

This paper is dedicated to Professor David Shugar on the occasion of his 80th birthday

NAD-analogues as potential anticancer agents: Conformational restrictions as basis for selectivity*

Krzysztof W. Pankiewicz^{a**}, Andrzej Zatorski^b and Kyoichi A. Watanabe^a

^aOncorPharm, 200 Perry Parkway, Gaithersburg, MD 20877, U.S.A.

^bSloan-Kettering Institute, Memorial Sloan-Kettering Cancer Center, Sloan-Kettering Division of Graduate School of Medical Sciences, Cornell University, New York, New York 10021, U.S.A.

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Cofactor type inhibitors (NAD-analogues) of IMP-dehydrogenase (IMPDH) were synthesized and their application as potential anticancer agents are discussed. C-nucleoside isosteres of NAD, C-NAD and C-PAD, showed an effective competitive inhibition of IMPDH. C-NAD but not C-PAD caused extremely potent inhibition of alcohol dehydrogenase. We also synthesized compounds in which nicotinamide riboside was replaced with tiazofurin (TAD-analogues) and the 2' and 3'-positions of adenosine part were fluorinated. The ribose ring of 2'-deoxy-2'-fluoroadenosine is in the C3'-endo conformation whereas 3'-deoxy-3'-fluoroadenosine favors the C2'-endo sugar pucker. These derivatives are good inhibitors of IMPDH type II, the isoenzyme dominant in neoplastic cells. In contrast, all these analogues showed rather weak inhibitory activity against alcohol dehydrogenase. Nicotinamide riboside derivatives in which the base and the sugar are linked through an oxygen or a methylene bridge were synthesized. NAD-analogues containing such conformationally restricted nicotinamide nucleoside moiety (*syn* or *anti*) are expected to be selective inhibitors of B-specific (IMPDH) or A-specific dehydrogenases, respectively.

IMP-dehydrogenase (IMPDH), the enzyme which catalyzes the NAD-dependent conversion of inosine 5'-monophosphate to guanosine 5'-monophosphate, is positioned at the branch point in the *de novo* synthesis of guanine nucleotides [1]. Inhibition of IMPDH has been shown to have antiviral [2, 3], immunosuppressive [4, 5], antiparasitic [6, 7] and anticancer effects [8, 9]. IMPDH has been suggested [1, 8, 9] as a key enzyme in neoplasia and one of the most sensitive targets for cancer chemotherapy. The level of IMP-dehydrogenase activity was found to be

much greater in several tumors as compared to normal tissues [10]. Human IMPDH cDNA was first cloned and sequenced by Collart & Huberman [11] showing that the protein contained 514 amino acids. Increased IMPDH gene expression has been then found in several tumor cell lines [12]. It has been recently discovered [13] that human IMP-dehydrogenase exists as two isoforms, types I and II. The two distinct cDNAs have been characterized and it has been found that the two isoforms are differently regulated [14-16]. Type I is constitutively ex-

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**To whom correspondence should be addressed.

Abbreviations: ADH, alcohol dehydrogenase; C-NAD, C-glycosidic nicotinamide adenine dinucleotide; C-PAD, C-glycosidic picolinamide adenine dinucleotide; IMP, inosine 5'-monophosphate; IMPDH, IMP-dehydrogenase; TAD, thiazole-4-carboxamide adenine dinucleotide.

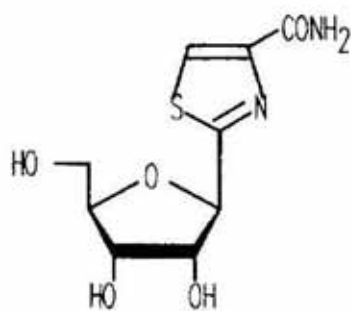
pressed and is the preponderant isoform in normal cells, while type II is selectively up-regulated in neoplastic and replicating cells and emerges as the dominant species. Interestingly, when neoplastic cells are induced to differentiate, the type II transcript is selectively down-regulated to a level below that of type I. Thus, the selective inhibition of type II IMPDH is expected to provide significant therapeutic advantage by eliminating or reducing potential toxicity caused by inhibition of type I isoform [13–16].

Tiazofurin, 2-(β -D-ribofuranosyl)thiazole-4-carboxamide, is metabolically converted into the NAD analogue, thiazole-4-carboxamide adenine dinucleotide (TAD, Scheme 1) which was found to be a competitive inhibitor of IMP-dehydrogenase [17]. Tiazofurin exhibits potent activity against various murine and human tumors [18, 19], and has been evaluated as an anticancer agent in Phase 1 and Phase 2 clinical studies [20]. In recent years, the ability of tiazofurin to decrease guanine nucleotide pool created new hope and excitement. In 1991 the compound was on the cover of the journal *Blood* [21, 22] as an inducer of myeloid differentiation. Due to the inhibition of G-protein mediated transmembrane signaling [23], cells had undergone maturation and differentiation. Weber [24] noted that patients with chronic granulocytic leukemia in blast crisis treated with tiazofurin demonstrated no depletion in bone marrow. Examination of bone marrow showed decrease in blast cells and increase in more mature forms of granulocytes. These results appear to indicate drug-induced cell differentiations in humans [24–27]. More importantly this treatment has induced complete remissions in 77% of patients [24]. It was also found

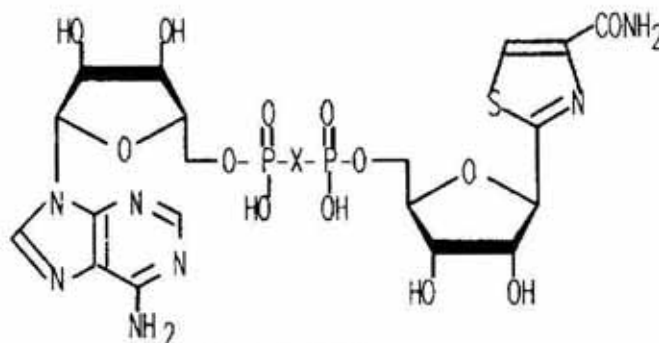
that tiazofurin treatment modulates gene expression. Rapid decrease in *c-Ki-ras* messenger RNA levels and *c-myc* expression in K-562 cells were reported, whereas the mRNA level of *c-abl* gene was not affected [27–29].

Tiazofurin requires a unique metabolic activation. It is phosphorylated by adenosine kinase or other kinase(s) to the 5'-monophosphate [30] and then converted to TAD by NAD-pyrophosphorylase [31]. TAD has been synthesized [31, 32] and has been found to be an even more potent inhibitor of IMP-dehydrogenase than tiazofurin. TAD mimics NAD but can not function as the coenzyme. It was found to interfere with NAD synthesis *in vitro*, and is a weak inhibitor of poly(ADP-ribose) synthase [33]. Accumulation of TAD, however, is regulated by cellular phosphodiesterase called TAD-ase, which is responsible for TAD breakdown and development of cell lines resistant to tiazofurin [34, 35]. In order to circumvent the resistance development, a TAD analogue containing the phosphonate ($-P-CH_2-P-$, Scheme 1) instead of pyrophosphate ($-P-O-P-$) moiety was synthesized [36]. This analogue (β - CH_2 -TAD) is not susceptible to enzymic hydrolysis, and is the most potent inhibitor of IMP-dehydrogenase [36].

Since the discovery of oncolytic properties of tiazofurin, other C-nucleoside analogues have been studied. Selenazofurin (5, Scheme 2), an analogue containing selenium instead of sulfur, was synthesized [37] and its corresponding dinucleotide anabolite was found to be even more cytotoxic than tiazofurin. In contrast, isosteric replacement of the sulfur or selenium with oxygen, as in oxazofurin (6) [38] abolished the antitumor activity. Arabino (7) and xylo (8) congeners of tiazofurin show no cytotoxicity



1. Tiazofurin



2. X=O; TAD

3. X=CH₂; β -methylene TAD

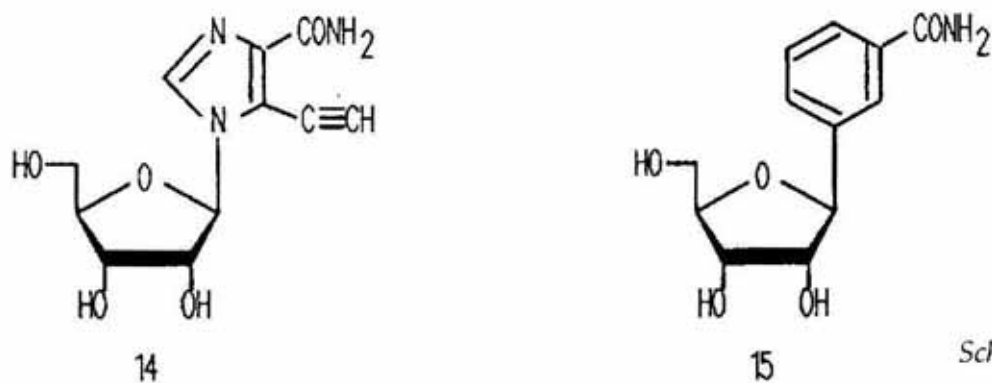
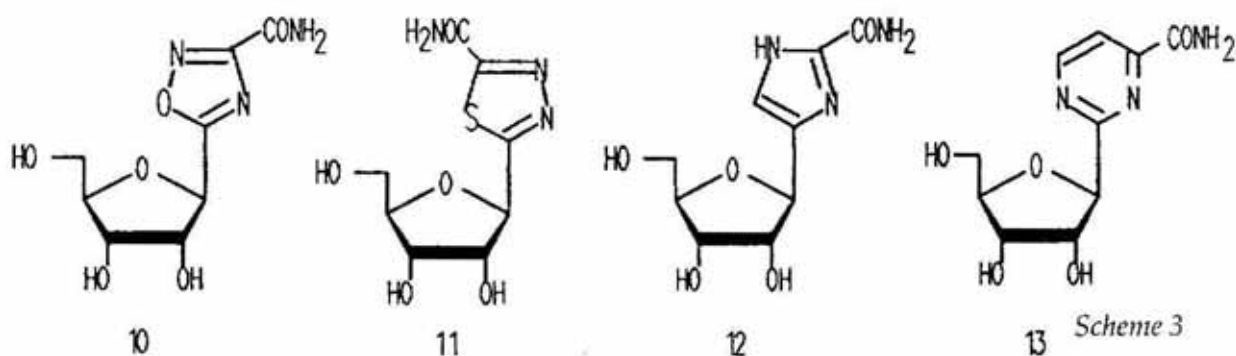
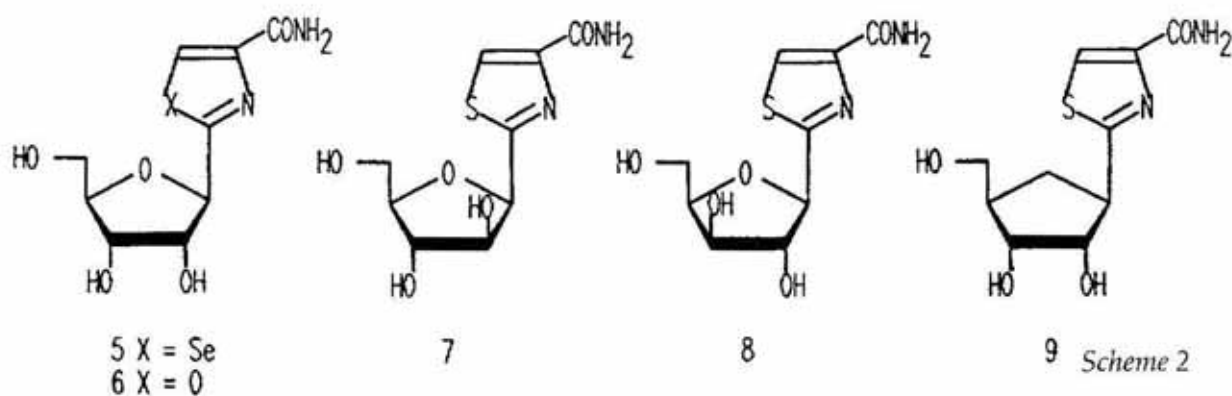
4. X=CF₂; β -difluoromethylene TAD

[39]. Recently, carbocyclic tiazofurin (**9**) and its enantiomer have been reported to display cytotoxicity against a breast carcinoma cell line [40]. Another nucleoside, the 5-(β -D-ribofuranosyl)-1,2,4-oxadiazole-3-carboxamide (**10**, Scheme 3) caused only weak inhibition of leukemia L1210 and P388 lines [41]. No activity was reported for 1,3,4-thiadiazole C-nucleoside analogue **11** of tiazofurin [42]. An imidazole C-nucleoside, 4-(2-deoxy- β -D-ribofuranosyl)imidazole-2-carboxamide (**12**) has also been prepared recently [43]. The synthesis of the 2-(β -D-ribofuranosyl)pyrimidine-4-carboxamide (**13**) and its NAD analogue has been published, but the antitumor activity of these compounds has not been reported [44].

Two novel, potent inhibitors of IMPDH have recently been reported. First, 5-ethynyl-1- β -D-

ribofuranosylimidazole-4-carboxamide (**14**, Scheme 4, synthesized by Matsuda *et al.* [45]) and its 5'-monophosphate derivative inhibit IMPDH from L1210 cells competitively with respect to IMP [46]. Second, 3- β -D-ribofuranosylbenzamide (**15**) showed toxicity in a nanomolar concentration to S49.1 lymphoma cells [47] and leukemia K562 cells [48]. Interestingly, benzoic acid riboside is almost nontoxic to the lymphoma cells [47]. Benzamide adenine dinucleotide has been prepared enzymatically and found to be an active metabolite of benzamide riboside [49].

The known inhibitors of IMPDH can be broadly classified into three groups based upon the mode of enzyme binding [50]. The first group consists of IMP/XMP (substrate) analogues such as 6-chloropurine ribotide (**16**, Scheme 5), riba-

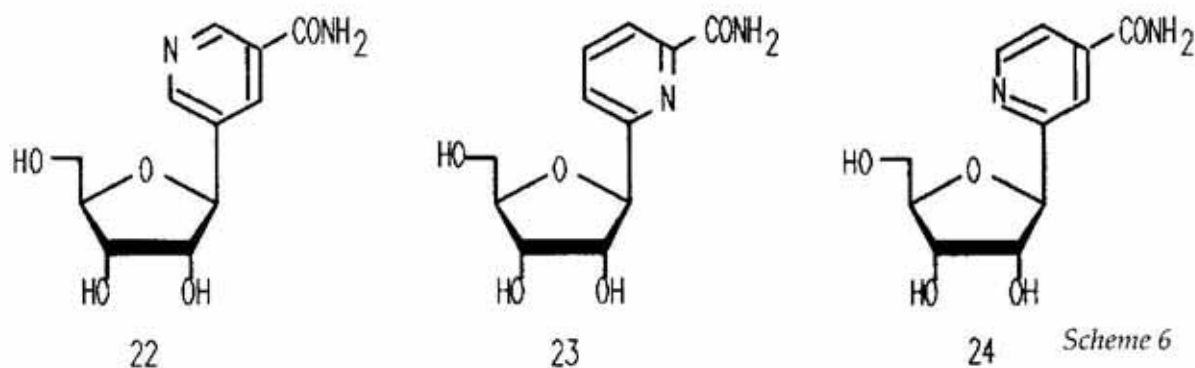
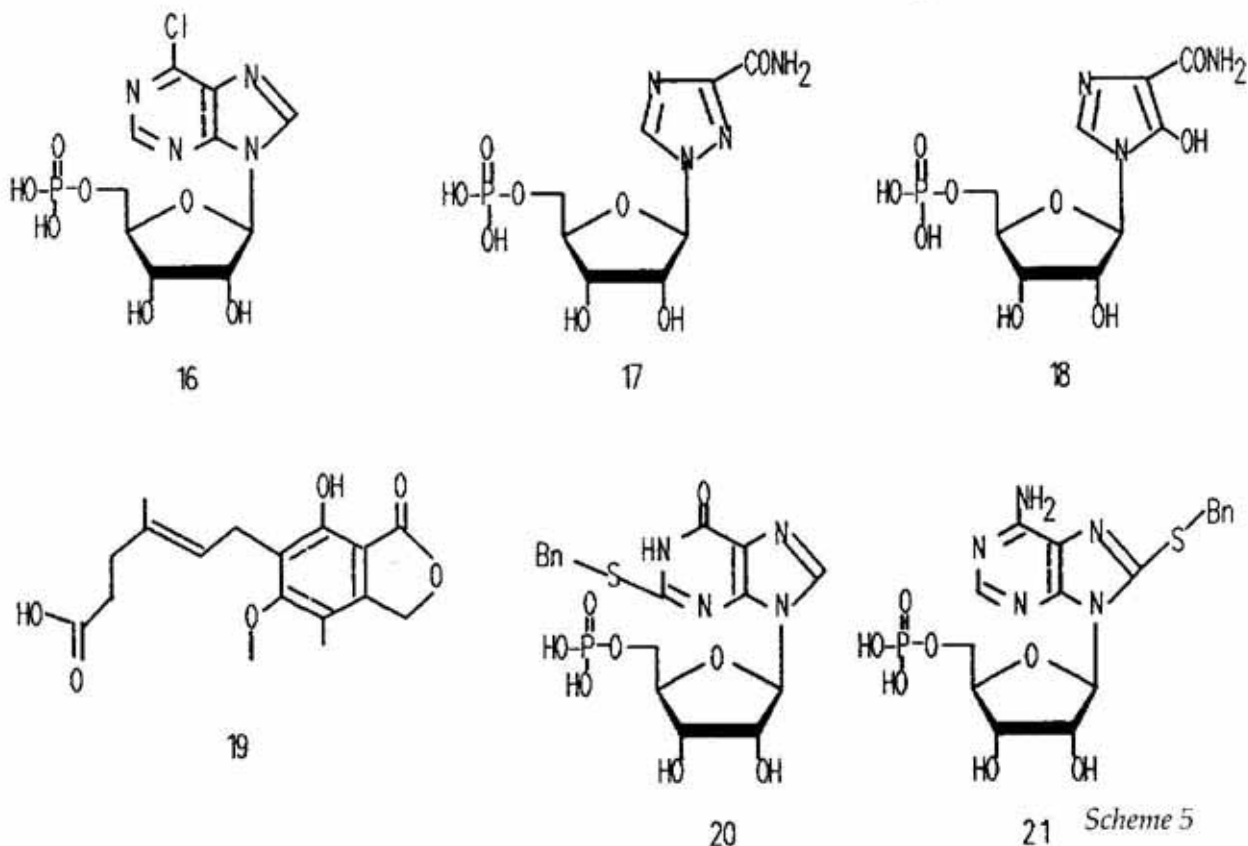


virin-5'-phosphate (17), and bredinin-5'-phosphate (18). The second group are the NAD⁺/NADH analogues, tiazofurin and mycophenolic acid (19). The third group of IMPDH inhibitors are 2-thiobenzyl IMP (20) and 8-thiobenzyl AMP (21), multisubstrate inhibitors that presumably occupy the IMP/XMP binding domain with groups projecting into NAD site to enhance binding. The importance of cofactor type inhibitors consists in their mode of action which offers specificity for IMPDH [50]. More general inhibitors of NAD-dependent enzymes would be expected to have greater toxicity [50].

We have reported [51–53] the synthesis of the closest structural analogues of nicotinamide riboside, i.e. C-nucleoside isosteres — nicotin-

amide 22 and picolinamide 23 (Scheme 6). Later, 2-β-D-ribofuranosyl-isonicotinamide (24) was synthesized by others [54]. In contrast to our expectation these compounds showed only weak inhibitory activity against L1210, P-815, HL-60, CCRF-CEM, MOLT/4F, and MT-4 cell lines [51–54].

Recently, we converted our nicotinamide C-nucleoside 22 and picolinamide C-nucleoside 23 into the corresponding NAD analogues, C-NAD (Scheme 7) and C-PAD, respectively [55]. We found [55] that C-NAD was an effective competitive inhibitor of IMPDH (ID₅₀ = 20 μM). Interestingly, C-NAD caused extremely potent inhibition of horse liver alcohol dehydrogenase (ADH, K_i = 1.1 nM), whereas C-PAD was found to be a much less potent inhibitor of ADH



($K_i = 20 \mu\text{M}$). It was later discovered by Goldstein *et al.* [56, 57] that the specificity and affinity of C-NAD for ADH are due to coordination of the zinc cation at the ADH catalytic site by the pyridine nitrogen of C-NAD. Displacement of the pyridine nitrogen to the opposite side of the ring, as in C-PAD, removes the specificity for ADH [56, 57].

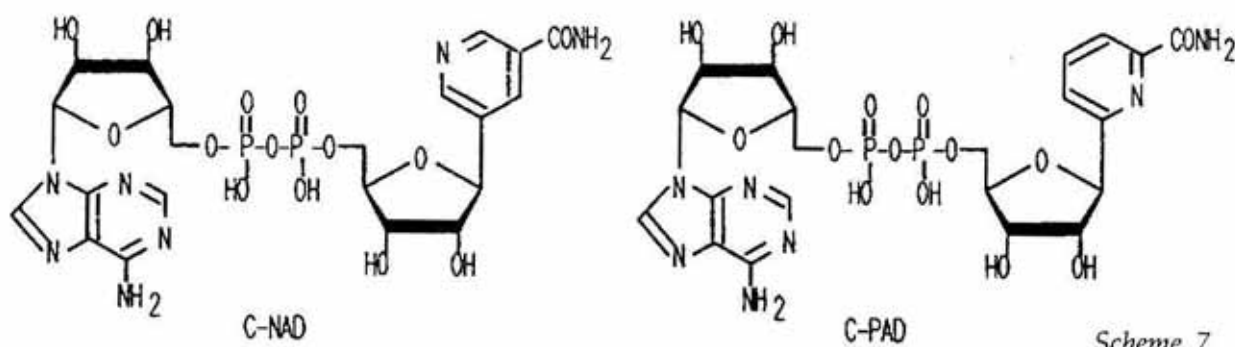
All these studies indicate that with the exception of tiazofurin, selenazofurin, and perhaps carbocyclic tiazofurin, other nucleosides related to nicotinamide riboside are not efficiently converted into corresponding NAD analogues by cellular enzymes. Therefore, NAD analogues, *not nucleosides*, that are able to penetrate cells membrane should be of therapeutic interest. These NAD analogues should be specific inhibitors of IMPDH (possibly type II), not toxic or less toxic towards cellular dehydrogenases. In contrast to nucleosides, NAD analogues do not require metabolic activation by cellular enzymes. NAD analogues containing C-nucleosides instead of nicotinamide riboside should be resistant to cellular catabolic enzymes, such as NAD-glycohydrolases. Methylene bisphosphonate analogue of NAD should be effective against tiazofurin resistant cell lines [34, 35] due to "built in" resistance to NAD-phosphodiesterases.

Since the X-ray structure of IMPDH is not known, we used well established ADH structure for general studies of enzyme-inhibitor (cofactor type) interactions. Our recent crystallographic studies [58] of a complex of ADH and β -methylene-TAD showed preservation of hydrogen bonds between donors O2' and O3' of the adenosine ribose and negatively charged Asp-223. In addition, the positively charged Lys-228 donates a hydrogen bond to adenosine O3'. These interactions are also observed in ADH-bound NAD, and are conserved in a number of dehydrogenases [59]. In IMPDH, as

in ADH, TAD displaces NAD binding at the cofactor site. Initial recognition of the cofactor appears to occur *via* binding to the adenosine end [60].

These findings prompted us to synthesize three TAD analogues containing fluorine atom at the C2'-ribo, C3'-ribo, and C2'-arabino positions of the adenine nucleoside (25, 26, and 27, respectively, Scheme 8) as probes of the stereochemical requirements of the adenine end on IMPDH [61]. Fluorine can mimic hydroxyl group in terms of size and polarity [62] and is able to form an even stronger hydrogen bond than a hydroxyl group, acting as an acceptor. Some dehydrogenases require the C2'-endo conformation of the adenosine ribose, others require C3'-endo. Interestingly, the ribose ring of 2'-deoxy-2'-fluoro-adenosine is in a C3'-endo conformation due to the highly electronegative 2'-substituent [63]. In contrast, 3'-deoxy-3'-fluoro-adenosine favors a C2'-endo sugar pucker [64]. Thus, TAD analogues 25-27 are expected to bind tightly to enzymes which are able to provide a good donor type hydrogen bonding.

We found [61] (Table 1) that all our compounds showed good inhibitory activity against IMPDH-II, the isoenzyme predominant in neoplastic cells [16]. Analogues 25 ($K_i = 0.5 \mu\text{M}$), 26 ($K_i = 0.7 \mu\text{M}$), and 27 ($K_i = 2.9 \mu\text{M}$) were a little less effective than TAD ($K_i = 0.2 \mu\text{M}$). Interestingly, the level of inhibition of ADH was found to be much lower (0.1 mM for 25 and 26, and no inhibition up to 10 mM for 27). Thus, new compounds are potent inhibitors of IMPDH-II, show some selectivity, and may be of therapeutic potential. TAD-25 and TAD-27 do not contain the 2'-hydroxyl function of the adenosine moiety and therefore can not be converted into the corresponding NAD(P) analogue by cellular enzyme(s). Thus, these compounds are expected to be less toxic than TAD itself, since the former should be



harmless toward numerous NAD(P)-dependent cellular enzymes.

We also synthesized the β -difluoromethylene-TADs (β -CF₂-TADs, 28 and 29, Scheme 9) [65] containing -CF₂- group as replacement of pyrophosphate oxygen, and found [61] that β -CF₂-TAD-28 ($K_i = 0.17 \mu\text{M}$) was almost equally effective to β -CH₂-TAD ($K_i = 0.11 \mu\text{M}$) against IMPDH-II. Transport properties, metabolism, and toxicity of these fluoro-substituted analogues are to be studied.

We assumed that even subtle modifications in the structure of inhibitor of cofactor type, might alter selectivity against targeted dehydrogenases. Thus, changes in conformation of adenine sugar moiety, *vide supra*, or restrictions in *syn-anti* positioning of nicotinamide (rotation around the glycosyl bond), *vide infra*, should affect drug-enzyme interactions.

Restricted rotation around the glycosyl bond of tiazofurin has been suggested as an important factor in its biological activity [66]. It was speculated that good IMPDH inhibitory activity of tiazofurin was due to its favorable conformation for binding to enzyme(s) that converts tiazofurin to TAD, or due to tight binding of TAD to IMPDH. Goldstein and his group found that close intramolecular contact between the sulfur and the ribose ring oxygen in tiazofurin (responsible for restriction in rotation) [66] is maintained in β -CH₂-TAD bound to ADH [58]

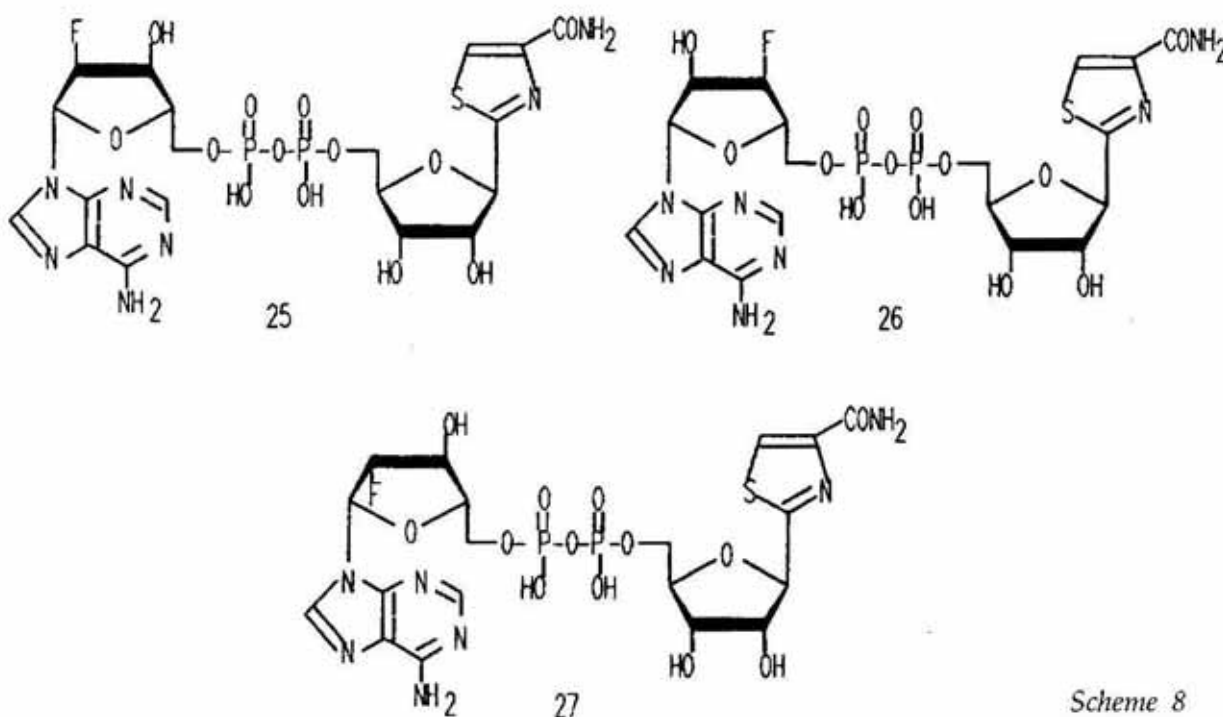
The importance of rigid molecules that have fixed glycosyl torsion angle X is probably most dramatically expressed in the case of uridine phosphorylase inhibition. The K_i value of the 2,2'-anhydro-5-ethyluridine, one of the most potent inhibitors of this enzyme, is more than three orders of magnitude lower than that of arabinofuranosyl-5-ethyluracil, the corresponding unrestricted analogue [67].

Table 1
Inhibition of human IMP-dehydrogenase type II by TAD analogues.

NAD is the variable substrate, with IMP constant at 100 μM . The pattern of inhibition in each case is non-competitive. Values were obtained at 37 °C.

Inhibitor	K_{ij} (μM)	K_{is} (μM)
25	0.5 ± 0.1	6.0 ± 12.0
26	0.7 ± 0.1	1.5 ± 0.6
27	2.9 ± 0.5	13 ± 10
β -CF ₂ -TAD	0.17 ± 0.03	0.3 ± 0.2
β -CH ₂ -TAD	0.11 ± 0.02	0.2 ± 0.1
TAD	0.19 ± 0.03	0.3 ± 0.1
NADH	120 ± 7	175 ± 22

ADH inhibition: K_i 's for 25, 26 and β -CF₂-TAD no less than 0.1 mmol; for 27 no inhibition up to 10 mmol.



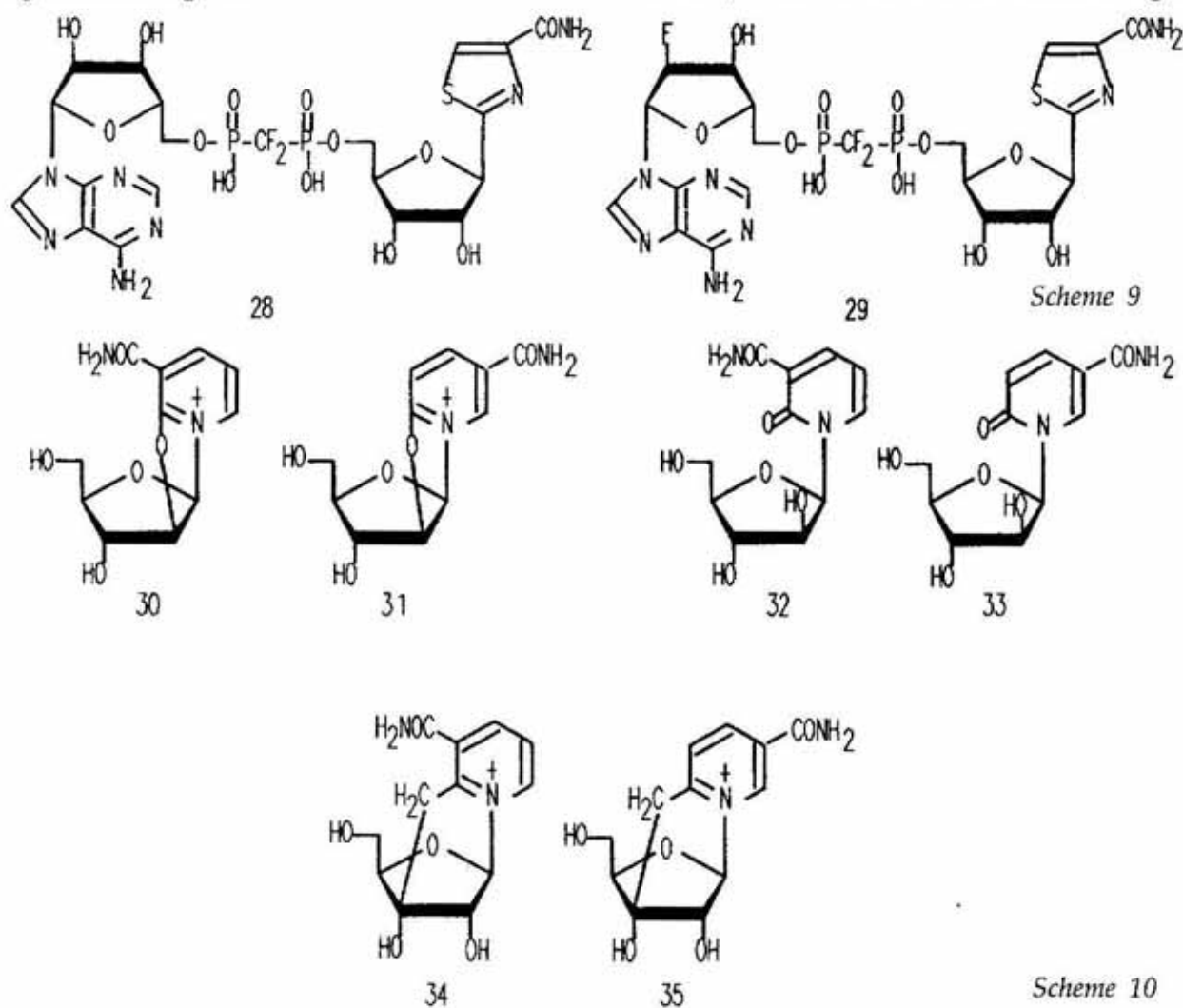
Scheme 8

Enzymic oxidation-reduction by all known dehydrogenases are stereospecific processes [68], i.e., only pro-R hydrogen of the dihydropyridine ring in NADH is transferred by some dehydrogenases (A-specific, as ADH), whereas others (B-specific, as IMPDH) transfer exclusively pro-S hydrogen. The X-ray studies of various complexes of dehydrogenases and NAD revealed that those enzymes that bind the NADH molecule in *syn* conformation transfer pro-S hydrogen, whereas transfer of pro-R hydrogen requires that NADH is bound to the enzyme in the *anti* conformation [69]. If conformation of nicotinamide nucleoside moiety of NAD analogue is restricted to *syn*, such NAD analogue may serve as coenzyme only for B-specific but not for A-specific dehydrogenases. Therefore, at least in principle, the *syn* conformationally restricted NAD analogues are expected to function as inhibitors only for B-specific enzymes. Consequently, NAD analogue containing *anti* restricted nicotinamide nu-

cleoside should exhibit inhibitory activity against A-specific dehydrogenases. Thus, it should be expected that the value of X (defined as the angle O4', C1', N1, C2, with *syn* and *anti* as the borderlines) would dramatically affect enzyme-inhibitor interaction and consequently the selectivity of enzymes inhibition.

In order to test the above hypothesis, we synthesized 2,2'-anhydro-2-hydroxy- and 6,2'-anhydro-6-hydroxy-1-(β -D-arabinofuranosyl)nicotinamide (30 and 31, respectively, Scheme 10) [70]. Our attempted conversion of 30 and 31 into their corresponding 5'-monophosphates was, however, unsuccessful due to instability of the anhydro linkage of 30 and 31 resulting in the formation of *arabino* derivatives 32 and 33 [70].

Recently, we have prepared analogues of nicotinamide mononucleotide in which the base and sugar are linked through a methylene bridge, 34 and 35, making the conformation restricted to the *syn* and *anti*, respectively [71, 72]. We found, however, that due to severe strain in their rigid



Scheme 10

structures, these compounds were susceptible to hydrolysis of the glycosyl bond. Therefore, we are now synthesizing conformationally restricted C-nucleosides with stable glycosyl bond. If the concept of conformationally restricted NAD analogous is valid, approximately half of the cellular dehydrogenases (pro-S or pro-R) will be unaffected (lack of toxicity) by our synthetic targets.

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