Abiotic synthesis of ATP from AMP in the gas phase: implications for the origin of biologically important molecules from small molecular clusters

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Abstract

In spite of decades of research, little is known about the processes leading to the prebiotic formation of some of the key molecular components involved in the origin of life. We have undertaken a systematic investigation of small noncovalent clusters in the gas phase, using collisional activation to effect reactions between the cluster components, with the goal of identifying processes that lead to the production of chemical species which may have been key players at the threshold of life. Here, we report that collisional activation of a trimer of adenosine 5′-monophosphate (AMP) leads exclusively to formation of the important species adenosine 5′-triphosphate (ATP). The AMP trimer can be either cationic or anionic with each AMP present as the sodium salt. An additional sodium ion yields the cationic cluster, while the anionic cluster lacks one sodium ion. The energetics of this reaction are explored using PM5 semi-empirical calculations and density functional theory. The generation of polyphosphate bonds is not exclusive to the AMP trimer. Sodiated clusters of other nucleotides, ribose phosphate, or phosphate itself lead to the generation of other biologically important polyphosphates. The implications of this chemistry for chemical and biological evolution are discussed.
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Keywords: ATP; Phosphate; Salt bridge; Origin of life; Aerosol

1. Introduction

Adenosine 5′-triphosphate (ATP) is a ubiquitous biological molecule used for storing energy in the form of chemical bonds. The cellular machinery for creating ATP has been found in the oldest forms of life, yet it is still found in all groups of present day organisms [1]. Despite this ancient lineage and the extreme importance of ATP in many biological processes, it is surprising that the molecular mechanism for the synthesis of ATP is still open to debate [2,3]. Even more puzzling is how ATP originally became the primary energy currency for all of life [4]. In order to develop a more complete picture of the role of ATP in the origin of life, it is of interest to identify prebiotic routes that may have possibly contributed to the generation of ATP.

Several explanations for the terrestrial synthesis of thermodynamically unstable phosphoanhydride bonds have been given [5,6] and disputed [7]. The primary difficulty associated with the production of ATP on earth is that this type of chemistry is unlikely

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ATP
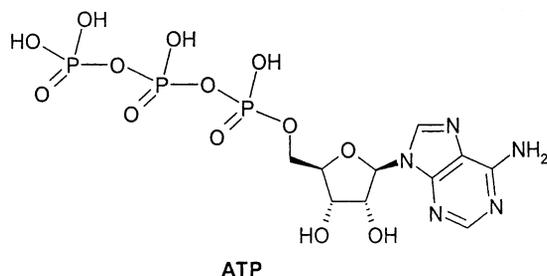
to occur in water. Not only is the hydrolysis of ATP exothermic [8], but the oceans of the primitive world contained pyrite, which catalyses the hydrolysis of phosphoanhydride bonds [9]. Rigorously non-aqueous organic solvents may be used for the synthesis of phosphoanhydride bonds [10], but the availability of such environments may have been limited in the early earth. Perhaps the best approach to phosphoanhydride synthesis involves chemistry in which no solvent is present, a suggestion which will be explained shortly.

Space is an environment which can easily accommodate conditions which are devoid of solvent. The presence of organic molecules in comets and meteorites and the possible role of these organics in the origin of life have drawn much attention over the years [11]. Related work has shown that it is possible to synthesize nucleotides such as adenosine 5′-monophosphate (AMP) by simply exposing the precursor molecules on a thin film to open space [12]. Interestingly, all of the components necessary to synthesize ATP have been detected in meteorites. A wide variety of sugars and their acid and alcohol counterparts have been detected in the Murchison and Murray meteorites [13]. Phosphates and alkyl phosphonic acids in many forms have been detected in various meteorites [14,15]. Nucleobases and other nitrogen heterocycles, including adenine, have been detected in meteorites as well [16], and they have been synthesized in experiments approximating the conditions in space [17].

Terrestrial phenomena can also account for the extraction of key chemical components into a solvent free state. For example, it has been suggested that aerosol particles produced by sea spray may have formed cell-like reactors in which geochemistry may

have evolved into biochemistry [18]. Interestingly, organics from meteorites have been observed to transfer into these aerosol particles in high concentrations [19]. However, if the organic content of an aerosol particle is low and thus insufficient to form a protective monolayer, then as the particle travels to a region of low humidity the water will evaporate away. This could eventually lead to small, desolvated salt clusters. This process is fundamentally analogous to sonic spray ionization (SSI) experiments which can be monitored by mass spectrometry in a controlled environment [20,21]. It is interesting to note that SSI experiments lead to a great enhancement in the formation of clusters when compared to other ionization methods [22].

The gas phase study of non-covalently bound clusters is a rapidly expanding area in the field of mass spectrometry [23–25]. The chemistry of small, non-covalently bound clusters and the role that they may have played in the prebiotic origin of biopolymers is the subject of the present work. It is demonstrated that trimers of AMP will yield ATP exclusively under the appropriate conditions. Collision-activated dissociation (CAD) is utilized to initiate the reactions [26]. Furthermore, this chemistry is general in nature and can be applied to the generation of other polyphosphates which are also of biological significance. It is postulated that other linear polymers of biological importance could be generated from small clusters in a similar manner.

2. Experimental

Mass spectra were obtained using a Finnigan LCQ ion trap quadrupole mass spectrometer without modification. The signal was optimized using the automatic tuning capabilities of the LCQ. To enhance cluster formation, the following settings were used for cationic clusters: source voltage 4.15 kV, capillary voltage 5.0 V, capillary temperature 200 °C, and tube lens offset −40 V. CAD was performed on isolated ions by applying a resonance excitation RF voltage of between 0.98 and 2.45 V for a period of 30 ms.

Adenosine (267 Da)   AMP (347 Da)   Adenine (135 Da)
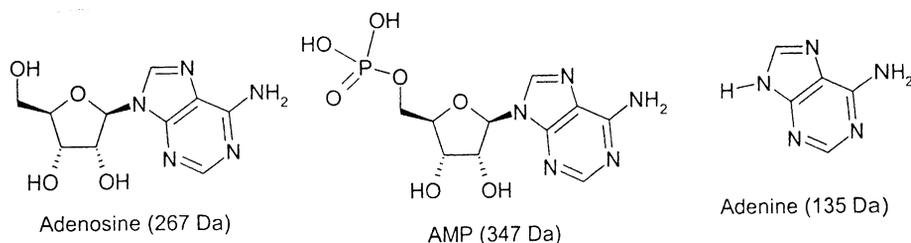
Sample concentrations were in the range of ~50 μM. Samples were electrosprayed using a 80:20 methanol/water mixture at a flow rate of 3 μl/min from a 250 μl Hamilton syringe. Silica tubing with an inner diameter of 12.7 μm inside a stainless steel sheath was used for electrospraying the mixture. Compounds were purchased commercially and used without further purification.

3. Results

3.1. Anionic clusters

The negative ion ESI-MS spectrum for the sodium salt of AMP is shown in Fig. 1a. The two major peaks correspond to [AMP-H]$^-$ and [2AMP-H]$^-$. Clusters incorporating one or more sodium ions replacing protons are present in smaller abundance. Isolation of [3AMP-H]$^-$ followed by CAD leads exclusively to the loss of neutral AMP, as shown in Fig. 1b. The results differ marginally when one hydrogen is replaced by sodium. The CAD spectrum for [3AMP-2H + Na]$^-$ is shown in Fig. 1c. The primary product is the loss of neutral AMP. However, there is also a minor loss of 267 Da, which corresponds to the loss of an adenosine nucleoside. In other words, one AMP has been fragmented, releasing neutral adenosine and leaving PO$_3$ with the cluster. The structures for AMP and two important fragments are given above.

Fig. 2 shows the results for the sequential CAD of the [3AMP-3H + 2Na]$^-$. The loss of adenosine leads to the base peak in Fig. 2a. Some additional loss of a second adenosine is observed in Fig. 2a as well. The primary product is isolated in Fig. 2b. Further CAD in Fig. 2c leads exclusively to the loss of the sec-

ond adenosine. Tentatively, the results in these spectra correspond to the transformation of 3AMP into AMP + ADP, which is then converted into ATP, as will be discussed in greater detail below.

Three additional collisional activation steps are performed on the primary product from Fig. 2c in Fig. 3. The primary peak in Fig. 3a results from the loss of 18 Da (water). An additional loss of 102 Da is also observed. In Fig. 3b, the adenine base is lost, leading to the base peak in the spectrum. This loss is accompanied by the pickup of water and methanol, yielding peaks with masses that are 18 and 32 Da higher than the daughter fragment, respectively.[1] Isolation of the daughter fragment does not eliminate these peaks, confirming that they are picked up from residual solvent in the trap. A competitive loss of 102 Da is also observed. Isolation of the most abundant product, followed by collisional activation yields primarily a loss of 102 Da in the MS^6 spectrum shown in Fig. 3c. An additional minor peak corresponding to the loss of a second 102 Da is also present in Fig. 3c. This 102 Da most likely corresponds to neutral loss of NaPO$_3$.

The sequential CAD of an authentic sample of ATP is shown in Fig. 4. In Fig. 4a, isolation of [ATP-3H + 2Na]$^-$ is accompanied by the isobaric doubly charged dimer [2ATP-6H + 4Na]$^{2-}$. Therefore, several of the daughter peaks in the CAD spectrum in Fig. 4a result from the dissociation of the doubly charged dimer and are marked with a double asterisk. The loss of water leads to the base peak, just as in Fig. 3a. In Fig. 4b, interference from doubly charged peaks has been eliminated, and the spectrum is nearly identical

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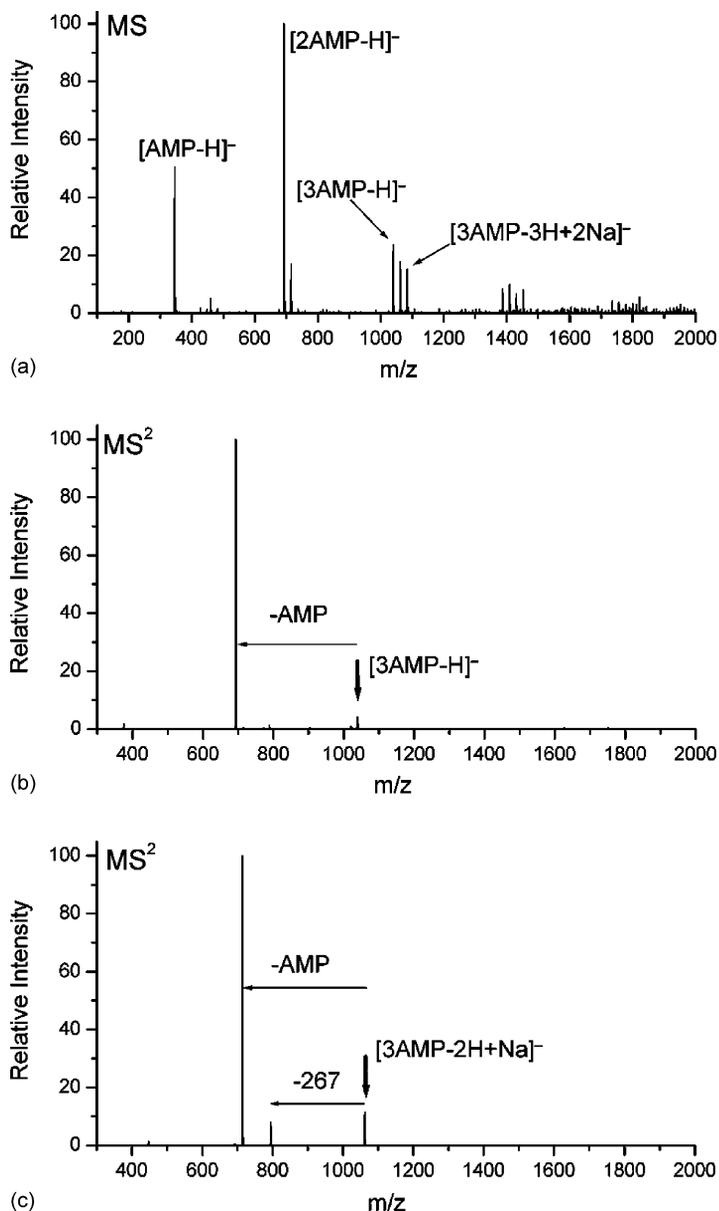[1] This phenomena is often observed in ion traps, particularly with transition metal ions.

Fig. 1. (a) Anionic clusters of AMP with sodium. (b) CAD spectrum of the deprotonated AMP trimer showing that no covalent bonds are broken. (c) CAD spectrum of AMP trimer with inclusion of one sodium ion, in which a minor peak corresponding to the fragmentation of one AMP (namely, loss of adenosine) is observed.

to that shown in Fig. 3b. Similarly, Fig. 4c is very comparable to Fig. 3c. In fact, even the majority of the minor peaks in both Fig. 4b and c are identical to those found in Fig. 3b and c.

3.2. Cationic clusters

Fig. 5 shows the positive ion ESI spectrum for the sodium salt of AMP. Clusters with a sodium cation
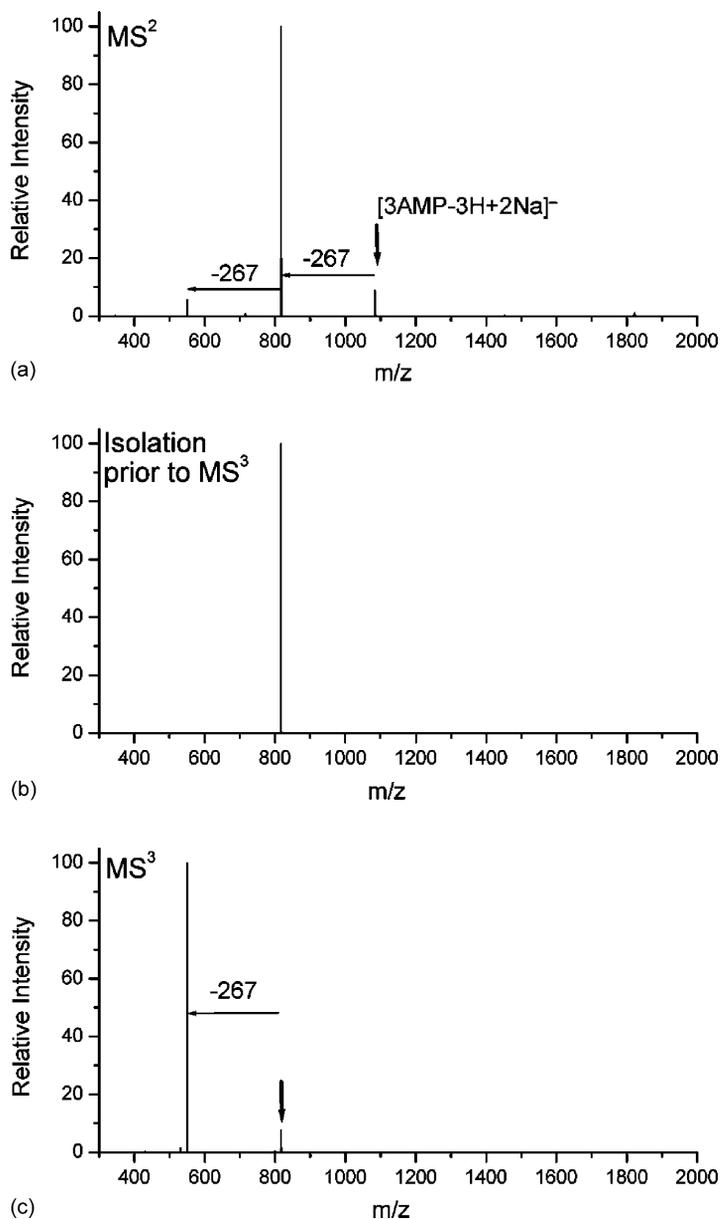
Fig. 2. (a) CAD spectrum of the anionic sodiated salt cluster of the AMP trimer. The loss of adenosine is predominant and no dissociation of the cluster by loss of AMP is observed. (b) Isolation of the product peak. (c) MS^3 spectrum demonstrating the loss of a second adenosine and the generation of ATP.

providing the net charge are easily generated. Clusters incorporating up to five AMP molecules are observed. The maximum intensity for each cluster distribution corresponds to the peak fitting $[xAMP - xH + (x + 1)Na]^+$, where $x = 1$–5. Formally then, each AMP is deprotonated and coupled with a sodium cation, forming salt clusters.

In Fig. 5b, results are presented for the collisional activation of the cluster $[3AMP-3H + 4Na]^+$. The loss of adenosine yields the most abundant product.
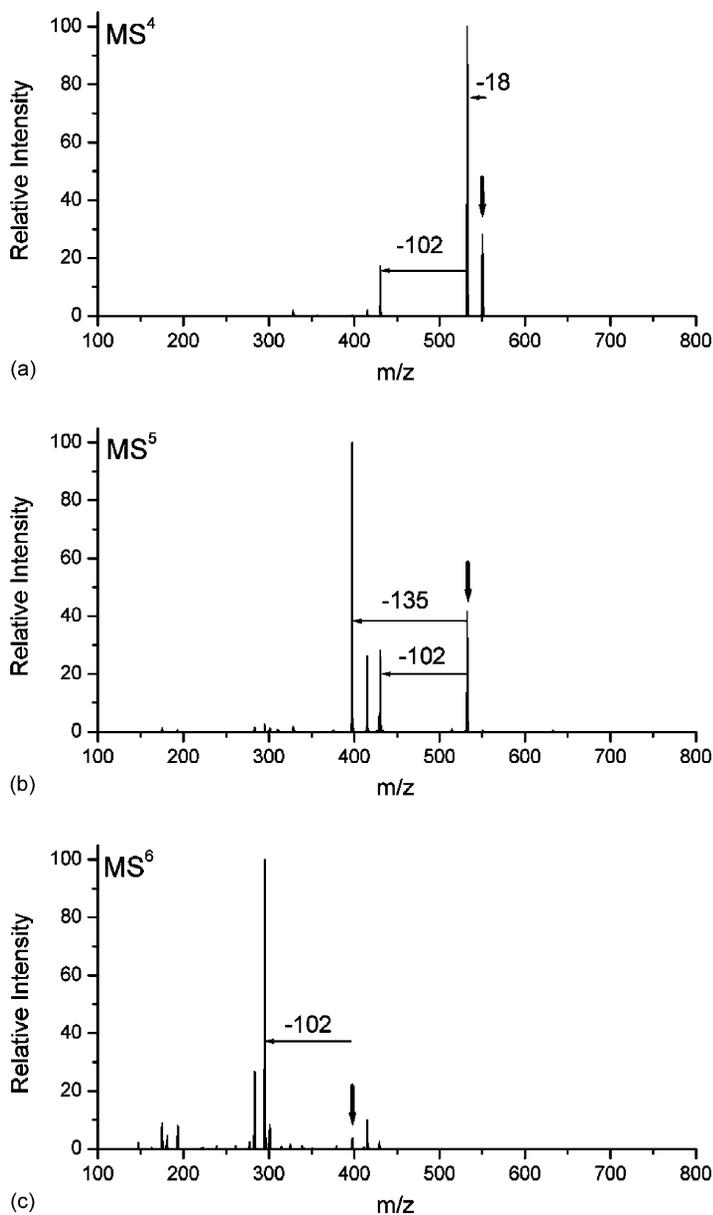
Fig. 3. Three further collisional activation steps on the product (ATP) generated in Fig. 2c. (a) The lowest energy pathway leads to the loss of water. (b) Adenine is lost next. (c) Sequential losses of 102 Da or $NaPO_3$ are the final losses that are observed in this MS^6 spectrum.

Isolation of this product, followed by further collisional activation, leads primarily to the loss of another adenosine as seen in Fig. 5c. These losses are identical to those observed in Fig. 2 for the anionic clusters. The base peak produced in Fig. 5c is isobaric with $[ATP-3H + 4Na]^+$. Further CAD of this peak yields products that are indistinguishable from those produced by CAD of $[ATP-3H + 4Na]^+$ using an authentic ATP sample. The results for the cationic clusters closely parallel those obtained for the anionic clusters.
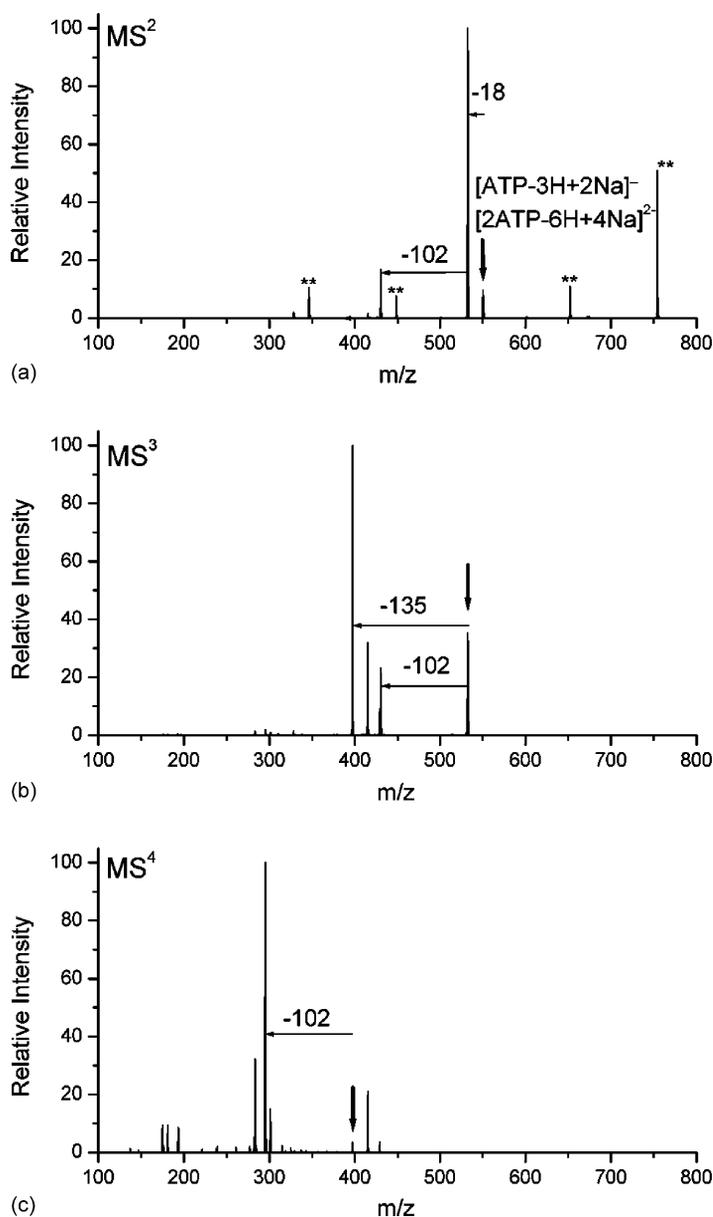
Fig. 4. Three CAD steps on authentic ATP, [ATP-3H + 2Na]⁻. (a) Some interference from the doubly charged dimer leads to additional peaks. (b) The loss of adenine is very abundant, as is also observed in Fig. 3b. (c) Loss of 102 Da is observed in a similar fashion as Fig. 3c, suggesting that the product produced in Fig. 2c is in fact ATP. **Daughter ions from the doubly charged dimer are indicated by a double asterisk.

3.3. Phosphate clusters

The CAD spectra for clusters of phosphate are shown in Fig. 6. The sodiated cluster $[3H_2PO_4 + Na]^-$ loses water exclusively as shown in Fig. 6a. Subsequent activation of the daughter ion results in the loss of another water and the formation of triphosphate as shown in Fig. 6b. This loss is accompanied by the
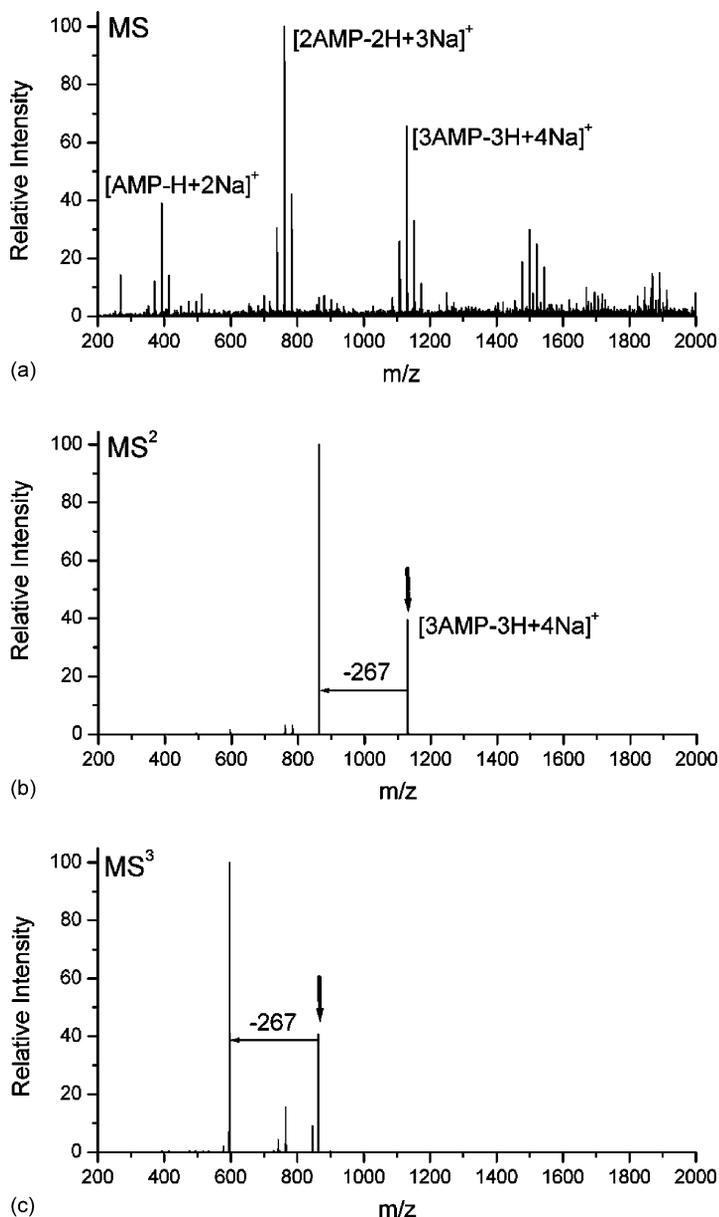
Fig. 5. (a) Sodiated cationic salt clusters of AMP. (b) CAD of the trimer leads to the formation of ADP + AMP. (c) A second adenosine is lost, implying that ATP is generated in high yield by a reaction similar to that observed for the anions.

pick up of water and methanol in low abundance. It is unclear whether the added solvent molecules react with the cluster or are merely coordinated to it. In Fig. 6c, $[3H_3PO_4–H]^-$ is subjected to CAD. In the absence of the sodium ions, the only observed product corresponds to a loss of H_3PO_4. These results closely parallel those observed for the AMP clusters.
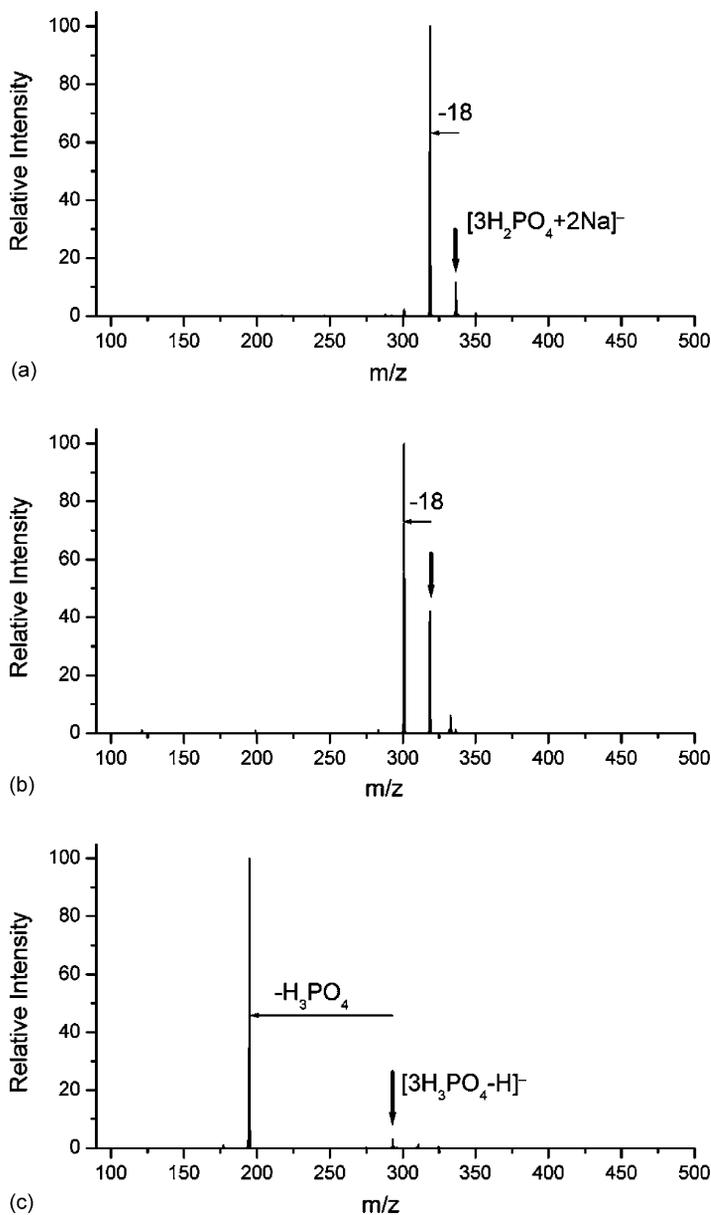
Fig. 6. (a) CAD spectrum of $[3H_2PO_4 + Na]^-$ which loses water to form a phosphoanhydride. (b) CAD of the daughter ion from (a), yielding triphosphate from the loss of second water. (c) CAD of $[3H_3PO_4–H]^-$. The hydrogen bound cluster simply dissociates by the loss of H_3PO_4.

3.4. Calculations

The enthalpic contributions to reaction 1 were calculated at the PM5 semi-empirical level and utilizing density functional theory at the BLYP/DZVP and B3LYP/6-31G** levels. The results are given in Table 1. Methylphosphate is used instead of AMP to evaluate the binding energies of the clusters in order to allow for higher levels of theory to be utilized in a reasonable time. This approximation is supported

Dissociation
Threshold
51 kcal/mol
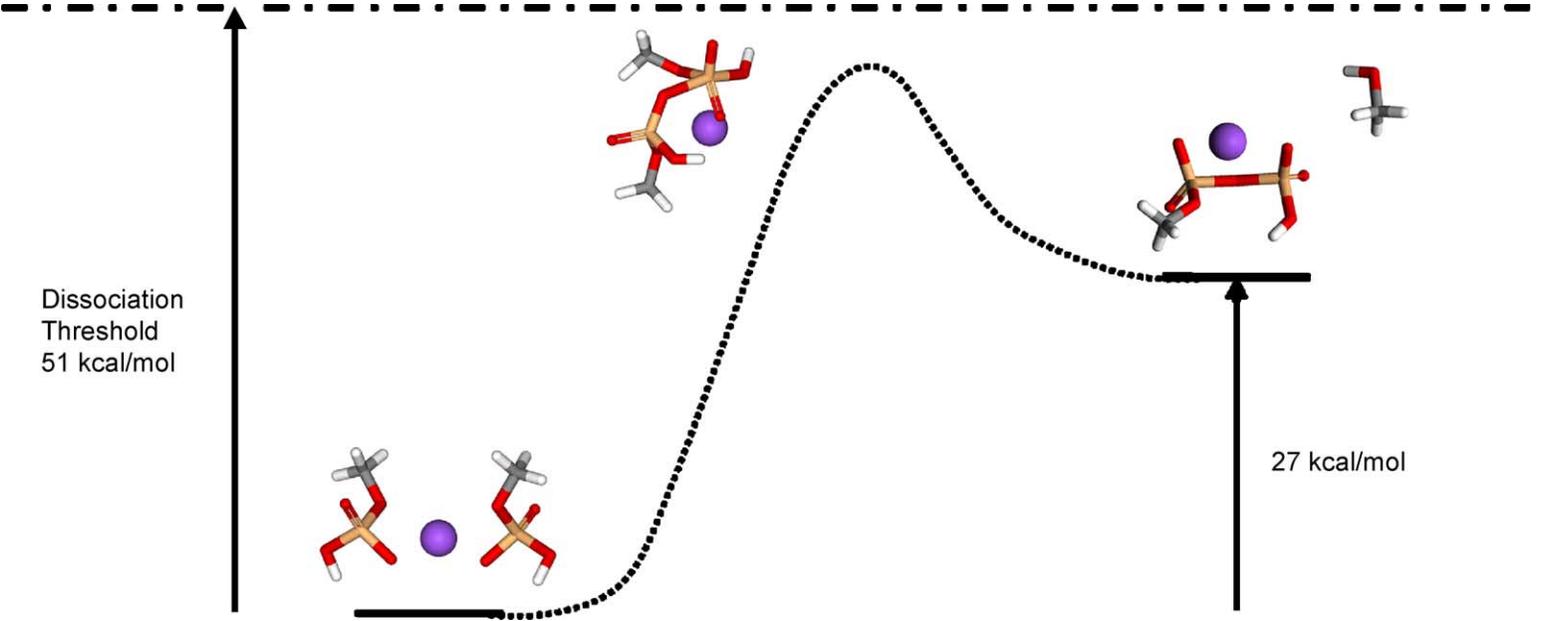
27 kcal/mol

Scheme 1. Reaction coordinate diagram for model chemical system.

Table 1
Calculated energetics in kcal/mol

Cluster	Binding energy PM5	Binding energy BLYP/DZVP	Binding energy B3LYP/6-31G**
$(CH_3OPO_3H)_2H^-$	−22.36	−34.2309	−40.2867
$(CH_3OPO_3H)_2Li^-$	−46.37	−48.1216	−54.807
$(CH_3OPO_3H)_2Na^-$	−44.95	−44.9093	−50.7288
$(CH_3OPO_3H)_2K^-$	−46.71	−46.0386	−46.6662[a]
$(CH_3OPO_3H)_2Rb^-$	−59.17	−41.0822	−38.2897[a]

[a] Calculated using the LACVP basis set.

by the experimental results shown in Fig. 6 which suggest that most of the chemistry takes place in the phosphate/sodium salt portion of the clusters. The calculations reveal that metal bound phosphate salt clusters are more strongly bound in the gas phase than the proton bound counterparts. The PM5 methodology is sufficient to qualitatively predict the results, although the energetic ordering among the alkali metals is not correct.

$$M \cdot CH_3HPO_4 + CH_3HPO_4^-$$
$$\rightarrow M \cdot CH_3HPO_4 \cdot CH_3HPO_4^-$$
$$(M = H, Li, Na, K, Rb) \qquad (1)$$
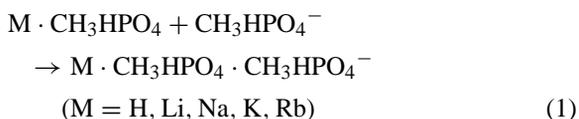
The calculated energetics and structures for the collision-induced reaction of $[2CH_3HPO_4 + Na]^-$ to yield $[CH_3HPO_3PO_4 + Na]^-$ and methanol are given in Scheme 1 in kcal/mol utilizing DFT at the B3LYP/6-31G** level. The reaction is uphill, and proceeds via a metastable penta-coordinate phosphorous intermediate.

4. Discussion

The binding energy of a noncovalent cluster must be greater than the activation barrier for a reaction to occur by collisional activation. In CAD experiments with clusters, rearrangement processes involving the cleavage of covalent bonds are usually not competitive with dissociation of the complex [27]. This is well illustrated in Fig. 1, where the hydrogen bonded clusters are shown to simply dissociate into the con-

stituent molecules. The addition of a single sodium ion increases the binding energy sufficiently to allow some covalent bond cleavage, as seen in Fig. 1c. However, the formation of a true salt cluster in Fig. 2 lends greater binding energy to the cluster, enabling reactions with higher activation barriers. The greater binding energy is quantified by the computational results, which predict the salt cluster to be more strongly bound than the hydrogen bound cluster by roughly ∼10 kcal/mol for each substitution. As a result, in Fig. 2 the sequential loss of two adenosine nucleosides which results from the fragmentation of covalent bonds is observed almost exclusively. The final peak produced in Fig. 2c appears to be ATP, but the fact that it has the same mass is not necessarily proof of structure.

In order to address that question, further CAD experiments and calculations were performed. The results for the calculations are shown in Scheme 1 as a reaction coordinate diagram. Theory predicts that the ΔH for the reaction of a sodium bound dimer is significantly uphill, which is to be expected. The exact energy for the transition state was not calculated, but the transition state must lie below the dissociation threshold for the cluster and probably involves a penta-coordinate phosphorous as shown in Scheme 1.

The quasimolecular ion produced in Fig. 2c and actual ATP were both subjected to further steps of CAD for comparison. The results from these experiments are presented in Figs. 3 and 4. With the exception of the peaks originating from the doubly charged cluster of ATP in Fig. 4a, the spectra in both Figures are nearly identical. The peaks that are produced and the relative intensity of those peaks are very similar, including peaks that only constitute minor contributions to the total ion intensity. Although this is not absolute proof that the structure of the product in Fig. 2c is ATP, it provides strong support for the conjecture.

Further support for this notion is obtained from the cationic clusters. Despite a different number of sodium ions and presence of net positive charge, the same chemistry is observed in sodiated clusters of AMP.

Fig. 5 clearly shows a favorable route to what appears to be ATP. Again, comparison of the subsequent CAD of this molecule with a verified sample of ATP reveals that the two yield identical daughter ions. There is no interference from a dimer in this case, and the peaks and relative intensities of the daughter ions are nearly identical. This strongly suggests that ATP is indeed formed from the salt cluster trimer. The notion that two different products would coincidentally produce spectra that overlap exactly with ATP is unlikely.

Other important observations are related to the fact that the same chemistry operates in cationic and anionic clusters. This suggests that the net charge does not play a significant role in the mechanism that leads to the generation of ATP. Neutral clusters would probably yield similar results, but unfortunately neutral clusters are not well suited for the experiments performed in the present work. This does not preclude the formation of such clusters in the atmosphere, either from meteorites or sea spray, or the subsequent chemistry that follows. The results obtained in the present work suggest a viable prebiotic route to the synthesis of ATP.

Furthermore, the chemistry is not limited to the formation of ATP. In fact, it was possible to synthesize adenosine pentaphosphate, although the structure of this compound was not verified for lack of a sample for comparison. Other nucleotides and deoxynucleotides were basically interchangeable with AMP, allowing for the synthesis of other polyphosphates. Sodiated clusters of 5'-ribose-phosphate also yielded ribose polyphosphate upon collisional activation. Polyphosphate, another important and ubiquitous biopolymer [5], can be synthesized from sodium salt clusters of phosphate itself. The facile generation of polyphosphates by this mechanism is therefore applicable to a wide range of biologically relevant molecules.

5. Conclusion

ATP is easily generated by collisional activation of a sodiated salt cluster containing three AMP precursors. The sequential loss of two adenosines leads to the formation of ATP. Comparison of several subsequent steps of CAD dissociation of ATP generated by this method with that of actual ATP reveals that the two ions produce identical CAD spectra. This is true for ATP generated from both cationic and anionic sodiated salt clusters of AMP. Polyphosphates can also be generated by collisional activation of sodiated clusters of phosphate, ribose phosphate, and other nucleotides and deoxynucleotides.

This type of chemistry represents a feasible pathway for the generation of biologically important polymers. A potential prebiotic source for ATP or other polyphosphates is outlined in the present work, but conceivably other biopolymers may be generated by similar reactions. It is interesting to note that small clusters of biomolecules can have an inherent preference for homochirality, as is demonstrated by the serine octamer [28]. It appears quite possible that the chemistry of these small clusters may be more important than has been previously realized.

Acknowledgements

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