



Review

The antiviral activity and mechanism of action of (S)-[3-hydroxy-2-(phosphonomethoxy)propyl] (HPMP) nucleosides

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ABSTRACT

One class of compounds that has shown promise as antiviral agents are the (S)-[3-hydroxy-2-(phosphonomethoxy)propyl] (HPMP) nucleosides, members of the broader class of acyclic nucleoside phosphonates. These HPMP nucleosides are nucleotide analogs and have been shown to be effective inhibitors of a wide range of DNA viruses. Prodrugs of these compounds, which achieve higher levels of the active metabolites within the cell, have an expanded activity spectrum that also includes RNA viruses and retroviruses. Because they are analogs of natural nucleotide substrates, HPMP nucleosides are predicted to target polymerases (DNA polymerases, RNA polymerases and reverse transcriptases), resulting in the inhibition of viral genome replication. Previous work using the replicative enzymes of different viruses including human cytomegalovirus (HCMV) and vaccinia virus DNA polymerases and human immunodeficiency virus type 1 (HIV-1) reverse transcriptase has shown that the activated forms of these compounds are substrates for viral polymerases and that incorporation of these compounds into either the primer strand or the template strand inhibits, but does not necessarily terminate, further nucleic acid synthesis. The activity of these compounds against other viruses that do not encode their own polymerases, like polyoma viruses and papilloma viruses, suggests that host cell DNA polymerases are also targeted. This complex mechanism of action and broad activity spectrum has implications for the development of resistance and host cell genome replication, and suggests these compounds may be effective against other viruses such as influenza virus, respiratory syncytial virus and Dengue virus. This class of nucleotide analogs also points to a potential avenue for the development of newer antivirals.

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1. Acyclic nucleoside phosphonates

The acyclic nucleoside phosphonates are a class of antiviral agents that are nucleotide, rather than nucleoside, analogs. As a group, these compounds have shown activity against a range of DNA viruses, RNA viruses, and retroviruses (De Clercq and Holy, 2005; De Clercq et al., 1986; Wyles et al., 2009). The structure of these compounds consists of an acyclic nucleoside moiety attached to a phosphonate group through a phosphate–carbon–oxygen bond. This type of bond is not cleaved by cellular esterases, making these compounds more stable than those that contain the phosphate–oxygen–carbon bond of a phosphate group (De Clercq, 1997; De Clercq and Holy, 2005). The phosphonate group on this class of compounds acts as a phosphate mimic, making these antiviral agents analogs of deoxynucleoside monophosphates (dNMPs). As a result of this structure, only two phosphorylation steps are required to activate these compounds to their active diphosphoryl metabolites, which then act as analogs of deoxynucleoside triphosphates (dNTPs).

Several acyclic nucleoside phosphonates have been described, and these generally fall into three categories: the phosphonomethoxyethyl (PME) purine and pyrimidine derivatives, the phosphonomethoxypropyl (PMP) purine and pyrimidine derivatives, and the (S)-[3-hydroxy-2-(phosphonomethoxy)propyl] (HPMP) purine and pyrimidine derivatives (Fig. 1). The PME and PMP derivatives, including 9-(2-phosphonomethoxyethyl)adenine (PMEA, adefovir) and 9-(2-phosphonomethoxypropyl)adenine (PMPA, tenofovir), are obligate chain terminators because they lack a hydroxyl group that can act as a 3'-hydroxyl as found in the natural dNTP structure. In contrast, the HPMP derivatives do possess a 3'-hydroxyl group in the correct position and can theoretically be incorporated into DNA and extended (De Clercq and Holy, 2005; De Clercq and Neyts, 2009).

2. HPMP nucleosides

The first of the acyclic nucleoside phosphonates to be described was HPMP-adenine (HPMPA) (De Clercq et al., 1986) (Fig. 2). This compound originated as a merger of two other antiviral agents, phosphonoformate and (S)-9-(2,3-dihydroxypropyl)adenine [(S)-DHPA] (De Clercq and Holy, 2005; De Clercq et al., 1987). Other HPMP nucleosides have subsequently been developed (reviewed in De Clercq, 2007, 2011; De Clercq and Holy, 2005); each of these compounds has the same backbone structure and differ only in the

base moiety. This review will focus on those HPMP nucleosides that possess the natural bases adenine, cytosine, guanine and thymine: HPMPA, HPMP-cytosine (HPMPC, cidofovir), HPMP-guanine (HPMPG) and HPMP-thymine (HPMPT), analogs of dAMP, dCMP, dGMP and dTMP, respectively (Fig. 2). We have focused on these compounds as they exhibit a more interesting and complex mode of action than do the chain terminating drugs.

2.1. Uptake and metabolism of HPMP nucleosides

Much of the work on the uptake and metabolism of the HPMP nucleosides has been performed using the two most well studied compounds, HPMPA and HPMPC. Both of these analogs are taken up into cells by fluid-phase endocytosis (Connelly et al., 1993; Palú et al., 1991), where they are converted to various products including their monophosphoryl and diphosphoryl derivatives; the latter are the active metabolites (Fig. 3A). Cellular enzymes catalyze the metabolism of these compounds, as shown by data demonstrating that the phosphorylation pattern of HPMPA is similar in uninfected cells and in cells infected with herpes simplex virus type 1 (HSV-1) (Votruba et al., 1987), and the production of HPMPC metabolites is unchanged by infection with HSV-1 (Ho et al., 1992), murine cytomegalovirus (MCMV) (Okleberry et al., 1997) or with cowpox virus (Smee et al., 2002). The fact that cellular enzymes are used to catalyze these steps means that viral enzymes are not required, thus expanding the range of potential viral targets to include kinase-deficient and mutant viruses.

The metabolites formed after the uptake of HPMPA are HPMPA monophosphate (HPMPAp) and HPMPA diphosphate (HPMPApp) (Votruba et al., 1987). AMP(dAMP) kinase has been shown to catalyze the two-step phosphorylation of HPMPA to HPMPApp (Merta et al., 1992). Creatine kinase is also able to convert HPMPAp to HPMPApp (Merta et al., 1992).

In contrast to HPMPA, the intracellular metabolism of HPMPC yields three major products: HPMPC monophosphate (HPMPCp), HPMPC diphosphate (HPMPCpp), and HPMPC phosphocholine (HPMPCp-choline; Fig. 3B) (Aduma et al., 1995; Ho et al., 1992). Pyrimidine nucleoside monophosphate kinase has been shown to catalyze the first phosphorylation step, and one of three enzymes, pyruvate kinase, creatine kinase or nucleoside diphosphate kinase, catalyze the second step (Cihlar and Chen, 1996). Interestingly, Cihlar and Chen (1996) showed that human cytomegalovirus (HCMV) infection resulted in an elevation of the intracellular levels of HPMPCp and HPMPCpp relative to uninfected cells. These ele-

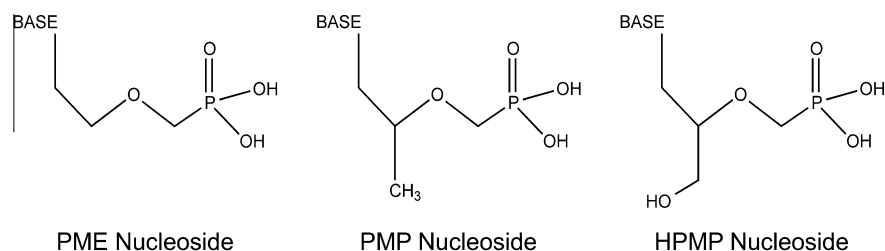


Fig. 1. Categories of acyclic nucleoside phosphonates. The phosphonomethoxyethyl (PME) nucleosides and the phosphonomethoxypropyl (PMP) nucleosides lack a hydroxyl group equivalent to the 3'-hydroxyl of a natural nucleoside and as such are obligate chain terminators. In contrast, the (S)-[3-hydroxy-2-(phosphonomethoxy)propyl] (HPMP) nucleosides do possess a hydroxyl group in the appropriate position.

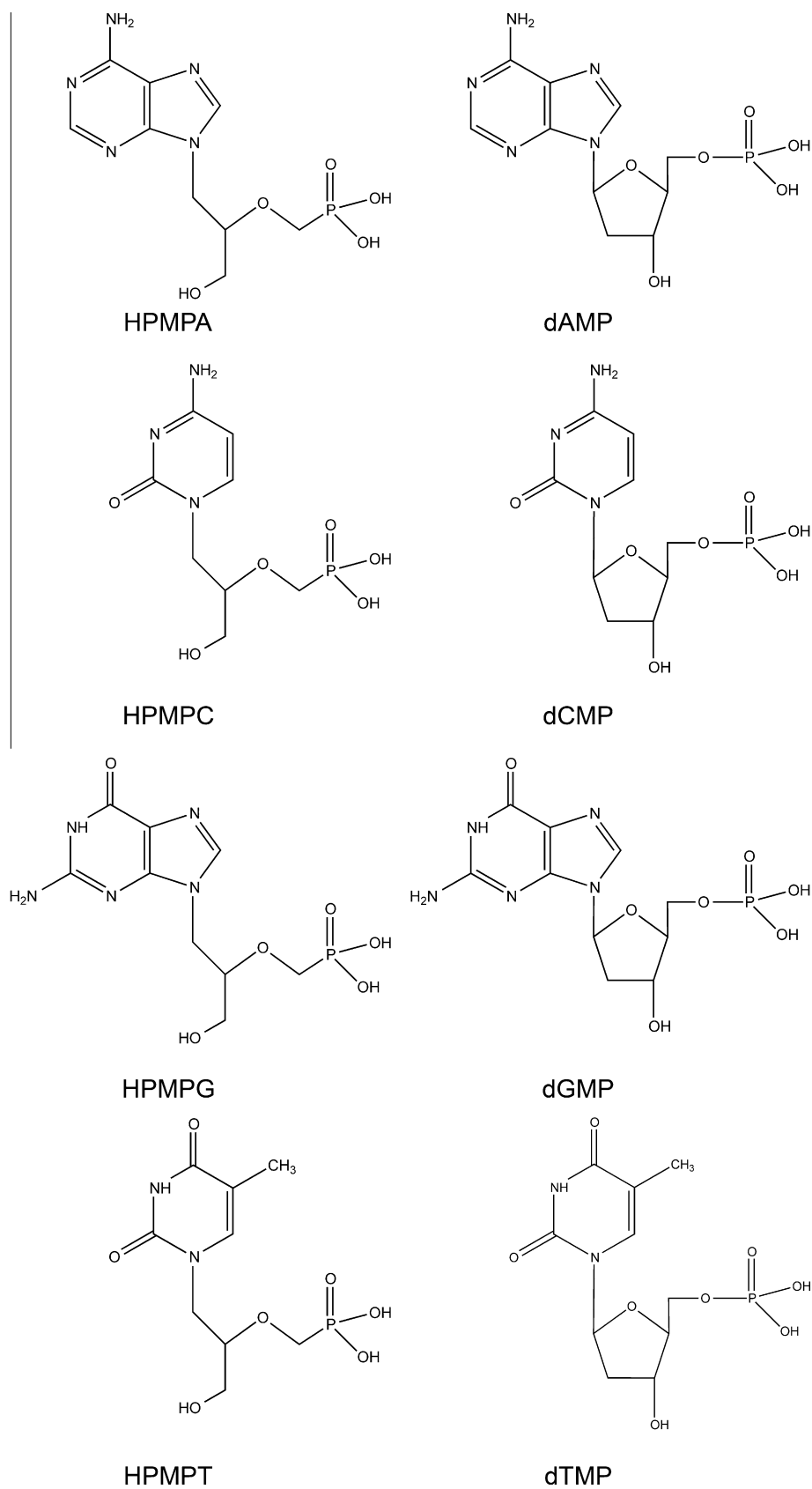


Fig. 2. Structures of the HPMP nucleosides. HPMPA, HPMP, HPMPG and HPMP are analogs of the natural nucleotides dAMP, dCMP, dGMP and dTMP, respectively. The acyclic backbone structures of these compounds are identical; only the structures of the base moieties differ.

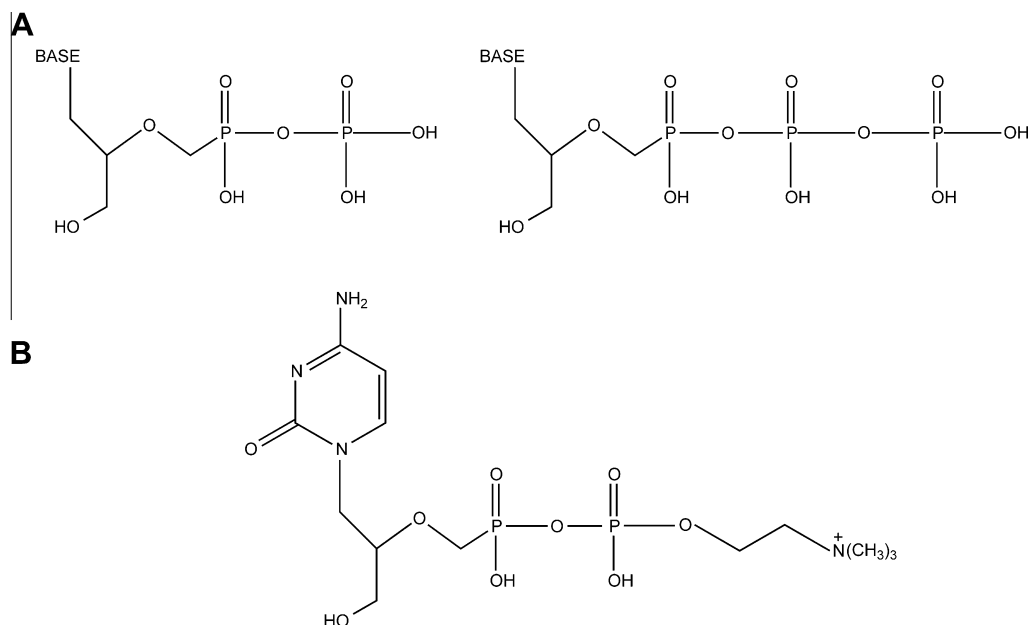


Fig. 3. Metabolites of HPMP nucleosides. (A) Structures of the monophosphoryl (left) and diphosphoryl (right) derivatives of the HPMP nucleosides. The HPMP nucleoside diphosphates are the active intracellular compounds. (B) Structure of HPMP phosphocholine (HPMPCp-choline), a third metabolite formed upon treatment of cells with HPMP.

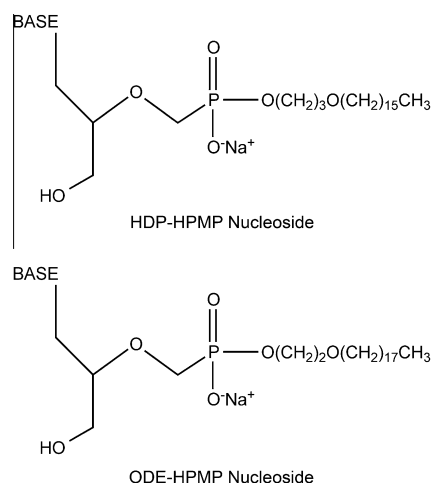


Fig. 4. Alkoxyalkyl ester derivatives of HPMP nucleosides. Structures of the hexadecyloxypropyl (HDP; top) and octadecyloxyethyl (ODE; bottom) esters are shown.

vated levels were due to higher activities of the cellular kinases in infected cells, rather than the induction of virus-encoded enzymes capable of phosphorylating HPMP (Cihlar and Chen, 1996).

The formation of HPMPCp-choline, an analog of the phospholipid synthesis intermediate cytidine 5'-diphosphocholine (Cihlar et al., 2001), from HPMPCpp is catalyzed by choline phosphate cytidyltransferase (Cundy, 1999). HPMPCp-choline has a half-life of greater than 48 h (Ho et al., 1992) and acts as a slow-release reservoir for HPMPCpp. The production of this metabolite is thought to be the source of the prolonged antiviral state (>7 days) induced by HPMP pretreatment of cells (Aduma et al., 1995). This effect is presumably also responsible for the infrequent dosing that can be used to treat established virus infections (Bronson et al., 1990; Flores-Aguilar et al., 1994; Quenelle et al., 2003; Smee et al., 2000). In comparison, the antiviral state induced by pretreat-

ment of cells with HPMPA is much shorter (24–72 h); the latter compound is not metabolized to a long-lasting reservoir like HPMPCp-choline (Aduma et al., 1995).

Although an analysis has not been performed of the metabolites produced upon treatment of cells with HPMPG, Terry et al. (1988) showed that chemically-synthesized (*R,S*)-HPMPG could be enzymatically activated to HPMPCpp. Guanylate kinase catalyzes the conversion of HPMPG to HPMPC monophosphate (HPMPGp), and nucleoside 5'-diphosphate kinase catalyzes the second phosphorylation step to HPMPC diphosphate (HPMPGpp).

A series of alkoxyalkyl ester prodrugs of the HPMP nucleosides designed to resemble natural lipids have also been developed (Painter and Hostetler, 2004); these conjugates include the hexadecyloxypropyl (HDP) and octadecyloxyethyl (ODE) esters of these compounds (Fig. 4). Treatment of cells with the HPMPA or HPMP prodrugs results in greater intracellular levels of the respective diphosphoryl derivatives compared to treatment with the unmodified compounds (Aldern et al., 2003; Magee et al., 2008). This increase in active metabolite concentration appears to be a result of a rapid cellular uptake of the prodrugs based on their interaction with the cell membrane relative to slow uptake of the parent compound by endocytosis (Aldern et al., 2003). As shown with HDP-HPMP (CMX001), the prodrug forms a large pool in the cell membrane that is first metabolized by cellular phospholipase C and phosphodiesterases to release the parent compound, which is then further metabolized to the diphosphoryl derivative (Aldern et al., 2003).

2.2. Antiviral activity spectrum

Collectively, the HPMP nucleosides inhibit the replication of a number of virus families in cell culture, including the DNA viruses herpesviruses [HSV-1, herpes simplex virus type 2 (HSV-2), CMV, Epstein-Barr virus (EBV), varicella zoster virus, human herpesvirus 6 (HHV6), human herpesvirus 7, human herpesvirus 8] (Table 1), poxviruses (vaccinia virus, cowpox virus, ectromelia virus, monkeypox virus, orf virus, variola virus, camelpox virus) (Table 2), adenoviruses, asfarviruses [African swine fever virus (ASFV)], hep-

Table 1

Inhibition of herpesviruses by HPMP nucleosides.

Virus ^a	Antiviral activity				Selected references
	HPMPA	HPMPC	HPMPG	HPMPT	
HSV-1	+	+	+	+/- ^b	Balzarini et al. (1993), De Clercq et al. (1986, 1987), (1987), Terry et al. (1988)
HSV-2	+	+	+	+/-	Balzarini et al. (1993), De Clercq et al. (1986, 1987), Terry et al. (1988), Yu et al. (1992)
CMV	+	+	+	+/-	Balzarini et al. (1993), De Clercq et al. (1986, 1987), Shigeta et al. (1991), Snoeck et al. (1988), Terry et al. (1988), Yu et al. (1992)
EBV	+	+	Unk ^c	Unk	Lin et al. (1991)
VZV	+	+	+	+/-	Andrei et al. (1995), Balzarini et al. (1993), De Clercq et al. (1986), De Clercq et al. (1987), Terry et al. (1988)
HHV6	+	+	+	Unk	Naesens et al. (2006), Reymen et al. (1995), Yoshida et al. (1998)
HHV7	+	+	+	+	Yoshida et al. (1998), Zhang et al. (1999)
HHV8	+	+	Unk	Unk	Neyts and De Clercq (1997)

^a Virus abbreviations are HSV-1 (herpes simplex virus type 1), HSV-2 (herpes simplex virus type 2), CMV (cytomegalovirus), EBV (Epstein-Barr virus), VZV (varicella zoster virus), HHV6 (human herpes virus 6), HHV7 (human herpes virus 7), HHV8 (human herpes virus 8).

^b Weak or no activity.

^c Unknown or not tested.

Table 2

Inhibition of poxviruses by HPMP nucleosides.

Virus	Antiviral activity				Selected references
	HPMPA	HPMPC	HPMPG	HPMPT	
Vaccinia	+	+	+	+/- ^a	Baker et al. (2003), Balzarini et al. (1993), De Clercq et al. (1986), De Clercq et al. (1987), Quenelle et al. (2007), Smee et al. (2002), Terry et al. (1988)
Cowpox	+	+	Unk ^b	Unk	Baker et al. (2003), Quenelle et al. (2007), Smee et al. (2002)
Ectromelia	+	+	Unk	Unk	Buller et al. (2004)
Monkeypox	+	+	Unk	Unk	Baker et al. (2003), Smee et al. (2002)
Orf	+	+	Unk	Unk	Dal Pozzo et al. (2007)
Variola	+	+	Unk	Unk	Baker et al. (2003)
Camelpox	+	+	Unk	Unk	Duraffour et al. (2007), Smee et al. (2002)
Molluscum	+	+	Unk	Unk	contagiosum
Unk	+	Unk	Unk	Unk	Bachmeyer et al. (2009), Baxter and Highet (2004), Briand et al. (2008), Davies et al. (1999), Ibarra et al. (2000), Lin et al. (1991), Meadows et al. (1997), Toro et al. (2000), Zabawski and Cockerell (1999)

^a Weak or no activity.

^b Unknown or not tested.

adnaviruses [human hepatitis B virus (HBV), duck hepatitis B virus (DHBV)], and polyomaviruses (BK virus, murine polyomavirus, SV40), and the simple retrovirus murine sarcoma virus (Table 3). Further, HPMPC inhibits the proliferation of human papillomavirus infected cells (Andrei et al., 1998a; Johnson and Gangemi, 1999) and although activity has not been demonstrated in vitro, this compound has also been used to treat human molluscum contagiosum infections (see references in Table 2). In contrast, these compounds do not have activity (or lack selective activity) against RNA viruses and human immunodeficiency virus types 1 and 2 (HIV-1 and HIV-2) (Balzarini et al., 1993; De Clercq et al., 1986; Pauwels et al., 1988; Yu et al., 1992), although Srinivas et al. (1997) demonstrated that HPMPC could inhibit HIV-1 in HeLa-CD4 cells, but not in the MT2 lymphocyte cell line. Interestingly, the related compound HPMP-uracil (HPMPU), has no activity against HSV-1, HSV-2, CMV, VZV, vaccinia virus, adenovirus or ASFV (De Clercq et al., 1987; Gil-Fernandez et al., 1987).

The alkoxyalkyl ester derivatives of the HPMP nucleosides described in Section 2.1 exhibit increased antiviral activity relative to the parent compounds (Beadle et al., 2006; Buller et al., 2004; Dal Pozzo et al., 2007; Hartline et al., 2005; Keith et al., 2004; Kern et al., 2002; Lebeau et al., 2006; Morrey et al., 2009; Quenelle et al., 2007, 2004; Valiaeva et al., 2009). This increased activity is thought to be due to the increased levels of active diphosphoryl metabolites produced upon treatment of cells with these prodrugs (Aldern et al., 2003). A further advantage of these prodrugs is that they ex-

pand the antiviral activity spectrum of these compounds to HIV-1 (Hostetler et al., 2006; Magee et al., 2011), hepatitis C virus (Wyles et al., 2009) and JC virus (Jiang et al., 2010).

2.3. Inhibition of viral and cellular enzymes

The active diphosphoryl metabolites of the HPMP nucleosides are analogs of dNTPs and as such, are predicted to target DNA polymerases. Indeed, these compounds inhibit both viral and cellular DNA polymerases. This activity has not been well studied with a great diversity of isolated enzymes, due to the greater difficulty of purifying DNA polymerases and of obtaining the biochemically active diphosphoryl compounds, as well as the greater complexity of the assays. The HCMV and vaccinia virus DNA polymerases have been the best-studied of the purified virus enzymes and consequently we will emphasize the results of these studies in the discussions that follow.

2.3.1. HPMPApp

HPMPApp, an analog of deoxyadenosine triphosphate (dATP), has been shown to inhibit the DNA replication of HSV-1 (Votruba et al., 1987), EBV (Lin et al., 1991), vaccinia virus (Magee et al., 2008), adenovirus (Mul et al., 1989), HBV (Yokota et al., 1991), and DHBV (Yokota et al., 1990). HPMPApp is a competitive inhibitor of DNA polymerases with respect to dATP and exhibits some selectivity for viral DNA polymerases over cellular DNA polymer-

Table 3
Inhibition of other viruses by HPMP nucleosides.

Virus ^a	Antiviral activity				Selected references
	HPMPA	HPMPC	HPMPG	HPMPT	
Adenoviruses	+	+	+	+/- ^b	De Clercq et al. (1986, 1987), Gordon et al. (1991), Hartline et al. (2005), Naesens et al. (2005)
Asfarvirus ASFV	+	+	+	+/-	De Clercq et al. (1986), Gil-Fernandez et al. (1987)
Hepadnaviruses					
HBV	+	+	Unk ^c	Unk	Heijntink et al. (1994), Yokota et al. (1991)
DHBV	+	+	Unk	Unk	Yokota et al. (1990)
Polyomaviruses					
BK Virus	+	+	Unk	Unk	Topalis et al. (2011)
Murine Polyomavirus	+	+	Unk	Unk	Andrei et al. (1997)
SV40	+	+	Unk	Unk	Andrei et al. (1997)
Retroviruses					
MSV	+	Unk	+	-	De Clercq et al. (1986), Pauwels et al. (1988)
HIV-1	+/-	+/-	+/-	Unk	Balzarini et al. (1993), Hostetler et al. (2006), Pauwels et al. (1988), Srinivas et al. (1997), Yu et al. (1992)
HIV-2	+/-	Unk	+/-	Unk	Balzarini et al. (1993)

^a Virus abbreviations are ASFV (African swine fever virus) HBV (human hepatitis B virus), DHBV (duck hepatitis B virus), MSV (murine sarcoma virus), HIV-1 (human immunodeficiency virus type 1), HIV-2 (human immunodeficiency virus type 2).

^b Weak or no activity.

^c Unknown or not tested.

ases (Kramata et al., 1996; Merta et al., 1990; Mul et al., 1989). This selectivity is illustrated by the respective $K_i(\text{HPMPApp})/K_m(\text{dATP})$ values for HSV-1 DNA polymerase, adenovirus DNA polymerase, human DNA polymerase α , rat DNA polymerase α , human DNA polymerase β , rat DNA polymerase δ , and rat DNA polymerase ϵ : 1.9, 0.58, 3.1, 2.3, >170, 0.25 and 0.07, respectively (Kramata et al., 1996; Merta et al., 1990; Mul et al., 1989). The low K_i/K_m values (with the exception of human DNA polymerase β) obtained with the cellular enzymes are most likely the cause of the cytotoxic effects of HPMPA seen at higher drug doses (Bronson et al., 1989; Vesely et al., 1990).

Although we did not calculate a K_i value for HPMPApp and vaccinia virus DNA polymerase, we demonstrated that this compound is an efficient substrate for the viral enzyme. The K_m of HPMPApp was calculated to be $3.8 \pm 0.8 \mu\text{M}$, relative to a value of $4.6 \pm 0.5 \mu\text{M}$ for dATP (Magee et al., 2008). The V_{\max} for the two substrates were similar, with values of 2.1 ± 0.1 and $2.0 \pm 0.07 \text{ pmol/min}$, respectively (Magee et al., 2008).

2.3.2. HPMPGpp

The deoxycytidine triphosphate (dCTP) analog HPMPGpp also inhibits viral DNA replication, as shown for vaccinia virus (Magee et al., 2008), EBV (Lin et al., 1991) and HBV (Yokota et al., 1991). This active compound is a competitive inhibitor of DNA polymerases with respect to dCTP (Cherrington et al., 1994; Ho et al., 1992; Xiong et al., 1996), and interestingly, has greater selectivity for viral enzymes relative to cellular DNA polymerases than is seen with HPMPApp. HPMPGpp is a poor inhibitor of human DNA polymerases β and γ ($K_i(\text{HPMPGpp})/K_m(\text{dCTP})$ values are 121 and 1424, respectively) (Cherrington et al., 1994), a moderate inhibitor of human DNA polymerase α and HCMV DNA polymerase ($K_i(\text{HPMPGpp})/K_m(\text{dCTP})$ values are 10.8 and 9.2, respectively) (Ho et al., 1992; Xiong et al., 1996), and a strong inhibitor of HSV-1 and HSV-2 DNA polymerases ($K_i(\text{HPMPGpp})/K_m(\text{dCTP})$ values are 2.8 and 3.8, respectively) (Ho et al., 1992). Smee et al. (2002) also showed that HPMPGpp is a moderate inhibitor of partially purified cowpox virus DNA polymerase, with a $K_i(\text{HPMPGpp})/K_m(\text{dCTP})$ value of 11.3. Further, HPMPGpp inhibited the activity the molluscum contagiosum, cowpox virus and HSV-1 DNA polymerases expressed in an in vitro transcription and translation system, although kinetic data were not shown (Watanabe and Tamaki, 2008). As with HPMPApp, we

did not calculate a K_i value for HPMPGpp and vaccinia virus DNA polymerase, but did present data showing this compound is a less favored substrate for the viral enzyme than is dCTP. The K_m of HPMPGpp was determined to be $23 \pm 6 \mu\text{M}$, higher than the calculated K_m for dCTP ($3.8 \pm 0.7 \mu\text{M}$) (Magee et al., 2005). Also as seen for HPMPApp and vaccinia virus DNA polymerase, the V_{\max} of the two substrates were similar (3.0 ± 0.4 and $2.4 \pm 0.2 \text{ pmol/min}$, respectively) (Magee et al., 2005). These results are comparable to those previously obtained using HCMV DNA polymerase, where the K_m values for HPMPGpp and dCTP were 18.7 ± 1.4 and $0.67 \pm 0.07 \mu\text{M}$, respectively and the V_{\max} values for the two substrates were 0.27 ± 0.09 and $0.39 \pm 0.08 \text{ pmol/min}$, respectively (Xiong et al., 1996).

HPMPGpp has also inhibits HIV-1 reverse transcriptase (RT), with a $K_i(\text{HPMPGpp})/K_m(\text{dCTP})$ value of 4.6 for a DNA template and 5.9 for an RNA template (Cherrington et al., 1996). However, subsequent work has indicated that HPMPGpp is not active against HIV-1 RT (Frangeul et al., 2008). In determining the 50% inhibitory concentration of HPMPGpp, these authors showed that HIV-1 RT activity is not inhibited by concentrations of this compound up to 2 mM. This enzyme could however, use HPMPGpp as a substrate and incorporate it into DNA (Frangeul et al., 2008), a result that we have recently confirmed (Magee et al., 2011).

2.3.3. HPMPGpp

The addition of HPMPG to virus-infected cells inhibits HSV-1 DNA synthesis (Terry et al., 1988). This compound's active metabolite, HPMPGpp, is a competitive inhibitor of HSV-1 DNA polymerase with respect to dGTP and selectively inhibits this viral enzyme relative to human DNA polymerase α (Terry et al., 1988). Although $K_m(\text{dGTP})$ values were not provided, the $K_i(\text{HPMPGpp})$ values were determined to be 0.032 ± 0.015 and $2.3 \mu\text{M}$ for HSV-1 DNA polymerase and human DNA polymerase α , respectively (Terry et al., 1988). It is important to note that this study used the racemic mixture of (R,S)-HPMPGpp, and not (S)-HPMPGpp alone (Terry et al., 1988).

2.3.4. HPMPTpp

Few data have been obtained on HPMPT or its active metabolite HPMPTpp, most likely because this compound exhibits little to no antiviral activity (Table 1). The explanation for this lack of activity

is unclear but may be due to an inability of cellular kinases to activate HPMPPT to HPMPPTpp, an inability of HPMPPTpp to inhibit DNA polymerases, or both.

3. Mechanism of action of HPMP nucleosides

Evidence obtained using a number of virus polymerases (DNA polymerases and reverse transcriptase), as well as human DNA polymerases, indicates that HPMP nucleoside diphosphates (HPMPXpp, where X is adenine, cytosine, guanine or thymine) can be used as substrates by these enzymes and incorporated into the 3'-terminus of the growing DNA strand. Unlike many other antiviral nucleoside and nucleotide analogs however, the addition of HPMPX to the 3' end of the primer does not result in a termination of DNA synthesis. Rather, this incorporation results in a number of different effects on enzyme activity, as described below.

3.1. HSV-1 DNA polymerase

Merta et al. (1990) demonstrated that HPMPApp could replace dATP in reactions catalyzed by purified HSV-1 DNA polymerase using activated calf thymus DNA as the substrate. Significant DNA synthesis occurs in these reactions, although not to the levels observed when dATP is present. In contrast, PMEA diphosphate, an obligate chain terminator, cannot replace dATP to support DNA synthesis (Merta et al., 1990). Although HPMPApp substrate usage was not examined directly, these results suggest HPMPA is incorporated into DNA and where it can promote continued DNA synthesis.

3.2. HCMV DNA polymerase

The first studies to thoroughly examine the mechanism of action of HPMP nucleosides against DNA polymerases looked at the interactions between HPMPApp and HCMV DNA polymerase (Xiong et al., 1997, 1996). Xiong et al. (1997) showed with primer extension analyses using defined primer template pairs that HPMPApp can be used as an alternative substrate for this enzyme, and this compound is faithfully incorporated into DNA opposite a template dGMP. These authors suggested that the rate of incorporation of CDV and dCMP into DNA are similar, based on observations of the extension of the original primer strands (Xiong et al., 1997). The addition of a single HPMP molecule to the primer terminus does not lead to chain termination and HCMV DNA polymerase can use a synthetic primer terminating in HPMP or dCMP with similar affinities (K_m values of 165 ± 42 and 90 ± 8 nM, respectively) (Xiong et al., 1996). However, the incorporation of a single HPMP molecule does slow subsequent elongation (Xiong et al., 1997). In contrast, when two consecutive molecules are incorporated into DNA, or when two molecules of HPMP separated by a natural nucleotide are incorporated, elongation is dramatically reduced or terminated. Further, when present at the 3'-terminus of the primer strand, HPMP is not excised by the 3'-to-5' proofreading exonuclease activity of HCMV DNA polymerase (Xiong et al., 1997). In addition, when present in the template strand, pausing is observed at positions immediately prior to, at, and immediately after the HPMP molecule. However, HPMP in the template strand is not an absolute block to DNA synthesis as full-length extension past the lesion is also observed (Xiong et al., 1997). Interestingly, the degree of inhibition of HCMV DNA polymerase is much larger when HPMP is located in the template strand than with a template incorporating HPMP into the nascent strand. The incorporation of HPMP into DNA therefore results in several mechanisms of HCMV DNA polymerase inhibition.

3.3. Vaccinia virus DNA polymerase

We have previously studied the effects of HPMPApp and HPMPApp on primer extension reactions catalyzed by vaccinia virus DNA polymerase (Magee et al., 2008, 2005). As described in Sections 2.3.1 and 2.3.2, both compounds can be used as substrates by this enzyme. HPMPApp was shown to be a more efficient substrate for vaccinia virus DNA polymerase than was HPMPApp (Magee et al., 2008, 2005). Indeed, HPMPApp is as good as, if not a better substrate than, the natural substrate dATP. In contrast, HPMPApp is a less efficient substrate for this enzyme than dCTP. Both HPMPA and HPMP are incorporated into the 3'-terminus of the growing DNA strand and these HPMPA/C-terminated primers can also be extended by each polymerase, indicating that the hydroxymethyl moiety on each compound can act as a 3'-hydroxyl group to initiate nucleophilic attack on the incoming dNTP. The presence of either compound in the primer strand did however, lead to the slowing or pausing of continued DNA synthesis (Magee et al., 2008, 2005). The incorporation of a single HPMPA or HPMP residue by vaccinia virus DNA polymerase results in pausing at the N + 1 position, where N is the expected site of incorporation of the compound. When two consecutive molecules of HPMPA or HPMP are incorporated into DNA by this enzyme, a slowing of DNA synthesis was also observed relative to control reactions incubated with the natural nucleotides (dATP and dCTP, respectively); pause sites were located at positions N, N + 1, N + 2, and N + 3. This slowing of DNA synthesis following HPMP nucleoside incorporation is more profound when HPMP is incorporated in the primer strand than when HPMPA was present, particularly when two molecules are incorporated. In all of these reactions however, the production of full-length DNA strands was observed (Magee et al., 2008, 2005). These results indicate that HPMPA and HPMP are not simple chain terminators. They can be incorporated into DNA and this permits the continuation of DNA synthesis.

One of the issues these studies highlight is the danger of relying upon simplified biochemical assays as screens for drugs with antiviral properties. HPMPA is a highly effective inhibitor of vaccinia virus growth in vivo, with EC_{50} values ranging from 0.3 to 8.1 μ g/ml (Baker et al., 2003; De Clercq et al., 1986, 1987), and yet the HPMPApp metabolite is just as good a substrate for vaccinia virus DNA polymerase as is dATP in primer extension assays. As we note below, chain extension assays do not always accurately reflect the primary mode of action of this compound.

The effect of HPMPA and HPMP on the 3'-to-5' exonuclease activity of vaccinia virus DNA polymerase was also investigated. Although HPMPA and HPMP can be excised from DNA bearing these compounds at the primer terminus, oligonucleotides containing either compound at the penultimate position of this terminus, equivalent to the N + 1 structure described above, are refractory to exonuclease activity (Magee et al., 2008, 2005). These results are different from what is seen with HCMV DNA polymerase (Section 3.2). The HCMV enzyme's 3'-to-5' exonuclease activity cannot excise a molecule of HPMP located at the primer terminus (Xiong et al., 1997).

The ability of HPMPA and HPMP to be incorporated into DNA by vaccinia virus DNA polymerase without causing chain termination prompted an examination of the effects of these compounds when positioned in the template strand. Similar to the results observed using HCMV DNA polymerase (Section 3.2) (Xiong et al., 1997), the presence of HPMPA or HPMP in the template results in inhibition of DNA synthesis. However, unlike HCMV DNA polymerase, vaccinia virus DNA polymerase is completely inhibited by drug in the template; nucleotide can be incorporated opposite the HPMPA/C residue but further extension does not occur. This inhibition of extension is observed even if oligonucleotides terminating one nucleotide past the HPMPA/C residue are used to prime

DNA synthesis. Interestingly, these latter primers are rapidly degraded by one nucleotide by the vaccinia virus DNA polymerase 3'-to-5' exonuclease activity, suggesting that these structures are not favored substrates.

Jesus et al. (2009) examined the effects of HPMP on the vaccinia virus replicative cycle, and found that treatment of virus-infected cells with this compound inhibits genome encapsidation and virion morphogenesis. Further, analysis by atomic force spectroscopy of genomic DNA synthesized in the presence of HPMP showed aggregated structures that were not seen when this compound is absent. These authors suggested that the incorporation of HPMP into DNA alters the structure of the nucleic acid and prevents its packaging into virions (Jesus et al., 2009). More recently we have solved the solution structure of a DNA duplex containing a single molecule of HPMP (Julien et al., 2011). This compound is well accommodated in the DNA structure and comparison with an isosequential control DNA duplex shows a root mean square deviation of only 1.5 Å over all common atoms. The stability of the HPMP-containing DNA duplex however, is less than the control DNA with a lower melting temperature relative to the latter (Julien et al., 2011). We favor the hypothesis that once an HPMP moiety has been incorporated into DNA, it creates a subtle disturbance in the structure of the helix that then creates an impediment to the transit of the molecule through the DNA binding domains of a DNA polymerase or packaging helicase. A good analogy would be the difficulty encountered trying to screw a nut over a bolt bearing a stripped thread.

3.4. Adenovirus DNA polymerase

HPMPApp inhibits adenovirus DNA replication at the level of elongation, but not at initiation (Mul et al., 1989). These authors did not demonstrate HPMPA incorporation into DNA catalyzed by adenovirus DNA polymerase, but did present data showing HPMPApp inhibits, but does not stop elongation.

3.5. HIV-1 RT

Frangeul et al. (2008) first demonstrated the ability of HIV-1 RT to incorporate HPMP into DNA, although HPMPApp was also determined to be an inefficient substrate. More recently, we showed that HPMPApp and HPMPApp can replace dATP and dCTP, respectively, to support DNA synthesis catalyzed by HIV-1 RT (Magee et al., 2011). The incorporation of either compound into DNA slows, but does not absolutely terminate continued DNA synthesis. Interestingly, the incorporation of two consecutive molecules of HPMP, but not HPMPA, results in a severe inhibition of HIV-1 RT activity. The presence of either compound in the template strand also inhibits HIV-1 RT; this inhibition is more profound with HPMP than with HPMPA (Magee et al., 2011).

3.6. Human DNA polymerases

The ability of HPMP nucleosides to inhibit the growth of a variety of different tumors and transformed cell lines, including those with and without a known viral cause (Andrei et al., 1998a,b; Collette and Zechel, 2011; Johnson and Gangemi, 1999; Liekens et al., 1998, 2001a,b; Redondo et al., 2000; Sherman et al., 2002; Snoeck et al., 2000; Spanos et al., 2005; Yoshizaki et al., 2008), as well as their ability to inhibit the replication of viruses that use cellular DNA polymerases to replicate their genomes (Andrei et al., 1997; Topalis et al., 2011) suggests that cellular DNA polymerases are also inhibited by these compounds. Indeed, rat cellular DNA polymerases α , δ , and ϵ can use HPMPApp as a substrate and incorporate two to four consecutive HPMPA molecules into a growing DNA strand (Birkus et al., 2004; Kramata et al., 1996). Further, both DNA

polymerases δ and ϵ can excise HPMPA from a primer terminus, with more efficient removal of the drug by polymerase ϵ (Birkus et al., 2004). In addition, the mitochondrial abnormalities observed in biopsies from patients with BV virus nephropathy treated with HPMP (Talmon et al., 2010) suggests that DNA polymerase γ may use these compounds as substrates as well.

3.7. Model of the mechanism of action of HPMP nucleosides

The inhibition of polymerases, including DNA polymerases and reverse transcriptase by HPMP nucleosides occurs after their incorporation into DNA and is due to a number of different factors, including the inhibition of DNA synthesis by the presence of compound in the primer strand and inhibition of DNA synthesis by these compounds in the template strand; inhibition of 3'-to-5' exonuclease activity also plays a role (Fig. 5). The inhibition of DNA synthesis by the presence of these compounds in the template is of particular interest, as this mechanism is not relevant for most other antiviral nucleoside or nucleotide analogs because they act as either obligate or functional chain terminators. One notable exception is the antiviral drug entecavir, a 2'-deoxyguanosine analog that, like the HPMP nucleosides, possesses a 3'-hydroxyl moiety. Entecavir also has a complex mechanism of inhibition against HIV-1 RT, inhibiting DNA synthesis when present in either the primer or the template strands (Domaoal et al., 2008; Tchesnokov et al., 2008).

4. Resistance to HPMP nucleosides

The complex mechanism of action of HPMP nucleosides against DNA polymerases and reverse transcriptases has implications for the development of resistance to these drugs. The in vitro selection of resistance to HPMPA and/or HPMP has been described for HSV-1, HCMV, HHV6, vaccinia virus, cowpox virus, camelpox virus,

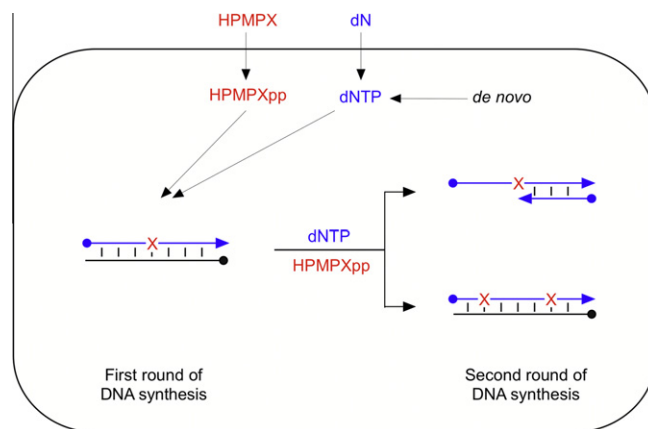


Fig. 5. The mechanism of action of the HPMP nucleosides. HPMP nucleosides (HPMPX) are taken up into cells by endocytosis and then converted to their active intracellular metabolites, HPMP nucleoside diphosphates (HPMPXpp), by cellular enzymes. As analogs of dNTPs, HPMPXpp can be used as substrates by polymerases (DNA polymerases and reverse transcriptases) and incorporated into DNA (X residues [red] embedded in newly synthesized ordinary DNA [blue]). This incorporation of HPMPX into the primer strand results in a slowing of subsequent DNA elongation, but not termination. Since these primer strands are eventually fully elongated, they can serve as template DNA in the second round of DNA synthesis (at right). In this case, the presence of HPMPX in what is now a new template strand (top right) also causes inhibition of DNA synthesis; this inhibition can either cause a slowing or termination of further DNA elongation, depending on the enzyme. The semi-conservative nature of DNA replication also results in further rounds of HPMPX incorporation opposite the old, original, DNA template (black strand, bottom right). HPMPX in the DNA primer strand can also inhibit the DNA polymerase 3'-to-5' exonuclease activity, potentially creating mutations as well as slowing replication (not shown).

Table 4

Virus resistance to HPMP nucleosides.

Virus ^a	Resistance described				References
	HPMPA	HPMPC	HPMPG	HPMPT	
HSV-1	+	+	Unk ^b	Unk	Andrei et al. (2007), Andrei et al. (2000)
HSV-2	Unk	+	Unk	Unk	Wyles et al. (2005)
HCMV	+	+	Unk	Unk	Cihlar et al. (1998), Scott et al. (2007), Smith et al. (1998), Snoeck et al. (1996)
HHV6	Unk	+	Unk	Unk	Bonnafous et al. (2008)
Vaccinia	+	+	Unk	Unk	Andrei et al. (2006), Becker et al. (2008), Kornbluth et al. (2006), Smee et al. (2002)
Cowpox	+	+	Unk	Unk	Smee et al. (2002)
Camelpox	+	+	Unk	Unk	Smee et al. (2002)
Monkeypox	+	+	Unk	Unk	Smee et al. (2002)
Adenovirus	Unk	+	Unk	Unk	Gordon et al. (1996), Kinchington et al. (2002)

^a Virus abbreviations are HSV-1 (herpes simplex virus type 1), HSV-2 (herpes simplex virus type 2), HCMV (human cytomegalovirus), HHV6 (human herpes virus 6).^b Unknown or not tested.

monkeypox virus and adenovirus (Table 4). In some cases resistance loci create cross-resistance to many different related nucleoside phosphonate compounds (Duraffour et al., 2012; Gammon et al., 2008). Resistance to HPMPC, the only member of the HPMP nucleoside class to be approved for use in humans, has also been found in clinical isolates of HSV-2 and HCMV (Table 4).

In many of these cases, resistance has been mapped to viral DNA polymerase genes and the specific residues conferring resistance have been located in both the polymerase domain and the 3'-to-5' exonuclease domain of these enzymes. Although more work is needed to examine the interactions of HPMPXpp with these drug-resistant polymerases, it is probable that mutated residues in the polymerase domain allow the enzyme to select against incorporation of the compounds into DNA, or to bypass HPMPC residues in the template strand, while mutations in the 3'-to-5' exonuclease domain facilitate removal of the compound after incorporation. For example, in vitro selection for vaccinia viruses that are resistant to HPMPC and related compounds repeatedly selects for an A314T substitution in the exonuclease domain and for an A684V mutation in the polymerase domain (Andrei et al., 2006; Gammon et al., 2008). The A314T mutation can be modeled to fall within an extended β -hairpin region in the homologous RB69 DNA polymerase structure (Shamoo and Steitz, 1999) and has been speculated to play some role in transferring DNA from the polymerase to exonuclease domains. This is supported by in vitro enzyme assays where polymerases encoding the A314T substitution excise HPMPC from the penultimate position in a primer strand far faster than wild type vaccinia DNA polymerase (Gammon and Evans, 2009). The A684V mutation falls in a highly conserved part of the polymerase domain active site and likely influences the positioning of another highly conserved nearby tyrosine residue that plays an important role in nucleotide selection (Andrei et al., 2006). This mutation could thus favor either selection against HPMPXpp nucleotides or promote trans-lesion bypass when the enzyme encounters a HPMPX moiety in the template. Viruses encoding the A684V mutation are weak mutators (Andrei et al., 2006), a phenotype compatible with either hypothesis.

The inhibitory effect caused by HPMP nucleosides in the template strand may explain why it is so difficult to select for viruses exhibiting high-level resistance to these compounds. The few mutants that have been identified (e.g. the vaccinia A314T and A684V mutations) are not highly resistant and exhibit reduced virulence in mice (Andrei et al., 2006). Although one can select for higher levels of resistance with continued passage, these mutant viruses exhibit even greater levels of impairment. In many ways these compounds create lesions resembling a form of DNA damage. This damage is not necessarily a problem for cells as it can be accommodated using bypass polymerases and repaired using cellular DNA repair systems. [However, the mitochondria may be less capa-

ble of repairing such damage, perhaps accounting for the mitochondrial toxicity reported for some of these compounds (Talmon et al., 2010).] It is much more of a problem for viruses, especially because no mutant polymerase could ever completely avoid incorporating these compounds into DNA and thus avoid creating an impediment to second strand synthesis. It appears to be difficult for viruses to transit a mutational landscape selecting for DNA polymerases that exhibit a combination of three complex traits (efficient drug excision, great selectivity against HPMPXpp incorporation, and a tolerance for damaged templates) without negatively affecting fitness. This is of special relevance in the bio-defense realm, where the studies with vaccinia virus shows that high virulence appears to be incompatible with drug resistance in an Orthopoxvirus (Andrei et al., 2006; Becker et al., 2008; Duraffour et al., 2012; Gammon et al., 2008; Kornbluth et al., 2006; Smee et al., 2002, 2005). This suggests that drug-based countermeasures will not be readily defeated through the malicious creation (or natural evolution) of drug-resistant strains and provides some support for the development of compounds like CMX001 as a smallpox therapeutic (Painter and Hostetler, 2004).

Although drug resistance has been more extensively evaluated using vaccinia virus, some of the same themes seem to be applicable to other viruses. The Hostetler laboratory has showed that the alkoxyalkyl ester derivatives of HPMP nucleosides retain activity against HIV-1 strains possessing a number of resistance mutations at clinically relevant concentrations (Hostetler et al., 2006; Magee et al., 2011). It will be of interest to determine the types of mutations that will develop upon the in vitro selection of resistance of HIV-1 to HPMP nucleosides. Since these compounds exert major effects on HIV-1 RT when present in the template strand, we predict that resistance will be more difficult to achieve than that obtained with other antiretroviral compounds that act as obligate chain terminators and that novel mutations will develop.

5. Conclusions

The complex mechanism of action that has been described for HPMP nucleosides, particularly the inhibition of enzyme activity by the presence of these compounds in the template strand, points to a new avenue for targeted antiviral drug design. Nucleoside and nucleotide analogs can be developed that are efficient and selective substrates for viral polymerases and that cause little inhibition of nucleic acid synthesis when being incorporated into DNA or RNA. These analogs would instead exert their effects in subsequent rounds of replication. Modifications to these drugs could then be made, if necessary, to improve oral bioavailability, uptake, and metabolism. Future work will also include an elucidation of the structural basis for the inhibition of DNA polymerases and reverse

transcriptases, and a determination of the mechanism of inhibition of the diphosphoryl derivatives of these compounds on the polymerases of HBV and HCV. These experiments will help to determine if the complex mechanism of action that has been observed against HCMV DNA polymerase, vaccinia virus DNA polymerase, and HIV-1 RT is more broadly applicable to these other virus systems. In addition, it will be of interest to test the efficacy of HPMP nucleosides, and their alkoxyalkyl ester derivatives, against other viruses, like influenza, measles, or respiratory syncytial virus, to determine the full antiviral activity spectrum of these drugs. Both the parent compounds and their derivatives should be tested, in case high levels of drug are required for activity, as is seen in the case of HIV-1 and hepatitis C virus.

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