

# *The Next Ten Stories on Antiviral Drug Discovery (Part E): Advents, Advances, and Adventures*

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**Abstract:** This review article presents the fifth part (part E) in the series of stories on antiviral drug discovery. The ten stories belonging to this fifth part are dealing with (i) aurotricarboxylic acid; (ii) alkenyldiarylmethanes; (iii) human immunodeficiency virus (HIV) integrase inhibitors; (iv) lens epithelium-derived growth factor as a potential target for HIV proviral DNA integration; (v) the *status presens* of neuraminidase inhibitors NAIs in the control of influenza virus infections; (vi) the *status presens* on respiratory syncytial virus inhibitors; (vii) tricyclic (1,*N*-2-ethenoguanine)-based acyclovir and ganciclovir derivatives; (viii) glycopeptide antibiotics as antivirals targeted at viral entry; (ix) the potential (off-label) use of cidofovir in the treatment of polyoma (JC and BK) virus infections; and (x) finally, thymidine phosphorylase as a target for both antiviral and anticancer agents. © 2009 Wiley Periodicals, Inc. *Med Res Rev*, 31, No. 1, 118–160, 2010

**Key words:** ATA; ADAM; HIV integrase inhibitors; LEDGF; influenza; neuraminidase inhibitors; RSV (respiratory syncytial virus) inhibitors; tricyclic guanosine derivatives; glycopeptide antibiotics; polyoma virus infections; thymidine phosphorylase

## **1. INTRODUCTION**

This review article corresponds to the fifth paper in the series of stories on antiviral drug discovery (part E), thus following up, in a similar, narrative style on the stories presented in part A (although, it was not labeled as such),<sup>1</sup> part B,<sup>2</sup> part C,<sup>3</sup> and part D.<sup>4</sup> The present article (part E) is entitled “Advents, Advances, and Adventures” because it heralds new opportunities for antiviral drug design, that have blossomed recently, and may represent important advances in our understanding of antiviral drug development, but considering the pitfalls involved they may also signal some adventures ahead.

The first story (story E1) concerns aurotricarboxylic acid (ATA), a compound accredited over the past 25 years with an ever-increasing number of biological, including antiviral, activities.

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From ATA originated the alkenyldiarylmethanes (ADAMs), which, as told in story E2, have been primarily pursued as anti-human immunodeficiency virus (HIV)-1 agents, alike the non-nucleoside reverse transcriptase inhibitors (NNRTIs) HEPT (A8), TIBO (A9), rilpivirine (C9), and TBZ (D3).

Integrase inhibitors (INIs) have, with raltegravir, only recently made their appearance on the anti-HIV drug scene (story E3), which is almost two decades after integrase was considered a potential target for anti-HIV drug design.

Whether the host cell factor LEDGF (lens epithelium-derived growth factor), which helps the HIV integrase in accomplishing the integration of the proviral DNA into the cellular genome could be considered as an attractive antiviral target remains an intriguing story (story E4).

For influenza (already covered in C3 and D5), the seasonal flu has continued to lead to its annual outbreaks (with, unexpectedly, an outbreak of influenza A (H1N1), tentatively called “Mexican”), but the so much feared bird flu (avian influenza A H5N1) has (fortunately) not (yet) arrived, while the stockpiled neuraminidase inhibitors (NAIs), particularly oseltamivir, are facing increasing emergence of drug-resistant influenza H1N1 variants, which would be particularly worrisome should this extend to the “Mexican” H1N1 and to the avian H5N1 (story E5).

Although considered as only second in importance to influenza, respiratory syncytial virus (RSV) infections have received relatively little attention from a chemotherapeutic viewpoint. Available options are ribavirin and monoclonal antibody (palivizumab), and a number of fusion inhibitors, as well as one nucleocapsid binder (RSV-604), are under scrutiny (story E6).

In the past, little, if any attention has been given to a class of tricyclic acyclovir or ganciclovir derivatives (containing 1,*N*-2-ethenoguanine) that besides being active as antiviral agents [for acyclovir, see C7] are also endowed with fluorescent properties (story E7).

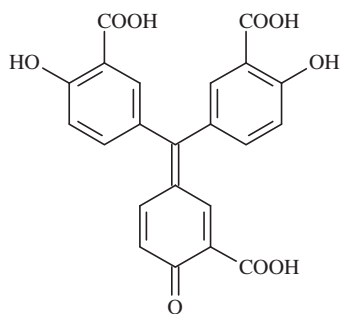
Antibiotics have been rarely linked to antiviral activity, except for aminoquinolone derivatives that, being transcription inhibitors, deploy some activity against HIV. In story E8, I review the inhibitory effects of aglycosylated glycopeptide antibiotics (i.e. vancomycin, teicoplanin, etc.) on the entry of a number of viruses [particularly HIV, but also coronaviruses (including the severe acute respiratory syndrome coronavirus (SARS CoV))].

Story E9 follows up on D8, where I described the potential of cidofovir (off-label) in the treatment of human papillomavirus (HPV)-associated infections. In story E9 I now further address the potential of cidofovir for the treatment of polyoma (JC and BK) virus infections.

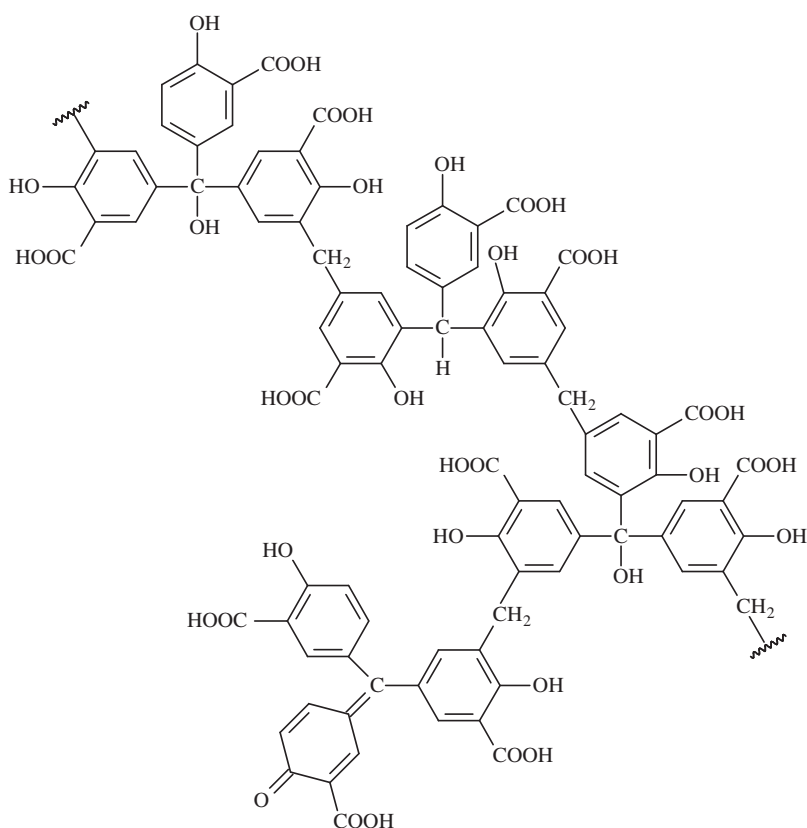
During our studies with (*E*)-5-(2-bromovinyl)-2'-deoxyuridine (BVDU) (A3), we observed that the compound is readily cleaved through thymidine (dThd) phosphorylase to yield (*E*)-5-(2-bromovinyl)uracil (BVU) and 2-deoxyribose-1-phosphate. In story E10 I discuss dThd phosphorylase as a target for antiviral as well as antitumor activity.

### **A. ATA, a Salicylic Acid Derivative, With a Myriad of Antiviral Effects**

ATA (Fig. 1) is a heterogeneous mixture of polymers formed when salicylic acid is treated with formaldehyde, sulfuric acid, and sodium nitrite.<sup>5,6</sup> ATA is often incorrectly represented as a triphenylmethane dye (Fig. 1) rather than as a polymer (Fig. 1).<sup>7</sup> While ATA had been known for a long time to interact with a number of enzymes including DNA polymerases, RNA polymerases, reverse transcriptase (RNA-dependent DNA polymerase), aminoacyl-tRNA synthetase, ribonucleotide reductase, and ribonucleases (as reviewed by Cushman et al.<sup>7</sup>), it was considered as a nonspecific enzyme inhibitor,<sup>8</sup> simply because ATA, being a polycarboxylate, would bind by electrostatic interactions to any protein that contains



ATA monomer



ATA polymer

**Figure 1.** Aurintricarboxylic acid (ATA) is often incorrectly represented as a triphenylmethane dye rather than as a polymer.

positively charged residues. [The use of ATA as an inhibitor of nucleases during nucleic acid isolation was advocated at a certain time.<sup>9</sup>]

Interest in ATA revived when it was shown to inhibit the cytopathogenicity of HIV in cell culture.<sup>10,11</sup> ATA was then shown to specifically interact with CD4, the cell's principal receptor for HIV, as it selectively prevented the binding of OKT4A/Leu-3a monoclonal

antibody to the CD4 cell receptor.<sup>12</sup> In further studies, ATA also appeared to block the binding of monoclonal antibody to the viral envelope glycoprotein gp120.<sup>13</sup> The lower molecular weight fractions of ATA bound to gp120, but not CD4, and, as they prevented the cytopathicity of HIV-1 and HIV-2, it could be concluded that the ATA fractions bound more avidly to gp120 than to CD4, and that the binding of ATA to gp120 in the absence of CD4 binding was sufficient for anti-HIV activity.<sup>7</sup> After a number of ATA monomer analogues (such as the triphenylcarbinol and triphenylmethane) had proven to be virtually inactive against HIV-1,<sup>14</sup> interest in ATA, as a potential lead compound for the treatment of HIV infections, largely waned.

Yet, new viruses emerged (or re-emerged) and so did the interest in ATA. After ATA had been reported as a potent inhibitor of the SARS CoV,<sup>15</sup> ATA was predicted, by using computer-aided analysis, to bind to and inhibit the SARS CoV RNA-dependent RNA polymerase (RdRp), thereby blocking viral transcription.<sup>16</sup> However, whether ATA inhibits SARS CoV replication through interference with the RdRp function has not been directly demonstrated [considering the negative charges of ATA, one could indeed question how easily this compound enters the cells].

Then followed the report that ATA inhibits (the early stage of) vaccinia virus replication,<sup>17</sup> as the result of two mechanisms of action: (i) blocking the extracellular signal-regulated kinase (ERK) phosphorylation [activation of the ERK cascade is essential for vaccinia virus replication], and (ii) inhibiting the phosphatase activity of the viral enzyme HIL [which plays an essential role in initiating viral early gene transcription]. In fact, ATA would be able to interfere with various signaling pathways: viz. the ERK pathway,<sup>18</sup> the JAK-STAT signaling pathway,<sup>19</sup> and the insuline-like growth factor 1 signaling pathway.<sup>20</sup>

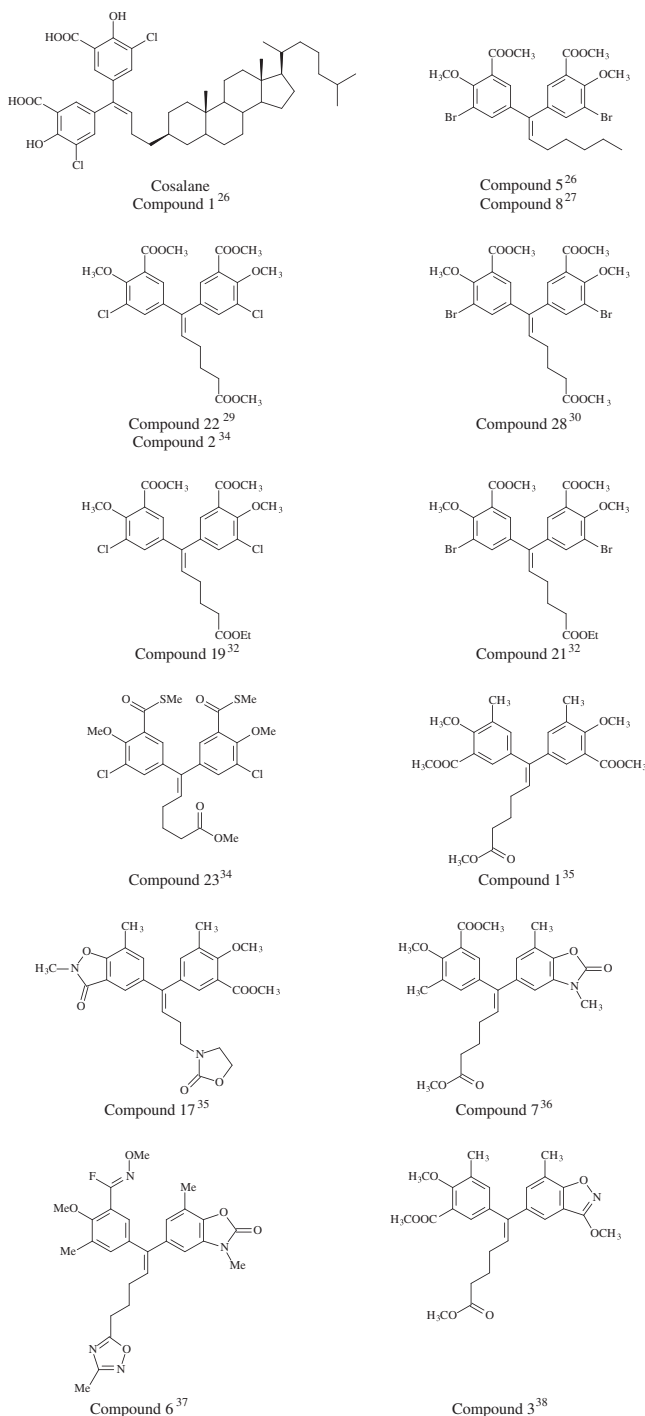
Given the current interest in new anti-influenza therapeutics, a battery of approximately 2,000 structurally diverse compounds were screened, and ATA emerged as one of the most promising NAIs.<sup>21</sup> Although ATA was originally assumed to inhibit influenza virus replication through the inhibition of the influenza viral polymerase (RdRp) activity,<sup>22,23</sup> Hung et al.<sup>21</sup> found that ATA did not inhibit the influenza polymerase, but, instead, interfered with the viral neuraminidase by competing with sialic acid (although ATA is structurally distinct from sialic acid) at the enzyme's substrate binding site. At least, this was suggested by molecular modeling experiments.<sup>21</sup>

Another virus and accompanying viral target, that has received increasing attention is the *S*-adenosylmethionine (SAM)-dependent methyltransferase (MTase)<sup>3</sup> associated with flaviviruses (i.e. dengue virus).<sup>4</sup> Here, again, ATA has been identified, as the result of a combined computational and experimental screening approach, as a potent flaviviral MTase inhibitor, even the most potent known to date, according to Milani et al.<sup>24</sup>

Thus, there is no shortage of potential therapeutic applications for ATA, but separately from its eventual therapeutic usefulness, what has to be resolved in each case, is whether the proposed molecular target, despite all the molecular docking and computation, really accounts for the observed antiviral activity, or, given the myriad of possible interaction sites of ATA, only represent(s) one of these many sites with which the compound could theoretically interact.

### ***B. ADAM: Non-Nucleoside HIV-1 Reverse Transcriptase Inhibitors that Should Go Beyond Just HIV-1***

Following up on the "ATA" lead (see supra in this series of stories), Mark Cushman and his collaborators then described cosalane (Fig. 2), which should conceptually be considered as an ATA derivative (missing one salicylic acid part) conjugated to cholestane: this compound acts primarily by the inhibition of viral (gp120-cellular) CD4 binding,<sup>25</sup> the cholestane part



**Figure 2.** Structures of alkenyl/diaryl methane (ADAM) non-nucleoside reverse transcriptase inhibitors (NNRTIs).

being responsible for anchoring the compound into the cell membrane. Why cholestane was, as such, not further developed for its anti-HIV properties, has remained unclear (to me), but, instead, it opened the way to the ADAM derivatives, as potential NNRTIs, thus joining the

wide array of NNRTIs (HEPT, TIBO, nevirapine, BHAP, TSAO,  $\alpha$ -APA, etc.) described at that time.<sup>26</sup>

The ADAM derivatives were from the very beginning recognized as genuine NNRTIs, as, for example, suggested for 3',3''-dibromo-4',4''-dimethoxy-5',5''-bis(methoxycarbonyl)-1,1-diphenyl-1-heptene (compound **8** in Cushman et al.<sup>27</sup>; or compound **5** in Cushman et al.<sup>26</sup>) (Fig. 2), although structurally little resemblance could be observed between the “classical” NNRTIs (HEPT, TIBO, ...) and the ADAMs, particularly with regard to the paradigmatic “butterfly” structure assigned to NNRTIs such as TBZs.<sup>4</sup>

Starting from the original ADAM derivatives,<sup>27</sup> new “ADAMs” were derived with optimized potency,<sup>28</sup> exhibiting an EC<sub>50</sub> of 0.013  $\mu$ M and a selectivity index of 2430 [i.e. methyl-3',3''-dichloro-4',4''-dimethoxy-5',5''-bis(methoxycarbonyl)-6,6-diphenylhexenoate].<sup>29</sup> This potency was even further increased to 0.0013  $\mu$ M (EC<sub>50</sub>) and the selectivity increased to 10,000, if the chlorines were replaced by bromines, as in methyl-3',3''-dibromo-4',4''-dimethoxy-5',5''-bis(methoxycarbonyl)-6,6-diphenyl-5-hexenoate (Fig. 2)<sup>30</sup> [solid-phase synthesis of the ADAMs was described by Xu et al.<sup>31</sup>].

Further improvement of anti-HIV-1 potency, especially against NNRTI resistance mutations (i.e. A98G) were noted with ADAM derivatives based on the ethyl (instead of methyl) 5-hexenoate, as in both the 3',3''-dichloro- and 3',3''-dibromo-derivatives (Fig. 2), so that the latter compounds were at the time (year 2001) considered as the most promising ADAM derivatives.<sup>32</sup> New methods were in the mean time devised for the preparation of new ADAM derivatives with nonidentical aromatic rings.<sup>33</sup>

The potential therapeutic usefulness of the ADAMs could be comprised by the metabolic instability that may be expected from their ester moieties that are likely to be hydrolyzed by nonspecific esterases present in human plasma.<sup>34</sup> Therefore, structural modifications (i.e. thioesters) introduced in the parent compound **2** (Fig. 2), as in compound **23** (Fig. 2), resulted in enhanced stability ( $t_{1/2}$  = 55 min), diminished cytotoxicity (CC<sub>50</sub> > 224  $\mu$ M), while maintaining reasonable potency (EC<sub>50</sub> = 1.8  $\mu$ M).<sup>34</sup>

Similarly, modifications of compound **1** (Fig. 2) (including introduction of a benzo[d]isoxazole as in compound **17**) (Fig. 2) enhanced stability ( $t_{1/2}$  = 22 min), diminished cytotoxicity (CC<sub>50</sub> = 16  $\mu$ M), while maintaining moderate antiviral potency (EC<sub>50</sub> = 2.7  $\mu$ M).<sup>35</sup> For the benzoxazolyl derivative (*E*)-5-[1-(3,7-dimethyl-2-oxo-2,3-dihydro-benzoxazol-5-yl)-5-methoxycarbonyl-pent-1-enyl]-2-methoxy-3-methylbenzoic acid methyl ester (compound **7**) (Fig. 2), EC<sub>50</sub> values of 30 and 90 nM against HIV-1 (RF and III<sub>B</sub> strains) were recorded.<sup>36</sup> For (*Z*)-5-((*E*)-1-(3,7-dimethyl-2-oxo-2,3-dihydrobenzo[d]oxazol-5-yl)-5-(3-methyl-1,2,4-oxadiazol-5-yl)pent-1-enyl)-*N*,2-dimethoxy-3-methylbenzimidoyl fluoride (compound **6**) this was accompanied by enhanced metabolic stability in rat plasma ( $t_{1/2}$  = 61 hr).<sup>37</sup> At present, the most promising compound within the ADAM series is compound **3**, (*Z*)-2-methoxy-5-[5-methoxycarbonyl-1-(3-methoxy-7-methylbenzo[d]isoxazol-5-yl)-pent-1-enyl]-3-methylbenzoic acid methyl ester (Fig. 2) with EC<sub>50</sub> values of 40 nM (HIV-1 RF) and 20 nM (HIV-1 III<sub>B</sub>).<sup>38</sup>

The primary objective has been to optimize the ADAMs by replacing the metabolically unstable methyl ester moieties with stable isosteres (i.e. benzo[d]isoxazole), while maintaining or enhancing the antiviral potency of the ADAMs as potential NNRTI-type drugs. However, other potential applications of the ADAMs should not be ignored. They may not have much potential as PDE4 inhibitors [phosphodiesterase 4 (PDE4) belongs to the phosphodiesterase family of hydrolases responsible for regulating cellular cAMP levels, PDE4 playing a role in HIV infections, and PDE4B2 being specifically involved in inflammation], as they have only weak, if any, activity against PDE4B2.<sup>39</sup> However, two distantly related ADAM derivatives, i.e. compounds **15** and **16**,<sup>40</sup> like colchicine, proved active as tubulin polymerization inhibitors (IC<sub>50</sub> of 3.7 and 2.8  $\mu$ M, respectively). This makes them worth pursuing as potential anticancer drug candidates.

### ***C. HIV INIs, the Fifth Class of HIV Inhibitors, Following N(t)RTIs, NNRTIs, PIs, and CRIs***

Exactly 25 years after the discovery of HIV as the causative agent of AIDS by Françoise Barré-Sinoussi [Nobel Prize in Medicine or Physiology (2008)] and her co-workers<sup>41</sup> and by Mikulas Popovic and his co-workers,<sup>42</sup> 25 compounds have been officially approved for the treatment of AIDS.<sup>43</sup> The most recent compounds to complete the anti-HIV drug armamentarium are the INIs,<sup>44,45</sup> presently only one (raltegravir)<sup>45</sup> but soon to be expected two (raltegravir and elvitegravir).<sup>44</sup> Raltegravir plus optimized background therapy provided better viral suppression than optimized background therapy alone for at least 48 weeks,<sup>46</sup> and raltegravir has been considered a valuable addition to the current armamentarium for the treatment of patients infected with multi-drug-resistant HIV-1.<sup>47</sup>

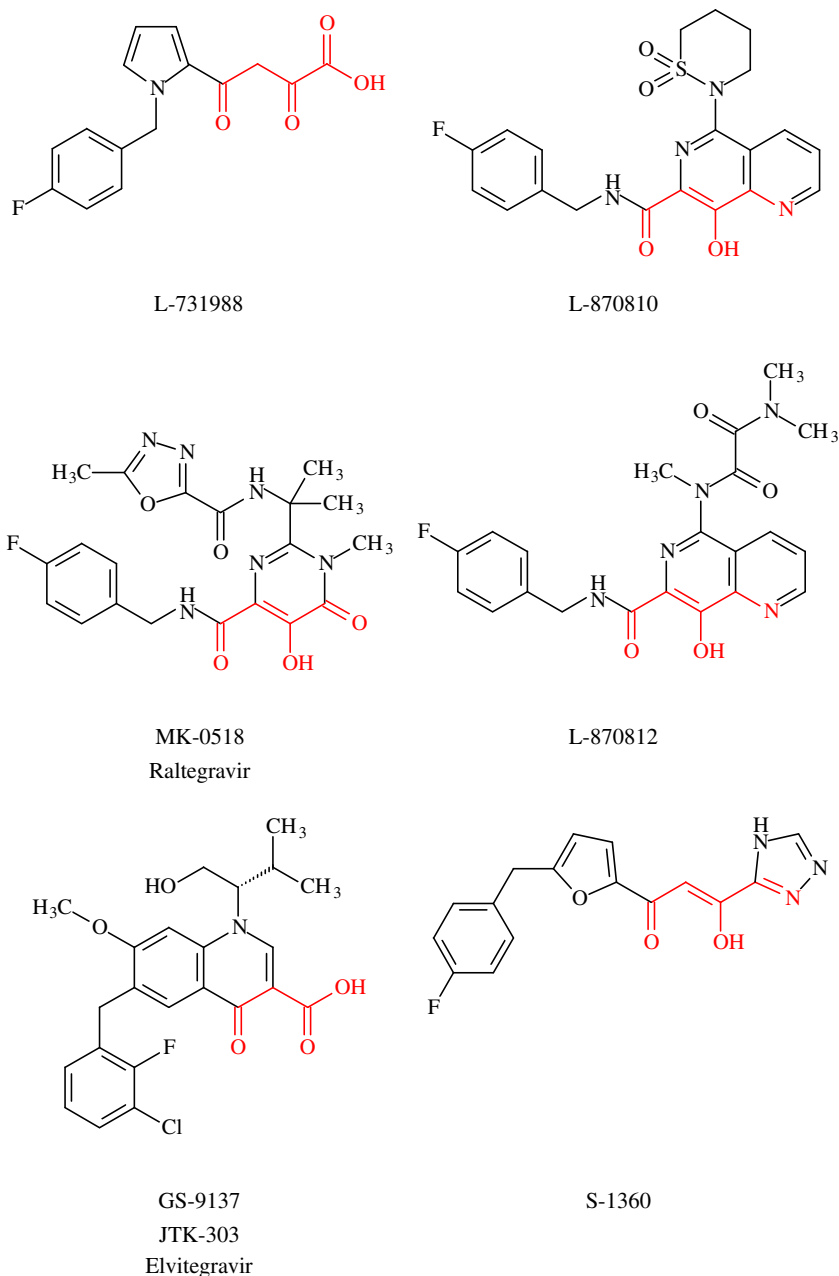
The era of the INIs started with Daria Hazuda's paper in *Science*<sup>48</sup> on the inhibitory effects of "diketo acids", i.e. L-731988, on integration and replication of HIV in cell culture, providing the first proof of concept for HIV-1 integrase (specifically, strand transfer) inhibitors to act as antiviral agents. While first demonstrated for L-731988,<sup>48</sup> this proof of principle has been extended to various related compounds, i.e. L-870810 (a naphthyridine carboxamide)<sup>49</sup> and L-870812<sup>50</sup> which proved efficacious *in vivo* against retroviral (simian-human immunodeficiency virus) replication *in vivo* (rhesus macaques).

Hence, it did not come as a surprise that the compound MK-0518 [4-[*N*-(4-fluorobenzyl)carbamoyl]-1-methyl-2-[1-methyl-1-(5-methyl-1,3,4-oxadiazol-2-ylcarboxamido)ethyl]-6-oxo-1,6-dihydropyrimidin-5-olate (raltegravir)] (Fig. 3) selected as the clinical candidate for further development<sup>51</sup> finally became licensed for clinical use as the first INI ever to be approved for the treatment of HIV infection. Meanwhile, L-870812 has been advocated as a microbicide to prevent cell-free and cell-associated HIV infection.<sup>52</sup> Should INIs ever be used topically as microbicides, they preferentially be combined, as is the rule for their systemic use as well, with other classes of HIV inhibitors, including reverse transcriptase inhibitors and viral entry inhibitors.

The runner-up INI to be licensed for clinical use is elvitegravir (GS-9137)<sup>53</sup> (Fig. 3). While raltegravir is currently dosed at 400 mg twice daily, elvitegravir is dosed once daily at 150 mg [currently in combination with 100 mg of ritonavir as its booster; on the horizon, however, is the combination of elvitegravir with Gilead's own booster (pharmacoenhancer) GS 9350 and Truvada (tenofovir disoproxil fumarate and emtricitabine), which together will make a one pill once-daily treatment (tentatively called "QUAD" as it will consist of four compounds)].

Structurally, elvitegravir (Fig. 3) is built upon the quinolone 3-carboxylic acid scaffold [this pharmacophore has also served as the starting point for the design of a second generation of HIV-1 INIs].<sup>54</sup> Quinolones, such as the fluoroquinolone K-12<sup>55</sup> and the 6-aminoquinolone (WM-5),<sup>56</sup> have been known as Tat-dependent transcription inhibitors. It was, therefore, interesting to ascertain whether elvitegravir, being both a "diketo acid" and a "quinolone" should behave as an INI or transcription inhibitor. Unlike WM-5, but like L-870810, elvitegravir behaved as a genuine INI.<sup>57</sup>

Elvitegravir is a more potent INI than raltegravir,<sup>58</sup> but its mechanism of action, i.e. inhibition of the strand transfer function of integrase must be similar, as both drugs exhibited a parallel resistance profile, the Q148 K and T66I mutations conferring the highest resistance to both drugs.<sup>58</sup> In addition to the Q148 K and T66I mutations, a few other mutations, i.e. E92Q, L74 M and S230N have been shown to confer resistance to elvitegravir and, where examined, other INIs (i.e. L-870810 and raltegravir) as well.<sup>59-61</sup> While, on the one hand, the clinical relevance of these mutations remains to be demonstrated, i.e. with regard to possible reduced fitness of the viral mutants, they should, on the other hand, help in gaining further insight in the mode of action of INIs and, eventually facilitate the design of new INIs.



**Figure 3.** HIV Integrase inhibitors (INIs) (in red: metal ( $Mg^{++}$  or  $Mn^{++}$ ) binding  $\beta$ -diketo acid or the like portion of the molecules).

#### ***D. LEDGF as Cellular Cofactor for HIV Integrase: An Attractive Target for anti-HIV Drug Design?***

The LEDGF “story” started in 2000, when Peter Cherepanov and his coworkers<sup>62</sup> demonstrated that using a synthetic gene, efficient expression of HIV-1 integrase was achieved in human cells. Cherepanov et al. then found in the nuclei of human cells stably expressing the HIV integrase that it formed stable tetramers associated with the LEDGF/p75, a protein



which had been implicated in the regulation of gene expression and cellular stress response.<sup>63</sup> Moreover, LEDGF was found to robustly enhance the strand transfer activity of HIV integrase *in vitro*, and Cherepanov et al.<sup>63</sup> concluded that LEDGF may constitute a novel target for anti-HIV drug therapy.

In subsequent studies it was further shown that LEDGF/p75 accounts for the karyophilic, i.e. nuclear targeting, and chromosomal targeting of HIV-1 integrase,<sup>64</sup> that LEDGF/p75 is essential for HIV DNA integration into chromosomal DNA,<sup>65</sup> and LEDGF/p75 is the first example of a cellular protein controlling the location of HIV integration in human cells.<sup>66</sup> In fact, LEDGF/p75 is able to interact with different lentiviral integrases, including those from bovine immunodeficiency virus, maedi-visna virus, and equine infectious anemia virus, but nonlentiviral integrases such as those from beta-, gamma-, deltaretroviruses or spumaviruses do not possess detectable affinity for LEDGF.<sup>67</sup>

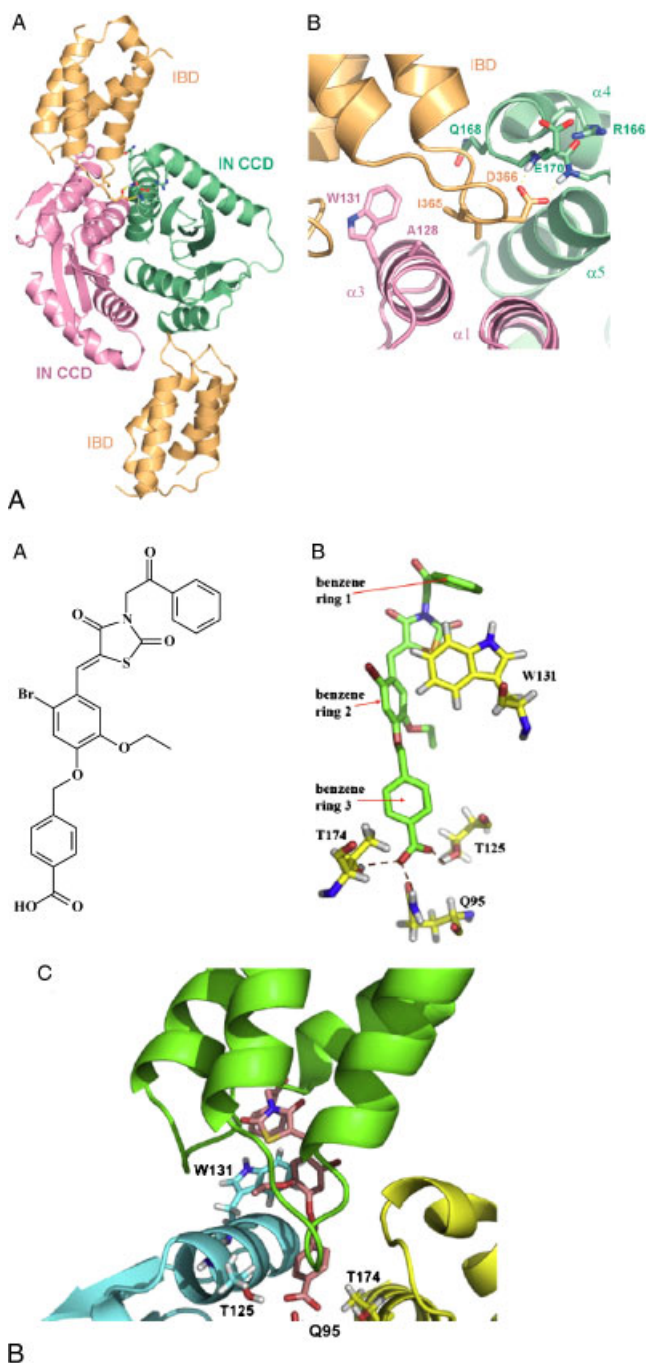
Poeschla thus concluded that LEDGF/p75 may act as a chromatin docking factor or receptor for lentiviral preintegration complexes; LEDGF/p75 tethers HIV integrase to chromatin, protects it from degradation, and strongly influences the genome-wide pattern of HIV integration.<sup>68</sup> Given the fact that LEDGF/p75 is a ubiquitous nuclear protein, tightly associated with chromatin throughout the cell cycle, and although an important but not strictly essential cofactor of lentiviral DNA integration,<sup>69</sup> could such ubiquitous protein (that is not strictly essential for retroviral DNA integration) serve as a target for the development of a novel class of antiretroviral drugs?

Not the LEDGF *per se*, but its interaction with the integrase (thus the protein–protein interaction (PPI)), which is further illustrated in Figure 4A<sup>70</sup> may serve as a potential target for drug discovery, but, therefore, several issues should be addressed<sup>71</sup>: (i) the target has to be validated as important for HIV-1 replication; (ii) inhibition of the specific PPI should not be associated with toxicity; (iii) structural information on the PPI should be available; and (iv) identification of genuine inhibitors should provide ultimate proof-of-concept.<sup>71</sup> The pursuit of LEDGF as a potential target for anti-HIV drugs may seem as a Sisyphean task as we not only have to take into account the PPI interaction between the integrase’s CCD and LEDGF’s IBD (Fig. 4A), but also the role this PPI has to fulfill, that is the integration of the proviral DNA into the target chromosomal DNA, which makes it, *stricto sensu*, a “menage à quatre.”

These tantalizing perspectives have not deterred several investigators to launch a search for small molecule PPI inhibitors (already nicknamed as SMPPIIs)<sup>70</sup> or to screen for small-molecule inhibitors of the HIV integrase-LEDGF/p75 interaction by a luminescent proximity assay.<sup>72</sup> A first (apparent) success has already been reported: that of D77, a benzoic acid derivative (Fig. 4B) that would function as a novel HIV-1 inhibitor targeting the interaction between the integrase’s CCD and the LEDGF’s IBD<sup>73</sup> [D77, as shown in Figure 4B, would (principally) interact with the CDD domain of the integrase]. This approach opens a wealth of possibilities, i.e. structure–activity relationship (SAR) studies, pharmacokinetics and toxicity, and, particularly, confirmatory studies to validate the presumed target of action and provide the ultimate proof-of-concept.

### ***E. While Waiting for the Avian H5N1 Influenza Pandemic, What is the Position of NAIs (such as Oseltamivir)?***

As put forward by Frederick Hayden,<sup>74</sup> new antiviral agents for the treatment of influenza are urgently needed, one of the limitations of current drugs (i.e. oseltamivir) being the emergence of resistance among influenza A (H1N1) strains and the fear it may also emerge among avian influenza A (H5N1) (“bird flu”) strains. Highly pathogenic avian influenza A (H5N1) viruses are entrenched among poultry in parts of Asia, Africa, and the Middle



**Figure 4.** (A) Interface between the HIV-1 integrase catalytic core domain (CCD) and the integrase-binding domain (IBD) of LEDGF/p75 using Protein Data Bank crystal structure file 2BJ4 (panel A). Panel B shows a close-up view of the interface. The integrase CCD monomers are colored purple and green, and the IBD subunit is orange. Integrase residues A128 and W131 are part of a hydrophobic patch, which accommodates the side chains of the LEDGF/p75 residues I365, F406 and V408. Data taken from Busschots et al.<sup>70</sup> (B) Molecular docking of D77 into the HIV-1 integrase catalytic core domain (CCD). Panel A. Chemical structure of D77, the benzoic acid derivative 4-[(5-bromo-4-[[2,4-dioxo-3-(2-oxo-2-phenylethyl)-1,3-thiazolidin-5-ylidene]methyl]-2-ethoxyphenoxy)methyl]benzoic acid. Panel B. View on D77 as docked into CCD (amino acid residues W131, T125, Q95, and T174). Panel C. Closer view on D77 as docked into the CCD/IBD complex (showing amino acid residues W131, T125, Q95, and T174 of the CCD domain). Data taken from Du et al.<sup>73</sup>

East,<sup>75</sup> and human infection with these viruses has been noted in many of countries including Vietnam,<sup>76</sup> Indonesia (three clusters!),<sup>77</sup> and Eastern Turkey<sup>78</sup> [Indonesia has had the most human cases of highly pathogenic avian influenza A (H5N1) and one of the highest case-fatality rates worldwide<sup>79</sup>].

It is re-assuring that person-to-person transmission of avian influenza A (H5N1) is rather limited,<sup>80</sup> and this is probably linked, as I explained before,<sup>81</sup> to the differences in receptor specificity: avian receptors prefer a  $\alpha(2-3)$  linkage, whereas humans prefer the  $\alpha(2-6)$  linkage so that productive infections may have difficulty in jumping from one host (avian) to the other (human).

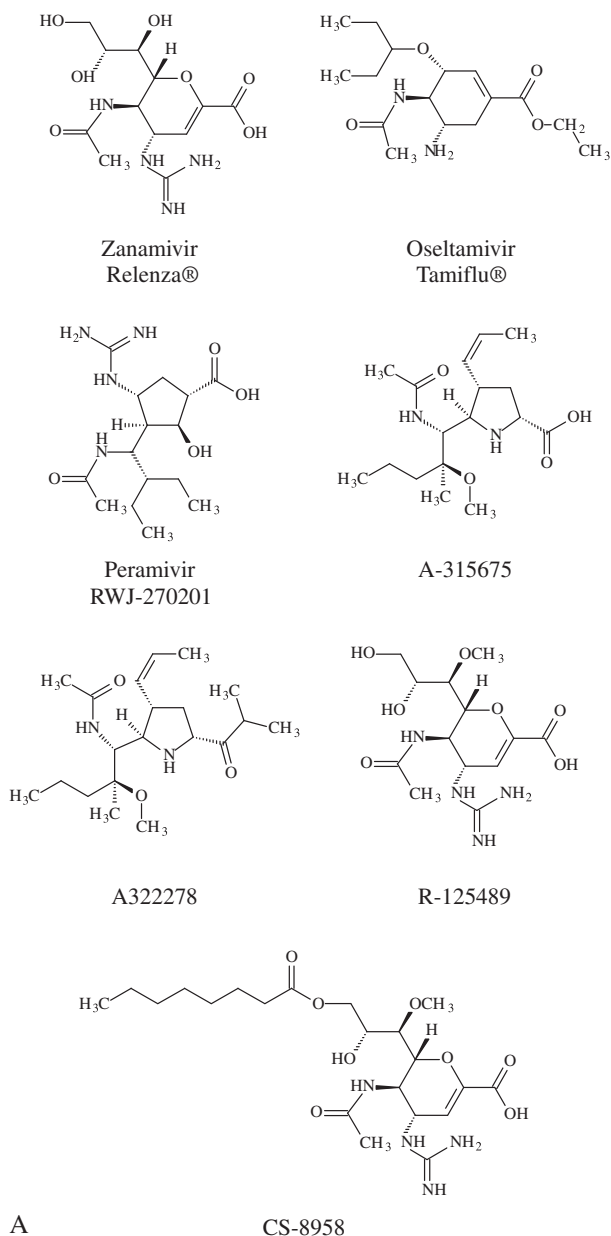
Oseltamivir has been considered as the golden grail and stockpiled for use should the need prevail: first, to treat infected individuals (to moderate disease severity); second, to protect family members of an index case (to interrupt transmission); and third, to all the people in an area surrounding an index case (ring prophylaxis).<sup>82</sup> As it can be administered orally, oseltamivir (Fig. 5A) is the drug of choice for the therapy of both seasonal and avian influenza<sup>83</sup> [with zanamivir (Fig. 5A) (that has to be inhaled via a rather sophisticated device) ranking second].

While waiting for a still hypothetical avian influenza A H5N1 pandemic, the world was recently confronted with a new influenza (first called “pig,” then “Mexican”) H1N1 variant, which spread rapidly over the five Continents and thus, in epidemiological terms, gave rise to a pandemic. Although highly contagious, the “Mexican” H1N1 influenza variant has so far not proven more pathogenic than other seasonal influenza A H1N1 variants (although it started its journey in the off-season). As all the other H1N1 variants, the “Mexican” H1N1 virus should normally be sensitive to the NAIs oseltamivir and zanamivir.

However, a worrying phenomenon that has been observed within the past few years is widespread oseltamivir resistance in influenza A (H1N1) in the United States,<sup>84,85</sup> Europe,<sup>86</sup> Japan,<sup>87</sup> Norway,<sup>88</sup> and South Africa.<sup>89</sup> It has been suggested that the H274Y mutation conferring resistance to oseltamivir leaves the influenza A (H5N1) virus severely compromised.<sup>90</sup> However, this oseltamivir-resistant virus can be pathogenic, and fatal, in an immunocompromised patient.<sup>91</sup> On the contrary, oseltamivir allowed complete recovery of a bone-marrow transplant recipient from an influenza A (H1N1) virus infection that did not respond to zanamivir.<sup>92</sup>

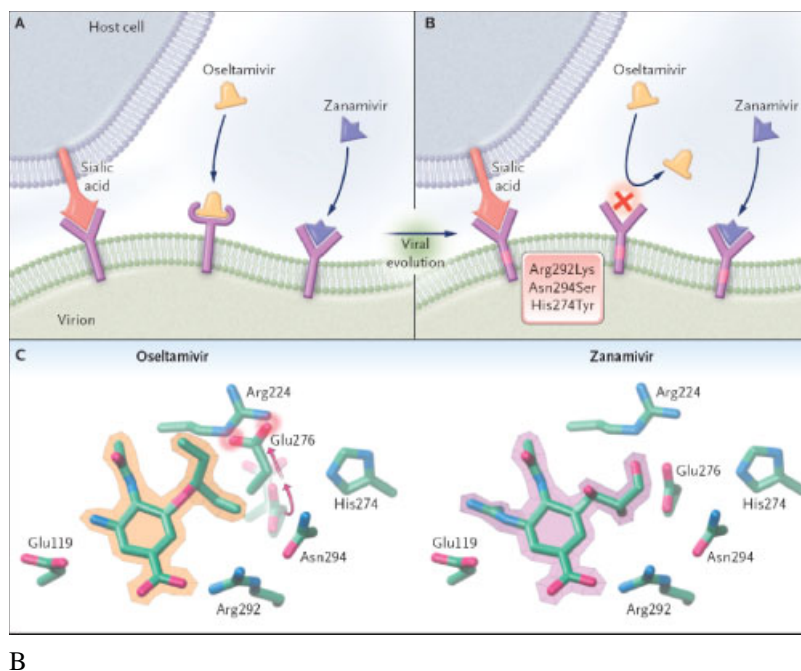
Influenza A virus strains resistant to NAIs (zanamivir, oseltamivir) would circulate at a stable and low level (1%) since these compounds were introduced in clinical practice.<sup>93</sup> Influenza B viruses with reduced sensitivity to NAIs may also arise, but not as frequently as for the influenza A viruses.<sup>94</sup> As originally pointed out by Peter Palese, influenza A may be effectively transmitted among guinea pigs (Personal communication, 8th International Symposium on NeuroVirology, San Diego, California, USA, 30 October-2, November 2007), and this was effectively demonstrated by Bouvier et al. for oseltamivir-resistant influenza A viruses.<sup>95</sup> This opens new avenues not only for the study of the transmission of influenza A virus infections, but also for the chemotherapy and chemoprophylaxis of these infections.

Although resistance to oseltamivir has also been noted with influenza A (H3N2) virus [i.e. due to a deletion of four amino acids ( $\Delta 245-248$ ) in the neuraminidase<sup>96</sup>], resistance to oseltamivir would emerge at a higher rate in influenza A (H1N1) virus than in influenza A (H3N2) virus or influenza B virus in children.<sup>97</sup> For influenza A (H5N1) virus there might be a significant natural variation in sensitivity to oseltamivir,<sup>98</sup> although reduced sensitivity of influenza A (H5N1) to oseltamivir has been noted at several occasions<sup>99,100</sup>; and NAI-resistant H5N1 influenza viruses (i.e. those carrying the H274Y mutation) may retain the high pathogenicity of the wild-type virus both in vitro and in vivo (in mice).<sup>101</sup> As learned from ferrets, it should be advisable to increase the dose of oseltamivir to treat the highly pathogenic avian influenza H5N1 infections in humans.<sup>102,103</sup>



**Figure 5.** (A) Structural formulae of zanamivir, oseltamivir, peramivir, A-322278, R-125489 and CS-8958. (B) Mechanism of development of resistance to oseltamivir. Binding of oseltamivir, but not zanamivir, to the neuraminidase active site requires a shape change that creates a pocket (Panel A), and mutations that prevent formation of this pocket may prevent binding of oseltamivir but permit binding of zanamivir (Panel B). Crystal structure analysis shows that binding of oseltamivir requires a rotation of the Glu276 residue away from the hydrophobic pentyloxy group of oseltamivir, which then makes hydrophobic contact with a methylene group on Glu276 (Panel C). However, binding of either sialic acid (the natural substrate) or zanamivir does not require a change in the position of Glu276. The His274Tyr substitution in the neuraminidase active site pushes the Glu276 further into the binding site and disrupts the hydrophobic pocket that is required only for oseltamivir binding. According to Moscona.<sup>117</sup>

Novel neuraminidase mutations D197E<sup>104</sup> and D198N<sup>105</sup> have been described that show resistance to both oseltamivir and zanamivir, but the clinical relevance of these resistance mutations remains to be further elucidated. Taking together all the resistance mutations



B

**Figure 5.** Continued.

collected for oseltamivir, Skehel and colleagues considered it prudent for pandemic stockpiles of oseltamivir to be augmented by additional antiviral drugs including zanamivir.<sup>106</sup>

To stockpile oseltamivir (tamiflu) as a precautionary measure against an influenza pandemic is both risky and costly. It is recomforting, therefore, that intensive efforts have been made to improve the synthesis of oseltamivir [the starting material is shikimic acid normally obtained from the Chinese star anise<sup>107</sup>] and these efforts have indeed allowed a more practical and, at the same time, high-yielding synthesis of oseltamivir.<sup>108–111</sup> An environmental issue, linked to its chemical stability, is that oseltamivir would not be easily removed or degraded in normal sewage water.<sup>112</sup> This not only points to the stability of the compound, but also its propensity to contribute to far-reaching anti-influenza virus drug resistance.

Are new NAIs forthcoming that may supersede the classical NAIs oseltamivir and zanamivir? A-322278 (Fig. 5A) certainly seems promising as it is active against the oseltamivir-resistant H274Y A (H1N1) influenza virus mutant in mice.<sup>113</sup> A-322278, the 2-methylpropanoyl prodrug of A-315675, would have comparable activity to oseltamivir in immunocompetent and -compromised murine models of influenza virus infection.<sup>114</sup> A-315675 has potent inhibitory activity against oseltamivir-resistant influenza viruses (N1 and N2 subtypes).<sup>115</sup> In addition to zanamivir and oseltamivir, peramivir, A-322278 and A-315675, also R-125489 and CS-8958 (Fig. 5A) show high promise as anti-influenza virus agents, the latter (CS-8958) demonstrating long-acting anti-influenza virus activity.<sup>116</sup>

The mechanism of development of resistance to oseltamivir (Fig. 5B) as proposed by Moscona<sup>117</sup> is paradigmatic, but is it correct? NAI inhibitors (whether zanamivir, oseltamivir or peramivir) are still trying to find their best way of application, prophylactic/therapeutic,<sup>118</sup> or would injectable peramivir offer the ideal, if not final, solution?<sup>119</sup> Drug-resistant virus will remain an important factor for NAIs for many years to come,<sup>120</sup> but the design, synthesis, and structure–activity relationship (SAR) studies will be continued to trying to cope with this challenge.<sup>121</sup>

Of primordial importance, if the avian H5N1 influenza virus, or any H1N1 influenza virus such as the “Mexican” variant, becomes a pandemic threat in humans, is that we are

“prepared” [as we are already supposed to be for many years]; and the combination of oseltamivir with, for example, amantadine,<sup>122</sup> or ribavirin,<sup>123</sup> or double combinations of amantadine, oseltamivir and ribavirin<sup>124</sup> may rank among the various measures that could be taken to curb influenza virus infections.

### ***F. RSV: Still in Search of Specific Antiviral Drugs***

In 1996, I reviewed the perspectives for the chemotherapy of RSV infections.<sup>125</sup> At the time, ribavirin (given as an aerosol at 20 mg/mL) was the only antiviral agent approved for the treatment of RSV infection. Several new possible drug candidates were considered.<sup>125</sup> What has happened with ribavirin and these drug candidates, and how does the chemotherapy (and/or–prophylaxis) of RSV infections look like nowadays?

Human RSV is the major cause of upper and lower respiratory tract infections in the pediatric population. These infections are particularly problematic in infants that are born prematurely or with congenital heart disease or chronic lung disease (CLD) or are otherwise immune compromised (ref.<sup>126</sup>; and references therein). Elderly and immunocompromised adults are also at increased risk for developing complications or even death associated with RSV infection (for a review, see Falsey & Walsh<sup>127</sup>).

Ribavirin still is the only antiviral agent approved for the treatment of RSV infection, but due to efficacy and toxicity issues, it has only limited utility.<sup>128</sup> Inhaled (nebulized) ribavirin has remained the standard therapy for the treatment of RSV infections, despite it is costly and cumbersome. Pelaez et al.<sup>129</sup> recently pointed to the efficacy of oral ribavirin in the treatment of lower respiratory tract infection after lung transplantation. They argued that further studies should be conducted to compare the long-term efficacy of oral ribavirin vs. nebulized therapy for RSV.

After more than 40 years of research, there is no approved vaccine, and the only prophylactic therapies available are RSV-IVIG, a polyclonal RSV immunoglobulin,<sup>130</sup> and palivizumab (Synagis<sup>®</sup>), a humanized monoclonal antibody targeting the RSV fusion protein.<sup>131</sup> While effective, this treatment is only administered to high-risk pediatric patients. There is a clear need for new anti-RSV therapeutics, with improved efficacy and safety for broader applications (for a review, see Meanwell & Krystal<sup>132</sup>).

RSV encodes three surface glycoproteins, the fusion protein F, the attachment glycoprotein G, and the small hydrophobic protein SH. Both the F and G glycoproteins are required for efficient infectivity *in vivo*, but the F protein alone is sufficient for virus binding and entry into cells *in vitro*.<sup>133</sup>

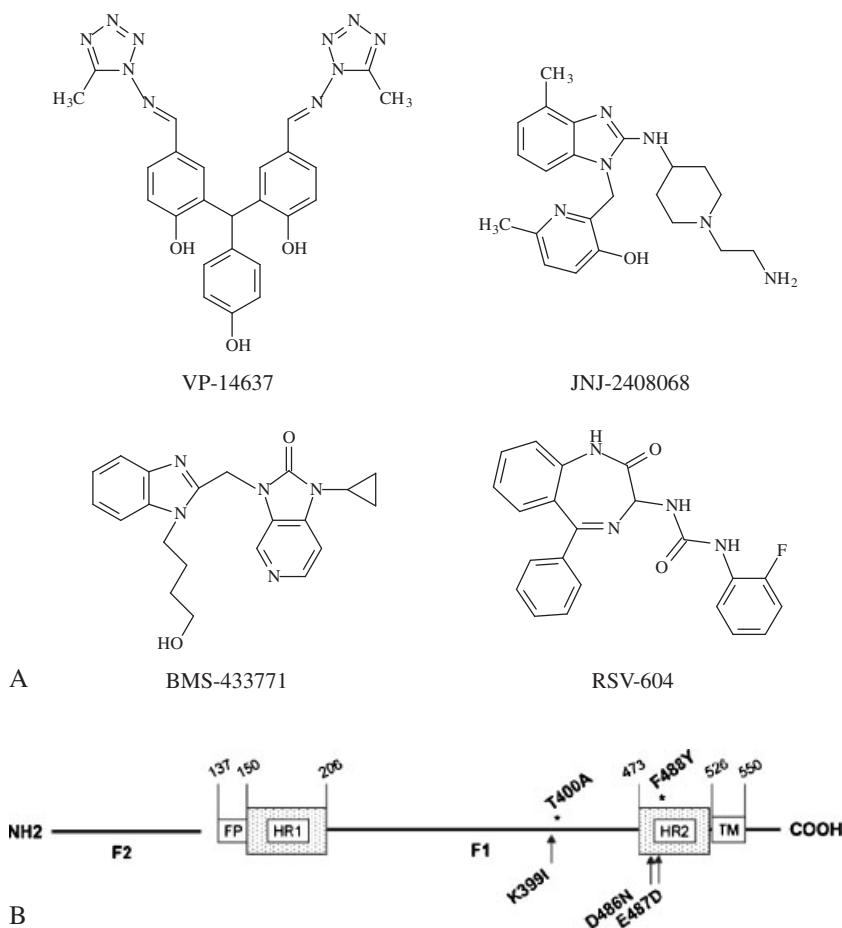
RSV is a major cause of hospitalization in preterm infants and infants with CLD.<sup>134</sup> Prophylaxis with palivizumab significantly reduced the incidence of RSV-related hospitalization relative to placebo and is generally well tolerated in high-risk infants aged <2 years, including those with prematurity and bronchopulmonary dysplasia/CLD or hemodynamically significant congenital heart disease, which are risk factors for early or serious RSV infection. Palivizumab has been approved for use in these patients.<sup>135</sup>

Children who experience RSV lower respiratory tract infections early in life have high rates of subsequent recurrent wheezing. Palivizumab, by ameliorating or preventing this lower respiratory tract infection in preterm infants may reduce subsequent recurrent wheezing.<sup>136</sup> Wu et al.<sup>137</sup> described the selection of ultra-potent anti-RSV antibodies for preventing RSV infection. They applied an iterative mutagenesis approach, and were able to identify palivizumab Fab variants with up to 1500-fold improvement and palivizumab IgG variants with up to 44-fold improvement in the ability to neutralize RSV. Recently, a new, ultra-potent antibody, motavizumab, has been developed for the prevention of RSV infections.<sup>138</sup> Motavizumab binds to RSV F protein 70-fold better than palivizumab, and it reduced pulmonary RSV titers to up to 100-fold lower levels than did palivizumab.

Motavizumab is currently being evaluated in pivotal clinical trials for RSV prophylaxis, and for the reduction of RSV-related asthma.

There is an experimental concern, originated from preclinical studies in cotton rats, which RSV escape mutants may emerge in immunosuppressed patients. Should palivizumab arise in humans, palivizumab may obviously be ineffective.<sup>139</sup> Zhao and Sullender<sup>140</sup> evaluated the potential for palivizumab-resistant RSV mutants to arise *in vivo*. Cotton rats were immunosuppressed with cyclophosphamide. Three of the five animals had mixed populations of lung virus, and over 50% of the clones from the three animals revealed F gene mutations associated with resistance to palivizumab. Thus, prolonged pulmonary replication of RSV in the presence of palivizumab was followed by the appearance of viruses resistant to palivizumab.

Using an RNA interference (RNAi) approach, inhibition of both RSV and parainfluenza virus can be achieved in the mouse by intranasally administered short interfering RNAs (siRNAs),<sup>141</sup> and a similar success has been obtained with intranasal siRNA nanoparticles targeting the viral NS1 gene (siNS1).<sup>142</sup> Initial clinical studies on safety, tolerability, and pharmacokinetics favor the further pursuit of siRNAs (i.e. ALN-RSV01) for RSV in humans.<sup>143</sup> Similarly



**Figure 6.** (A) Structural formulae of RSV inhibitors. (B) RSV inhibitors interacting with the F protein. Location of the resistance mutations selected in the presence of VP-14637 and JNJ-2408068. Schematic view of the RSV F polyprotein indicating salient features: F1 and F2 subunits; FP, fusion peptide; HR1, heptad repeat 1; HR2, heptad repeat 2; TM, transmembrane domain. The numbers represent amino acid designations, the asterisks above the line indicate the resistance mutations selected by VP-14637, and the arrows below the line indicate the resistance mutations selected by JNJ-2408068. According to Douglas et al.<sup>148</sup>

to siRNAs, RSV-targeted deoxyribozymes (DNA zymes) potentially present a therapeutic (or prophylactic) approach for RSV diseases. Their activity is based on the ability to bind and cleave complementary RNA sequences, thus inhibiting protein expression. D71133 is an example of such deoxyribozyme that targets the conserved genomic RNA sequence of the RSV nucleocapsid protein: it has been shown to block RSV infection both *in vitro*<sup>144</sup> and *in vivo*.<sup>145</sup>

Sidwell and Barnard<sup>146</sup> recently reviewed the prospects of the principal drug candidates for the control of RSV infections. Prominent among the cited compounds were VP-14637, JNJ-2408068, BMS-433771, and A-60444 (the latter compound became later known as RSV604) (Fig. 6A). Two RSV fusion inhibitors, VP-14637 (Fig. 6A) and JNJ-2408068 (Fig. 6A), have been identified,<sup>147,148</sup> which are both targeted at the F1 glycoprotein [as could be deduced from the resistance mutations selected by VP-14637 and JNJ-2408068 (Fig. 6B)].

VP-14637 (Fig. 6A) was first reported by D.C. Pevear (ViroPharma) [as mentioned by Wyde et al.<sup>149</sup>] before it was further evaluated by Cihlar and his colleagues<sup>148,150</sup> and found to be targeted at the F protein of RSV. The efficacy of VP-14637 against RSV was demonstrated *in vitro*, and *in vivo* in cotton rats following delivery by small droplet aerosol.<sup>151</sup> According to Sidwell and Barnard,<sup>146</sup> the compound was in phase I trials before a decision not to develop it further, in part due to developmental costs.

JNJ-2408068, originally named R170591 and for the first time reported by Andries and his colleagues., has a structure (2-[[2-[[1-(2-aminoethyl)-4-piperidinyl]amino]-4-methyl-1*H*-benzimidazol-1-yl]-6-methyl-3-pyridinonol) (Fig. 6A), which is fundamentally different from that of VP-14637, but nevertheless targeted at the F protein of HSV<sup>148</sup> (Fig. 6B). JNJ-2408068 is highly potent (EC<sub>50</sub>: 0.16 nM) but according to Douglas,<sup>150</sup> it has limited oral bioavailability. Like VP-14637, JNJ2408068 was found to protect cotton rats from experimental RSV infection.<sup>151</sup>

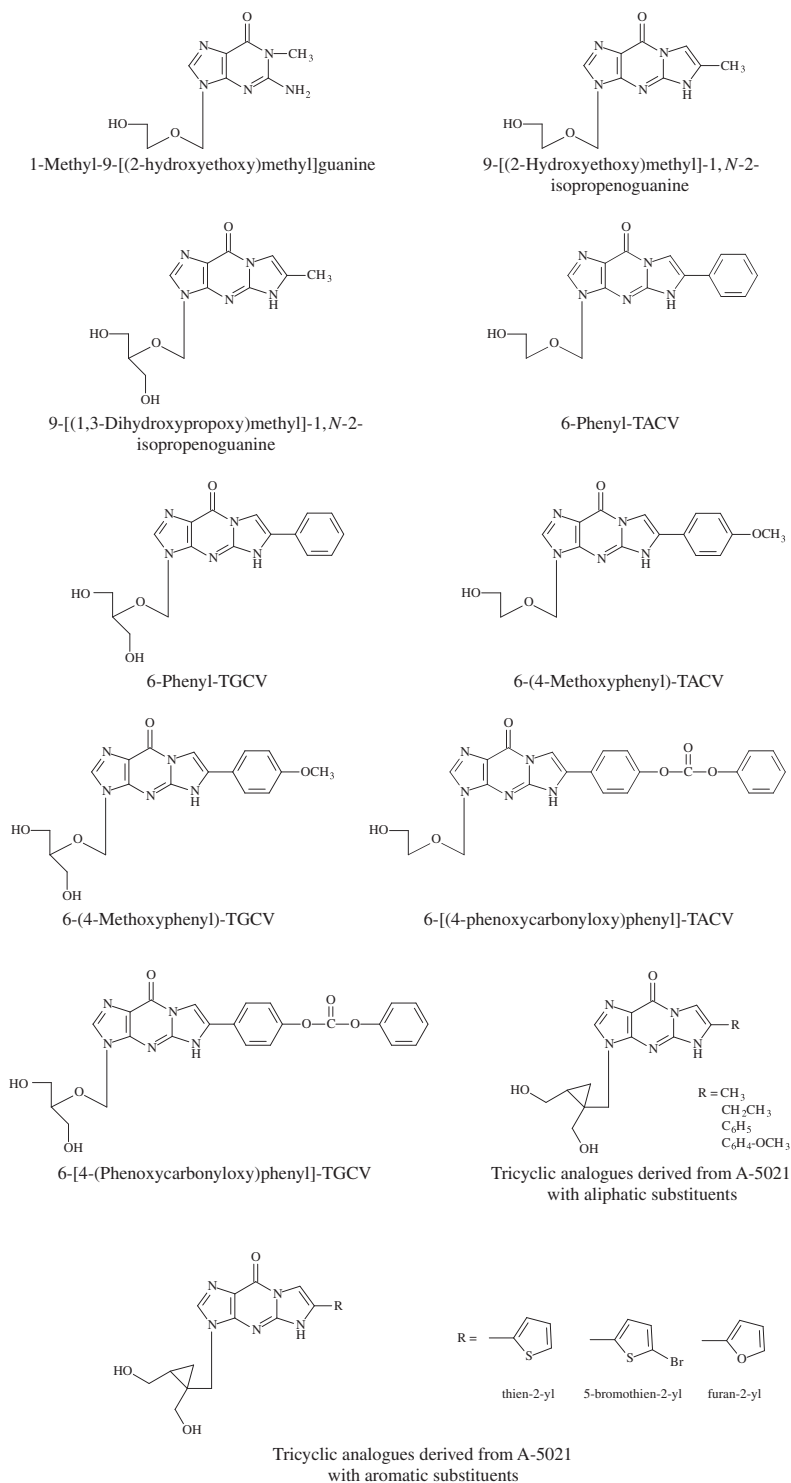
BMS-433771 [1-cyclopropyl-1,3-dihydro-3-[[1-(4-hydroxybutyl)-1*H*-benzimidazol-2-yl]-methyl]-2*H*-imidazo[4,5-*c*]pyridin-2-one] is another example of anti-RSV agents targeted at the F protein.<sup>152</sup> [To confirm that the mechanism of action was through inhibition of fusion, the K394R virus mutant was selected for resistance to BMS-433771 *in vitro*, and found refractory to BMS-433771 *in vivo*.] It proved orally efficacious against RSV infection in Balb/C mice and cotton rats.<sup>153,154</sup> Numerous structural variants of BMS-433771 have been synthesized<sup>155–160</sup> but, most likely, BMS-433771 still excelled in potency and oral bioavailability.

Powell and his colleagues have recently identified a new class of RSV inhibitors, namely that of 1,4-benzodiazepines,<sup>161</sup> which eventually led to the identification of RSV-604 (Fig. 6A) as the clinical candidate.<sup>162</sup> RSV-604 can be considered as a truly novel inhibitor of RSV replication in that, being a benzodiazinylurea derivative, it is targeted at the nucleocapsid, and, unlike all the F protein inhibitors (see *supra*) active against RSV after it has infected the cells.<sup>163</sup> In the conclusion of the latter paper, RSV-604 was hailed as the “most promising candidate to date for the treatment of RSV disease in humans.” In the wake of data on *in vivo* efficacy and oral bioavailability, this sounds as a daring statement.

### ***G. Fluorescent Tricyclic Analogues of Acyclovir and Ganciclovir Based on 1,*N*-2-ethenoguanine***

Following up on the work of Beauchamp et al. on a series of acyclovir derivatives with modifications in the heterocyclic base (which did not yield significantly active compounds),<sup>164</sup> we (Bozenna Golankiewicz, Jerzy Boryski, and I) started a new program around new acyclovir derivatives with one or more nitrogen centers blocked by methylation or incorporated into an additional ring structure. Marked activity against herpes simplex virus (HSV-1 and HSV-2), comparable to that of acyclovir, was obtained with 1-methyl-9-[(2-hydroxyethoxy)methyl]guanine and 9-[(2-hydroxyethoxy)methyl]-1,*N*-2-isopropeno-guanine (Fig. 7),





**Figure 7.** Tricyclic analogues of acyclovir, ganciclovir and A-5021, based on 1,N-2-ethenoguanine.

compounds **2** and **5**, respectively, in the original publication of Boryski et al.<sup>165</sup> While N-1 methylation allowed the antiviral activity of acyclovir to be preserved, it was virtually abolished following N-3 methylation.<sup>166</sup>

The earliest report on 1,*N*-2-etheno derivatives of guanine has been that of Sattangi et al.<sup>167</sup> on 1,*N*-2-ethenoguanosine. 1,*N*-2-ethenoguanine was not only successfully implemented in acyclovir but also in ganciclovir. The resulting tricyclic analogue of ganciclovir (compound **2b** in the original publication) (Fig. 7) proved markedly active not only against HSV-1 and HSV-2, but also varicella-zoster virus (VZV) and cytomegalovirus.<sup>168</sup> Further substitution of the methyl group in the 1,*N*-2-isopropeno moiety by a phenyl group as in 6-phenyl-TACV (tricyclic acyclovir) and 6-phenyl-TGCV (tricyclic ganciclovir) (Fig. 7), compounds **7** and **13** in the original publication<sup>169</sup> give rise to fluorescent antiviral compounds which, on the one hand, showed an antiviral activity profile similar to that of their parent compounds (acyclovir and ganciclovir), and, on the other hand, showed relatively strong fluorescence, making them useful for the noninvasive diagnosis of herpesvirus infections.<sup>169</sup> This line of research was then extended to 6-(4-methoxyphenyl)-TACV and 6-(4-methoxyphenyl)-TGCV (compounds **8** and **27** in the original publication) as fluorescent tricyclic analogues of acyclovir and ganciclovir.<sup>170</sup>

Yet further extension of the 6-side chain in TACV and TGCV yielded the 6-[4-(phenoxycarbonyloxy)phenyl]substituted derivatives (compounds **11** and **19** in the original publication)<sup>171</sup> (Fig. 7) with, again, combination of high antiherpetic activity and strong fluorescence. Similarly, the 6-(4-biphenyl)substituted TACV and TGCV derivatives showed high selectivity against HSV-1 together with fluorescent properties.<sup>172</sup> Introduction of a fluorine i.e. in the 6-phenyl part of the structure may make the tricyclic acyclovir and ganciclovir analogues amenable to <sup>19</sup>F NMR studies.<sup>173</sup>

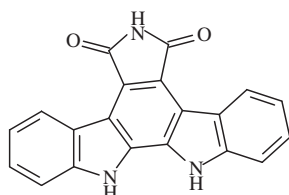
Meanwhile the fluorescent tricyclic acyclovir and ganciclovir derivatives had demonstrated a pronounced cytostatic activity (accompanied by a pronounced bystander effect) in HSV-1 thymidine kinase (TK) gene-transduced tumor cell lines.<sup>174</sup> The importance of this particular observation should be viewed in the context of our earlier observation that tumor cells transformed with the HSV-1 TK gene become highly sensitive to the cytostatic effects of anti-herpetic drugs such as ganciclovir.<sup>175</sup> In 1992 Culver et al. reported that rats bearing glioma tumors could be successfully treated with ganciclovir following *in situ* transfection by the HSV-1 TK gene.<sup>176</sup> This combined gene/chemotherapy approach for cancer has been further elaborated by Degève et al.<sup>177</sup> The fluorescent tricyclic acyclovir and ganciclovir may well fit in this approach as being fluorescent they could be readily monitored in biological fluids and tissues, and being more lipophilic than acyclovir or ganciclovir themselves they may better taken up from the blood into the central nervous system.<sup>174</sup>

Following acyclovir, ganciclovir, the relatively unknown 9-[[*cis*-1',2'-bis(hydroxymethyl)cycloprop-1'-yl]methyl]guanine and especially its 1'*S*,2'*R* enantiomer A-5021 served as the starting point for implanting the 1,*N*-2-etheno bridge. Although A-5021, quite unfortunately, has not been further pursued as an antiviral drug (candidate), its credentials have been known since the late 1990s.<sup>178-181</sup> I have, personally, always emphasized the potential of A-5021 as an antiviral drug,<sup>182</sup> and, therefore, I favor the use of the scaffold of A-5021 when building it up to tricyclic analogues, reminiscent of the tricyclic derivatives of acyclovir (TACV) and ganciclovir (TGCV) (see *supra*). The tricyclic analogues of A-5021 reported by Ostrowski et al.<sup>183</sup> confirm the potential of this approach.

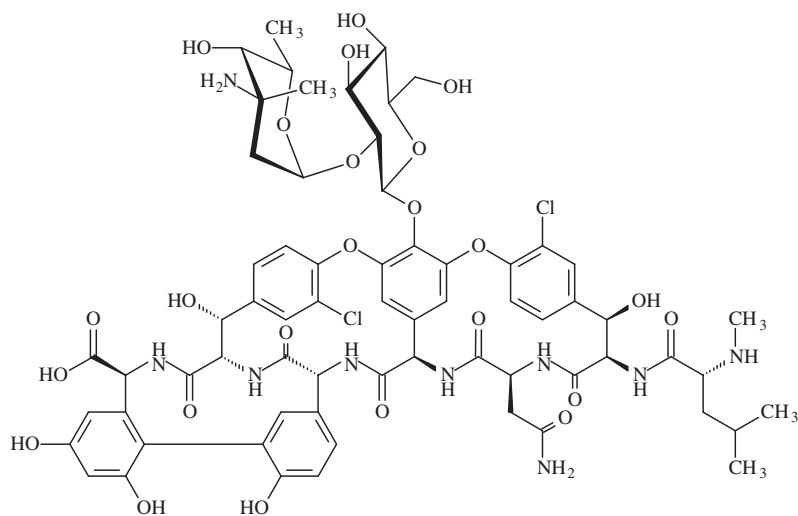
Prevailing also for the concept of the tricyclic acyclovir and ganciclovir approach is that their activity against VZV could be easily extended to a number of A-5021 derivatives, including its 6-thien-2-yl, 6-(5-bromothien-2-yl), and 6-furan-2-yl derivatives (Fig. 7).<sup>184</sup>

***H. Glycopeptide (and Other) Antibiotic Derivatives: an as Yet to be Fully Explored Opportunity to Inhibit Cellular Entry of a Wide Variety of (Enveloped) Viruses***

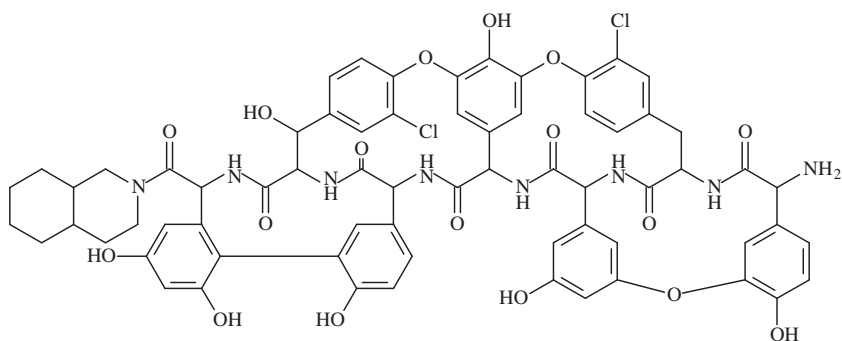
More than 20 ago, it must have been in the 1980's, I met with Dr. Maria Preobrazhenskaya [she went with me on a trip to our (remote) Campus of Kortrijk for a side visit]. She was (and still is) at the Gause Institute of New Antibiotics of the Russian Academy of Medical Sciences at Moscow (Russia). We decided to work on antibiotics, not as antibacterial agents,



Arcyriaflavin A

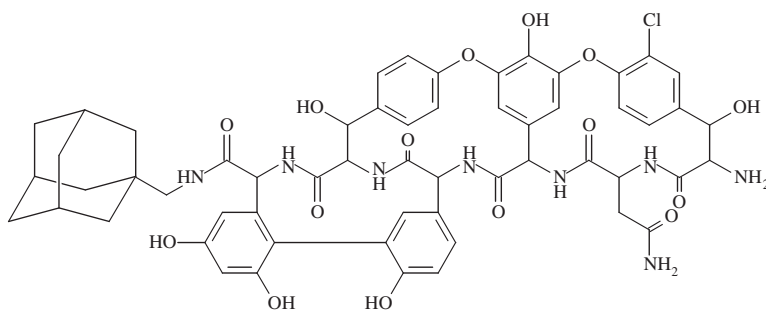
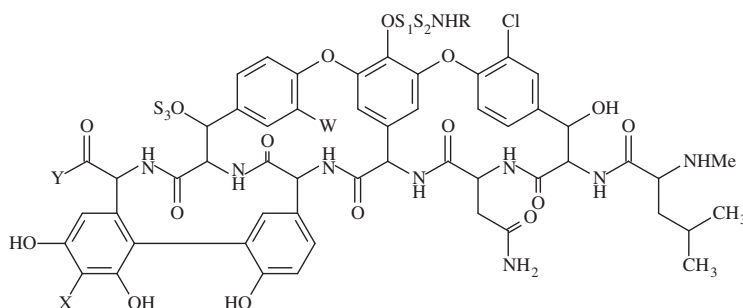


Vancomycin



Compound 62<sup>197</sup>

**Figure 8.** Glycopeptide antibiotic (e.g. vancomycin, teicoplanin, eremomycin) aglycon derivatives.

Compound 5a<sup>198</sup>

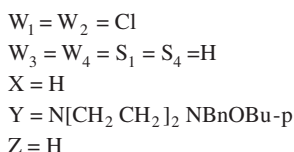
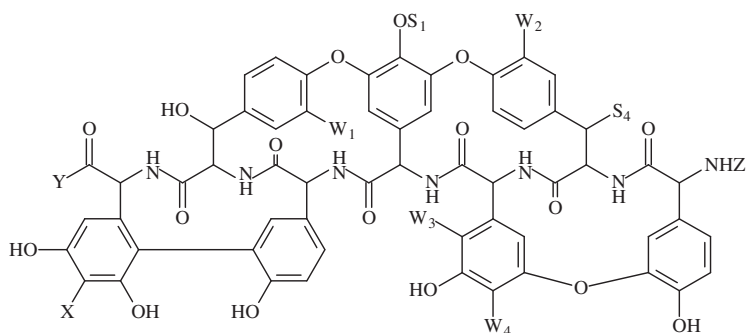
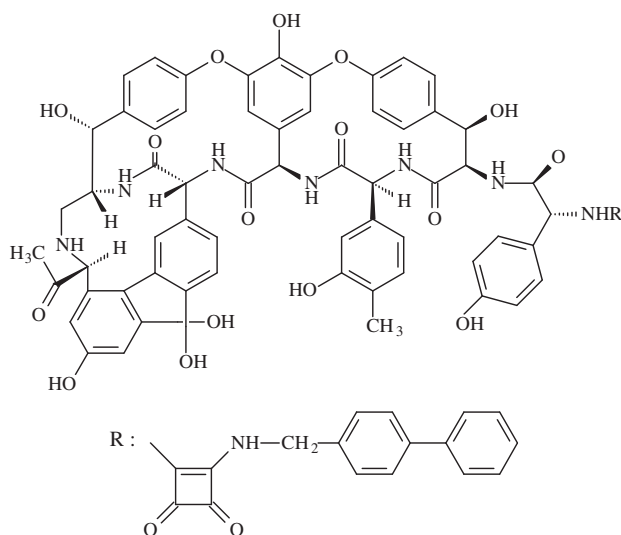
W = Cl, S<sub>1</sub> = Glc, S<sub>2</sub> = vancosamine, S<sub>3</sub> = H  
 X = CH<sub>2</sub>N[CH<sub>2</sub>CH<sub>2</sub>]<sub>2</sub>NBnBu-p  
 Y = NHMe  
 R = H

Compound 27<sup>206</sup>**Figure 8.** Continued.

but as cytostatic agents, and from our original collaboration resulted several papers describing the cytostatic activity of a number of anthracycline (i.e. daunorubicin) derivatives<sup>185–188</sup> as well as streptonigrin derivatives.<sup>189,190</sup> [Incidentally, Ferenc Sztaricskai (see *infra*) was a co-author on the latter paper.]

None of these compounds or any of their derivatives was further explored for their antiviral potential; and so was the antibiotic arcylriarubin.<sup>191</sup> Yet, antibiotics belonging to this class of compounds (such as arcylriaffavin) have been reported to exhibit potent and selective inhibition of human cytomegalovirus (HCMV) replication,<sup>192</sup> and bisindolylmaleimides such as arcylriaffavin A (Fig. 8) may even have potential usefulness as anti-hepatitis C virus agents.<sup>193</sup> In the mean time, novel derivatives of the antibiotic olivomycin<sup>194,195</sup> and novel anthracene-9,10-diones<sup>196</sup> were reported, although they were considered only as cytostatic, and, regrettably, not as (potential) antiviral agents.

Yet, the glycopeptide antibiotics vancomycin (Fig. 8), eremomycin, ristocetin and teicoplanin and their aglycons (= nonglycosylated parts) have served as the starting points for the synthesis of (semisynthetic) derivatives with antiviral activity, i.e. compound **62** (a teicoplanin type aglycon) (Fig. 8) exhibiting an EC<sub>50</sub> of 0.75 μM against HIV-1.<sup>197</sup> The mode of anti-HIV action of these glycopeptide antibiotics could be ascribed to an inhibition of the viral entry process.<sup>197</sup> From further work along these lines emanated several other glycopeptide derivatives, which, while inactive against bacteria, proved quite effective and selective as antiretroviral agents, i.e. compound **5a** (Fig. 8)<sup>198</sup> with an EC<sub>50</sub> of 1.6 μM against HIV-1.

Compound 115<sup>206</sup>Compound 8e<sup>207</sup>**Figure 8.** Continued.

As these compounds had lost their antibacterial activity (i.e. ability to interact with the peptidoglycan glycopeptidyl transferase) they may be expected not to lead to bacterial resistance even after prolonged administration. Glycopeptide aglycon derivatives such as **5a**<sup>198</sup> and **62**<sup>197</sup> may therefore be considered promising candidate drugs for the chemotherapy and/or prophylaxis of HIV infections.

Numerous glycopeptide antibiotics, primarily derived from eremomycin and teicoplanin, have been described for their antibacterial activity.<sup>199–205</sup> As shown earlier with regard to their anti-HIV activity,<sup>197,198</sup> many of these compounds also showed activity against the coronaviruses feline infectious peritonitis virus (FIPV) and human SARS CoV, i.e. compound

**27** [ $EC_{50}$ : 0.43  $\mu$ M (HIV-1), 5.4  $\mu$ M (FPIV), and 14  $\mu$ M (SARS CoV)] and compound **115** [ $EC_{50}$ : 0.7  $\mu$ M (HIV-1), 1.6  $\mu$ M (FIPV), and 8.0  $\mu$ M (SARS CoV)] (Fig. 8).<sup>206</sup> Although the mode of anti-coronavirus action could, like the anti-HIV-1 mode of action, be attributed to an inhibition of the viral entry process, there was, in general, no close correlation between the  $EC_{50}$  values of the compounds for HIV-1, FIPV, and SARS CoV.<sup>206</sup>

Whether any of the above mentioned glycopeptide aglycon derivatives would have any anti-influenza virus activity is not known (and has probably never been assessed). It therefore came as a surprise that one particular glycopeptide aglycon derivative, namely aglycoristocetin with cyclobutene dione carrying an hydrophobic side chain (Fig. 8), compound **8e** in the original publication, showed a specific, consistent and selective activity ( $EC_{50}$ : 0.4  $\mu$ M) against various influenza A and B viruses.<sup>207</sup> Again the mode of action of compound **8e** was assumed to be associated with the influenza virus entry process.<sup>207</sup> This signals that (many of) the glycopeptide antibiotic derivatives reported by Maria Peobrazhenskaya and her co-workers should be revisited for their potential anti-influenza virus activity, and, vice versa, the compounds from Ferenc Sztaricskai and his coworkers should also be looked at for their potential activity against HIV and SARS CoV.

### ***I. Is There a Role for Cidofovir in the Treatment of Polyomavirus (JC or BK) Infection: a Continuing Question?***

Since we (Graciela Andrei, Robert Snoeck, Michel Vandeputte and I) described (in 1997) the inhibitory effects of acyclic nucleoside phosphonates, such as cidofovir, against the in vitro replication of polyomaviruses,<sup>208</sup> cidofovir has been used with (anecdotal) success in the treatment of various diseases associated with the polyomaviruses JC [progressive multifocal leukoencephalopathy (PML)], BK (hemorrhagic cystitis (HC) and BK virus nephropathy (BKN)]. In 2003, I reviewed clinical data obtained with cidofovir in the treatment of polyomavirus (JC and BK)-associated PML and HC, respectively.<sup>209</sup> Here, I address further case reports that have appeared within the past 5-year period on the potential usefulness of cidofovir in the treatment of polyomavirus-associated infections.

The most remarkable report, perhaps, is that of a dual infection with polyomavirus BK and acyclovir-resistant herpes simplex virus (HSV) in a hematopoietic stem cell transplant (HSCT) recipient that was successfully treated with cidofovir (intravenously at a dose of 5 mg/kg weekly, three times, and then every other week for three additional times).<sup>210</sup> That cidofovir may preferentially inhibit HSV, as well as HCMV infection could be rationalized on the basis of its specific inhibitory effect on the viral DNA polymerase,<sup>211</sup> but polyomaviruses do not induce their own DNA polymerase, which makes it more difficult to explain the favorable results reported for cidofovir in the treatment of polyomavirus (BK) virus infections,<sup>212–214</sup> as reviewed by Rinaldo and Hirsch.<sup>215</sup> [Incidentally, in the latter review, the structure of cidofovir was not correctly presented.] As to its mechanism of action, cidofovir would inhibit polyomavirus BK replication in human renal tubular cells at a transcription of late viral mRNA, downstream of viral early gene expression.<sup>216</sup> Esterification of cidofovir with an ether lipid group, such as hexadecyloxopropyl, octadecyloxyethyl, or oleyloxyethyl, would further enhance the in vitro inhibitory effect of cidofovir on polyomavirus BK replication by 3 log's.<sup>217</sup>

BKN is an important cause of renal graft dysfunction in kidney transplant recipients [BK virus was first identified in 1971 in a kidney transplant patient with ureteric stenosis<sup>218</sup>] BK virus has been found in 50% of normal healthy kidneys but, when reactivated in a transplanted kidney, consequently to immunosuppression (i.e. with tacrolimus mycophenolate mofetil, and prednisolone) it may be causing interstitial nephritis nephropathy, which may

lead to irreversible graft failure.<sup>219,220</sup> Polyomavirus has been considered a “hot problem” in renal transplantation.<sup>221</sup>

Cidofovir that is highly concentrated in tubuli and renal tissue, the primary sites of BK virus infection, has proven to have a beneficial effect on BKN, even if administered weekly at a low dose (0.5–1.0 mg/kg intravenously) for 4–10 weeks.<sup>212</sup> This and other similarly favorable results obtained for cidofovir (reviewed by Bonvoisin et al.<sup>221</sup>) also including the first case in Japan of a kidney transplant recipient successfully treated with cidofovir for BKN<sup>222</sup> should be followed up by randomized controlled studies to prospectively evaluate the effects of cidofovir in graft survival as well as BK viral load.<sup>221</sup>

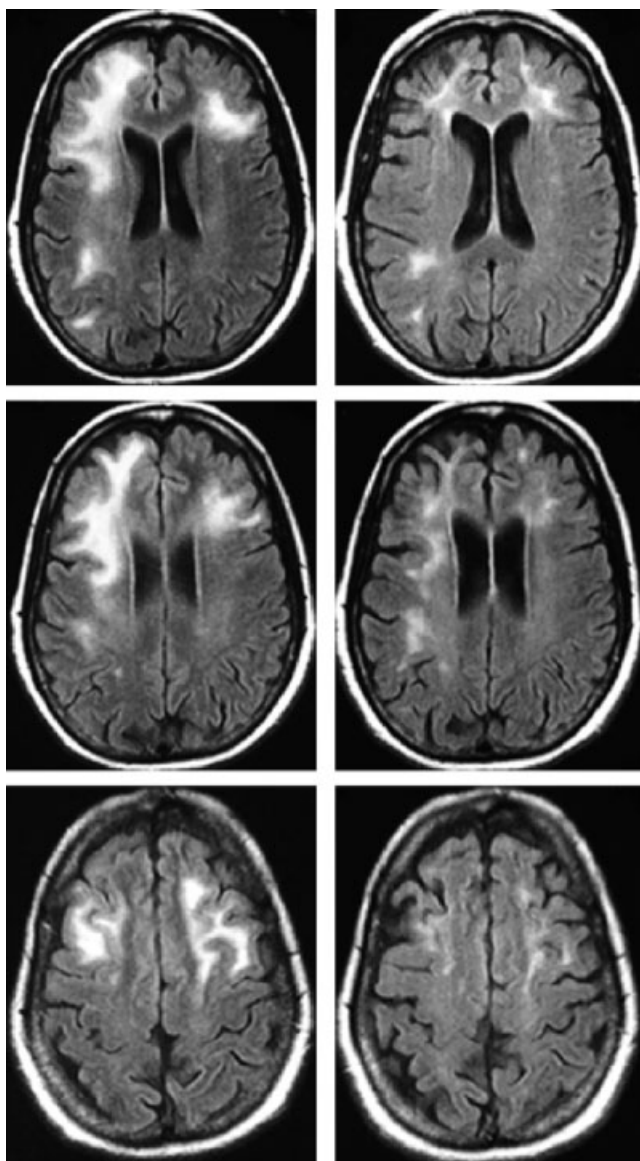
Low-dose cidofovir (weekly, 1 mg/kg intravenously) has also proven successful in the treatment of BK virus-associated HC in recipients of HSCTs,<sup>223</sup> and this observation, again, necessitates followup by a prospective trial. In the treatment of HC after allogeneic stem cell transplantation, cidofovir may even to be instilled directly into the bladder, as this regimen has been shown to decrease BK viral load and significantly improve the clinical outcome.<sup>214</sup>

Whether cidofovir, in addition to highly active anti-retroviral therapy (HAART), would give an additional benefit, as compared with HAART alone in the treatment of PML in HIV-infected patients could not be proven.<sup>224</sup> The usefulness of cidofovir in the treatment of PML has continued to be controversial since my previous review in 2003.<sup>209</sup> Cidofovir would be helpful in the treatment of PML according to some reports<sup>225,226</sup> but not according to others.<sup>227</sup> The favorable response of PML, as visualized by magnetic resonance imaging of the brain, to treatment with cidofovir (5 mg/kg every 2 weeks) in combination with cytarabine (2 mg/kg/day for 5 days every 3 weeks) is illustrated in Figure 9. In this case an obvious regression of the lesions was noted.<sup>226</sup>

### ***J. Thymidine Phosphorylase, a Target for Both Antiviral and Anticancer Agents***

Thymidine phosphorylase (TPase)<sup>228</sup> is one of the key enzymes involved in both the salvage and catabolism of pyrimidine 2'-deoxynucleosides: TPase catalyzes the reversible phosphorolysis of 2'-deoxythymidine, with formation of 2-deoxy- $\alpha$ -D-ribose-1-phosphate and thymine (Fig. 10A). TPase recognizes as substrate not only 2'-deoxythymidine and 2'-deoxyuridine but also a variety of 5-substituted 2'-deoxyuridine analogues, including BVDU (brivudin).<sup>229</sup> [BVDU has in the mean time been licensed in several European countries for the treatment of herpes zoster, where it has proved efficacious as a single oral dose of 125 mg once daily for 7 days: see ref.<sup>1</sup>.] It is in fact the TPase, which converts BVDU into its free base BVU. BVU is an inhibitor of DPD (dihydropyrimidine dehydrogenase), the enzyme responsible for the hydrogenation of the antitumor agent 5-fluorouracil (FU), and has been shown to enhance the toxicity of FU. Combination of BVDU with TPase inhibitors (see *infra*) should by preventing its degradation to BVU, in principle, enhance the antiviral activity of BVDU, and, concomitantly therewith, reduce the toxic side effects of FU, should the latter (or prodrugs thereof) be used (inadvertently) in combination with BVDU. Despite their attractive potential for increasing the antiviral activity of BVDU or other 5-substituted 2'-deoxyuridines (such as IDU or TFT: see ref.<sup>3</sup>). TPase inhibitors (see *infra*) have not been fully explored as (adjunct) antiviral agents.

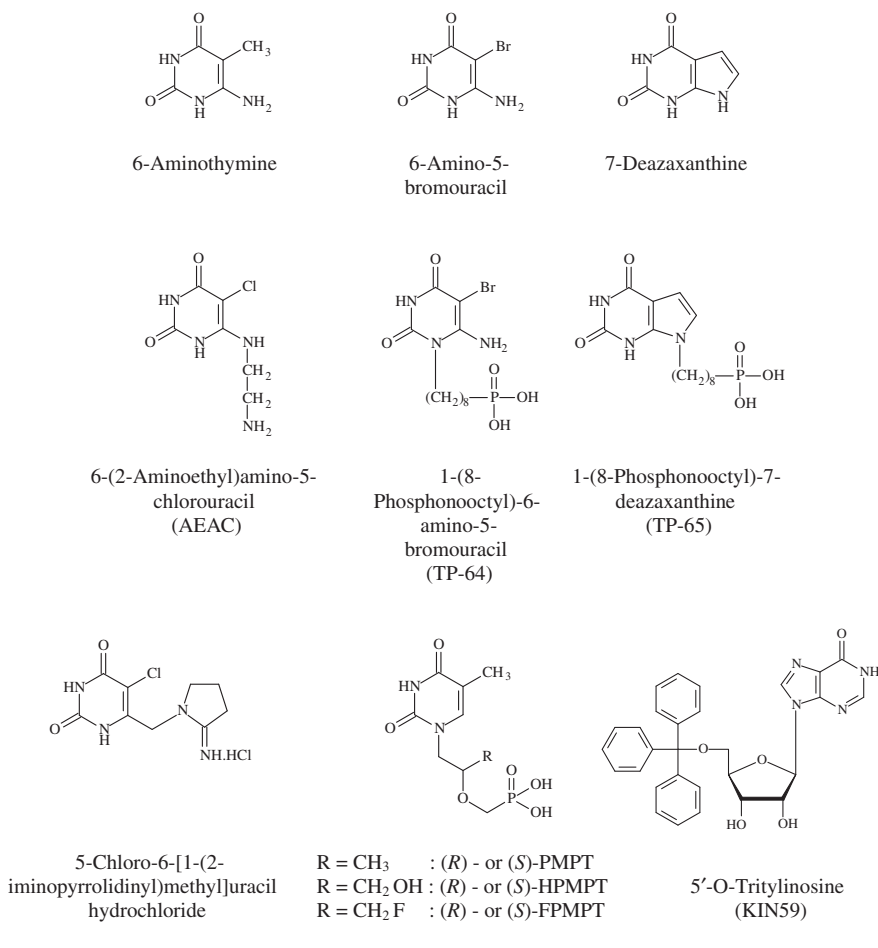
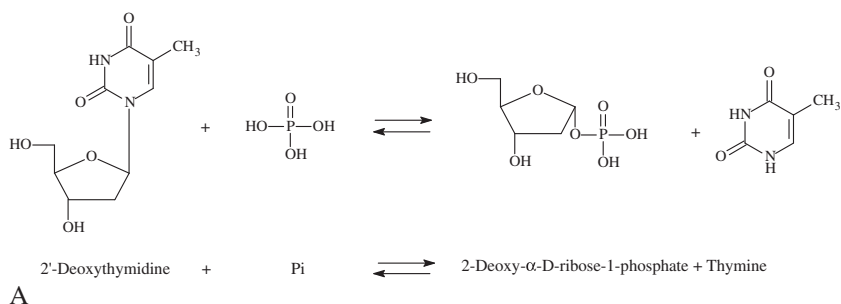
Instead, after it had become clear that TPase is identical to platelet-derived endothelial cell growth factor (PD-ECGF, a factor that had been known since the mid 1980s,<sup>230–233</sup> the sole angiogenic (angiogenesis-inducing) factor present in platelets,<sup>234</sup> interest in TPase inhibitors shifted toward their potential as anticancer agents blocking angiogenesis. As reviewed previously,<sup>235</sup> TPase is overexpressed in many solid tumors including breast, ovarian, colorectal, and pancreatic cancers, pointing to a role for this enzyme in tumor vascularization.



**Figure 9.** Magnetic resonance imaging (MRI) of the brain before (3 panels on the left) and after (3 panels on the right) six regimens of cidofovir (5 mg/kg every 2 weeks, intravenously) in combination with cytarabine (2 mg/kg/day for 5 days every 3 weeks, intravenously) in a non-AIDS patient with PML (progressive multifocal leukoencephalopathy). Data taken from Terrier et al.<sup>226</sup>

Which are the compounds that have been reported as TPase inhibitors? The first ones, and this dates from long before TPase was recognized to be an angiogenic factor, must have been 6-amino-5-bromouracil and 6-aminothymine (Fig. 10B), described in 1967 by Peter Langen and his coworkers.<sup>236</sup> It took until 1998 before 7-deazaxanthine (Fig. 10B), which could be viewed as a uracil derivative with a ring closure between C5 and C6, was described as a TPase inhibitor,<sup>237</sup> and in 2000, a new approach, based on “multisubstrate” inhibitors of TPase, was announced.<sup>238,239</sup> This “multisubstrate” concept based upon 1-(8-phosphooctyl)-7-deazaxanthine (TP-65) was, however, not followed up (except for its anti-angiogenic activity<sup>240</sup>) in further studies.





**Figure 10.** (A) Reversible phosphorolysis of 2'-deoxythymidine by thymidine phosphorylase (TPase). (B) Inhibitors of thymidine phosphorylase (TPase).

Novel 6-substituted uracil derivatives that have been described as TPase inhibitors with anti-angiogenic properties include 6-(2-aminoethyl)amino-5-chlorouracil (AEAC) (Fig. 10B).<sup>241</sup> One of the most promising TPase inhibitors described for their inhibitory effects on angiogenesis has been the thymidine phosphorylase inhibitor (TPI) 5-chloro-6-[1-(2-iminopyrrolidinyl)methyl]uracil hydrochloride (Fig. 10B).<sup>242</sup> Characteristically, this compound potentiated the antitumor activity of IDU and TFT.<sup>243</sup> In the light of what I explained

above, it would not be too farfetched to postulate that TPI may potentiate the antiviral activity of IDU, TFT and... BVDU as well.

Acyclic nucleoside phosphonates, such as cidofovir, adefovir, and tenofovir,<sup>211</sup> also referred by Holý as phosphonomethoxyalkyl analogues of nucleotides<sup>244</sup> have gained such a wide notoriety, that it would almost seem “odd” if they (or at least some of them) would not be active against TPase (PD-ECGF), and, indeed, substantial inhibition of PD-ECGF (from Sprague–Dawley lymphoma) has been observed with a series of phosphonomethoxyalkyl thymines (in order of decreasing activity): (*R*)-FPMPT > (*S*)-FPMPT ≥ (*R*)-HPMPT > (*S*)-PMPT > (*S*)-HPMPT ≥ (*R*)-PMPT.<sup>245</sup> [Inhibition of 2-deoxy-D-ribose-1-phosphate release from the endothelial cells may represent the potential anti-angiogenic target in this case.<sup>246</sup>] (*R*)- and (*S*)-PMPT, -HPMPT and -FPMPT (Fig. 10B)<sup>247,248</sup> should be further followed up as TPase (PD-ECGF) inhibitors for their anti-angiogenic activity in particular, and anticancer activity in general.

As for the reverse transcriptase (RT), both nucleoside type (i.e. TPI) and nucleotide type (i.e. (*R*)-FPMPT) of TPase inhibitors have been described. This analogy can be extended to non-nucleoside type of TPase inhibitors such as KIN59 (5'-O-tritylinosine) (Fig. 10B), which like the non-nucleoside RT inhibitors (NNRTIs), inhibit TPase via a non-competitive mechanism of action<sup>249</sup> and thus act as allosteric inhibitors of TPase.<sup>250</sup> As recently demonstrated,<sup>251</sup> 5'-O-tritylinosine would fit into a cavity (with aspartic acid residue –203 being critical) located at a distance of about 11 Å from the substrate-binding site of TPase.

## 2. CONCLUSIONS

The ten stories (E1–E10) that are subject of the present review offer a mix of premises and promises: they have done so over a period spanning one or two decades, and they are likely to continue to do so for the next 10 years or so.

E1. ATA analogues have continued to find new applications, one of the last being influenza virus, but given their lack of specificity, would they ever find their final “niche”?

E2. The ADAMs originated from the ATAs and are still looked upon as NNRTIs, but are they genuine NNRTIs (as are nevirapine, efavirenz, etravirine, rilpivirine, and others), or may they have, as yet uncovered, “side” effects worth pursuing?

E3. Among the HIV INIs, raltegravir was the first to cross the (finish) line, thereby fulfilling a long-vested premise. Will elvitegravir be the second crossing the line? What are the prospects for a quadruple drug combination consisting of tenofovir, emtricitabine, elvitegravir, and a booster (pharmacoenhancer)?

E4. In the aftermath of the INIs, LEDGF may be considered as a fascinating but risky target, and the development of LEDGF inhibitors may be a roller coaster ride.

E5. There is, at present, more talking (and writing) about influenza A (H1N1) virus strains that are resistant to oseltamivir, than about an imminent avian influenza A (H5N1) pandemic. [Meanwhile, the influenza A H5N1 Mexican pandemic swept through the five Continents.] Oseltamivir should have been stockpiled by now, and its synthesis has been improved. It should still be the first choice drug to be used if needed to curb any influenza A pandemic (whether H1N1 or H5N1).

E6. In the meantime RSV has remained a real threat, at least in the very young and (not so very) old, and besides the monoclonal antibody palivizumab and the (questionable) use of ribavirin, we are still waiting for an effective and versatile anti-RSV agent.

E7. From a purely chemical viewpoint it is surprising that tricyclic guanosine derivatives (based on 1*H*,2-ethenoguanine) has received so little attention from the biomedical world,

despite their unique combination: an anti-herpesvirus activity combined with marked fluorescent properties.

E8. Antibiotics and antivirals, often envisaged as antipodes, have been reconciled with the demonstration that glycopeptide antibiotics (or their aglycons) are able to block the entry of enveloped viruses such as HIV. This unique property may well extend, and should be further explored for other enveloped viruses, such as coronaviruses (already done), myxoviruses (only barely touched upon), and herpesviruses (not even yet considered).

E9. Cidofovir (off-label), and other acyclic nucleoside phosphonates have as yet unexplored potential for the treatment of what was once called papovaviruses. Their potential in the treatment of HPV infections has already been highlighted,<sup>4</sup> but their equally promising potential for the treatment of polyomavirus infections should be further explored as well.

E10. TPase as target to increase the antiviral activity of compounds like BVDU against HSV-1 and VZV that are readily degraded by dThd phosphorylase deserves (much) more attention than received so far and in its own right the dThd phosphorylase should be considered as a target for antitumor agents, given its angiogenic prowess thus offering the design of anti-angiogenic compounds with antitumor potential.

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