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Quantification of isomeric equilibria formed by metal ion complexes of 8-[2-(phosphonmethoxy)ethyl]-8-azaadenine (8,8aPMEA) and 9-[2-(phosphonmethoxy)ethyl]-8-azaadenine (9,8aPMEA). Derivatives of the antiviral nucleotide analogue 9-[2-(phosphonmethoxy) ethyl]adenine (PMEA)

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Abstract The acidity constants of the two-fold protonated acyclic 9-[2-(phosphonmethoxy)ethyl]-8-azaadenine, $H_2(9,8aPMEA)^{\pm}$, and its 8-isomer, 8-[2-(phosphonmethoxy)ethyl]-8-azaadenine, $H_2(8,8aPMEA)^{\pm}$, both abbreviated as $H_2(PA)^{\pm}$, as well as the stability constants of their $M(H;PA)^+$ and $M(PA)$ complexes with the metal ions $M^{2+} = Mg^{2+}, Ca^{2+}, Sr^{2+}, Ba^{2+}, Mn^{2+}, Co^{2+}, Ni^{2+}, Cu^{2+}, Zn^{2+}$ or Cd^{2+} , have been determined by potentiometric pH titrations in aqueous solution at $I=0.1$ M ($NaNO_3$) and 25 °C. Application of previously determined straight-line plots of $\log K_{M(R-PO_3)}^M$ versus $pK_{H(R-PO_3)}^H$ for simple phosphonate ligands, $R-PO_3^{2-}$, where R represents a residue without an affinity for metal ions, proves that for all $M(PA)$ complexes a larger stability is observed than is expected for a sole phosphonate coordination of the metal ion. This increased stability is attributed to the formation of five-membered chelates involving the ether oxygen present in the aliphatic residue ($-CH_2-O-CH_2-PO_3^{2-}$) of the ligands. The formation degrees of these chelates were calculated; they vary between about 13% for $Ca(8,8aPMEA)$ and 71% for

$Cu(8,8aPMEA)$. The adenine residue has no influence on complex stability except in the $Cu(9,8aPMEA)$ and $Zn(9,8aPMEA)$ systems, where an additional stability increase attributable to the adenine residue is observed and equilibria between four different isomers exist. This means (1) an open isomer with a sole phosphonate coordination, $M(PA)_{op}$, where $PA^{2-} = 9,8aPMEA^{2-}$, (2) an isomer with a five-membered chelate involving the ether oxygen, $M(PA)_{cl/O}$, (3) an isomer which contains five- and seven-membered chelates formed by coordination of the phosphonate group, the ether oxygen and the N3 site of the adenine residue, $M(PA)_{cl/O/N3}$, and finally (4) a macrochelated isomer involving N7, $M(PA)_{cl/N7}$. For $Cu(9,8aPMEA)$ the formation degrees are 15, 30, 48 and 7% for $Cu(PA)_{op}$, $Cu(PA)_{cl/O}$, $Cu(PA)_{cl/O/N3}$ and $Cu(PA)_{cl/N7}$, respectively; this proves that the macrochelate involving N7 is a minority species. The situation for the $Cu(PMEA)$ system, where $PMEA^{2-}$ represents the parent compound, i.e. the dianion of 9-[2-(phosphonmethoxy)ethyl]adenine, is quite similar. The relationship between the antiviral activity of acyclic nucleoside phosphonates and the structures of the various complexes is discussed and an explanation is offered why 9,8aPMEA is biologically active but 8,8aPMEA is not.

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Abbreviations (d)ATP⁴⁻: (2'-deoxy)adenosine 5'-triphosphate · PMEA: 9-[2-(phosphonmethoxy)ethyl]adenine · 8,8aPMEA: 8-[2-(phosphonmethoxy)ethyl]-8-azaadenine · 9,8aPMEA: 9-[2-(phosphonmethoxy)ethyl]-8-azaadenine · *I*: ionic strength · K_a : acidity constant · M^{2+} : divalent metal ion

Introduction

The idea to use nucleotide analogues as therapeutic agents has a long tradition [1, 2] and all building blocks of a nucleotide have been altered and varied over the years (see, for example, [3] and references therein). Within the large group of acyclic nucleotide analogues ([4, 5, 6] and references therein), two series deserve recognition, i.e. the (*S*)-3-hydroxy-2-(phosphonomethoxy)propyl (HPMP) derivatives and the (phosphonomethoxy)ethyl (PME) derivatives [7, 8]^{1,2}. Representative compounds of these series are (*S*)-9-[3-hydroxy-2-(phosphonomethoxy)propyl]adenine (HPMPA) and 9-[2-(phosphonomethoxy)ethyl]adenine (PMEA) [4, 5, 6, 8], both of which are active against a wide range of viruses, including herpesviruses, poxviruses [9], hepadnaviruses and retroviruses [4, 5, 6, 10]. This discovery [10] goes back to 1986 and in 2002 one of these compounds, namely PMEAs, now known as Adefovir [4, 5, 6], was approved by the US Food and Drug Administration (FDA)³ in its oral prodrug form, i.e. its bis(pivaloyloxymethyl)ester (Adefovir dipivoxil [4, 5, 6]), for the treatment of hepatitis B patients who suffer from the infection of a DNA virus. For the same treatment, the same compound but under the name Hepsersa was also approved in March 2003 for "Community Marketing" by the European Agency for the Evaluation of Medicinal Products (EMEA)⁴.

The dianion of PMEAs, which can be considered as an analogue of (2'-deoxy)adenosine 5'-monophosphate [(d)AMP²⁻] (Fig. 1 [11, 12, 13]) is converted in the cells [4, 5, 6] into its diphosphorylated form, PMEApp⁴⁻, which is an analogue of (2'-deoxy)adenosine 5'-triphosphate [(d)ATP⁴⁻]. PMEApp⁴⁻ is initially recognized by nucleic acid polymerases as a substrate and incorporated in the growing nucleic acid chain, which is then terminated due to the lack of a 3'-hydroxy group. In the polymerase reaction, two metal ions are involved [14, 15, 16]: one needs to be coordinated to the β,γ -phosphate units and the other to the α -phosphate group [17, 18] to promote the transfer of a nucleotidyl residue [19]. The observation that PMEApp⁴⁻ is initially a better substrate than the parent dATP⁴⁻ [20, 21] was rationalized by the suggestion [19, 22, 23] that the ether oxygen atom present in PMEAs facilitates the M²⁺/ α -phosph(on)ate coordination by the formation of a five-membered

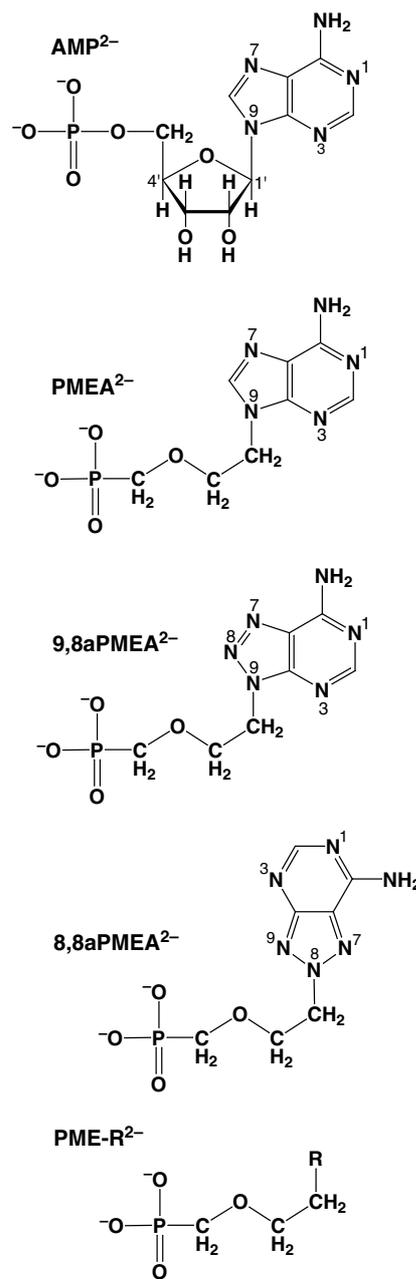


Fig. 1 Chemical structures of adenosine 5'-monophosphate (AMP²⁻) and of the dianions of 9-[2-(phosphonomethoxy)ethyl]adenine (PMEAs²⁻ = Adefovir) [4, 5, 6], 9-[2-(phosphonomethoxy)ethyl]-8-azaadenine (9,8aPMEAs²⁻) and of 8-[2-(phosphonomethoxy)ethyl]-8-azaadenine (8,8aPMEAs²⁻), together with the structure of PME-R²⁻, where R is a non-interacting residue, which represents the metal ion-coordinating properties of the ether-phosphonate chain occurring in PMEAs²⁻, 9,8aPMEAs²⁻ and 8,8aPMEAs²⁻. A further ligand to be considered in this study is 9-(4-phosphonobutyl)adenine, which is abbreviated as dPMEAs²⁻ (= 3'-deoxa-PMEAs²⁻) to indicate that its structure corresponds to that of PMEAs²⁻ except that the ether O atom is replaced by a CH₂ group. It should be noted that AMP²⁻ is shown in its dominating *anti* conformation [11, 12] and that the orientation of PMEAs²⁻ in solution [13] resembles this *anti* conformation

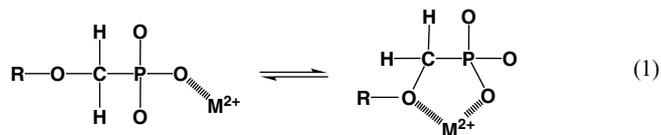
¹For further abbreviations see the legends to Figs. 1 and 2

²Species written without a charge either do not carry one or represent the species in general (i.e. independent of their protonation degree); which of the two possibilities applies is always clear from the context. In formulas like M(H;PA)⁺, the H⁺ and PA²⁻ are separated by a semicolon to facilitate reading, yet they appear within the same parenthesis to indicate that the proton is at the ligand without defining its location

³Information regarding US FDA: (2002) Chem Rundschau (CH-4501 Solothurn, Switzerland) no. 19 (Oct 8), p 68

⁴Information regarding EMEA as downloaded from the World Wide Web in December 2003: <http://www.emea.eu.int/humandocs/PDFs/EPAR/hepsersa/610202en1.pdf>

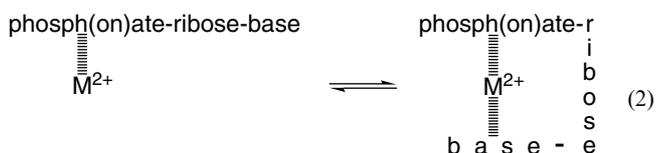
chelate ring, as is expressed in a simplified manner in equilibrium 1:



In fact, it is well known that this ether oxygen is crucial for the biological activity of PMEAs and that its replacement by an S atom [24] or a CH₂ unit [20] results in inactive compounds. Similarly, the position of this O atom within the aliphatic chain is crucial; already the insertion of one additional CH₂ unit deprives the resulting analogue of any useful antiviral activity [21, 25]; indeed, the six-membered chelate-ring analogue of equilibrium 1 is considerably less stable with metal ions like Mg²⁺ and Zn²⁺ or does even not form at all (Fernández-Botello A, Griesser R, Holý A, Moreno V, Sigel H, to be published).

With the mentioned observations in mind and with the aim to delineate further the structure–function relationship, we studied the metal ion-binding properties of the PMEA analogue 9-[2-(phosphonmethoxy)ethyl]-8-azaadenine (9,8aPMEA) and its isomer 8-[2-(phosphonmethoxy)ethyl]-8-azaadenine (8,8aPMEA) (see Fig. 1). This nucleobase modification, i.e. the replacement of (C8)H by N [26], was initiated by the more than 50-years-old discovery of the antibacterial [27] and antitumor [28] properties of 8-azaguanine, which has led to a continued interest in 8-azapurines and related nucleosides [29, 30, 31, 32]. Interestingly, the present study reveals that the metal ion-binding properties of 9,8aPMEA²⁻ are very similar to those of PMEAs²⁻ whereas the ones of 8,8aPMEA²⁻ differ, and these differences are also reflected in their biological activity [26, 33].

A further point to be emphasized is that the presence of the ether oxygen atom (Fig. 1) in all three nucleotide analogues, i.e. PMEAs²⁻ [22, 23, 34, 35, 36], 8,8aPMEAs²⁻ and 9,8aPMEAs²⁻, gives rise to the already mentioned equilibrium 1 with all the metal ion complexes studied. This property distinguishes the complexes of these analogues from those of their parent nucleotide AMP²⁻, which does not have this possibility and therefore forms either simply complexes with the phosphate residue or, as in the case of several divalent 3d metal ions, in addition macrochelates involving N7 [37, 38, 39, 40]; this situation is indicated in a simplified way in the intramolecular equilibrium 2:



As we shall see below, for the M(8,8aPMEA) complexes, equilibrium 1 is of relevance, whereas for some of the

M(9,8aPMEA) species the situation is considerably more complicated [41] and both equilibria 1 and 2 are to some extent of relevance.

Materials and methods

Materials

Twofold protonated 9-[2-(phosphonmethoxy)ethyl]-8-azaadenine, i.e. H₂(9,8aPMEA)[±], and its 8-isomer, 8-[2-(phosphonmethoxy)ethyl]-8-azaadenine, H₂(8,8aPMEA)[±], were synthesized by alkylation of 8-azaadenine with a synthon carrying the structural constituents of the required side chain [26]. For the present study, compounds from the same lots used previously [41, 42] were applied. The aqueous solutions of the ligands were freshly prepared daily just before the experiments by dissolving the substance in deionized, ultrapure (MILLI-Q185 PLUS; from Millipore, Molsheim, France) CO₂-free water, adjusted to pH about 8.5 by adding 2 equiv of 0.1 NaOH.

The disodium salt of 1,2-diaminoethane-*N,N,N',N'*-tetraacetic acid (Na₂H₂EDTA), potassium hydrogen phthalate, HNO₃, NaOH (Titrisol), and the nitrate salts of Na⁺, Mg²⁺, Ca²⁺, Sr²⁺, Ba²⁺, Mn²⁺, Co²⁺, Ni²⁺, Cu²⁺, Zn²⁺ and Cd²⁺ (all pro analysi) were from Merck (Darmstadt, Germany). All solutions for the potentiometric pH titrations were prepared with the mentioned ultrapure CO₂-free water. The buffer solutions (pH 4.00, 7.00, 9.00, based on the NBS scale; now NIST) for calibration were from Metrohm (Herisau, Switzerland).

The exact concentrations of the stock solutions of the divalent metal ions were determined by potentiometric pH titrations via their EDTA complexes by measuring the equivalents of protons liberated from H(EDTA)²⁻ upon complex formation with M²⁺. The exact concentration of the ligand solutions was newly determined for each experiment by evaluation of the corresponding titration pairs, i.e. the differences in NaOH consumption between solutions with and without ligand (see below).

Potentiometric pH titrations

The pH titration curves for the determination of the equilibrium constants in aqueous solution were recorded with a Metrohm E536 potentiograph connected to a Metrohm E655 dosimat and a Metrohm 6.0222.100 combined macro glass electrode. The pH calibration of the instrument was carried out with the mentioned buffer solutions of pH 4.00, 7.00 and 9.00. The titer of the NaOH used was determined with potassium hydrogen phthalate.

The direct pH meter readings were used in the calculations of the acidity constants, i.e. these constants determined at *I*=0.1 M (NaNO₃) and 25 °C are so-called practical, mixed or Brønsted constants [43].

They may be converted into the corresponding concentration constants by subtracting 0.02 from the listed pK_a values; this conversion term contains both the junction potential of the glass electrode and the hydrogen ion activity [43, 44]. It should be emphasized that the ionic product of water (K_w) and the mentioned conversion term do not enter into our calculation procedure because we always evaluate the differences in NaOH consumption between a pair of solutions, i.e. with and without ligand. The stability constants determined are concentration constants.

All equilibrium constants were calculated by curve-fitting procedures in the way and with the equipment described previously [45, 46].

Determination of equilibrium constants

The acidity constants $K_{H_2(9,8aPMEA)}^H$ (Eq. 4b) and $K_{H(9,8aPMEA)}^H$ (Eq. 5b) of $H_2(9,8aPMEA)^\pm$, where one proton is at N1 of the adenine moiety and the other at the phosphonate group, were determined recently [41] and these results were also confirmed now (25 °C; $I=0.1$ M, $NaNO_3$). The stability constants $K_{M(H;9,8aPMEA)}^M$ (Eq. 6b) and $K_{M(9,8aPMEA)}^M$ (Eq. 7b) of the $M(H;9,8aPMEA)^+$ and $M(9,8aPMEA)$ complexes, respectively, were determined under the same conditions as the acidity constants [41]. This means that 30 mL of aqueous 0.83 mM HNO_3 were titrated in the presence and absence of 0.4 mM deprotonated ligand under N_2 with 1 mL of 0.03 M NaOH (the differences in NaOH consumption between such a pair of titrations were used for the calculations), but $NaNO_3$ was partly or fully replaced by $M(NO_3)_2$ (25 °C; $I=0.1$ M). The M^{2+} /ligand ratios were 83:1 (Mg^{2+} , Ca^{2+} , Sr^{2+} , Ba^{2+} , Mn^{2+}), 75:1 (Mg^{2+}), 73.4:1 (Mn^{2+}), 55.6:1 (Mg^{2+} , Ca^{2+}), 52:1 (Ca^{2+}), 50:1 (Mn^{2+} , Co^{2+}), 49:1 (Co^{2+}), 41.7:1 (Sr^{2+} , Ba^{2+}), 28:1 (Cd^{2+}), 25:1 (Co^{2+}), 14:1 (Cd^{2+}), 13:1 (Cd^{2+}), 11:1 (Cu^{2+}) and 5.5:1 (Cu^{2+}).

The acidity constants $K_{H_2(8,8aPMEA)}^H$ (Eq. 4b) and $K_{H(8,8aPMEA)}^H$ (Eq. 5b) of $H_2(8,8aPMEA)^\pm$ were determined by titrating 30 mL of aqueous 1.27 mM HNO_3 in the presence and absence of 0.4 mM deprotonated ligand with 1.3 mL of 0.03 M NaOH (25 °C; $I=0.1$ M, $NaNO_3$). The differences in NaOH consumption between such a pair of titrations were used in the calculations. The pH range from 3.6 to 8.0 was evaluated, which corresponds to an initial formation degree of 48% for $H_2(8,8aPMEA)^\pm$ and to a final deprotonation degree of 94% for $8,8aPMEA^{2-}$. The results for the two acidity constants are the averages of 13 pairs of independent titrations.

The stability constants $K_{M(H;8,8aPMEA)}^M$ (Eq. 6b) and $K_{M(8,8aPMEA)}^M$ (Eq. 7b) of the $M(H;8,8aPMEA)^+$ and $M(8,8aPMEA)$ complexes were determined under the same conditions as the acidity constants, but $NaNO_3$ was partly or fully replaced by $M(NO_3)_2$ (25 °C;

$I=0.1$ M). The M^{2+} /ligand ratios were 83:1 (Mg^{2+} , Ca^{2+} , Sr^{2+} , Ba^{2+} , Mn^{2+}), 55.6:1 (Mg^{2+} , Ca^{2+}), 50:1 (Co^{2+} , Ni^{2+}), 41.7:1 (Sr^{2+} , Ba^{2+} , Mn^{2+}), 28:1 (Zn^{2+} , Cd^{2+}), 25:1 (Co^{2+} , Ni^{2+}), 14:1 (Cd^{2+}), 11:1 (Cu^{2+} , Zn^{2+}) and 5.5:1 (Cu^{2+}).

The stability constants were calculated by considering H^+ , $H_2(PA)^\pm$, $H(PA)^-$, PA^{2-} , M^{2+} , $M(H;PA)^+$ and $M(PA)$, where $PA^{2-}=9,8aPMEA$ or $8,8aPMEA^{2-}$. However, the formation of the monoprotonated $M(H;PA)^+$ complexes was usually rather small and therefore values for the corresponding constants had to be estimated. These estimations were based on the relationship between complex stability and ligand donor group basicity [47], i.e. $\log K_{\text{stability}}$ versus pK_a results in a straight line for families of closely related ligands and their complexes. This line is defined by the equation $y=mx+b$, where x represents the pK_a value, y the logarithm of the calculated stability constant of the corresponding M^{2+} -ligand complex, and m the slope of the straight line, b being the intercept with the y -axis. For the slope we used the average of the straight-line equations defined previously for benzimidazole- [48], imidazole- [49] and pyridine-type [50] ligands and their corresponding complexes. The stability constants for $M(H;PA)^+$ species were provided by two structurally related adenine derivatives, i.e. 9-[2-(phosphonomethoxy)ethyl]adenine (PMEA; $pK_{H_2(PMEA)}^H=4.16$ [51]) and 9-[2-(phosphonomethoxy)ethyl]-2,6-diaminopurine (PMEDAP; $pK_{H_2(PMEDAP)}^H=4.82$ [45]); the corresponding stability constant values have been tabulated [45, 51]. Since the $pK_{H_2(PA)}^H$ values are known, \log stability constants for $M(H;PA)^+$ complexes could be calculated from the PMEAs as well as from the PMEDAP data; the average of these two results obtained for each metal ion was then used in the final calculations.

For the 9,8aPMEA systems the experimental data were collected every 0.1 pH unit, beginning at about 1.5% of complex formation for $M(H;9,8aPMEA)^+$ to a neutralization degree of about 90% with respect to the species $H(9,8aPMEA)^-$ or until the beginning of the hydrolysis of $M(aq)^{2+}$, which was evident from the titrations without ligand. The maximal formation degree for the $M(H;9,8aPMEA)^+$ and $M(9,8aPMEA)$ species varies between 1.6–8.7% and 25.8–89.9%, respectively, depending on the metal ion considered. For the corresponding complexes with Ni^{2+} , Cu^{2+} and Zn^{2+} , reference [41] should be consulted.

For the 8,8aPMEA systems the experimental data were also collected every 0.1 pH unit from a formation degree of about 0.9% for the $M(H;8,8aPMEA)^+$ complex to a neutralization degree of about 90% with regard to the $H(8,8aPMEA)^-$ species or until the beginning of the hydrolysis of $M(aq)^{2+}$, which was again evident from the titrations in the absence of ligand. The maximal formation degree for the $M(H;8,8aPMEA)^+$ and $M(8,8aPMEA)$ complexes varied between 1.5–12.4% and 22.8–89.5%, respectively.

It needs to be emphasized that the results showed no dependence on the excess of metal ion concentration employed in the various experiments. The final results for the stability constants of the complexes listed below are the averages of at least five independent titrations for each system.

Results and discussion

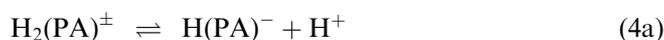
It is well known that nucleobases and their derivatives can undergo self-association via π stacking [11, 52, 53]. Therefore, the experimental conditions for the determination of the equilibrium constants by potentiometric pH titrations (25 °C; $I=0.1$ M, NaNO_3) were selected such that the results summarized below refer to monomeric species. This is ascertained with ligand concentrations of 0.4 mM, as has been shown previously for PME A [51].

Acidity constants of $\text{H}_3(8,8\text{aPME A})^+$ and $\text{H}_3(9,8\text{aPME A})^+$

The ligands $8,8\text{aPME A}^{2-}$ and $9,8\text{aPME A}^{2-}$, abbreviated as PA^{2-} , may accept three protons, two at the phosphonate group and one at the N1 site of the adenine residue [13] (see Fig. 1). Further protonations of the adenine residue are possible at N7 and N3, but these protons are released at $\text{p}K_{\text{a}} < 0$ [54, 55], and therefore they are not considered in this study. At $\text{pH} > 0$ the strongest acid that can be derived in aqueous solution from PA^{2-} is $\text{H}_3(\text{PA})^+$. Hence, the following three deprotonation reactions need to be considered:



$$K_{\text{H}_3(\text{PA})}^{\text{H}} = [\text{H}_2(\text{PA})^{\pm}][\text{H}^+]/[\text{H}_3(\text{PA})^+] \quad (3\text{b})$$



$$K_{\text{H}_2(\text{PA})}^{\text{H}} = [\text{H}(\text{PA})^-][\text{H}^+]/[\text{H}_2(\text{PA})^{\pm}] \quad (4\text{b})$$



$$K_{\text{H}(\text{PA})}^{\text{H}} = [\text{PA}^{2-}][\text{H}^+]/[\text{H}(\text{PA})^-] \quad (5\text{b})$$

However, the first proton according to equilibrium 3a is released from the $\text{P}(\text{O})(\text{OH})_2$ group of $\text{H}_3(\text{PA})^+$, and from the results obtained previously for $\text{H}_3(\text{PME A})^+$ it is known that $\text{p}K_{\text{H}_3(\text{PME A})}^{\text{H}} = 1.22 \pm 0.13$ [13]. The same value may be surmised for the corresponding acidity constant of $\text{H}_3(8,8\text{aPME A})^+$ and $\text{H}_3(9,8\text{aPME A})^+$. Hence, for the present study, for which all potentiometric pH titrations were carried out in the pH range above 3 (see ‘‘Determination of equilibrium constants’’ and [41, 42]), equilibrium 3a is actually also not of

relevance. Here, only the release of the protons from $\text{H}_2(\text{PA})^{\pm}$ need to be considered, i.e. the first one from the (N1) H^+ site (Eq. 4a) and the next one from the $\text{P}(\text{O})_2(\text{OH})^-$ group (Eq. 5a). The corresponding acidity constants are listed in Table 1, together with some related data [56, 57, 58, 59, 60, 61, 62].

The data in Table 1 offer comparisons for many conclusions; a few are given below:

1. Comparison of entries 1 and 2 with entry 3 confirms the conclusion indicated above that the first proton in $\text{H}_2(9,8\text{aPME A})^{\pm}$ is released from the (N1) H^+ site of the 8-azaadenine residue [56] and the second one from the monoprotonated phosphonate group.
2. Similarly, the combination of the values from entries 6 and 2 gives those of entry 5 and confirms the corresponding site attributions of the protons in $\text{H}_2(\text{PME A})^{\pm}$ [13, 51].
3. Comparison of entries 1 and 3 with 5 and 6 demonstrates that replacement of (C8)H by N acidifies the (N1) H^+ site by about 1.4 p K units; however, this substitution at position 8 of the purine ring has no remarkable effect on the basicity of the phosphonate group.
4. Similarly, the phosphonate group remains largely unaffected if the aliphatic side chain with the ether oxygen atom is moved from position 9 to 8 of the purine ring but the (N1) H^+ site is now acidified only by $\Delta \text{p} K_{\text{a}} = 0.6$, as a comparison of entries 4 and 5 demonstrates.

Table 1 Negative logarithms of the acidity constants of $\text{H}_2(\text{PA})^{\pm}$, where $\text{PA}^{2-} = 8,8\text{aPME A}^{2-}$ or $9,8\text{aPME A}^{2-}$ (Eqs. 4a,b and 5a,b), as determined by potentiometric pH titrations in aqueous solution at 25 °C and $I=0.1$ M (NaNO_3), together with the corresponding values of some related systems^{a,b}

No	Protonated species ^c	$\text{p}K_{\text{H}_2(\text{PA})}^{\text{H}}$ (N1) H^+	$\text{p}K_{\text{H}(\text{PA})}^{\text{H}}$ $\text{P}(\text{O})_2(\text{OH})^-$	Ref
1	$\text{H}(9\text{Me}8\text{azaAde})^+$	2.70 ^d		[56]
2	$\text{H}(\text{PME-R})^-$		6.99 ± 0.04^e	$-\frac{e}{-}$
3	$\text{H}_2(9,8\text{aPME A})^{\pm}$	2.73 ± 0.02	6.85 ± 0.02	$-\frac{f,g}{-}$
4	$\text{H}_2(8,8\text{aPME A})^{\pm}$	3.56 ± 0.02	6.79 ± 0.01	$-\frac{f}{-}$
5	$\text{H}_2(\text{PME A})^{\pm}$	4.16 ± 0.02	6.90 ± 0.01	[13, 51]
6	$\text{H}(9\text{MeAde})^+$	4.10 ± 0.01		[54, 55, 59]
7	$\text{H}_2(\text{dPME A})^{\pm}$	4.17 ± 0.02	7.69 ± 0.01	[46]
8	$\text{CH}_3\text{P}(\text{O})_2(\text{OH})^-$		7.51 ± 0.01	[60, 61]
9	$\text{H}_2(\text{AMP})^{\pm}$	3.84 ± 0.02	6.21 ± 0.01	[37, 62]

^aThe error limits given are three times the standard error of the mean value or the sum of the probable systematic errors, whichever is larger

^bSo-called practical, mixed or Brønsted constants are listed [43]

^c9Me8azaAde = 9-methyl-8-azaadenine; 9MeAde = 9-methyladenine

^dValue based on ^1H NMR shift experiments ($I=0.5$ M, KNO_3) [56]; a very similar result was obtained in an early spectrophotometric study [57]

^eAverage of the values due to (phosphonomethoxy)ethane (PME^{2-} ; $\text{R} = \text{H}$) [51] and 1-[2-(phosphonomethoxy)ethyl]cytosine (PMEC^{2-} ; $\text{R} = \text{cytosine residue}$) [58]

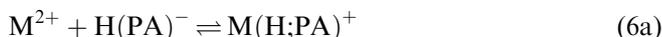
^fThis study

^gThe values given in [41] are now also confirmed

5. On the other hand, replacement of the ether oxygen by a methylene group in the alkyl chain has no effect on the basicity of N1 (see entries 5 and 7), but it increases the basicity of the phosphonate group by about 0.8 p*K* units. This conclusion is further confirmed by comparing entries 2–5 with 7 and 8.
6. Of further interest is a comparison of the values for H₂(AMP)[±] (entry 9) with some other data. It is evident that the (N1)H⁺ site in H₂(AMP)[±] is only a bit more acidic than the same site in H(9MeAde)⁺ or H₂(PMEA)[±] (entries 5, 6). This means that the replacement of the ribosyl residue in AMP by the ether oxygen-containing aliphatic chain has little effect on the aromatic ring system. However, the phosphate residue, -OP(O)₂(OH)⁻, of H(AMP)⁻ is by about 0.7 p*K* units more acidic than the phosphonate group, -P(O)₂(OH)⁻ (entries 5, 9). This means that the phosphate group is practically completely deprotonated at physiological pH (ca. 7.5), whereas the phosphonate group still carries a proton to some extent (about 20%) [13].

Stability constants of M(H;PA)⁺ and M(PA) complexes for PA²⁻ = 8,8aPMEA²⁻ and 9,8aPMEA²⁻

The experimental data of the potentiometric pH titrations of all the M²⁺/PA systems can be described completely by equilibria 4a, 5a, 6a and 7a,



$$K_{M(H;PA)}^M = [M(H;PA)^+]/([M^{2+}][H(PA)^-]) \quad (6b)$$



$$K_{M(PA)}^M = [M(PA)]/([M^{2+}][PA^{2-}]) \quad (7b)$$

if the evaluation is not carried into the pH range where hydroxo complexes form. The acidity constant of equilibrium 8a may be calculated with Eq. 9:



$$K_{M(H;PA)}^H = [M(PA)][H^+]/[M(H;PA)^+] \quad (8b)$$

$$pK_{M(H;PA)}^H = pK_{H(PA)}^H + \log K_{M(H;PA)}^M - \log K_{M(PA)}^M \quad (9)$$

The results are listed in Table 2; the stability constants given for the M(H;PA)⁺ complexes are only estimates since the formation degree of these species was low (see ‘‘Determination of equilibrium constants’’). The stability constants of the M(PA) complexes show the usual trends. For the alkaline earth ions the stability decreases with increasing ionic radii, indicating that metal ion binding at the phosphonate group is (at least) in part inner-sphere. For the

Table 2 Logarithms of the stability constants of the M(H;PA)[±] (Eq. 6a,b) and M(PA) complexes (Eq. 7a,b), where PA²⁻ = 8,8aPMEA²⁻ or 9,8aPMEA²⁻, as determined by potentiometric pH titrations, together with the negative logarithms of the acidity constants of the monoprotonated M(H;PA)⁺ complexes (Eqs. 8a,b and 9) (aqueous solution; 25 °C; I = 0.1 M, NaNO₃)^{a,b}

PA ²⁻	M ²⁺	log K _{M(H;PA)} ^M ^c	log K _{M(PA)} ^M	pK _{M(H;PA)} ^H
8,8aPMEA ²⁻	Mg ²⁺	0.2	1.83 ± 0.07	5.16 ± 0.26
	Ca ²⁺	0.15	1.59 ± 0.03	5.35 ± 0.25
	Sr ²⁺	0.1	1.37 ± 0.03	5.52 ± 0.25
	Ba ²⁺	0.0	1.34 ± 0.04	5.45 ± 0.25
	Mn ²⁺	0.3	2.45 ± 0.02	4.64 ± 0.25
	Co ²⁺	0.55	2.28 ± 0.04	5.06 ± 0.25
	Ni ²⁺	0.9	2.27 ± 0.03	5.42 ± 0.25
	Cu ²⁺	1.3	3.68 ± 0.06	4.41 ± 0.26
	Zn ²⁺	1.0	2.64 ± 0.09	5.15 ± 0.27
	Cd ²⁺	1.0	2.89 ± 0.08	4.90 ± 0.26
9,8aPMEA ²⁻	Mg ²⁺	0.2	1.84 ± 0.04	5.21 ± 0.25
	Ca ²⁺	0.15	1.62 ± 0.08	5.38 ± 0.26
	Sr ²⁺	0.1	1.41 ± 0.04	5.54 ± 0.25
	Ba ²⁺	0.0	1.38 ± 0.05	5.47 ± 0.26
	Mn ²⁺	0.15	2.49 ± 0.04	4.51 ± 0.25
	Co ²⁺	0.4	2.33 ± 0.04	4.92 ± 0.25
	Ni ²⁺	0.7	2.25 ± 0.08	5.30 ± 0.26
	Cu ²⁺	0.95	3.98 ± 0.04	3.82 ± 0.25
	Zn ²⁺	0.8	2.82 ± 0.09	4.83 ± 0.27
	Cd ²⁺	0.75	2.93 ± 0.06	4.67 ± 0.26

^aFor the error limits, see footnote (a) of Table 1. The error limits (3σ) of the derived data, in the present case for column 5, were calculated according to the error propagation after Gauss

^bThe values for the Ni²⁺, Cu²⁺ and Zn²⁺ systems of 9,8aPMEA are taken from [41]. The titrations with Zn²⁺ were hampered by precipitation, i.e. the pH range suitable for the evaluation of the stability constants was restricted [41]

^cThe stability constants listed for the M(H;PA)⁺ complexes as well as the error limits (±0.25 log units) are estimates

divalent 3d metal ions the long-standing experience [51, 63] is confirmed that the stabilities of phosph(on)ate–metal ion complexes often do not strictly follow [34, 38, 39, 45, 62, 64, 65, 66] the Irving–Williams sequence [67], an observation in accord with the fact that in ligands of this kind the phosph(on)ate group is always the main binding site [35, 36, 38, 39, 62, 65] in M(PA) complexes (see the next section and Fig. 2 below).

As far as the M(H;PA)⁺ complexes are concerned, it is evident that the evaluation of potentiometric pH titration data only allows the determination, or in the present case estimation, of their stability constants. Further information is required to detect the binding sites of the proton and the metal ion. At first, one may ask where the proton is located because binding of a metal ion to a protonated ligand commonly leads to an acidification of the ligand-bound proton [68, 69]. Indeed, the acidity constants of the M(H;PA)⁺ complexes given in column 5 of Table 2 are about 1.3–3 p*K* units smaller than pK_{H(PA)}}^H (Table 1), but about 0.9–2.8 log units larger than pK_{H₂(PA)}}^H. This comparison shows that the proton in M(H;PA)⁺ is clearly located at the phosphonate group; hence, one may tentatively

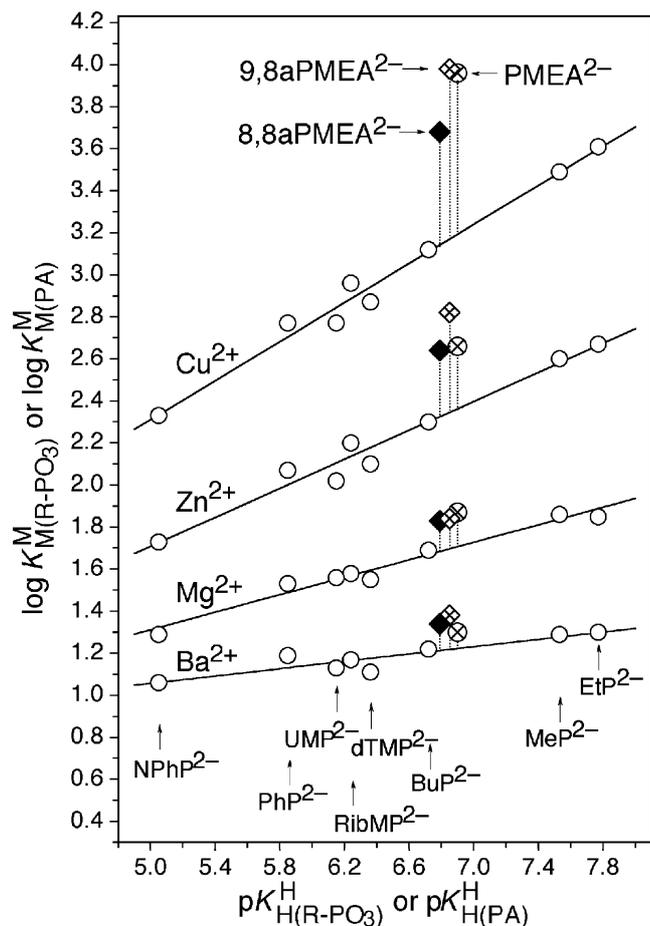


Fig. 2 Evidence for an enhanced stability of several $M(9,8aPMEA)$ (crossed diamonds), $M(8,8aPMEA)$ (solid diamonds) and $M(PMEA)$ (crossed circles) complexes, based on the relationship between $\log K_{M(R-PO_3)}^M$ and $pK_{H(R-PO_3)}^H$ for $M(R-PO_3)$ complexes of some simple phosphate monoester and phosphonate ligands ($R-PO_3^{2-}$) (open circles): 4-nitrophenyl phosphate ($NPhP^{2-}$), phenyl phosphate (PhP^{2-}), uridine 5'-monophosphate (UMP^{2-}), D-ribose 5-monophosphate ($RibMP^{2-}$), thymidine [$=1-(2'-deoxy-\beta-D-ribofuranosyl)thymine$] 5'-monophosphate ($dTMP^{2-}$), *n*-butyl phosphate (BuP^{2-}), methanephosphonate (MeP^{2-}) and ethanephosphonate (EtP^{2-}) (from left to right). The least-squares lines (Eq. 10) are drawn through the corresponding eight data sets (open circles) taken from [73] for the phosphate monoesters and from [51] for the phosphonates. The points due to the equilibrium constants for the $M^{2+}/9,8aPMEA$ (crossed diamonds) and $M^{2+}/8,8aPMEA$ (solid diamonds) systems are based on the values listed in Tables 1 and 2; those for the $M^{2+}/PMEA$ systems (crossed circles) are from [51]. The vertical broken lines emphasize the stability differences from the reference lines; they equal $\log \Delta_{M/9,8aPMEA}$ and $\log \Delta_{M/8,8aPMEA}$, as defined in Eqs. 11a and 11b, for the $M(9,8aPMEA)$ and $M(8,8aPMEA)$ complexes, respectively. All the plotted equilibrium constants refer to aqueous solutions at 25 °C and $I=0.1$ M ($NaNO_3$)

assume that the metal ion is bound preferentially to the nucleobase, since a monoprotonated phosphonate group is only a weak binding site. Indeed, this suggestion agrees with evidence obtained previously for other related $M(H;PA)^+$ species [35, 36, 51]. It is further supported by the observation that the estimated stabilities of the

$M(H;PA)^+$ complexes follow the Irving–Williams sequence [67], as is typical for metal ion binding to nitrogen donor sites [63], the preferred site most likely being N7 [70, 71].

Evidence for an enhanced stability of the $M(8,8aPMEA)$ and $M(9,8aPMEA)$ complexes

The $8,8aPMEA^{2-}$ and $9,8aPMEA^{2-}$ ligands offer four potential binding sites for the coordination of metal ions: the two-fold negatively charged phosphonate group, the ether oxygen of the $-CH_2CH_2-O-CH_2-PO_3^{2-}$ chain (see Fig. 1) and the adenine residue with its N7 and N3 sites; the N1 is not accessible by a phosphonate-bound metal ion [34, 51]. The phosphonate group is clearly the primary binding site for all metal ions considered in this study and therefore any participation in metal ion binding of one (or more) of the other potential sites has to be reflected in a relative stability increase [47]. Hence, it is necessary to define the stability of a pure PO_3^{2-}/M^{2+} interaction. This can be done by applying the previously defined [34, 65, 72] straight-line correlations which are based on $\log K_{M(R-PO_3)}^M$ versus $pK_{H(R-PO_3)}^H$ plots for simple phosphate monoesters [73] and phosphonates [51]; these ligands are abbreviated as $R-PO_3^{2-}$, where R represents a non-coordinating residue. The parameters for the corresponding straight-line equations, which are defined by Eq. 10,

$$\log K_{M(R-PO_3)}^M = m \times pK_{H(R-PO_3)}^H + b \quad (10)$$

have been tabulated [34, 51, 65, 72], i.e. the slopes m and the intercepts b with the y axis. Hence, with a known pK_a value for the deprotonation of a $P(O)_2(OH)^-$ group an expected stability constant can be calculated for any phosph(on)ate–metal ion complex.

Corresponding plots of $\log K_{M(R-PO_3)}^M$ versus $pK_{H(R-PO_3)}^H$ according to Eq. 10 are shown in Fig. 2 for 1:1 complexes of Ba^{2+} , Mg^{2+} , Zn^{2+} and Cu^{2+} , as examples, with the data points (open circles) of the eight simple ligand systems used [51, 73] for the determination of the straight reference lines [51]. The data points for the M^{2+} complexes of $PMEA^{2-}$ and its 8-aza analogues are for all four metal ion systems clearly above the reference lines, thus proving an increased stability for all 12 complexes considered though the stability enhancements for the Mg^{2+} and Ba^{2+} complexes are relatively small.

Furthermore, it is evident that the stability increase for the Mg^{2+} and Ba^{2+} complexes is within the error limits identical for all three ligands and this indicates that here equilibrium 1 is of relevance, since the affinity of the alkali earth metal ions toward N sites is small [63]. This observation contrasts with that made for the Cu^{2+} and Zn^{2+} complexes since their stability enhancement evidently differs for $8,8aPMEA^{2-}$ and

9,8aPMEA²⁻, indicating that here at least to some extent the adenine residue is (also) involved in metal ion binding. This is confirmed by the data points for the Cu²⁺ complexes of PME²⁻ and 9,8aPMEA²⁻, which indicate the same stability enhancement, and for Cu(PMEA) the involvement in metal ion binding of the adenine residue has been proven. The apparent difference for the corresponding Zn²⁺ complexes should not be interpreted, since the stability constant for Zn(PMEA) could not be measured due to precipitation [51]; it is an estimate only [22, 23] and, in fact, the corresponding stability constants are within their (large) error limits identical (see below).

Stability enhancements like those seen in Fig. 2 can be quantified by the differences between the experimentally (exptl) measured stability constants and those calculated (calcd) according to Eq. 10; this difference is defined in Eq. 11a,b:

$$\log \Delta_{M/PA} = \log K_{M(PA)exptl}^M - \log K_{M(PA)calcd}^M \quad (11a)$$

$$= \log K_{M(PA)}^M - \log K_{M(PA)op}^M (= \log \Delta) \quad (11b)$$

where the expressions $\log K_{M(PA)calcd}^M$ and $\log K_{M(PA)op}^M$ are synonymous because the calculated value equals the stability constant of the “open” isomer, M(PA)_{op}

Table 3 Stability constant comparisons for the M(PA) complexes, where PA²⁻ = 8,8aPMEA²⁻ or 9,8aPMEA²⁻, according to Eq. 11a,b; that is, between the experimentally measured (exptl) and the calculated (calcd) log stability constants, the latter being based on the reference-line equations (Eq. 10) [34, 51, 72] and the pK_{H(PA)}^H values (Table 1) of the monoprotonated H(PA)⁻ species (aqueous solution; 25 °C; I = 0.1 M, NaNO₃)^a

PA ²⁻	M ²⁺	log K _{M(PA)} ^M		log Δ _{M/PA}
		exptl ^b	calcd	
8,8aPMEA ²⁻	Mg ²⁺	1.83 ± 0.07	1.68 ± 0.03	0.15 ± 0.08
	Ca ²⁺	1.59 ± 0.03	1.53 ± 0.05	0.06 ± 0.06
	Sr ²⁺	1.37 ± 0.03	1.29 ± 0.04	0.08 ± 0.05
	Ba ²⁺	1.34 ± 0.04	1.21 ± 0.04	0.13 ± 0.06
	Mn ²⁺	2.45 ± 0.02	2.30 ± 0.05	0.15 ± 0.05
	Co ²⁺	2.28 ± 0.04	2.07 ± 0.06	0.21 ± 0.07
	Ni ²⁺	2.27 ± 0.03	2.09 ± 0.05	0.18 ± 0.06
	Cu ²⁺	3.68 ± 0.06	3.14 ± 0.06	0.54 ± 0.08
	Zn ²⁺	2.64 ± 0.09	2.33 ± 0.06	0.31 ± 0.11
	Cd ²⁺	2.89 ± 0.08	2.63 ± 0.05	0.26 ± 0.09
	9,8aPMEA ²⁻	Mg ²⁺	1.84 ± 0.04	1.70 ± 0.03
Ca ²⁺		1.62 ± 0.08	1.53 ± 0.05	0.09 ± 0.09
Sr ²⁺		1.41 ± 0.04	1.29 ± 0.04	0.12 ± 0.06
Ba ²⁺		1.38 ± 0.05	1.22 ± 0.04	0.16 ± 0.06
Mn ²⁺		2.49 ± 0.04	2.31 ± 0.05	0.18 ± 0.06
Co ²⁺		2.33 ± 0.04	2.08 ± 0.06	0.25 ± 0.07
Ni ²⁺		2.25 ± 0.08	2.10 ± 0.05	0.15 ± 0.09
Cu ²⁺		3.98 ± 0.04	3.17 ± 0.06	0.81 ± 0.07
Zn ²⁺		2.82 ± 0.09	2.35 ± 0.06	0.47 ± 0.11
Cd ²⁺		2.93 ± 0.06	2.65 ± 0.05	0.28 ± 0.08

^aFor the error limits, see footnote (a) of Table 2

^bFrom column 4 of Table 2

(see, for example, equilibria 1 and 2), in which only a PO₃²⁻/M²⁺ interaction occurs. In columns 3–5 of Table 3 the values for the three terms of Eq. 11a,b are listed. All log Δ_{M/PA} values being positive proves that at least one further binding site, next to the phosphonate group, must be involved in all these M(PA) complexes.

Evidence for an ether oxygen interaction in the M(PA) complexes and quantification of the corresponding intramolecular equilibrium

It is obvious that the vertical distance from the marked data points in Fig. 2 for the M(PA) complexes to their reference lines reflects the “intensity” of the participation of any other binding site, next to the phosphonate group, in metal ion binding and that this “intensity” is quantified by Eqs. 11a,b. To learn which of the above-mentioned additional binding sites (see also Fig. 1) is responsible for the observed stability enhancements, we have listed in column 2 of Table 4 the log Δ_{M/PME-R} values (according to Eqs. 11a,b) which are solely attributable to the formation of five-membered chelates as seen in equilibrium 1 because the ligand PME-R²⁻ does not offer any other binding site except the ether oxygen, aside from the phosphonate group itself (see Fig. 1). The corresponding data log Δ_{M/8,8aPMEA} and log Δ_{M/9,8aPMEA} for the M(8,8aPMEA) and M(9,8aPMEA) complexes [=M(PA)] are given in columns 3 and 4 of Table 4, respectively, and a comparison with those in column 2 is best done by calculating the differences according to Eq. 12:

$$\Delta \log \Delta_{M/PA/PME-R} = \log \Delta_{M/PA} - \log \Delta_{M/PME-R} \quad (12)$$

The corresponding results are listed in columns 6 and 7 for 8,8aPMEA²⁻ and 9,8aPMEA²⁻, respectively. It is obvious that all the values in column 6 are zero within the error limits, meaning that the stability enhancements log Δ_{M/8,8aPMEA} correspond to those of log Δ_{M/PME-R} and that therefore equilibrium 1 operates in all the M(8,8aPMEA) systems and that only a M²⁺-ether oxygen interaction is responsible for the increased complex stability. Exactly the same is true for the M(9,8aPMEA) systems, with two exceptions: in Cu(9,8aPMEA) and Zn(9,8aPMEA) a further interaction next to the one with the ether oxygen must occur, as is evident from the results listed in column 7.

Since it has been shown previously [41, 46] that in the Cu(PMEA) species not only equilibrium 1 with the ether oxygen interaction operates but that also the adenine residue is involved, mainly via N3 and to a small extent also via N7 [46], we have summarized the log Δ_{M/PMEA} values in column 5 of Table 4 and the differences according to Eq. 12 in column 8 at the

Table 4 Comparison of the stability enhancements according to Eq. 11a,b as observed for the M(8,8aPMEA) and M(9,8aPMEA) complexes (Table 3, column 5), with the corresponding values determined earlier for the related M(PME-R) [58] and M(PMEA) complexes [22, 23, 51] (aqueous solution; 25 °C; $I=0.1$ M, NaNO₃)^{a,b}

M ²⁺	log $\Delta_{M/PME-R}$	log $\Delta_{M/8,8aPMEA}$	log $\Delta_{M/9,8aPMEA}$	log $\Delta_{M/PMEA}$	Δ log $\Delta_{M/8,8aPMEA/PME-R}$	Δ log $\Delta_{M/9,8aPMEA/PME-R}$	Δ log $\Delta_{M/PMEA/PME-R}$
Mg ²⁺	0.16 ± 0.04	0.15 ± 0.08	0.14 ± 0.05	0.16 ± 0.05	-0.01 ± 0.09	-0.02 ± 0.06	0.00 ± 0.06
Ca ²⁺	0.12 ± 0.05	0.06 ± 0.06	0.09 ± 0.09	0.11 ± 0.07	-0.06 ± 0.08	-0.03 ± 0.10	-0.01 ± 0.09
Sr ²⁺	0.09 ± 0.05	0.08 ± 0.05	0.12 ± 0.06	0.07 ± 0.05	-0.01 ± 0.07	0.03 ± 0.08	-0.02 ± 0.07
Ba ²⁺	0.11 ± 0.05	0.13 ± 0.06	0.16 ± 0.06	0.08 ± 0.06	0.02 ± 0.08	0.05 ± 0.08	-0.03 ± 0.08
Mn ²⁺	0.19 ± 0.06	0.15 ± 0.05	0.18 ± 0.06	0.21 ± 0.08	-0.04 ± 0.08	-0.01 ± 0.08	0.02 ± 0.10
Co ²⁺	0.20 ± 0.06	0.21 ± 0.07	0.25 ± 0.07	0.28 ± 0.07	0.01 ± 0.09	0.05 ± 0.09	0.08 ± 0.09
Ni ²⁺	0.14 ± 0.07	0.18 ± 0.06	0.15 ± 0.09	0.30 ± 0.07	0.04 ± 0.09	0.01 ± 0.11	0.16 ± 0.10
Cu ²⁺	0.48 ± 0.07	0.54 ± 0.08	0.81 ± 0.07	0.77 ± 0.07	0.06 ± 0.11	0.33 ± 0.10	0.29 ± 0.10
Zn ²⁺	0.29 ± 0.07	0.31 ± 0.11	0.47 ± 0.11	0.30 ± 0.10 ^c	0.02 ± 0.13	0.18 ± 0.13	0.01 ± 0.12 ^c
Cd ²⁺	0.30 ± 0.05	0.26 ± 0.09	0.28 ± 0.08	0.33 ± 0.06	-0.04 ± 0.10	-0.02 ± 0.09	0.03 ± 0.08

^aFor the error limits, see footnote (a) of Table 2

^bThe values in columns 6, 7 and 8 for Δ log $\Delta_{M/8,8aPMEA/PME-R}$, Δ log $\Delta_{M/9,8aPMEA/PME-R}$ and Δ log $\Delta_{M/PMEA/PME-R}$ (Eq. 11a,b) result from the comparison between the stability enhancements

observed for the M(8,8aPME-R), M(9,8aPME-R) or M(PMEA) complexes and the ones for the M(PME-R) complexes, respectively ^cThis value is an estimate since the stability constant for the Zn(PMEA) complex could not be measured due to precipitation [51]

right. Indeed, all the Δ log $\Delta_{M/PMEA/PME-R}$ values are again zero within their error limits, pointing again to the importance of equilibrium 1, but there are also two exceptions, i.e. the Ni²⁺ and Cu²⁺ systems. In fact, the Δ log $\Delta_{Cu/PMEA/PME-R}$ value is identical with the one observed for Δ log $\Delta_{Cu/9,8aPMEA/PME-R}$; the meaning of this result will be discussed further below in the section “Cu(9,8aPMEA) and related systems. A four isomer problem”.

For the present we shall concentrate on those 18 M(PA) systems for which Δ log $\Delta_{M/PA/PME-R}$ is zero within the error limits (see Table 4, columns 6 and 7) and for which the intramolecular equilibrium 1 operates.

The dimensionless intramolecular equilibrium constant, $K_{I/O}$, is defined by Eq. 13,

$$K_{I/O} = [M(PA)_{cl/O}]/[M(PA)_{op}] \quad (13)$$

and values for $K_{I/O}$ can be calculated following known procedures [47, 51, 62, 65, 72], i.e. via Eq. 14:

$$K_{I/O} = 10^{\log \Delta} - 1 \quad (14)$$

Knowledge of $K_{I/O}$ then allows us, according to Eq. 15,

$$\% M(PA)_{cl/O} = 100 \times K_{I/O}/(1 + K_{I/O}) \quad (15)$$

to obtain the percentage of the closed isomers, M(PA)_{cl/O}, present in equilibrium 1, i.e. their formation degree. The corresponding results for the M(8,8aPMEA) and M(9,8aPMEA) complexes are summarized in Table 5, where the values for the 9,8aPMEA²⁻ systems with Cu²⁺ and Zn²⁺ are given in parentheses for the reasons outlined above.

It is remarkable to observe that for a given metal ion the formation degree of the five-membered chelate involving the ether oxygen according to equilibrium 1 is,

within the error limits, identical for the complexes with both ligands, i.e. 8,8aPMEA²⁻ and 9,8aPMEA²⁻. This means that the 8-azaadenine residue has no significant influence on the position of equilibrium 1 and that in all the complexes formed with Mg²⁺, Ca²⁺, Sr²⁺, Ba²⁺, Mn²⁺, Co²⁺, Ni²⁺ and Cd²⁺ these two ligands simply

Table 5 Extent of chelate formation according to equilibrium 1 as expressed by the dimensionless equilibrium constants $K_{I/O}$ (Eqs. 13 and 14) and the percentages of M(PA)_{cl/O} (Eq. 15), based on the log $\Delta_{M/PA}$ values listed in column 5 of Table 3 for the M(8,8aPMEA) and M(9,8aPMEA) systems (aqueous solution; 25 °C; $I=0.1$ M, NaNO₃)^a

PA ²⁻	M ²⁺	log $\Delta_{M/PA}$	$K_{I/O}$	% M(PA) _{cl/O}
8,8aPMEA ²⁻	Mg ²⁺	0.15 ± 0.08	0.41 ± 0.26	29 ± 13
	Ca ²⁺	0.06 ± 0.06	0.15 ± 0.16	13 ± 12
	Sr ²⁺	0.08 ± 0.05	0.20 ± 0.14	17 ± 10
	Ba ²⁺	0.13 ± 0.06	0.35 ± 0.19	26 ± 10
	Mn ²⁺	0.15 ± 0.05	0.41 ± 0.16	29 ± 8
	Co ²⁺	0.21 ± 0.07	0.62 ± 0.26	38 ± 10
	Ni ²⁺	0.18 ± 0.06	0.51 ± 0.21	34 ± 9
	Cu ²⁺	0.54 ± 0.08	2.46 ± 0.64	71 ± 5
	Zn ²⁺	0.31 ± 0.11	1.04 ± 0.52	51 ± 12
	Cd ²⁺	0.26 ± 0.09	0.82 ± 0.38	45 ± 11
9,8aPMEA ²⁻	Mg ²⁺	0.14 ± 0.05	0.38 ± 0.16	28 ± 8
	Ca ²⁺	0.09 ± 0.09	0.23 ± 0.26	19 ± 17
	Sr ²⁺	0.12 ± 0.06	0.32 ± 0.18	24 ± 10
	Ba ²⁺	0.16 ± 0.06	0.45 ± 0.20	31 ± 10
	Mn ²⁺	0.18 ± 0.06	0.51 ± 0.21	34 ± 9
	Co ²⁺	0.25 ± 0.07	0.78 ± 0.29	44 ± 9
	Ni ²⁺	0.15 ± 0.09	0.41 ± 0.29	29 ± 15
	Cu ²⁺	0.81 ± 0.07	(5.46 ± 1.04) ^b	(85 ± 2) ^b
	Zn ²⁺	0.47 ± 0.11	(1.95 ± 0.75) ^b	(66 ± 9) ^b
	Cd ²⁺	0.28 ± 0.08	0.91 ± 0.35	48 ± 10

^aFor the error limits, see footnote (a) of Table 2

^bThe parentheses indicate that the stability increase cannot solely be attributed to equilibrium 1 since in these systems a contribution from a nucleobase-metal ion interaction also exists [41]. See also the section “Cu(9,8aPMEA) and related systems. A four isomer problem”

behave as a PME-R^{2-} ligand, $\text{R-CH}_2\text{CH}_2\text{-O-CH}_2\text{-PO}_3^{2-}$, where R does neither participate in metal ion binding nor affect it in a negative sense, e.g. by steric inhibition.

Cu(9,8aPMEA) and related systems. A four isomer problem

The values for $\Delta \log \Delta_{\text{Cu/PA/PME-R}}$ due to the Cu(9,8aPMEA) and Cu(PMEA) complexes are identical (Table 4, columns 7 and 8) and in both instances the increased stability is clearly beyond that attributable to equilibrium 1 with the ether oxygen interaction and it is further clear that the additional interaction must be with the adenine residue. The additional stability increase due to $\Delta \log \Delta_{\text{Zn/9,8aPMEA/PME-R}}$ for the Zn(9,8aPMEA) system is clearly also beyond the error limits (Table 4, column 7). The apparent lack of such an increased stability for the Zn(PMEA) system (Table 4, column 8) is no surprise, as the data for this system are based on an estimation [22, 23, 51], though one should note that, owing to the relatively large error limits of the values for the Zn(PMEA) and Zn(9,8aPMEA) systems, the $\Delta \log \Delta_{\text{Zn/PA/PME-R}}$ values overlap, hence, the properties of the two systems are probably similar.

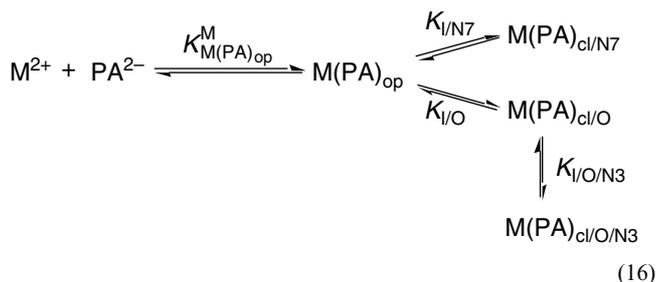
For the Ni(9,8aPMEA) and Ni(PMEA) systems (Table 4, columns 7 and 8) the differing situation is more complicated to rationalize, since there were no experimental difficulties. The $\Delta \log \Delta_{\text{Ni/PMEA/PME-R}}$ value for the latter system and, hence, a Ni^{2+} -adenine residue interaction, appears as being certain. The difference between these two systems, 0.15 ± 0.15 $[=(0.16 \pm 0.10) - (0.01 \pm 0.11)]$, seems also to be real, especially when one takes into account that the given error is based on 3σ ; the value for 1σ , i.e. 0.15 ± 0.05 , still corresponds to a confidence interval of about 70%. Hence, it appears that the Ni^{2+} -(N7) interaction is hampered in the Ni(9,8aPMEA) system, whereas in the Ni(PMEA) complex exactly this isomer occurs to a considerable formation degree [41]. It should be noted that in the Cu^{2+} and Zn^{2+} systems the macrochelated isomer involving N7 is a minority species (see below) and therefore no differences in the complexing

basicity of N7 in 9,8aPMEA $^{2-}$ which is due to the effect of the neighbouring N8 atom.

To complete the picture, the results regarding the previously established [41, 46] equilibrium scheme 16 need to be summarized shortly. The two isomeric species M(PA)_{op} and $\text{M(PA)}_{\text{cl/O}}$ have been discussed and defined above; they correspond to equilibrium 1. The species $\text{M(PA)}_{\text{cl/N7}}$ represents a third isomer and corresponds to the macrochelate seen in equilibrium 2 which is formed via N7. Finally, the fourth isomer in which not only a five-membered chelate involving the phosphate group and the ether oxygen exists, but in which in addition also a seven-membered chelate involving the N3 site is formed, is abbreviated as $\text{M(PA)}_{\text{cl/O/N3}}$.

The most thoroughly studied system of this kind is Cu(PMEA) [46]: the formation degrees for any Cu(PMEA) complex concentrations are $17 \pm 3\%$, $34 \pm 10\%$, $41 \pm 12\%$ and $7.7 \pm 5.3\%$ (errors 3σ) for $\text{Cu(PMEA)}_{\text{op}}$, $\text{Cu(PMEA)}_{\text{cl/O}}$, $\text{Cu(PMEA)}_{\text{cl/O/N3}}$ and $\text{Cu(PMEA)}_{\text{cl/N7}}$, respectively, showing that the macrochelated isomer involving N7 is a minority species; the important isomers involve the ether oxygen either in the form of $\text{Cu(PMEA)}_{\text{cl/O}}$ or together with N3 as $\text{Cu(PMEA)}_{\text{cl/O/N3}}$. Interestingly, the isomeric distribution for the Cu(9,8aPMEA) system is within the error limits (3σ) identical with that for the mentioned Cu(PMEA) system; the values for $\text{Cu(9,8PMEA)}_{\text{op}}$, $\text{Cu(9,8aPMEA)}_{\text{cl/O}}$, $\text{Cu(9,8aPMEA)}_{\text{cl/O/N3}}$ and $\text{Cu(9,8aPMEA)}_{\text{cl/N7}}$ are $15 \pm 3\%$, $30 \pm 10\%$, $48 \pm 11\%$ and $6.8 \pm 4.7\%$, respectively [41]. This demonstrates how closely related PMEA^{2-} and 9,8aPMEA $^{2-}$ are in their metal ion-binding properties, whereas 8,8aPMEA $^{2-}$ is clearly quite different in this respect.

Of further interest is the Zn(9,8aPMEA) system [41] (see Table 5, the second entry from below), of which $34 \pm 8\%$ exist as the $\text{Zn(9,8aPMEA)}_{\text{op}}$ isomer, whereas $32 \pm 13\%$, $24 \pm 26\%$ and $10 \pm 21\%$ are present as $\text{Zn(9,8aPMEA)}_{\text{cl/O}}$, $\text{Zn(9,8aPMEA)}_{\text{cl/O/N3}}$ and $\text{Zn(9,8aPMEA)}_{\text{cl/N7}}$, respectively. Two points need to be made here: (1) the total amount of chelated species encompasses 66% ($= 32 + 24 + 10$), as already given in parentheses in Table 5; (2) the large error limits (3σ ; confidence limits 99.7%) need to be put into perspective, i.e. if the error limits are given with one standard deviation (1σ), which corresponds to a confidence limit of about 70%, the results for $\text{Zn(9,8aPMEA)}_{\text{cl/O}}$, $\text{Zn(9,8aPMEA)}_{\text{cl/O/N3}}$ and $\text{Zn(9,8aPMEA)}_{\text{cl/N7}}$ are $32 \pm 4\%$, $24 \pm 9\%$ and $10 \pm 7\%$, respectively. Again, the chelated isomers involving the ether oxygen are the dominating ones among the "closed" species and the macrochelated isomer involving N7 is also in this case a minority species.



properties between PMEA^{2-} and 9,8aPMEA $^{2-}$ show up for these two metal ions. With Ni^{2+} this is different [41], as mentioned, but it may be explained with the reduced

Conclusions

The present results show that the metal ion-binding properties of 9,8aPMEA $^{2-}$ very closely resemble those of PMEA^{2-} itself, including the extent of the M^{2+} -ether

oxygen interaction. It is thus not surprising that 9,8aPMEA also shows antiviral activity [26, 33, 74], although it appears to be somewhat less active but also less toxic [74]. This contrasts with 8,8aPMEA which shows no useful biological activity [26, 33] and this despite the fact that also in its complexes an ether oxygen–metal ion interaction occurs and that it occurs with biologically meaningful metal ions like Mg^{2+} , Ca^{2+} or Mn^{2+} to the same extent as with 9,8aPMEA²⁻ or PMEAs²⁻ itself. However, since in the 8-aza isomer the orientation of the adenine residue is different than in PMEAs, whereas in the 9-aza isomer it is identical, this observation regarding the biological activity is no surprise.

Indeed, owing to this different orientation of the nucleobase moiety, also its binding properties towards metal ions with an N affinity, like Cu^{2+} , are different. In 9-substituted PAs a nucleobase interaction is possible and with ions like Cu^{2+} it actually occurs. In contrast, 8,8aPMEA²⁻ behaves in its complexing properties like a PME-R²⁻ ligand, which offers no binding site in its residue R. Since in the active site cavity the anchoring process of a substrate is important [22, 23], and because this process involves the nucleobase residue and dictates thus the orientation of the substrate, one may conclude that at least one of the reasons for the inactivity of 8,8aPMEA is a presumably different orientation in the active site of the polymerase.

In fact, if one considers the structure–function relationship for nucleotide analogues, one may conclude the following: if one assumes that 8,8aPMEA and 9,8aPMEA are transported to the cell and also diphosphorylated like PMEAs [75, 76, 77, 78], then it becomes understandable why 9,8aPMEA shows antiviral activity and 8,8aPMEA does not [26, 33]. The 9-aza isomer is structure-wise so similar to PMEAs itself (see Fig. 1) that a facilitated $M(P_{\alpha})$ binding (via the ether oxygen) is possible with 9,8aPMEAapp⁴⁻ and thus also the formation of the $M(P_{\alpha})$ - $M(P_{\beta}, P_{\gamma})$ binding mode, which is crucial for the transfer of a nucleotidyl unit in the polymerase reaction [22, 23]. In contrast, 8,8aPMEA is anchored with a different orientation (due to hydrogen bonding and stacking [42]) in the active site cavity of the polymerase, which prevents formation of the indicated reactive binding mode. To conclude, for biological activity of an acyclic nucleoside phosphonate derivative the ether oxygen is compulsory, but it must also be correctly orientated in space to become effective.

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