 6.09 (d, J = 3.1 Hz, 1H), 5.76 (s, 1H), 5.46 (M, 1H), 5.00 (m, 1H), 4.28 – 4.23 (m, 1H), 3.73 (s, 2H), 2.75 - 2.66 (m, 2H), 1.54 (s, 3H), 1.31 (s, 3H). ¹³C NMR (101 MHz, DMSO- d_6) δ 166.9, 156.5, 153.1, 149.3, 145.3, 140.4, 135.6, 128.9, 128.3, 127.9, 127.6, 126.7, 126.5, 119.7, 113.7, 89.7, 85.3, 83.1, 82.6, 69.9, 55.3, 53.0, 50.8, 27.5, 25.7. HRMS (ESI): calculated for $C_{40}H_{39}N_7O_4$ [M+Na]⁺ 704.2961, found 704.2975.

Methyl-3-(((((3aR,4R,6R,6aR))-6-((6-amino-9H-purin-9-yl)-2,2-dimethyltetrahydrofuro[3,4-d] [1,3]dioxol-4-yl)methyl)amino)benzoate (30). Following the procedure described for compound 29, coupling methyl 3-formylbenzoate 8 (0.51 g, 3.12 mmol) and 2'-3'-*O*-isopropylidene-6aminomethyl-adenosine (1.00 g, 3.43 mmol) afforded compound 30 as a white powder (0.92 g, 65% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.08 (s, 1H), 7.92 (s, 1H), 7.90 – 7.83 (m, 2H), 7.44 (d, *J* = 7.6 Hz, 1H), 7.32 (t, *J* = 7.6 Hz, 1H), 6.37 (d, *J* = 5.7 Hz, 2H), 5.95 (d, *J* = 3.1 Hz, 1H), 5.45 (m, 1H), 5.04 (m, 1H), 4.40 – 4.34 (m, 1H), 3.86 (s, 3H), 3.79 (s, 2H), 2.90-2.83 (m, 2H),

1.58 (s, 3H), 1.35 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 167.1, 155.8, 155.8, 153.0, 149.2, 140.4, 140.4, 139.8, 132.6, 132.6, 130.1, 129.1, 129.1, 128.4, 128.4, 128.2, 120.2, 114.4, 91.0, 85.5, 83.2, 83.2, 82.2, 82.2, 53.3, 52.1, 52.1, 50.6, 27.3, 27.2, 25.4, 25.3. HRMS (ESI): calculated for C₂₂H₂₆N₆O₅ [M+H]⁺ 455.2043, found 455.2050.

tert-Butyl-3-(((((3aR,4R,6R,6aR))-6-(6-amino-9H-purin-9-yl)-2,2-dimethyltetrahydrofuro[3,4d][1,3]dioxol-4-yl)methyl)amino)methyl)benzoate (31). Following the procedure described for compound 29, coupling *tert*-butyl 3-formylbenzoate 9 (0.64 g, 3.12 mmol) and 2'-3'-*O*isopropylidene-6-aminomethyl-adenosine (1.00 g, 3.43 mmol) afforded compound 31 as a white powder (0.77 g, 50% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.11 (s, 1H), 7.89 (s, 1H), 7.86-7.83 (m, 2H), 7.43 (d, *J* = 7.7 Hz, 1H), 7.31 (t, *J* = 7.6 Hz, 1H), 6.36 – 6.27 (m, 2H), 5.96 (d, *J* = 3.3 Hz, 1H), 5.46 (m, 1H), 5.04 (m, 1H), 4.38 (m, 1H), 3.80 (s, 2H), 2.94-2.81 (m, 2H), 1.58 (s, 3H), 1.55 (s, 9H), 1.36 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 165.7, 155.8, 155.8, 153.0, 149.3, 140.2, 139.8, 132.0, 132.0, 129.0, 128.2, 128.1, 120.3, 114.5, 91.0, 85.4, 83.2, 82.2, 80.9, 53.4, 50.6, 28.1, 27.3, 25.4. HRMS (ESI): calculated for C₂₅H₃₂N₆O₅ [M+H]⁺ 497.2512, found 497.2511.

9-((3aR,4R,6R,6aR)-2,2-dimethyl-6-(((naphthalen-2-ylmethyl)amino)methyl)tetrahydrofuro-

[3,4-d][1,3]dioxol-4-yl)-9*H*-purin-6-amine (32). Following the procedure described for compound 29, coupling 2-naphthaldehyde (0.49 g, 3.12 mmol) and 2'-3'-*O*-isopropylidene-6-aminomethyl-adenosine (1.00 g, 3.43 mmol) afforded compound 32 as a white powder (1.03 g, 74% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.11 (s, 1H), 7.88 (s, 1H), 7.78 (m, 3H), 7.70 (s, 1H), 7.48 – 7.38 (m, 3H), 6.05 (s, 2H), 5.99 (d, J = 3.3 Hz, 1H), 5.48 (m, 1H), 5.06 (m, 1H), 4.45 – 4.39 (m, 1H), 3.95 (s, 2H), 3.01 – 2.87 (m, 2H), 2.33 (br, 2H), 1.61 (s, 3H), 1.38 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 155.7, 153.0, 149.3, 139.9, 137.4, 133.3, 132.6, 128.0, 127.6 126.4,

3-(((((3aR,4R,6R,6aR)-6-(6-amino-9H-purin-9-yl)-2,2-dimethyltetrahydrofuro[3,4-d][1,3]dioxol-4-yl)methyl)(4-oxo-4-(tritylamino)butyl)amino)methyl)-N-tritylbenzamide (33). Oxo-N-tritylbutanamide 16 (62 mg, 0.18 mmol), compound 29 (100 mg, 0.15 mmol) and AcOH (1 drop) were dissolved in 1,2-dichloroethane (DCE, 10 mL) and stirred at room temperature under a N₂ atmosphere. After 3 hours, NaBH₄ (49 mg, 0.23 mmol) was added and the reaction mixture was stirred overnight at room temperature. The reaction was guenched by adding 1 N NaOH (10 mL), and the product was extracted with CH₂Cl₂. The combined organic layers were washed with brine and dried over Na_2SO_4 . The solvent was evaporated and the crude product was purified by column chromatography (10% MeOH in CH₂Cl₂) to give compound 33 as a white powder (83 mg, 55% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.15 (s, 1H), 7.69 (s, 1H), 7.67 (s, 1H), 7.53 (d, J = 7.1 Hz, 2H), 7.39 – 7.09 (m, 32H), 6.61 (s, 1H), 5.95 (d, J = 1.9 Hz, 1H), 5.65 (s, 2H), 5.36 (m, 1H), 4.89 (m, 1H), 4.40 - 4.34 (m, 1H), 3.56 (d, J = 3.4 Hz, 2H), 2.68 (d, J =6.8 Hz, 2H), 2.46 (m, 2H), 2.26 (m, 2H), 1.81 - 1.69 (m, 2H), 1.52 (s, 3H), 1.30 (s, 3H). ^{13}C NMR (101 MHz, CDCl₃) δ 171.5, 166.7, 155.4, 152.9, 149.0, 144.8, 144.7, 140.0, 139.9, 135.3, 131.5, 128.8, 128.7, 128.0, 127.9, 127.0, 126.9, 125.3, 114.1, 90.8, 85.7, 83.8, 83.4, 70.7, 70.4, 58.6, 56.0, 53.5, 34.9, 27.0, 25.3, 22.7. HRMS (ESI): calculated for $C_{63}H_{60}N_8O_5$ [M+H]⁺ 1009.4765, found 1009.4765.

3-(((((3aR,4R,6R,6aR)-6-(6-amino-9H-purin-9-yl)-2,2-dimethyltetrahydrofuro[3,4-

d][1,3]dioxol-4-yl)methyl)(5-oxo-5-(tritylamino)pentyl)amino)methyl)-N-tritylbenzamide

(34). Following the procedure described for compound 33, coupling compound 29 (100 mg, 0.15 mmol) with 5-oxo-*N*-tritylpentanamide 17 (64 mg, 0.18 mmol) afforded compound 34 as a white

powder (88 mg, 57% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.16 (s, 1H), 7.67 (s, 2H), 7.57 (br, 1H), 7.52 (s, 1H), 7.41 – 7.13 (m, 32H), 6.62 (s, 1H), 5.96 (d, *J* = 1.8 Hz, 1H), 5.83 (br, 2H), 5.38 (m, 1H), 4.92 (m, 1H), 4.40 – 4.34 (m, 1H), 3.54 (s, 2H), 2.65 (d, *J* = 6.9 Hz, 2H), 2.46 – 2.38 (m, 2H), 2.13 (m, 2H), 1.56 (s, 3H), 1.42 – 1.33 (m, 2H), 1.30 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 171.6, 166.6, 155.5, 152.9, 149.1, 144.8, 144.8, 140.3, 140.0, 135.2, 131.6, 128.8, 128.7, 128.2, 128.0, 127.9, 127.5, 127.0, 127.0, 125.4, 114.1, 90.9, 85.9, 83.8, 83.4, 77.3, 70.7, 70.4, 58.6, 56.1, 53.9, 37.1, 27.1, 26.3, 25.4, 23.1. HRMS (ESI): calculated for C₆₄H₆₂N₈O₅ [M+H]⁺ 1023.4921, found 1023.4918.

tert-Butyl 4-((((3a*R*,4*R*,6*R*,6a*R*)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(3-(tritylcarbamoyl)benzyl)amino)butanoate (35). Following the procedure described for compound 33, coupling *tert*-butyl 4-oxobutanoate 22 (29 mg, 0.18 mmol) and compound 29 (100 mg, 0.15 mmol) afforded compound 35 as a white powder (61 mg, 49% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.16 (s, 1H), 7.70 (d, *J* = 5.8 Hz, 2H), 7.58 (d, *J* = 7.6 Hz, 2H), 7.38 – 7.15 (m, 17H), 5.97 (d, *J* = 2.0 Hz, 3H), 5.36 (m, 1H), 4.93 (m, 1H), 4.35 (m, 1H), 3.63 – 3.52 (m, 2H), 2.76 – 2.63 (m, 2H), 2.47 (t, *J* = 7.1 Hz, 2H), 2.23 – 2.12 (m, 2H), 1.75 – 1.65 (m, 2H), 1.55 (s, 3H), 1.36 (s, 9H), 1.29 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 172.7, 166.7, 155.5, 152.9, 149.0, 144.8, 144.8, 140.0, 139.9, 131.6, 128.8, 128.7, 128.0, 128.0, 127.0, 125.5, 120.1, 90.8, 85.8, 83.9, 83.4, 80.1, 70.7, 61.9, 58.8, 55.9, 53.4, 33.0, 32.4, 28.0, 27.1, 25.3, 22.4. HRMS (ESI): calculated for C₄₈H₅₃N₇O₆ [M+H]⁺ 824.4136, found 824.4142.

tert-Butyl-5-((((3aR,4R,6R,6aR)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(3-(tritylcarbamoyl)benzyl)amino)pentanoate (36). Following the procedure described for compound 33, coupling *tert*-butyl 5-oxopentanoate 23 (31 mg, 0.18 mmol) and compound 29 (100 mg, 0.15 mmol) afforded compound 36 as a white powder (67 mg, 53% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.17 (s, 1H), 7.70 (d, J = 8.7 Hz, 2H), 7.58 (d, J = 7.8 Hz, 1H), 7.51 (s, 1H), 7.42 – 7.27 (m, 14H), 7.19 (t, J = 7.7 Hz, 1H), 5.98 (s, 1H), 5.72 (s, 2H), 5.38 (m, 1H), 4.94 (s, 1H), 4.40 – 4.33 (m, 1H), 3.57 (s, 2H), 2.70 – 2.65 (m, 2H), 2.47 (t, J = 7.0 Hz, 2H), 2.13 (t, J = 7.2 Hz, 2H), 1.57 (s, 3H), 1.52 – 1.45 (m, 2H), 1.41 (s, 9H), 1.32 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 172.9, 166.6, 155.4, 152.9, 149.1, 144.8, 140.2, 139.9, 135.2, 131.6, 128.7, 128.2, 128.0, 127.4, 127.0, 125.4, 120.2, 114.1, 90.8, 85.8, 83.9, 83.4, 80.0, 70.7, 58.7, 56.0, 54.0, 35.2, 28.1, 27.1, 26.3, 25.3, 22.7. HRMS (ESI): calculated for C₄₉H₅₅N₇O₅ [M+H]⁺ 838.4292, found 838.4298.

tert-Butyl-(S)-4-((((3aR,4R,6R,6aR)-6-(6-amino-9H-purin-9-yl)-2,2-

dimethyltetrahydrofuro[3,4-d][1,3]dioxol-4-yl)methyl)(3-(tritylcarbamoyl)benzyl)amino)-2-((*tert*-butoxycarbonyl)amino)butanoate (37). Following the procedure described for compound 33, coupling *tert*-butyl (R)-2-((*tert*-butoxycarbonyl)amino)-4-oxobutanoate 27 (49 mg, 0.18 mmol) and compound 29 (100 mg, 0.15 mmol) afforded compound 37 as a white powder (94 mg, 67% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.15 (s, 1H), 7.70 (m, 2H), 7.57 (d, *J* = 9.9 Hz, 2H), 7.41 – 7.14 (m, 15H), 5.98 (s, 1H), 5.59 (s, 2H), 5.37 (m, 2H), 4.91 (s, 1H), 4.36 (s, 1H), 4.17 (s, 1H), 3.62 (d, *J* = 13.8 Hz, 1H), 3.54 (d, *J* = 13.8 Hz, 1H), 2.76 – 2.48 (m, 4H), 1.99 (d, *J* = 6.2 Hz, 1H), 1.76 (br, 1H), 1.57 (s, 3H), 1.39 (m, 15H), 1.32 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 171.6, 166.6, 155.4, 152.9, 149.0, 144.8, 140.0, 139.5, 135.4, 131.6, 128.8, 128.0, 127.0, 125.5, 114.2, 90.7, 85.6, 83.9, 83.4, 81.7, 79.4, 70.7, 58.9, 58.2, 55.9, 52.7, 50.9, 50.6, 36.5, 29.7, 28.3, 28.2, 28.0, 27.9, 27.1, 25.3. HRMS (ESI): calculated for C₅₃H₆₁N₈O₈ [M+H]⁺ 939.4749, found 939.4784.

3-((((((3a*R*,4*R*,6*R*,6a*R*)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro[3,4d][1,3]dioxol-4-yl)methyl)(*iso*propyl)amino)methyl)-*N*-tritylbenzamide (38). Following the

procedure described for compound **33**, coupling 5 mL dry acetone (large excess) and compound **29** (100 mg, 0.15 mmol) afforded compound **38** as a white powder (68 mg, 63% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.22 (s, 1H), 7.75 (s, 1H), 7.60 (d, J = 10.4 Hz, 2H), 7.49 – 7.40 (m, 2H), 7.33 – 7.21 (m, 15H), 5.91 (s, 1H), 5.34 (m, 2H), 4.92 – 4.87 (m, 1H), 4.23 (d, J = 3.2 Hz, 1H), 3.57 (s, 2H), 2.92 – 2.83 (m, 1H), 2.76 – 2.68 (m, 1H), 2.59 (m, 1H), 1.49 (s, 3H), 1.25 (s, 3H), 1.01 (d, J = 6.6 Hz, 3H), 0.90 (d, J = 6.6 Hz, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 166.7, 155.7, 152.9, 149.0, 144.8, 141.4, 139.9, 135.2, 131.4, 128.7, 128.3, 128.0, 127.1, 125.4, 120.2, 114.0, 90.9, 86.5, 83.7, 83.3, 70.7, 54.4, 51.8, 50.5, 27.1, 25.4, 18.7, 17.2. HRMS (ESI): calculated for C₄₃H₄₅N₇O₄ [M+H]⁺724.3611, found 724.3618.

Methyl 3-(((((3aR, 4R, 6R, 6aR)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(4-oxo-4-(tritylamino)butyl)amino)methyl)benzoate (39). Following the procedure described for compound 33, coupling 4-oxo-N-tritylbutanamide 16 (62 mg, 0.18 mmol) and compound 30 (68 mg, 0.15 mmol) afforded compound 39 as a white powder (63 mg, 54% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.17 (s, 1H), 7.90 (s, 1H), 7.84 – 7.79 (m, 1H), 7.77 (s, 1H), 7.40 (d, *J* = 7.7 Hz, 1H), 7.31 – 7.08 (m, 17H), 6.60 (s, 1H), 5.97 (d, *J* = 2.2 Hz, 1H), 5.67 (s, 2H), 5.34 (m, 1H), 4.86 (m, 1H), 4.34 (m, 1H), 3.85 (s, 3H), 3.57 (m, 2H), 2.74 – 2.62 (m, 2H), 2.45 (t, *J* = 7.0 Hz, 2H), 2.31-2.16 (m, 2H), 1.75 (m, 2H), 1.53 (s, 3H), 1.32 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 171.5, 167.0, 155.4, 153.0, 149.1, 144.7, 139.9, 139.6, 133.3, 129.9, 129.7, 128.7, 128.2, 128.1, 127.9, 127.0, 126.9, 120.2, 114.2, 90.8, 85.4, 83.7, 83.4, 70.4, 58.5, 55.7, 53.6, 52.1, 34.9, 27.0, 25.2, 22.7. HRMS (ESI): calculated for C₄₅H₄₈N₇O₆ [M+H]⁺782.3666, found 782.3666.

 $\label{eq:methyl} Methyl 3-(((((3aR,4R,6R,6aR)-6-(6-amino-9H-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(5-oxo-5-(tritylamino)pentyl)amino)methyl)benzoate (40).$

Following the procedure described for compound **33**, coupling 5-oxo-*N*-tritylpentanamide **17** (64 mg, 0.18 mmol) and compound **30** (68 mg, 0.15 mmol) afforded compound **40** as a white powder (67 mg, 53% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.17 (s, 1H), 7.89 (s, 1H), 7.83 (d, *J* = 7.8 Hz, 1H), 7.76 (s, 1H), 7.44 (d, *J* = 7.6 Hz, 1H), 7.26-7.17 (m, 14H), 6.69 (s, 1H), 6.35 (br, 2H), 5.99 (d, *J* = 1.8 Hz, 1H), 5.39 (m, 1H), 4.90 (m, 1H), 4.36 (m, 1H), 3.83 (s, 3H), 3.62 – 3.48 (m, 2H), 2.65 (m, 2H), 2.41 (t, *J* = 6.9 Hz, 2H), 2.14 (p, *J* = 8.2 Hz, 2H), 1.56 (s, 3H), 1.33 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 171.7, 167.1, 155.8, 152.9, 149.1, 144.8, 139.9, 139.8, 133.4, 129.9, 129.8, 128.7, 128.2, 128.2, 127.9, 126.9, 114.1, 90.8, 85.6, 83.7, 83.4, 70.4, 58.5, 55.8, 53.8, 52.1, 37.1, 27.1, 26.3, 25.3, 23.1. HRMS (ESI): calculated for C₄₆H₄₉N₇O₆ [M+H]⁺796.3823, found 796.3814.

Methyl 3-(((((3a*R*,4*R*,6*R*,6a*R*)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(4-(*tert*-butoxy)-4-oxobutyl)amino)methyl)benzoate (41). Following the procedure described for compound 33, coupling *tert*-butyl 4-oxobutanoate 22 (29 mg, 0.18 mmol) and compound 30 (68 mg, 0.15 mmol) afforded compound 41 as a white powder (65 mg, 73% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.13 (s, 1H), 7.85 (s, 1H), 7.80 (d, J = 8.1 Hz, 2H), 7.41 (d, J = 7.6 Hz, 1H), 7.23 (t, J = 7.6 Hz, 1H), 6.47 (s, 2H), 5.98 (d, J = 1.9 Hz, 1H), 5.33 (d, J = 6.4 Hz, 1H), 4.87 (m, 1H), 4.30 (m, 1H), 3.84 (s, 3H), 3.61-3.48(m, 2H), 2.75 – 2.69 (m, 2H), 2.43 (m, 2H), 2.16-2.10 (m, 2H), 1.53 (s, 3H), 1.39 – 1.30 (m, 15H). ¹³C NMR (101 MHz, CDCl₃) δ 172.8, 167.0, 155.8, 152.9, 148.9, 139.7, 139.6, 133.3, 129.9, 129.7, 128.2, 128.1, 120.0, 114.1, 114.1, 90.7, 85.4, 83.7, 83.3, 80.0, 61.6, 61.1, 58.6, 55.6, 53.4, 52.0, 32.9, 32.3, 28.0, 27.5, 27.0, 25.3, 22.3. HRMS (ESI): calculated for C₃₀H₄₀N₆O₇ [M+H]⁺ 597.3037, found 597.3037.

Methyl 3-(((((3aR, 4R, 6R, 6aR)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(5-(*tert*-butoxy)-5-oxopentyl)amino)methyl)benzoate (42). Following the procedure described for compound 33, coupling *tert*-butyl 5-oxopentanoate 23 (31 mg, 0.18 mmol) and compound 30 (68 mg, 0.15 mmol) afforded compound 42 as a white powder (56 mg, 61% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.12 (s, 1H), 7.84 (s, 1H), 7.79 (d, *J* = 7.8 Hz, 2H), 7.39 (d, *J* = 7.5 Hz, 1H), 7.21 (t, *J* = 7.7 Hz, 1H), 6.52 (s, 2H), 5.97 (s, 1H), 5.36 – 5.30 (m, 1H), 4.86 (m, 1H), 4.33 – 4.26 (m, 1H), 3.82 (s, 3H), 3.58 (d, *J* = 13.8 Hz, 1H), 3.47 (d, *J* = 13.7 Hz, 1H), 2.69-2.56 (m, 2H), 2.43 – 2.35 (m, 2H), 2.07 (t, *J* = 6.8 Hz, 2H), 1.52 (s, 3H), 1.48 – 1.42 (m, 2H), 1.34 (s, 9H), 1.30 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 172.8, 167.0, 155.8, 152.9, 149.0, 139.7, 139.7, 133.3, 129.9, 129.7, 128.2, 128.1, 128.1, 120.1, 114.1, 90.8, 85.5, 83.6, 83.5, 83.3, 79.9, 58.6, 55.6, 53.9, 52.0, 35.2, 28.0, 27.1, 26.2, 25.3, 22.7. HRMS (ESI): calculated for C₃₁H₄₂N₆O₇ [M+H]⁺611.3193, found 611.3182.

Methyl 3-(((((3aR,4R,6R,6aR)-6-(6-amino-9H-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)((S)-4-(*tert*-butoxy)-2-((*tert*-butoxycarbonyl)amino)-4-

oxobutyl)amino)methyl)benzoate (43). Following the procedure described for compound 33, coupling *tert*-butyl (*R*)-2-((*tert*-butoxycarbonyl)amino)-4-oxobutanoate 27 (49 mg, 0.18 mmol) and compound 30 (68 mg, 0.15 mmol) afforded compound 43 as a white powder (62 mg, 58% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.19 (s, 1H), 7.87 (d, *J* = 6.5 Hz, 2H), 7.82 (s, 1H), 7.48 (d, *J* = 7.7 Hz, 1H), 7.30 (t, *J* = 7.9 Hz, 1H), 6.01 (s, 1H), 5.73 (s, 2H), 5.38 (m, 2H), 4.89 (m, 1H), 4.35 (m, 1H), 4.20 – 4.11 (m, 1H), 3.90 (s, 3H), 3.71-3.52 (m, 2H), 2.78 (m, 1H), 2.65 (m, 2H), 2.51 (m, 1H), 1.96 (s, 2H), 1.76 (m, 1H), 1.59 (s, 3H), 1.40 (m, 18H), 1.37 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 171.6, 167.0, 155.4, 155.4, 153.0, 149.1, 139.9, 139.2, 133.4, 130.0, 129.8,

128.4, 128.3, 120.3, 114.3, 90.7, 85.3, 83.8, 83.3, 81.7, 79.4, 58.6, 55.7, 52.7, 52.1, 50.5, 29.5, 28.3, 27.9, 27.1, 25.3. HRMS (ESI): calculated for C₃₅H₄₉N₇O₉ [M+H]⁺ 712.3670, found 712.3682.

Methyl 3-(((((3a*R***,4***R***,6***R***,6***aR***)-6-(6-amino-9***H***-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d**][**1,3**]dioxol-4-yl)methyl)(*iso*propyl)amino)methyl)benzoate (44). Following the procedure described for compound **33**, coupling dry acetone (5 mL, large excess) and compound **30** (68 mg, 0.15 mmol) afforded compound **44** as a white powder (42 mg, 57% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.22 (s, 1H), 7.94 (s, 1H), 7.84 (d, *J* = 7.7 Hz, 1H), 7.77 (s, 1H), 7.52 (d, *J* = 7.6 Hz, 1H), 7.29 (t, *J* = 7.7 Hz, 1H), 5.96 (d, *J* = 2.4 Hz, 2H), 5.36 (dd, *J* = 6.4, 2.4 Hz, 1H), 4.87 (dd, *J* = 6.4, 3.0 Hz, 1H), 4.26 – 4.20 (m, 1H), 3.88 (s, 3H), 3.62 (d, *J* = 14.2 Hz, 1H), 3.54 (d, *J* = 14.2 Hz, 1H), 2.88 (p, *J* = 6.6 Hz, 1H), 2.73-2.59(m, 2H), 1.53 (s, 3H), 1.33 (s, 3H). ¹³C NMR (101 MHz, Chloroform-*d*) δ 167.1, 155.6, 152.9, 149.2, 141.0, 139.9, 133.2, 129.9, 129.6, 128.1, 128.1, 120.2, 114.0, 91.0, 86.1, 83.5, 83.2, 60.3, 54.3, 52.0, 51.4, 50.3, 27.1, 25.3, 21.0, 18.9, 16.8, 14.2. HRMS (ESI): calculated for C₃₅H₃₂N₆O₅ [M+H]⁺497.2512, found 497.2510.

tert-Butyl 3-(((((3a*R*,4*R*,6*R*,6a*R*)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(4-oxo-4-(tritylamino)butyl)amino)methyl)benzoate (45). Following the procedure described for compound 33, coupling 4-oxo-*N*-tritylbutanamide 16 (62 mg, 0.18 mmol) and compound 31 (75 mg, 0.15 mmol) afforded compound 45 as a white powder (93 mg, 75% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.23 (s, 1H), 7.90 (s, 1H), 7.84 – 7.79 (m, 2H), 7.43 (d, *J* = 7.6 Hz, 1H), 7.31 – 7.16 (m, 17H), 6.68 (s, 1H), 6.07 (s, 2H), 6.01 (d, *J* = 2.2 Hz, 1H), 5.39 (m, 1H), 4.89 (m, 1H), 4.37 (m, 1H), 3.66 (d, *J* = 13.9 Hz, 1H), 3.55 (d, *J* = 13.9 Hz, 1H), 2.76 (m, 1H), 2.66 (m, 1H), 2.47 (t, *J* = 6.8 Hz, 2H), 2.28 (m, 2H), 1.77 (m, 2H), 1.59 (s, 9H), 1.56 (s, 3H), 1.35 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 171.6, 165.7, 155.6, 153.0, 149.1, 144.8, 139.7, 139.4, 132.8, 131.9, 129.6, 128.7, 128.1, 128.0, 127.9, 126.9, 120.2, 114.2,

90.8, 85.4, 83.6, 83.4, 80.9, 70.4, 58.6, 55.7, 53.6, 34.9, 28.2, 27.1, 25.3, 22.7. HRMS (ESI): calculated for $C_{49}H_{53}N_7O_6 [M+H]^+$ 824.4136, found 824.4123.

tert-Butyl-3-(((((3*aR*,4*R*,6*R*,6*aR*)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(5-oxo-5-(tritylamino)pentyl)amino)methyl)benzoate (46). Following the procedure described for compound 33, coupling 5-oxo-*N*-tritylpentanamide 17 (64 mg, 0.18 mmol) and compound 31 (75 mg, 0.15 mmol) afforded compound 46 as a white powder (97 mg, 77% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.23 (d, *J* = 7.6 Hz, 1H), 7.90 (d, *J* = 9.6 Hz, 1H), 7.85 – 7.79 (m, 2H), 7.47 (d, *J* = 7.7 Hz, 1H), 7.32 – 7.15 (m, 16H), 6.71 (d, *J* = 8.4 Hz, 1H), 6.35 (d, *J* = 14.9 Hz, 2H), 6.03 (d, *J* = 2.1 Hz, 1H), 5.43 (m, 1H), 4.93 (m, 1H), 4.41-4.37 (m, 1H), 3.65 (d, *J* = 13.8 Hz, 1H), 3.54 (d, *J* = 13.8 Hz, 1H), 2.75 -2.62 (m, 1H), 2.48 – 2.39 (m, 2H), 2.16 (t, *J* = 7.2 Hz, 2H), 1.59 (s, 12H), 1.45 – 1.37 (m, 2H), 1.36 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 171.7, 165.8, 155.8, 153.0, 149.1, 144.8, 144.8, 139.9, 139.6, 132.9, 131.8, 129.7, 128.7, 128.2, 128.1, 128.0, 128.0, 127.9, 126.9, 120.2, 114.1, 90.9, 90.8, 85.6, 83.7, 83.6, 83.4, 80.9, 70.4, 58.6, 58.6, 55.7, 53.8, 53.6, 37.1, 34.9, 28.2, 27.1, 27.1, 26.3, 25.3, 25.3, 23.1, 22.7. HRMS (ESI): calculated for C₄₉H₅₅N₇O₆ [M+H]⁺ 838.4292, found 838.4314.

tert-Butyl-3-((((((3a*R*,4*R*,6*R*,6a*R*)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(4-(*tert*-butoxy)-4-oxobutyl)amino)methyl)benzoate (47). Following the procedure described for compound 33, coupling *tert*-butyl 4-oxobutanoate 22 (29 mg, 0.18 mmol) and compound 31 (75 mg, 0.15 mmol) afforded compound 47 as a white powder (64 mg, 67% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.21 (s, 1H), 7.86 – 7.77 (m, 3H), 7.42 (d, *J* = 7.6 Hz, 1H), 7.26 (d, *J* = 7.7 Hz, 1H), 6.15 (s, 2H), 5.99 (d, *J* = 2.2 Hz, 1H), 5.36 (m, 1H), 4.88 (m, 1H), 4.32 (m, 1H), 3.65 (d, *J* = 13.8 Hz, 1H), 3.50 (d, *J* = 13.8 Hz, 1H), 2.78-2.73 (m, 1H), 2.64-2.59 (m,1H), 2.42 (m, 2H), 2.22 – 2.09 (m, 2H), 1.55 (s, 12H), 1.36 (s, 9H), 1.33 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 172.8, 165.7, 155.7, 153.0, 149.1, 139.8, 139.5, 132.8, 131.8, 129.6, 128.1, 128.0, 120.2, 114.2, 90.8, 85.5, 83.7, 83.3, 80.8, 80.0, 58.7, 55.6, 53.4, 33.0, 28.2, 28.0, 27.1, 25.3, 22.4. HRMS (ESI): calculated for $C_{33}H_{46}N_6O_7$ [M+H]⁺ 639.3506, found 639.3506.

tert-Butyl-3-(((((3aR,4R,6R,6aR)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(5-(*tert*-butoxy)-5-oxopentyl)amino)methyl)benzoate (48). Following the procedure described for compound 33, coupling *tert*-butyl 5-oxopentanoate 23 (31 mg, 0.18 mmol) and compound 31 (75 mg, 0.15 mmol) afforded compound 48 as a white powder (72 mg, 73% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.20 (s, 1H), 7.84 – 7.77 (m, 3H), 7.42 (d, *J* = 7.6 Hz, 1H), 7.24 (t, *J* = 7.6 Hz, 1H), 6.19 (s, 2H), 5.99 (d, *J* = 2.2 Hz, 1H), 5.37 (m, 1H), 4.88 (m, 1H), 4.35 – 4.30 (m, 1H), 3.65 -3.48 1H), 2.71-2.59 (m, 1H), 2.46 – 2.38 (m, 2H), 2.10 (t, *J* = 7.1 Hz, 2H), 1.55 (s, 12H), 1.44 (m, 2H), 1.37 (s, 9H), 1.33 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 172.9, 165.7, 155.7, 153.0, 149.1, 139.7, 139.5, 132.9, 131.8, 129.6, 128.0, 128.0, 120.2, 114.1, 90.8, 85.5, 83.6, 83.3, 80.8, 79.9, 58.7, 55.6, 53.9, 35.2, 28.2, 28.1, 27.1, 26.2, 25.3, 22.7. HRMS (ESI): calculated for C₃₄H₄₈N₆O₇ [M+H]⁺ 653.3663, found 653.3669.

tert-Butyl-3-((((((3a*R*,4*R*,6*R*,6a*R*)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)((*S*)-4-(*tert*-butoxy)-3-((*tert*-butoxycarbonyl)amino)-4 oxobutyl)amino)methyl)benzoate (49). Following the procedure described for compound 33,

coupling *tert*-butyl (*R*)-2-((*tert*-butoxycarbonyl)amino)-4-oxobutanoate **27** (49 mg, 0.18 mmol) and compound **31** (75 mg, 0.15 mmol) afforded compound **49** as a white powder (85 mg, 75% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.18 (s, 1H), 7.79 (d, *J* = 6.7 Hz, 3H), 7.44 (s, 1H), 7.28 – 7.23 (m, 1H), 6.20 (s, 2H), 5.99 (s, 1H), 5.50 – 5.43 (m, 1H), 5.34 (d, *J* = 5.6 Hz, 1H), 4.86 (m, 1H), 4.31 (m, 1H), 4.15 – 4.07 (m, 1H), 3.67 (br, 1H), 3.47 (br, 1H), 2.76 (br, 2H), 2.59 (m, 2H), 2.44 (m, 2H), 1.93 (m, 1H), 1.73 (m, 1H), 1.54 (s, 12H), 1.35 (m, 21H). ¹³C NMR (101 MHz,

 CDCl₃) δ 171.6, 165.6, 155.7, 155.7, 155.3, 153.0, 149.1, 139.8, 139.0, 132.9, 131.8, 129.6, 128.2, 128.2, 120.2, 114.3, 90.6, 85.3, 83.7, 83.3, 81.6, 80.9, 79.3, 58.8, 55.7, 52.7, 50.5, 29.4, 28.3, 28.2, 27.9, 27.1, 25.4. HRMS (ESI): calculated for C₃₈H₅₅N₇O₉ [M+H]⁺ 754.4140, found 754.4129.

tert-Butyl 3-(((((3a*R*,4*R*,6*R*,6a*R*)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro[3,4d][1,3]dioxol-4-yl)methyl)(isopropyl)amino)methyl)benzoate (50). Following the procedure described for compound 33, coupling 5 mL dry acetone (large excess) and compound 31 (75 mg, 0.15 mmol) afforded compound 50 as a white powder (85 mg, 79% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.26 (s, 1H), 7.90 (s, 1H), 7.80 (m, 2H), 7.51 (d, *J* = 7.6 Hz, 1H), 7.28 (t, *J* = 7.7 Hz, 1H), 5.97 (d, *J* = 2.4 Hz, 1H), 5.92 (s, 2H), 5.37 (m, 1H), 4.86 (m, 1H), 4.26 – 4.20 (m, 1H), 3.64 (br, 1H), 3.54 (br, 1H), 2.87 (m, 1H), 2.73-2.56 (br, 2H), 1.57 (s, 9H), 1.53 (s, 3H), 1.33 (s, 3H), 1.03 (d, *J* = 6.6 Hz, 3H), 0.89 (d, *J* = 6.5 Hz, 3H). ¹³C NMR (101 MHz, CDCl3-*d*) δ 165.8, 155.5, 153.0, 149.2, 140.8, 139.9, 132.7, 131.8, 129.4, 128.0, 127.9, 120.2, 114.0, 91.0, 86.1, 83.4, 83.2, 80.8, 54.5, 51.3, 50.4, 28.2, 27.1, 25.3, 19.0, 16.7. HRMS (ESI): calculated for C₂₈H₃₈N₆O₅ [M+H]⁺ 539.2982, found 539.2982.

4-(((((3aR,4R,6R,6aR)-6-(6-amino-9H-purin-9-yl)-2,2-dimethyltetrahydrofuro[3,4-

d][1,3]dioxol-4-yl)methyl)(naphthalen-2-ylmethyl)amino)-*N*-tritylbutanamide (51). Following the procedure described for compound **33**, coupling 4-oxo-*N*-tritylbutanamide **16** (62 mg, 0.18 mmol) and compound **32** (67 mg, 0.15 mmol) afforded compound **51** as a white powder (85 mg, 73% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.10 (s, 1H), 7.79 – 7.73 (m, 2H), 7.72 – 7.66 (m, 2H), 7.61 (s, 1H), 7.46 – 7.38 (m, 3H), 7.28 – 7.12 (m, 15H), 6.59 (s, 1H), 5.97 (d, *J* = 2.2 Hz, 1H), 5.81 (s, 2H), 5.30 (m, 1H), 4.83 (m, 1H), 4.38 (s, 1H), 3.77 (d, *J* = 13.7 Hz, 1H), 3.63 (d, *J* = 13.7 Hz, 1H), 2.78-2.64 (m, 2H), 2.51 (t, *J* = 6.9 Hz, 2H), 2.30 – 2.20 (m, 2H), 1.79 (m, 2H), 1.52 (s, 3H), 1.31 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 171.6, 155.5, 153.0, 144.8, 139.7, 136.6, 133.2, 132.7, 128.7, 127.9, 127.8, 127.6, 127.6, 127.4, 127.2, 126.9, 126.0, 125.6, 114.2, 90.8, 85.4, 83.7, 83.4, 70.4, 59.1, 55.8, 53.8, 35.0, 27.0, 25.3, 22.8. HRMS (ESI): calculated for C₄₇H₄₇N₇O₄ [M+H]⁺774.3768, found 774.3769.

5-(((((3aR,4R,6R,6aR)-6-(6-amino-9H-purin-9-yl)-2,2-dimethyltetrahydrofuro[3,4-

d][1,3]dioxol-4-yl)methyl)(naphthalen-2-ylmethyl)amino)-*N*-tritylpentanamide (52).

Following the procedure described for compound **33**, coupling 5-oxo-*N*-tritylpentanamide **17** (64 mg, 0.18 mmol) and compound **32** (67 mg, 0.15 mmol) afforded compound **52** as a white powder (85 mg, 64% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.12 (s, 1H), 7.84 – 7.55 (m, 5H), 7.42 (m, 3H), 7.28-7.16 (m, 14H), 6.57 (s, 1H), 5.98 (s, 1H), 5.76 (s, 2H), 5.35 (d, *J* = 6.3 Hz, 1H), 4.88 (d, *J* = 6.2 Hz, 1H), 4.39 (s, 1H), 3.75 (d, *J* = 13.6 Hz, 1H), 3.62 (d, *J* = 13.6 Hz, 1H), 2.77 – 2.62 (m, 2H), 2.52 – 2.37 (m, 2H), 2.16-2.09 (m, 2H), 1.56 (s, 3H), 1.45-1.38 (m, 2H), 1.32 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 171.7, 155.4, 152.9, 149.2, 144.8, 139.9, 136.8, 133.2, 132.7, 128.7, 128.4, 127.9, 127.8, 127.6, 127.4, 127.2, 127.0, 125.9, 125.5, 120.2, 114.1, 90.9, 85.7, 83.7, 83.5, 70.4, 59.1, 55.8, 53.8, 37.2, 27.1, 26.3, 25.3, 23.2. HRMS (ESI): calculated for C₄₈H₄₉N₇O₄ [M+H]⁺788.3924, found 788.3932.

tert-Butyl-4-(((((3a*R*,4*R*,6*R*,6a*R*)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(naphthalen-2-ylmethyl)amino)butanoate (53). Following the procedure described for compound 33, coupling *tert*-butyl 4-oxobutanoate 22 (29 mg, 0.18 mmol) and compound 32 (67 mg, 0.15 mmol) afforded compound 53 as a white powder (67 mg, 76% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.12 (s, 1H), 7.82 (s, 1H), 7.78 – 7.60 (m, 4H), 7.48 – 7.35 (m, 3H), 6.74 (s, 2H), 6.02 (s, 1H), 5.34 (m, 1H), 4.89 (m, 1H), 4.38 (m, 1H), 3.79 (d, *J* = 13.6 Hz, 1H), 3.60 (d, *J* = 13.6 Hz, 1H), 2.80 (m, 1H), 2.65 (m, 1H), 2.57-2.44 (m, 2H), 2.29-2.13 (m,

2H), 1.75 (p, J = 7.3 Hz, 2H), 1.57 (s, 3H), 1.40 – 1.08 (m, 12H). ¹³C NMR (101 MHz, CDCl₃) δ 172.9, 156.0, 152.9, 152.9, 149.0, 139.6, 136.7, 133.2, 132.7, 127.8, 127.6, 127.3, 127.2, 125.9, 125.5, 120.1, 114.1, 90.8, 85.5, 83.7, 83.5, 80.0, 59.2, 55.7, 53.6, 33.0, 28.0, 27.5, 27.1, 25.4, 22.4. HRMS (ESI): calculated for C₃₂H₄₀N₆O₅ [M+H]⁺ 589.3138, found 589.3143.

tert-Butyl-5-((((3aR, 4R, 6R, 6aR)-6-(6-amino-9*H*-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(naphthalen-2-ylmethyl)amino)pentanoate (54). Following the procedure described for compound 33, coupling *tert*-butyl 5-oxopentanoate 23 (31 mg, 0.18 mmol) and compound 32 (67 mg, 0.15 mmol) afforded compound 54 as a white powder (62 mg, 69% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.11 (s, 1H), 7.80 (d, J = 24.4 Hz, 2H), 7.73 (d, J = 8.4 Hz, 2H), 7.64 (s, 1H), 7.47 – 7.38 (m, 3H), 6.42 (s, 2H), 6.02 (s, 1H), 5.34 (d, J = 6.3 Hz, 1H), 4.90 (m, 1H), 4.42 – 4.36 (m, 1H), 3.80 (d, J = 13.6 Hz, 1H), 3.62 (d, J = 13.6 Hz, 1H), 2.77 (m, 1H), 2.70 – 2.62 (m, 1H), 2.54 (s, 2H), 2.15 (t, J = 6.7 Hz, 2H), 1.58 (s, 3H), 1.48 (d, J = 9.8 Hz, 2H), 1.41 (s, 9H), 1.35 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 172.9, 155.8, 152.9, 149.1, 139.7, 136.8, 133.2, 132.7, 127.7, 127.6, 127.5, 127.3, 127.2, 125.9, 125.5, 120.2, 114.1, 90.9, 85.5, 83.7, 83.6, 83.4, 79.9, 59.1, 55.7, 54.1, 35.3, 28.1, 28.0, 27.1, 26.3, 25.3, 22.8. HRMS (ESI): calculated for C₃₃H₄₂N₆O₅ [M+H]⁺603.3295, found 603.3311.

tert-Butyl-(R)-4-(((((3aR,4R,6R,6aR)-6-(6-amino-9H-purin-9-yl)-2,2-

dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(naphthalen-2-ylmethyl)amino)-2-((*tert*-butoxycarbonyl)amino)butanoate (55). Following the procedure described for compound 33, coupling *tert*-butyl (*R*)-2-((*tert*-butoxycarbonyl)amino)-4-oxobutanoate 27 (49 mg, 0.18 mmol) and compound 32 (67 mg, 0.15 mmol) afforded compound 55 as a white powder (72 mg, 68% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.08 (s, 1H), 7.87 – 7.67 (m, 4H), 7.61 (s, 1H), 7.54-7.39 (m, 3H), 6.27 (d, *J* = 11.3 Hz, 2H), 6.00 (s, 1H), 5.71 – 5.61 (m, 1H), 5.30 (d, *J* = 5.1 Hz, 1H), 4.84 (m, 1H), 4.39 – 4.34 (m, 1H), 4.23 – 4.14 (m, 1H), 3.84 (d, J = 13.5 Hz, 1H), 3.59 (d, J = 13.5 Hz, 1H), 2.82 (br, 2H), 2.65 (br, 2H), 2.57-2.51 (m, 1H), 2.06 – 1.94 (m, 1H), 1.86 – 1.78 (m, 1H), 1.57 (s, 3H), 1.50-1.18 (m, 21H). ¹³C NMR (101 MHz, CDCl₃) δ 171.7, 155.4, 152.9, 149.0, 139.7, 136.2, 133.2, 132.7, 127.9, 127.6, 127.6, 127.5, 127.2, 127.1, 125.9, 125.6, 120.1, 114.3, 90.7, 85.3, 83.7, 83.4, 81.6, 79.3, 59.2, 55.7, 52.9, 50.8, 29.4, 28.3, 27.9, 27.9, 27.1, 25.4. HRMS (ESI): calculated for C₃₃H₄₂N₆O₅ [M+H]⁺704.3772, found 704.3777.

9-((3aR,4R,6R,6aR)-6-((isopropyl(naphthalen-2-ylmethyl)amino)methyl)-2,2-

dimethyltetrahydrofuro[3,4-d][1,3]dioxol-4-yl)-9*H*-purin-6-amine (56). Following the procedure described for compound 33, coupling 5 mL dry acetone (large excess) and compound 32 (67 mg, 0.15 mmol) afforded compound 56 as a white powder (35 mg, 48% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.21 (s, 1H), 7.83 – 7.68 (m, 5H), 7.53 (m, 1H), 7.47 – 7.39 (m, 2H), 6.26 (s, 2H), 5.98 (d, *J* = 2.3 Hz, 1H), 5.32 (m, 1H), 4.85 (m, 1H), 4.32-4.27 (m, 1H), 3.79 (d, *J* = 13.9 Hz, 1H), 3.64 (d, *J* = 14.0 Hz, 1H), 3.00-2.93 (m, 1H), 2.78 (m 1H), 2.64 (m, 1H), 1.53 (s, 3H), 1.31 (s, 3H), 1.09 (d, *J* = 6.6 Hz, 3H), 0.95 (d, *J* = 6.6 Hz, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 155.7, 152.9, 149.1, 139.7, 138.0, 133.2, 132.7, 127.7, 127.6, 127.5, 127.2, 127.0, 125.9, 125.4, 120.2, 113.9, 91.1, 86.2, 83.5, 83.3, 77.3, 54.9, 51.4, 50.4, 27.0, 25.3, 19.2, 16.6. HRMS (ESI): calculated for C₂₇H₃₂N₆O₃ [M+H]⁺489.2614, found 489.2611.

3-(((4-Amino-4-oxobutyl)((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-

dihydroxytetrahydrofuran-2-yl)methyl)amino)methyl)benzamide (57). To a solution of compound 33 (100 mg, 0.098 mmol) in 5 mL CH₂Cl₂ was added 5 mL TFA and the mixture was stirred at room temperature. After 2 hours, 2 mL H₂O was added and the mixture was stirred for 1 hour at room temperature. The mixture was concentrated and the crude product was purified by preparative HPLC affording compound 57 as a white powder. ¹H NMR (400 MHz, D₂O) δ 8.46

- 8.06 (m, 2H), 7.87 - 7.26 (m, 4H), 6.08 (br, 1H), 4.75 - 4.36 (m, 4H), 4.27 (br, 1H), 3.84 - 3.27 (m, 4H), 2.38 (br, 2H), 2.10 (br, 2H). ¹³C NMR (101 MHz, D₂O) δ 177.5, 162.8, 162.5, 149.6, 143.8, 134.8, 134.1, 132.7, 129.6, 129.1, 128.3, 118.9, 117.6, 114.7, 90.4, 77.7, 73.6, 71.5, 57.9, 54.8, 31.8, 19.0. HRMS (ESI): calculated for C₂₂H₂₈N₈O₅ [M+H]⁺ 485.2261, found 485.2265.

3-(((5-Amino-5-oxopentyl)((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-dihydroxy-

tetrahydrofuran-2-yl)methyl)amino)methyl)benzamide (58). Following the procedure described for compound 57, compound 34 (50 mg, 0.049 mmol) was deprotected to obtain compound 58 as a white powder (16 mg, 56% yield). ¹H NMR (400 MHz, D₂O) δ 8.43 – 8.12 (m, 2H), 7.84 – 7.26 (m, 4H), 6.08 (br, 1H), 4.65 – 4.21 (m, 5H), 3.63-3.48 (m, 2H), 3.34 (br, 2H), 2.35 (br, 2H), 1.84 (br, 2H), 1.58 (br, 2H). ¹³C NMR (101 MHz, D₂O) δ 177.6, 170.6, 162.8, 149.6, 147.3, 143.8, 134.8, 129.7, 129.6, 129.4, 129.1, 128.3, 117.6, 114.7, 90.5, 77.8, 77.4, 71.6, 71.4, 57.8, 54.6, 32.6, 22.4, 21.0. HRMS (ESI): calculated for C₂₃H₃₀N₈O₅ [M+H]⁺ 499.2417, found 499.2420.

4-((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-dihydroxytetrahydrofuran-2-

yl)methyl)(3-carbamoylbenzyl)amino)butanoic acid (59). Following the procedure described for compound 57, compound 35 (50 mg, 0.060 mmol) was deprotected to obtain compound 59 as a white powder (21 mg, 60% yield). ¹H NMR (400 MHz, D₂O) δ 8.38 – 8.06 (m, 2H), 7.71 – 7.26 (m, 4H), 6.05 (br, 1H), 4.64 – 4.21 (m, 5H), 3.53 (br, 2H), 3.35 (s, 2H), 2.41 (br, 2H), 2.02 (br, 2H). ¹³C NMR (101 MHz, D₂O) δ 176.4, 170.5, 149.6, 147.3, 143.8, 134.8, 132.7, 129.6, 129.5, 128.3, 117.5, 114.6, 90.4, 77.7, 73.5, 71.4, 57.8, 52.8, 38.6, 30.2, 18.4. HRMS (ESI): calculated for C₂₂H₂₈N₇O₆ [M+H]⁺486.2101, found 486.2103.

5-((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-dihydroxytetrahydrofuran-2-

yl)methyl)(3-carbamoylbenzyl)amino)pentanoic acid (60). Following the procedure described for compound 57, compound 36 (50 mg, 0.059 mmol) was deprotected to obtain compound 60 as a white powder (17 mg, 50% yield). ¹H NMR (400 MHz, D₂O) δ 8.29 (br, 2H), 7.84 – 7.58 (m, 3H), 7.46 (br, 1H), 6.13 (br, 1H), 4.70 – 4.33 (m, 5H), 3.66 (br, 2H), 3.48 – 3.31 (m, 2H), 2.42 (br, 2H), 1.88 (s, 2H), 1.65 (br, 2H). ¹³C NMR (101 MHz, D₂O) δ 177.7, 170.8, 163.0, 150.5, 147.6, 145.1, 143.1, 134.5, 132.9, 129.7, 129.3, 128.4, 119.0, 117.7, 90.3, 77.7, 73.3, 32.8, 22.4, 21.0. HRMS (ESI): calculated for C₂₃H₃₀N₇O₆ [M+H]⁺ 500.2258, found 500.2267.

3-((((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-dihydroxytetrahydrofuran-2-

yl)methyl)(isopropyl)amino)methyl)benzamide (61). Following the procedure described for compound 57, compound 38 (50 mg, 0.069 mmol) was deprotected to obtain compound 61 as a white powder (22 mg, 60% yield). ¹H NMR (400 MHz, Acetone- d_6) δ 8.48-8.39 (m, 2H), 8.26 (br, 1H), 7.94 (d, J = 7.8 Hz, 1H), 7.9-7.73 (m, 2H), 7.46 (m, 1H), 6.81 (br, 1H), 6.13 (d, J = 3.4 Hz, 1H), 4.74 (br, 2H), 4.65 (s, 1H), 4.53 (br, 1H), 4.46 (br, 1H), 3.93 – 3.69 (m, 3H), 3.31 (s, 1H), 1.49-1.45 (m, 6H). ¹³C NMR (101 MHz, Acetone- d_6) δ 152.7, 148.4, 146.1, 142.5, 135.0, 134.0, 130.5, 130.2, 119.9, 90.6, 79.3, 73.5, 72.5, 55.6, 54.3, 54.0, 51.4, 16.6, 15.0. HRMS (ESI): calculated for C₂₁H₂₈N₇O₄ [M+H]⁺ 442.2203, found 442.2203.

Methyl 3-(((4-amino-4-oxobutyl)(((2R,3S,4R,5R)-5-(6-amino-9*H*-purin-9-yl)-3,4dihydroxytetrahydrofuran-2-yl)methyl)amino)methyl)benzoate (62). Following the procedure described for compound 57, compound 39 (50 mg, 0.064 mmol) was deprotected to obtain compound 62 as a white powder (20 mg, 53% yield). ¹H NMR (400 MHz, D₂O) δ 8.38 – 7.98 (m, 2H), 7.88 – 7.50 (m, 3H), 7.35 (br, 1H), 6.05 (br, 1H), 4.64 – 4.32 (m, 4H), 4.20 (br, 1H), 3.78 (s, 3H), 3.55 (br, 1H), 3.47 – 3.30 (m, 2H), 2.39 (br, 2H), 2.08 (br, 2H). ¹³C NMR (101

MHz, D₂O) δ 177.5, 167.3, 149.5, 147.2, 143.7, 143.6, 135.8, 134.6, 131.3, 130.7, 129.9, 129.8, 129.4, 129.1, 118.8, 90.6, 77.8, 77.4, 73.8, 73.1, 71.7, 71.4, 57.6, 56.9, 55.2, 54.8, 53.6, 52.7, 31.8, 19.0. HRMS (ESI): calculated for C₂₃H₂₉N₇O₆ [M+H]⁺ 500.2258, found 500.2265.

Methyl 3-(((5-amino-5-oxopentyl)((((2R,3S,4R,5R))-5-(6-amino-9*H*-purin-9-yl)-3,4dihydroxytetrahydrofuran-2-yl)methyl)amino)methyl)benzoate (63). Following the procedure described for compound 57, compound 40 (50 mg, 0.063 mmol) was deprotected to obtain compound 63 as a white powder (21 mg, 55% yield).¹H NMR (400 MHz, D₂O) δ 8.44 – 8.07 (m, 2H), 7.96 – 7.34 (m, 4H), 6.12 (br, 1H), 4.50 (br, 4H), 4.32 (s, 1H), 3.86 (s, 3H), 3.62 (br, 1H), 3.52 – 3.34 (m, 2H), 2.32 (br, 2H), 1.89 (br, 2H), 1.68 (br, 2H). ¹³C NMR (101 MHz, D₂O) δ 178.7, 167.5, 162.6, 149.6, 143.8, 143.6, 135.9, 130.9, 130.0, 129.9, 129.5, 129.1, 117.6, 114.7, 111.8, 71.4, 52.7, 33.9, 22.4, 21.9. HRMS (ESI): calculated for C₂₄H₃₁N₇O₆ [M+H]⁺ 514.2414, found 514.2415.

4-((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-dihydroxytetrahydrofuran-2-

yl)methyl)(3-(methoxycarbonyl)benzyl)amino)butanoic acid (64). Following the procedure described for compound 57, compound 41 (50 mg, 0.084 mmol) was deprotected to obtain compound 64 as a white powder (24 mg, 49% yield). ¹H NMR (400 MHz, D₂O) δ 8.29 (s, 1H), 8.11 (s, 1H), 7.94 – 7.64 (m, 4H), 7.45 (t, *J* = 7.9 Hz, 1H), 6.09 (s, 1H), 4.62 (br, 4H), 4.49 (br, 1H), 3.88 (s, 3H), 3.67 (br, 2H), 3.55-3.41 (m, 6.9 Hz, 3H), 2.53 (t, *J* = 6.4 Hz, 2H), 2.18 – 2.09 (m, 2H). ¹³C NMR (101 MHz, D₂O) δ 176.8, 167.6, 150.3, 147.4, 144.8, 143.3, 130.1, 129.8, 129.5, 129.2, 90.5, 77.7, 71.6, 57.7, 52.8, 30.6, 18.5. HRMS (ESI): calculated for C₂₃H₂₈N₆O₇ [M+H]⁺ 501.2098, found 501.2097.

5-((((2*R*,3*S*,4*R*,5*R*)-5-(6-amino-9*H*-purin-9-yl)-3,4-dihydroxytetrahydrofuran-2vl)methyl)(3-(methoxycarbonyl)benzyl)amino)pentanoic acid (65). Following the procedure

described for compound **57**, compound **42** (50 mg, 0.082 mmol) was deprotected to obtain compound **65** as a white powder (30 mg, 59% yield).¹H NMR (400 MHz, D₂O) δ 8.22 (br, 2H), 7.86 (br, 2H), 7.67 (br, 1H), 7.46 (br, 1H), 6.08 (br, 1H), 4.55 (br, 4H), 4.35 (br, 1H), 3.88 (s, 3H), 3.64 (br, 1H), 3.43 (br, 2H), 2.45 (br, 2H), 1.90 (br, 2H), 1.67 (br, 2H). ¹³C NMR (101 MHz, D₂O) δ 178.1, 168.0, 150.3, 144.5, 143.8, 130.5, 130.2, 129.9, 129.6, 118.0, 115.1, 72.0, 53.1, 33.1, 23.6, 22.8, 21.4. HRMS (ESI): calculated for C₂₄H₃₀N₆O₇ [M+H]⁺ 515.2254, found 515.2257.

(S)-2-Amino-4-((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-

dihydroxytetrahydrofuran-2-yl)methyl)(3-(methoxycarbonyl)benzyl)amino)butanoic acid (66). Following the procedure described for compound 57, compound 43 (50 mg, 0.070 mmol) was deprotected to obtain compound 66 as a white powder (26 mg, 60% yield).¹H NMR (400 MHz, D₂O) δ 8.24 (s, 1H), 8.01 (s, 1H), 7.72 (br, 1H), 7.64 (d, *J* = 7.8 Hz, 1H), 7.54 (d, *J* = 7.8 Hz, 1H), 7.31 (t, *J* = 7.8 Hz, 1H), 6.03 (s, 1H), 4.56 – 4.52 (m, 1H), 4.50 – 4.35 (m, 4H), 4.07 (m, 1H), 3.74 (s, 3H), 3.69 – 3.54 (m, 4H), 2.55-2.45 (m, 1H), 2.41-2.33 (m, 1H). ¹³C NMR (101 MHz, D₂O) δ 170.8, 167.2, 162.7, 162.3, 149.5, 147.2, 143.6, 143.6, 131.0, 130.0, 129.5, 129.3, 129.1, 118.8, 117.6, 114.7, 90.8, 77.5, 73.5, 71.4, 53.9, 50.7, 24.4. HRMS (ESI): calculated for C₂₃H₂₉N₇O₇ [M+H]⁺ 516.2207, found 516.2206.

Methyl 3-((((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-dihydroxytetrahydrofuran-2-yl)methyl)(*iso* $propyl)amino)methyl)benzoate (67). Following the procedure described for compound 57, compound 44 (50 mg, 0.101 mmol) was deprotected to obtain compound 67 as a white powder (33 mg, 59% yield).¹H NMR (400 MHz, D₂O) <math>\delta$ 8.19 (d, J = 3.9 Hz, 1H), 7.99 – 7.54 (m, 4H), 7.29 (t, J = 7.6 Hz, 1H), 5.91 (s, 1H), 4.62 – 4.46 (m, 3H), 4.33 – 4.25 (m, 1H), 4.20 (br, 1H), 3.87 (s, 3H), 3.79-3.62 (m, 2H), 3.39 – 3.28 (m, 1H), 1.54 – 1.37 (m, 6H). ¹³C

NMR (101 MHz, D₂O) δ 167.6, 149.7, 147.3, 144.1, 143.3, 136.1, 131.3, 130.1, 129.7, 128.7, 118.7, 114.8, 90.6, 78.6, 73.6, 71.4, 58.7, 55.2, 52.7, 50.5, 16.2, 15.7. HRMS (ESI): calculated for C₂₂H₂₈N₆O₅ [M+H]⁺ 457.2199, found 457.2196.

3-(((4-Amino-4-oxobutyl)((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-

dihydroxytetrahydrofuran-2-yl)methyl)amino)methyl)benzoic acid (68). Following the procedure described for compound 57, compound 45 (50 mg, 0.061 mmol) was deprotected to obtain compound 68 as a white powder (15 mg, 42% yield). ¹H NMR (400 MHz, D₂O) δ 8.21 (br, 2H), 7.86 (br, 2H), 7.65 (d, *J* = 7.6 Hz, 1H), 7.43 (br, 1H), 6.08 (br, 1H), 4.70 – 4.24 (m, 5H), 3.63 (br, 1H), 3.52 – 3.38 (m, 2H), 2.48 (br, 2H), 2.12 (br, 2H). ¹³C NMR (101 MHz, D₂O) δ 177.5, 149.8, 147.4, 144.0, 143.6, 131.2, 130.3, 129.9, 129.1, 117.7, 114.8, 90.5, 77.7, 71.5, 31.8, 19.1. HRMS (ESI): calculated for C₂₂H₂₆N₇O₆ [M+H]⁺ 486.2101, found 486.2089.

3-(((5-Amino-5-oxopentyl)(((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-

dihydroxytetrahydrofuran-2-yl)methyl)amino)methyl)benzoic acid (69). Following the procedure described for compound 57, compound 46 (50 mg, 0.059 mmol) was deprotected to obtain compound 69 as a white powder (19 mg, 65 % yield). ¹H NMR (400 MHz, D₂O) δ 8.19 (br, 2H), 7.83(br, 2H), 7.63 (br, 1H), 7.40 (br, 1H), 6.03 (br, 1H), 4.58-4.41 (m, 4H), 4.31 (br, 1H), 3.63 (br, 1H), 3.42 (d, *J* = 7.8 Hz, 2H), 2.34 (br, 2H), 1.90 (br, 2H), 1.68 (br, 2H). ¹³C NMR (101 MHz, D₂O) δ 178.7, 149.7, 143.9, 143.5, 130.3, 129.8, 129.8, 129.1, 120.6, 117.7, 114.8, 105.0, 77.9, 71.5, 33.9, 21.9. HRMS (ESI): calculated for C₂₃H₃₀N₇O₆ [M+H]⁺ 500.2258, found 500.2253.

3-((((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-dihydroxytetrahydrofuran-2-

yl)methyl)(3-carboxypropyl)amino)methyl)benzoic acid (70). Following the procedure described for compound 57, compound 47 (50 mg, 0.078 mmol) was deprotected to obtain

compound **70** as a white powder (21 mg, 46% yield). ¹H NMR (400 MHz, D₂O) δ 8.27 (s, 1H), 8.14 (s, 1H), 7.85 (br, 2H), 7.64 (d, *J* = 7.7 Hz, 1H), 7.49 – 7.35 (m, 1H), 6.08 (br, 1H), 4.55 (br, 5H), 3.65 (br, 1H), 3.48-3,43 (m, 2H), 2.52 (br, 2H), 2.13 (br, 2H). ¹³C NMR (101 MHz, D₂O) δ 176.6, 168.8, 149.8, 147.4, 144.0, 143.5, 131.4, 130.4, 129.7, 129.1, 117.7, 114.8, 90.5, 77.7, 71.6, 30.4, 22.1, 18.5. HRMS (ESI): calculated for C₂₂H₂₇N₆O₇ [M+H]⁺ 487.1941, found 487.1945.

3-(((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-dihydroxytetrahydrofuran-2-

yl)methyl)(4-carboxybutyl)amino)methyl)benzoic acid (71). Following the procedure described for compound 57, compound 48 (50 mg, 0.076 mmol) was deprotected to obtain compound 71 as a white powder (24 mg, 52% yield). ¹H NMR (400 MHz, D₂O) δ 8.20 (br, 2H), 7.83 (br, 2H), 7.63 (d, *J* = 7.5 Hz, 1H), 7.42 (br, 1H), 6.05 (br, 1H), 4.53 (br, 4H), 4.31 (br, 1H), 3.63 (br, 1H), 3.49 – 3.32 (m, 2H), 2.42 (br, 2H), 1.88 (br, 2H), 1.66 (br, 2H). ¹³C NMR (101 MHz, D₂O) δ 177.7, 150.0, 144.3, 143.4, 131.4, 130.4, 130.1, 129.7, 129.1, 117.7, 114.8, 90.4, 77.7, 71.6, 32.7, 22.4, 21.0. HRMS (ESI): calculated for C₂₃H₂₉N₆O₇ [M+H]⁺ 501.2098, found 501.2096.

3-((((S)-3-Amino-3-carboxypropyl)(((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-

dihydroxytetrahydrofuran-2-yl)methyl)amino)methyl)benzoic acid (72). Following the procedure described for compound 57, compound 49 (50 mg, 0.066 mmol) was deprotected to obtain compound 72 as a white powder (24 mg, 61% yield). ¹H NMR (400 MHz, D₂O) δ 8.23 (bs, 1H), 8.06 (s, 1H), 7.76 (s, 1H), 7.69 (d, *J* = 7.6 Hz, 1H), 7.56 (d, *J* = 7.8 Hz, 1H), 7.33 (t, *J* = 7.8 Hz, 1H), 6.06 – 5.98 (m, 1H), 4.59 – 4.34 (m, 5H), 4.06 (m, 1H), 3.72 – 3.51 (m, 4H), 2.55 – 2.4m (m, 1H), 2.40-2.32 (m, 1H). ¹³C NMR (101 MHz, D₂O) δ 171.0, 168.4, 162.8, 162.4, 149.5, 147.3, 143.7, 143.5, 131.3, 130.3, 129.7, 129.5, 129.1, 118.9, 117.6, 114.7, 90.6, 77.5, 149.5, 147.3, 143.7, 143.5, 131.3, 130.3, 129.7, 129.5, 129.1, 118.9, 117.6, 114.7, 90.6, 77.5, 129.1, 118.9, 117.6, 114.7, 90.6, 77.5, 129.1, 118.9, 117.6, 114.7, 90.6, 77.5, 129.1, 118.9, 117.6, 114.7, 90.6, 77.5, 129.5, 129.1, 118.9, 117.6, 114.7, 90.6, 77.5, 114.7, 90.6, 77.5, 114.7, 90.6, 77.5, 114.7, 90.6, 77.5, 114.7, 90.6, 77.5, 114.7, 90.6, 77.5, 114.7, 90.6, 77.5, 114.7, 90.6

73.5, 71.5, 50.9, 24.4. HRMS (ESI): calculated for $C_{22}H_{27}N_7O_7 [M+H]^+$ 502.2050, found 502.2048.

3-((((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-dihydroxytetrahydrofuran-2-

yl)methyl)(isopropyl)amino)methyl)benzoic acid (73). Following the procedure described for compound 57, compound 50 (50 mg, 0.093 mmol) was deprotected to obtain compound 73 as a white powder (27 mg, 54% yield). ¹H NMR (400 MHz, D₂O) δ 8.21 (s, 1H), 8.16 (s, 1H), 7.90 (s, 1H), 7.58 (m, 2H), 7.27 (t, *J* = 7.3 Hz, 1H), 5.89 (s, 1H), 4.55 (t, *J* = 10.4 Hz, 2H), 4.31 – 4.15 (m, 2H), 3.98 – 3.90 (m, 1H), 3.78 (br, 1H), 3.32 (br, 1H), 1.50 (m, 6H). ¹³C NMR (101 MHz, D₂O) δ 215.3, 168.7, 149.9, 147.3, 144.4, 143.1, 136.1, 131.7, 129.9, 128.7, 118.7, 117.7, 114.8, 89.7, 78.5, 73.5, 71.4, 58.8, 55.3, 51.7, 30.1, 16.2, 15.7. HRMS (ESI): calculated for C₂₁H₂₆N₆O₅ [M+H]⁺ 443.2043, found 443.2040.

4-(((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-dihydroxytetrahydrofuran-2-

yl)methyl)(naphthalen-2-ylmethyl)amino)butanamide (74). Following the procedure described for compound 57, compound 51 (50 mg, 0.064 mmol) was deprotected to obtain compound 74 as a white powder (22 mg, 58% yield). ¹H NMR (400 MHz, D₂O) δ 8.11 (br, 1H), 7.57 – 7.25 (m, 4H), 7.09 (d, *J* = 8.0 Hz, 1H), 5.89 (br, 1H), 4.35 (br, 4H), 3.94 – 3.63 (m, 1H), 3.49 – 3.22 (m, 3H), 2.44 (br, 2H), 2.13 – 1.82 (br, 2H). ¹³C NMR (101 MHz, D₂O) δ 177.5, 148.4, 146.0, 143.3, 142.3, 132.1, 131.6, 129.5, 128.0, 127.7, 127.2, 127.0, 126.8, 126.7, 125.1, 118.0, 90.6, 77.8, 74.1, 71.2, 58.0, 55.9, 54.9, 32.0, 19.0. HRMS (ESI): calculated for C₂₅H₂₉N₇O₄ [M+H]⁺ 492.2359, found 492.2363.

5-((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-dihydroxytetrahydrofuran-2-

yl)methyl)(naphthalen-2-ylmethyl)amino)pentanamide (75). Following the procedure described for compound 57, compound 52 (50 mg, 0.063 mmol) was deprotected to obtain

compound 75 as a white powder (23 mg, 60% yield). ¹H NMR (400 MHz, D₂O) δ 8.11 (s, 1H), 7.72 - 7.61 (m, 2H), 7.55 - 7.48 (m, 2H), 7.46 - 7.39 (m, 1H), 7.37 - 7.15 (m, 2H), 5.89 (br, 1H), 4.66 – 4.48 (m, 2H), 4.42 – 4.11 (m, 3H), 3.61 – 3.33 (m, 4H), 2.43 – 2.27 (m, 2H), 1.89 (d, J = 9.8 Hz, 2H), 1.77 – 1.61 (m, 2H). ¹³C NMR (101 MHz, D₂O) δ 178.8, 148.6, 146.3, 143.8, 143.4, 142.7, 142.2, 132.4, 131.8, 129.9, 128.2, 127.8, 127.4, 126.8, 125.1, 90.7, 78.1, 74.2, 71.3, 58.6. 57.8. 55.8. 54.6. 34.0. 22.5. 22.0. HRMS (ESI): calculated for $C_{26}H_{31}N_7O_4 [M+H]^+ 506.2516$, found 506.2520.

4-((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-dihydroxytetrahydrofuran-2-

yl)methyl)(naphthalen-2-ylmethyl)amino)butanoic acid (76). Following the procedure described for compound 57, compound 53 (50 mg, 0.085 mmol) was deprotected to obtain compound 76 as a white powder (30 mg, 60% yield). ¹H NMR (400 MHz, D₂O) δ 8.14 (br, 1H), 7.75 – 7.23 (m, 8H), 5.95 (br, 1H), 4.62 – 4.54 (m, 1H), 4.43 – 4.26 (m, 4H), 3.69 – 3.53 (m, 2H), 3.45 (m, 2H), 2.54 (t, *J* = 6.5 Hz, 2H), 2.19 – 2.04 (m, 2H). ¹³C NMR (101 MHz, D₂O) δ 179.8, 179.8, 152.0, 149.1, 146.3, 145.7, 134.9, 134.4, 132.5, 130.6, 129.9, 129.8, 129.3, 121.0, 120.2, 117.3, 93.3, 80.2, 74.2, 60.8, 33.6, 21.4. HRMS (ESI): calculated for C₂₃H₃₀N₈O₅ [M+H]⁺ 493.2199, found 493.2199.

5-((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-dihydroxytetrahydrofuran-2-

yl)methyl)(naphthalen-2-ylmethyl)amino)pentanoic acid (77). Following the procedure described for compound 57, compound 54 (50 mg, 0.083 mmol) was deprotected to obtain compound 77 as a white powder (31 mg, 61% yield). ¹H NMR (400 MHz, D₂O) δ 8.27 (br, 2H), 7.86 – 7.55 (m, 5H), 7.52 – 7.47 (m, 1H), 7.40 (d, *J* = 7.7 Hz, 1H), 5.95 (s, 1H), 4.66-4.40 (m, 4H), 4.31 (br, 1H), 3.70 – 3.43 (m, 4H), 2.50 (br, 2H), 1.97 (br, 2H), 1.80 – 1.67 (br, 2H). ¹³C NMR (101 MHz, D₂O) δ 178.0, 163.0, 162.7, 149.5, 146.6, 143.8, 143.1, 132.4, 131.9, 130.0,

128.0, 127.3, 127.2, 126.8, 117.7, 38.6, 32.9, 21.2. HRMS (ESI): calculated for $C_{26}H_{28}N_6O_5 [M+H]^+ 507.2356$, found 507.2355.

(S)-2-Amino-4-((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-

dihydroxytetrahydrofuran-2-yl)methyl)(naphthalen-2-ylmethyl)amino)butanoic acid (78). Following the procedure described for compound **57**, compound **55** (50 mg, 0.071 mmol) was deprotected to obtain compound **78** as a white powder (28 mg, 65% yield). ¹H NMR (400 MHz, D₂O) δ 8.03 (bs, 2H), 7.54 – 6.97 (m, 7H), 5.83 (bs, 1H), 4.55 – 4.25 (m, 3H), 4.24 – 4.16 (m, 1H), 4.16 – 4.09 (m, 1H), 3.65 (s, 2H), 3.49 (s, 1H), 2.46 (br, 2H). ¹³C NMR (101 MHz, D₂O) δ 170.8, 148.3, 146.0, 143.3, 142.2, 132.0, 131.5, 129.7, 127.1, 126.9, 126.6, 126.3, 118.2, 117.6, 114.7, 90.9, 77.5, 73.8, 71.4, 50.7, 24.5. HRMS (ESI): calculated for C₂₅H₂₉N₇O₅ [M+H]⁺ 508.2308, found 508.2309.

(2R,3R,4S,5R)-2-(6-amino-9H-purin-9-yl)-5-((isopropyl(naphthalen-2-ylmethyl))

amino)methyl) tetrahydrofuran-3,4-diol (79). Following the procedure described for compound **57**, compound **56** (50 mg, 0.102 mmol) was deprotected to obtain compound **79** as a white powder (36 mg, 65% yield).¹H NMR (400 MHz, D₂O) δ 7.86 (s, 1H), 7.64 – 7.47 (m, 2H), 7.42 – 7.10 (m, 6H), 5.58 (s, 1H), 4.41 – 4.35 (m, 1H), 4.29 (br, 1H), 4.16 – 4.04 (m, 2H), 3.92 (br, 1H), 3.80-3.73(m, 1H), 3.58 (br, 1H), 3.24 (m, 1H), 1.35 (m, 6H). ¹³C NMR (101 MHz, D₂O) δ 148.7, 146.3, 143.6, 143.0, 142.6, 132.2, 131.7, 130.4, 128.3, 127.5, 127.4, 118.0, 90.0, 77.5, 73.9, 71.9, 59.1, 51.7, 49.6, 17.4, 15.9. HRMS (ESI): calculated for C₂₄H₂₈N₆O₃ [M+H]⁺ 449.2301, found 449.2299.

tert-Butyl (S)-4-(((((3aR,4R,6R,6aR)-6-(6-amino-9H-purin-9-yl)-2,2-dimethyltetrahydrofuro-[3,4-d][1,3]dioxol-4-yl)methyl)(naphthalen-2-ylmethyl)amino)-2-bis(*tert*butoxycarbonyl)amino)pentanoate (80). Following the procedure described for compound 33, *tert*-butyl (*R*)-2-((*tert*-butoxycarbonyl)amino)-5-oxopentanoate **28** (312 mg, 0.80 mmol) and compound **32** (300 mg, 0.67mmol) to obtain compound **80** as a white powder (319 mg, 58% yield). ¹H NMR (400 MHz, CDCl₃) δ 8.05 (s, 1H), 7.79 (s, 1H), 7.73 m, 1H), 7.68 (d, *J* = 8.6 Hz, 2H), 7.61 (s, 1H), 7.45 – 7.34 (m, 3H), 6.59 (s, 2H), 5.98 (d, *J* = 2.2 Hz, 1H), 5.27 (dd, *J* = 6.4, 2.2 Hz, 1H), 4.84 (dd, *J* = 6.4, 3.1 Hz, 1H), 4.69 (dd, *J* = 9.6, 5.2 Hz, 1H), 4.37 (m, 1H), 3.80 (br, 1H), 3.57 (br, 1H), 2.76 (m, 1H), 2.64 – 2.47 (m, 3H), 2.04 (m, 1H), 1.83 (m, 1H), 1.54 (s, 3H), 1.43 (s, 16H), 1.41 (s, 8H), 1.31 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 155.9, 152.9, 152.5, 139.5, 127.7, 127.5, 127.3, 127.2, 125.8, 125.5, 90.8, 85.4, 83.6, 83.41, 82.6, 81.0, 59.0, 55.6, 54.0, 28.0, 27.1, 27.0, 25.3, 23.8. HRMS (ESI): calculated for C₄₃H₅₉N₇O₉ [M+H]⁺ 818.4453, found 818.4458.

(S)-2-Amino-5-((((2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-3,4-

dihydroxytetrahydrofuran-2-yl)methyl)(naphthalen-2-ylmethyl)amino)pentanoic acid (81). Following the procedure described for compound 57, compound 80 (120 mg, 0.15 mmol) was deprotected to obtain compound 81 as a white powder (58 mg, 63% yield). ¹H NMR (600 MHz, D₂O) δ 8.14 (br, 1H), 7.69–6.93 (m, 8H), 5.93 (br, 1H), 4.59–4.43 (m, 2H), 4.27 (br, 2H), 4.15– 3.73 (m, 2H), 3.47 (m, 4H), 2.16–1.90 (m, 4H). ¹³C NMR (151 MHz, D₂O) δ 171.6, 162.9, 162.7, 148.6, 131.8 127.3, 127.1, 126.8, 119.2, 117.3, 115.3, 90.7, 78.0, 74.3, 71.3, 58.8, 52.3, 26.9, 19.4. HRMS (ESI): calculated for C₂₆H₃₁N₇O₅ [M+H]⁺ 522.2465, found 522.2468.

Inhibition studies: Expression and purification of full-length wild type NNMT protein (NNMTwt) were performed as previously described.³² The purity of the enzyme was confirmed using SDS-PAGE with Coomassie blue staining and NNMT identity was confirmed using SDS-PAGE and Western blotting. Catalytic activity of the recombinant protein was evaluated with 1

unit of enzyme activity representing the formation of 1 nmol of MNA per hour of incubation at 37°C. The specific activity of the batch used in the inhibitory activity assays was 18,665 units per mg of protein at a protein concentration of 0.56 mg/mL. NNMT was used at a final concentration of 100 nM diluted in assay buffer (50 mM Tris buffer (pH 8.4) and 1 mM DTT). The compounds were dissolved in DMSO and diluted with water to concentrations ranging from 0.1 μ M to 500 μ M (DMSO was kept constant at 1.25% final concentration). The compounds were incubated with the enzyme for 10 minutes at 37°C before initiating the reaction with a mixture of NA and AdoMet at their K_M values of 200 µM and 8.5 µM respectively. The formation of MNA was measured after 30 minutes at 37 °C. The reaction was guenched by addition of 15 μ L sample to 70 μ L acetonitrile containing 50 nM deutero-methylated nicotinamide as internal standard. The enzymatic activity assays were performed using UHP-HILIC-MS/MS as previously described with minor modifications.²⁴ The UHP-HILIC-MS/MS system consisted of a binary UHPLC system, consisting of two LC-30AD pumps, a SIL30-ACmp auto-sampler, a CTO-20AC column oven, and a DGU-20A5R degasser, (all from Shimadzu, 's-Hertogenbosch, The Netherlands). Isocratic elution was performed after 1 µL injections on a Waters Acquity BEH Amide HILIC column (3.0 x 100 mm, 1.7 µm particle size, Waters, Milford, USA), using water containing 300 μ M formic acid and 550 μ M NH₄OH (pH 9.2) at 40% v/v and acetonitrile at 60% v/v, with a run-time of 3 min. Calibration samples were prepared using 75 μ L internal standard d₃-MNA at 50 nM in acetonitrile and 25 μ L of an aqueous solution of reference standard MNA with concentrations ranging from 2500 nM to 1.221 nM. For detection, a Sciex QTRAP® 5500 triple quadrupole mass spectrometer, with Analyst 1.6.2, and MultiQuant 3.0.1 software (Sciex, Ontario, Canada) was used. Settings used for the ionization source were: curtain gas, 40 psi; collision gas, 'medium'; ionspray voltage,

5000 V; temperature, 600°C; ion source gas 1, 60 psi; ion source gas 2, 80 psi. Dwell times were 10 msec, and entrance potential was set to 10 V; compound specific parameters can be found in Table 2. The whole eluate was transferred to the electrospray probe from 1.0 till 2.8 min using the MS diverter valve. Ratios of the sums of the MNA and d_3 -MNA transitions were calculated and plotted versus concentration.

Compound	Q1 (m/z)	Q3 (m/z)	DP	CE	СХР
MNA	137.101	94.0	136	27	12
		92.0	136	29	12
		78.0	136	35	10
MNA- d_3	140.128	97.1	121	29	12
		95.1	121	31	12
		78.0	121	35	10

 Table 2. Tuned MS/MS parameters for all quantified components.

The entrance potential was set at 10 V for all compounds, dwell-time was 10 msec. Q1: quadrupole 1, Q3: quadrupole 3, z: charge, DP: declustering potential, CE: collision energy, CXP: collision cell exit potential.

Isothermal Titration Calorimetry: Expression and purification of full-length wild type NNMT protein (NNMTwt) were performed as previously described.³¹ Isothermal titration calorimetry (ITC) measurements were made at 25°C on a MicroCal ITC200 Instrument (Malvern Instruments) with 2 μ L injections. NNMTwt was diluted at 200 μ M in ITC buffer (50 mM Tris (pH 8.0), 150 mM NaCl) supplemented with 4% DMSO. Compounds were dissolved in DMSO at 50 mM and diluted to 2 mM in ITC buffer with a final DMSO concentration of 4%. Binding constants were calculated by fitting the data using the ITC data analysis module in Origin 7.0 (OriginLab Corp.).

Modeling studies: Docking computations were performed using Autodock 4.2.⁴⁹ Compounds 1, 2, 78, and 81 were docked into the catalytic pocket of the structure taken from PDB ID: 3ROD.³² Four molecular dynamic simulations were performed with GROMACS 2018.2⁵⁰ using the AMBER03 force field⁵¹. Each structure was immersed in a cubic box using TIP3P water molecules⁵² and neutralized with counter ions. A production step of 250 ns was carried out using the Parrinello-Rahman algorithm⁵³ for temperature and pressure control, with coupling constants of T=0.1 ps and P=2.0 ps, for compounds 1, 2, 81 and extended to 450 ns for compound 78, in order to reach equilibrium of the system. Coordinates were saved every 200 ps and the protein/ligand binding energy was estimated using g_mmpbsa calculations^{54,55} on the last 50 ns of each trajectory. The conformation of minimal energy in these 50 ns was extracted from the simulations and minimized in order to represent the interactions between the ligands and NNMT protein.

Enzyme assay for selectivity: Methyltransferase inhibition assays were performed as described⁵⁶ by using commercially available chemiluminescent assay kits for PRMT1 and NSD2 (purchased from BPS Bioscience). The enzymatic reactions were conducted in duplicate at room temperature for 1 h (PRMT1) or 2 h (NSD2) in substrate-coated well plates at a final reaction volume of 50 μ L containing: the manufacturer's proprietary assay buffer, AdoMet (at a concentration of 5 times the respective Km value for each enzyme), the methyltransferase enzyme: PRMT1 (100 ng per reaction) and NSD2 (500 ng per reaction), and inhibitor **78**. Before addition of AdoMet, the enzyme was first incubated with the inhibitor for 15 min at room temperature. Positive controls were performed in the absence of inhibitor using water to keep the final volume consistent. Blanks and substrate controls were performed in the absence of the enzyme and AdoMet, respectively. Following the enzymatic reactions, 100 μ L of primary

antibody (recognizing the respective immobilized methylated product) was added to each well, and the plate was incubated at room temperature for an additional 1 h. Then, 100 μ L of secondary horseradish peroxidase (HRP)-conjugated antibody was added to each well, and the plate was incubated at room temperature for additional 30 min. Finally, 100 μ L of an HRP substrate mixture was added to the wells, and the luminescence was measured directly by using a standard microplate reader. The luminescence data were normalized with the positive controls defined as 100% activity and blank defined as 0%.

Cell culture and treatment with compounds: The HSC-2 human oral cancer cell line was purchased from the American Type Culture Collection (ATCC, Rockville, MD, USA), and cultured in DMEM/F12 medium, supplemented with 10% fetal bovine serum and 50 μ g/ml gentamicin, at 37°C in a humidified 5% CO₂ incubator. Compounds 1, 2, 78, and 81 were tested for their inhibitory effect on cell proliferation of HSC-2 cells. Each compound was dissolved in DMSO at 100mM concentration. This stock solution was then diluted in culture medium to final concentration values ranging between 1 μ M and 100 μ M. For each sample, DMSO was kept constant at 0.1% final concentration.

The day before starting treatment, cells were seeded in 96-well plates, at a density of 1×10^3 cells/well. Cells were allowed to attach overnight and then incubated with compounds at different final concentrations, or with DMSO only, for 24, 48 and 72 hours. All experiments were performed in triplicate.

MTT assay: Cell proliferation was determined using a colorimetric assay with 3-(4,5dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT). The MTT assay measures the

conversion of MTT to insoluble formazan by dehydrogenase enzymes of the intact mitochondria of living cells. HSC-2 cell proliferation was evaluated by measuring the conversion of the tetrazolium salt MTT to formazan crystals upon treatment with compounds or DMSO only for 24, 48 and 72 hours. Briefly, cells were incubated for 2 hours at 37° C with 100 µl fresh culture medium containing 5µl of MTT reagent (5mg/ml in PBS). The medium was removed and 200µl isopropanol were added. The amount of formazan crystals formed correlated directly with the number of viable cells. The reaction product was quantified by measuring the absorbance at 540nm using an ELISA plate reader. Experiments were repeated three times. Results were expressed as percentage of the control (control equals 100% and corresponds to the absorbance value of each sample at time zero) and presented as mean values \pm standard deviation of three independent experiments performed in triplicate. Data were analysed using GraphPad Prism software (GraphPad Software, San Diego, CA). Significant differences between groups were determined using the one-way analysis of variance (ANOVA). A p value <0.05 was considered as statistically significant.

Quantitative measurements of MNA levels in cultured cells: The analysis was performed as previously described⁵⁷ with minor modifications. Cellular MNA levels were determined using the same UHP-HILIC-MS/MS employed for the inhibition studies as described above. To determine the effect of compound **78** on NNMT activity in the HSC-2 oral cancer cell line used cells were treated with **78** at 100 μ M (final DMSO content 0.1%) and incubated for 24, 48, or 72 hours. The day prior to starting treatment, cells were seeded in 6-well plates, at a density of $3x10^4$ cells/well. Cells were allowed to attach overnight and were then incubated with compound **78**. All experiments were performed in duplicate. Following treatment, medium was removed, and adherent cells were trypsinized and harvested by centrifugation at 1,000xg for 3 min at 4°C.

Supernatant was then discarded and cell pellets were stored at -80°C until further use. The extraction of MNA from the cell pellets was performed as previously described.⁵⁸ Briefly, 100 μ L acetonitrile containing 50 nM d₃-MNA (as internal control) was added to the cell pellets and the cells were lyzed for 20 minutes at room temperature with mild shaking. 50 μ L of purified water was added, followed by mixing and the resulting cell debris centrifuged for 10 minutes at 5,000 rpm. 100 μ L of the resulting supernatant was then transferred to a 96-well plate and analysed for MNA content.

ASSOCIATED CONTENT

Supporting Information

Analytical data for all new compounds including: ¹H and ¹³C NMR spectra and molecular formula strings. Supporting figures for: enzyme inhibition curves, molecular modelling results, and cellular MNA determination. Supporting tables for: selectivity assays and estimated binding energies. This material is available free of charge via the Internet at http://pubs.acs.org.

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ABBREVIATIONS

COMT, catechol *O*-methyltransferase; DCE, 1,2-dichloroethane; DIBAL-H, diisobutylaluminium hydride; DMAP, 4-dimethylaminopyridine; K_d , dissociation constant; IC₅₀, half maximal inhibitory concentration; ITC, isothermal titration calorimetry; MNA, 1-methyl-nicotinamide; NA, nicotinamide; NNMT, nicotinamide *N*-methyltransferase; NNMTwt, wild type NNMT; PDC, pyridinium dichromate; QTOF-MS, Quadrupole Time-Of-Flight Mass Spectrometry; RP-HPLC, reversed phase HPLC; SAH, *S*-Adenosyl-L-homocysteine; SAM, *S*-adenosyl-L-methionine; Trt, triphenylmethyl (trityl); UHP-HILIC, Ultra High Performance Hydrophilic Liquid Interaction Chromatography.

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