



Cite this: *Chem. Commun.*, 2026, 62, 2633

Received 27th November 2025,
Accepted 23rd December 2025

DOI: 10.1039/d5cc06746d

rsc.li/chemcomm

We report a robust method for C7-sulfonamide installation on 7-deazaadenosines, enabled by an unprecedentedly stable sulfonyl chloride. This general transformation introduces a new functionalization site and expands nucleoside chemical space, offering a powerful platform for constructing tailored probes and analogues in synthetic and bioorganic chemistry.

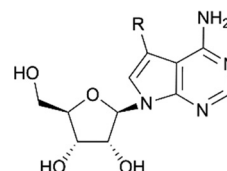
Nucleoside analogues are indispensable tools in medicinal chemistry and chemical biology, serving as both therapeutic agents and molecular probes. Structural modification of nucleosides has yielded numerous antiviral and anticancer drugs.¹ Pyrrolo[2,3-*d*]pyrimidine (7-deazapurine) nucleosides form a class of compounds closely mimicking purine nucleosides, and naturally occurring 7-deazaadenosine (tubercidin) (1), sangivamycin (2) and toyocamycin (3) (Fig. 1) display potent cytotoxic activities, making this scaffold highly attractive for further derivatization.²

Replacement of N7 by carbon in 7-deazapurines generates a unique substitution site at C7. Halo, (het)aryl, alkynyl, alkenyl, and alkyl substituents at this position have been extensively explored and shown to modulate the biological properties of the respective molecules.^{3–5} Our group has recently demonstrated that C7 substitution can be utilized to enhance activity and selectivity against several methyltransferase targets.^{6–10} In contrast, heteroatom-linked substituents remain underexplored. While a C7-carboxamide is occasionally present in sangivamycin analogues, it has rarely been used as a linker for further functionalization.¹¹

Sangivamycin itself has attracted longstanding interest. Initially studied as an anticancer nucleoside,^{12,13} it was later shown to possess broad antiviral activity,^{14–16} and to inhibit protein kinase C.¹⁷ More recently, it was identified as a potent

C7-Sulfonamide functionalization of 7-deazaadenosines: sangivamycin analogues with Haspin inhibitory activity

Milan Štefek, ^{ab} Milan Dejmek, ^a Michal Šála ^a and Radim Nencka *^a



- 1: R = H, tubercidin
- 2: R = CONH₂, sangivamycin
- 3: R = CN, toyocamycin

Fig. 1 Natural bioactive 7-deazaadenosines.

inhibitor of Haspin kinase,¹⁸ a serine/threonine kinase essential for mitotic chromosome alignment, which is overexpressed in malignant tissue and is responsible for cancer cell proliferation.¹⁹ Inhibition of Haspin by sangivamycin has been linked to pronounced anticancer effects, including induction of cell death in pancreatic cancer cell lines and tumor regression *in vivo*.¹⁸

In our exploration of 7-substituted-7-deazaadenosine derivatives, we identified C7-sulfonamides as an unexplored yet desirable modification. Sulfonamides are privileged motifs in drug discovery, valued for their polarity, hydrogen-bonding capacity, and metabolic stability, and they occur in numerous clinical agents.²⁰ Despite their widespread use in medicinal chemistry,^{21,22} C7-sulfonamide substitution of 7-deazapurines has not been reported. We therefore set out to establish a general method for their synthesis, demonstrate its applicability across diverse nucleoside scaffolds, and assess the biological activity of the resulting analogues.

Sulfonamides have been introduced to other positions of purine nucleosides, notably C2 and C6, yielding derivatives with antitumor and antiviral activities.^{23–25} However, C7 substitution of 7-deazapurines with sulfonamides has not been described, leaving this modification unexplored.

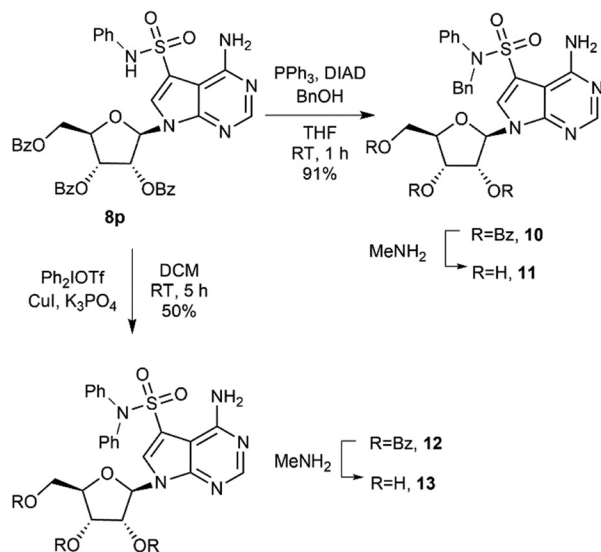
Despite the availability of numerous approaches to sulfonamide synthesis, the use of sulfonyl chlorides remains to be the preferred method because of their reliable reactivity and broad substrate compatibility.^{26–28} The C2 and C6 chlorosulfonyl-purines have previously been described as unstable, prone to hydrolysis or substitution by chloride anions. Yet, they are still

^a Institute of Organic Chemistry and Biochemistry of the Czech Academy of Sciences, Flemingovo náměstí 2, Prague 6 166 10, Czech Republic.

E-mail: radim.nencka@uochb.cas.cz

^b Department of Organic Chemistry, Faculty of Science, Charles University, Prague 128 00, Czech Republic



Scheme 2 N-Functionalization of sulfonamide **8p**.

exclusively to sulfonic acid, most likely *via* reaction of the sulfonyl chloride with the solvent to form a transient iminium salt.³² A biphasic DCM/water system with K_2CO_3 gave acceptable yields only after prolonged stirring, underscoring the hydrolytic stability of the sulfonyl chloride.

While most tested amines provided the corresponding sulfonamides in good to excellent yields, several substrates proved incompatible with sulfonyl chloride **7**. Electron-poor amines such as 3-aminopyridine, 4-aminopyridine, and 2-aminothiazole afforded only complex mixtures (data not shown), whereas diphenylamine was insufficiently reactive and did not yield any detectable sulfonamide product (see SI, Table S1). These observations indicate that strongly electron-deficient or weakly nucleophilic amines represent practical limitations of the current methodology.

Prepared secondary sulfonamides were amenable to further derivatization, as demonstrated with compound **8p** (Scheme 2). Alkylation under Mitsunobu conditions furnished benzylated compound **10** in excellent yield, while **12**, unavailable *via* direct sulfonation of diphenylamine, was obtained through Cu-catalysed coupling with Ph_2IOTf .³³ Final deprotection using $MeNH_2$ provided the free nucleosides.

Importantly, the methodology proved highly general. Beyond the sangivamycin analogue, it was successfully extended to multiple nucleoside scaffolds, including 2-substituted-7-deazaadenosines (**14**, **15**), pseudoadenosine with a pyrrolo[2,1-*f*]triazin core (**16**), 2'-modified nucleosides (**17**, **18**), and a carbocyclic nucleoside (**19**) (Fig. 2). This highlights the robustness of the transformation and establishes C7-sulfonamides as a versatile platform for nucleoside diversification.

Selected compounds were evaluated for inhibition of Haspin kinase (Table 1). Four analogues showed double-digit nanomolar IC_{50} values. The most potent, **9a**, inhibited Haspin at 21 nM, which is comparable to 7 nM value for sangivamycin. The IC_{50} data also allowed a basic SAR analysis. Changing the nucleobase core (compound **16**) afforded an analogue with similar

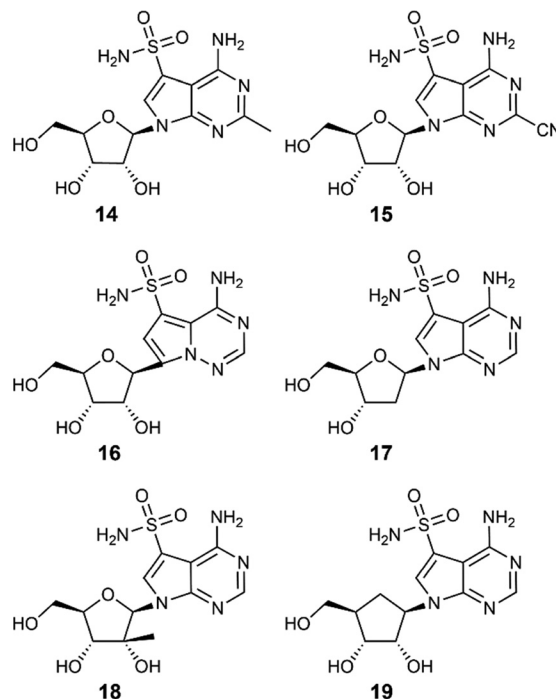


Fig. 2 Scope of C7-sulfonamide nucleosides. The transformation is compatible with diverse scaffolds.

Table 1 Inhibition of Haspin kinase by selected derivatives

Compound	Haspin IC_{50} (nM)
Sangivamycin	7
9a	21
9b	27
9c	55
14	> 10000
15	> 10000
16	30
17	354
18	344
19	1708

potency, suggesting that the central heterocycle can tolerate certain modifications. Introduction of substituents on the sulfonamide nitrogen (**9b**, **9c**) resulted in only a minor loss of activity, indicating that this moiety does not engage in crucial interactions with the protein. In contrast, modifications of the ribose portion of the nucleoside (**17**, **18**, **19**) led to substantially reduced inhibitory activity, and substitution at the C2 position produced compounds (**14**, **15**) with no detectable inhibition of Haspin kinase. These observations imply that the binding site is highly constrained around the C2 region and cannot accommodate additional substituents. To assess its selectivity, **9a** was profiled in a 63-kinase panel. Comparison with sangivamycin revealed that **9a** possesses an altered selectivity profile. Whereas sangivamycin inhibited several kinases in the panel, including $PKC\delta$, YSK4, KDR, DYRK1A, and DYRK2, compound **9a** showed reduced activity toward these off-targets but displayed measurable inhibition of AMPK α 1 and RSK1 instead (SI, Table S2). Overall, **9a** retains potent Haspin inhibition while



exhibiting a distinct and somewhat narrower pattern of off-target interactions (see SI, Table S2).

In summary, we report the first synthesis of C7-sulfonamide-7-deazaadenosines enabled by a uniquely stable sulfonyl chloride intermediate. The transformation is general, applicable to a wide range of amines and extending across multiple nucleoside scaffolds to provide derivatives readily suited for further modification. Several of the synthesized compounds exhibited potent low-nanomolar Haspin inhibition with high selectivity against other kinases. These findings establish C7-sulfonamides as a new and versatile entry point for nucleoside diversification with direct relevance to medicinal chemistry.

Conflicts of interest

There are no conflicts to declare.

Data availability

The data supporting this article have been included as part of the supplementary information (SI). Supplementary information is available. See DOI: <https://doi.org/10.1039/d5cc06746d>.

Acknowledgements

This research was funded by the project New Technologies for Translational Research in Pharmaceutical Sciences/NET-PHARM, project ID CZ.02.01.01/00/22_008/0004607, which is co-funded by the European Union. Also, the project was supported by the Academy of Sciences of the Czech Republic as part of the Strategy AV 21 Virology and Antiviral Therapy programme; the Czech Academy of Sciences (RVO: 61388963).

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