

## Prebiotic Adenine Synthesis from HCN—Evidence for a Newly Discovered Major Pathway

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Three new imidazole derivatives have been isolated and characterized from oligomerizing HCN solutions. On the basis of these results as well as the earlier identification of a new precursor to adenine, a new and major pathway leading to the formation of adenine is suggested. The route accounts for the synthesis of adenine-8-carboxamide from *cis*-diaminomaleonitrile, without requiring an isomerization to the *trans* configuration or reactions with formamidine. The formation of previously reported imidazoles is also explained.

### INTRODUCTION

The formation of biologically important molecules via the oligomerization of HCN is well established. Oro (1) and Oro and Kimball (2) isolated and identified the purine adenine from aqueous ammonium cyanide solutions (1–15 M). This first result has been supported and explored further (3, 4). Subsequently, several different classes of compounds have been reported including purines, pyrimidines, and imidazoles as well as amino acids, urea, oxalic acid, and guanidine (5–11). In this context, the oligomerization of HCN is considered to have been a major process in chemical evolution on the primitive Earth (12, 13).

However, the pathways for the formation of these compounds in the HCN oligomerization are not understood entirely. Suggested mechanisms for the formation of adenine are summarized in Fig. 1 (3, 11, 12).

The pathway from the trimer, aminomalononitrile **1**, is now considered to be unlikely in comparison to reactions involving the tetramer diaminomaleonitrile **2** (12). High steady-state concentrations of formamidine are also thought not to be plausible except in solutions with a high concentration of ammonia. The route to adenine *via* **2** is dependent on a photochemical isomerization and ring-closure reaction, and further conversion to adenine has been modeled by means of reaction with formamidine.

We have reported previously the isolation and preliminary characterization of a new precursor of adenine in the HCN oligomerization prior to hydrolysis. This compound—adenine-8-carboxamide (**6**), Fig. 5—was identified spectroscopically and by synthesis ((8) and unpublished results). The isolation of this precursor suggested a new pathway to adenine which was proposed tentatively (8). We have now isolated and identified several new compounds which support this mechanism, which we propose as a major route to purines in chemical evolution.

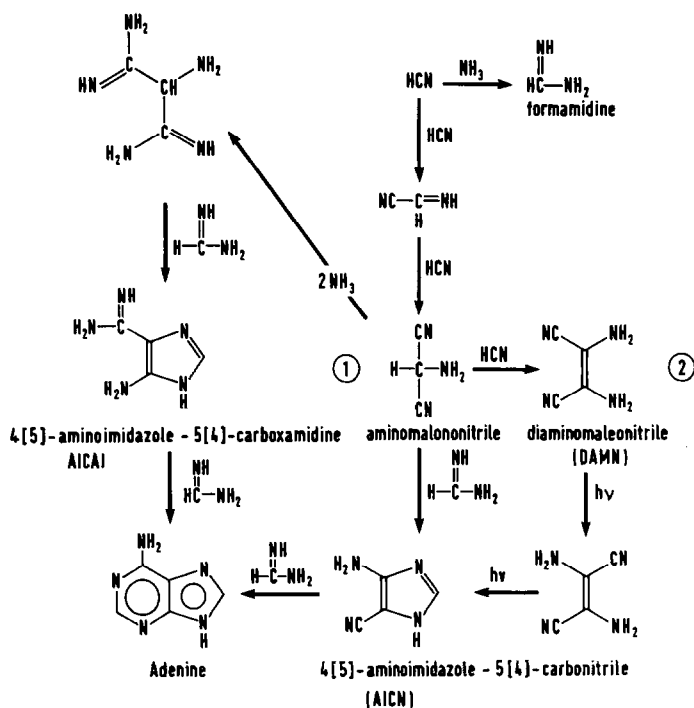


FIG. 1. Summary of earlier proposed pathways for the formation of adenine in aqueous ammonium cyanide solutions.

## EXPERIMENTAL

<sup>13</sup>C nmr spectra were determined on a Bruker Model WM-250 (62.89 MHz) with DMSO as solvent and as internal standard. <sup>1</sup>H nmr spectra were recorded in DMSO on either a Bruker Model WH-90, WM-250, or WM-400 (TMS as internal standard). Mass spectral data were obtained with either a Varian MAT 711 double-focusing mass spectrometer with combined EI/FI/FD ion source or a Finnigan 3100 quadrupole GC/MS (solid probe method). IR spectra were determined on a Perkin-Elmer 457 spectrophotometer as potassium bromide micropellets. UV spectra were obtained on a Cary-15 spectrophotometer. HPLC was performed on Aminex A-25 resins as described previously (14). Sephadex G-15 (40–120 μm) was purchased from Pharmacia Fine Chemicals, and was used in column chromatography for fractionation of HCN oligomers (90 × 1.6 cm) as well as for desalting purposes (30 × 1.4 cm). In both cases H<sub>2</sub>O was used as eluting solvent and the collected fractions were monitored continuously at 254 nm (LKB fraction collector connected to a Uvicord optical unit).

*General procedure for the preparation of HCN solutions.* HCN was freshly prepared by adding a saturated solution of NaCN to 75% aqueous H<sub>2</sub>SO<sub>4</sub> at 70°C with continuous distillation of the formed HCN. For a 1 M solution, 40 ml of HCN was added to 900 ml of H<sub>2</sub>O. The solution was then adjusted to pH 9.2 with

$\text{NH}_4\text{OH}$  and made up to 1 liter with  $\text{H}_2\text{O}$ . Solutions were allowed to stand at least 7 months at room temperature.

4[5]-Amino-5[4]-carboxamide-2-cyano-imidazole (3). Ammonium cyanide (1 M, 350 ml, reaction conditions: 8 months, RT, and pH 9.2) was passed through a Millipore filter ( $0.45\ \mu\text{m}$ ) to separate insoluble azulmic acid. The filtrate was evaporated to dryness *in vacuo* below  $30^\circ\text{C}$  and the residue was extracted with ethyl acetate ( $3 \times 100\ \text{ml}$ ). The residue was dissolved in 0.1 M HCl (20 ml) and the solution was extracted with EtOAc ( $3 \times 100\ \text{ml}$ ). The combined, dried ( $\text{Na}_2\text{SO}_4$ ) extracts were concentrated *in vacuo*. An aqueous solution of the residue was then applied to an Aminex A-25 HPLC column for fractionation. A peak with a retention time of 38.6 min was collected, desalted on Sephadex G-15, and evaporated *in vacuo* to yield 10–15 mg of product (0.09–0.14%). Mass:  $m/e$  151( $\text{M}^+$ ), 134(100), 106, 81, 55, 54, and 53. IR:  $3300, 1665\ \text{cm}^{-1}$  (amide);  $2215\ \text{cm}^{-1}$  ( $\text{C}\equiv\text{N}$ ). UV: 295, 240 nm (1 M HCl); 296, 238 nm (1 M NaOH); 298, 242 nm ( $\text{H}_2\text{O}$ ).  $^1\text{H}$  nmr (DMSO):  $\delta = 8.73$  and  $8.42$  ppm (see Results); 6.87 ppm (amino  $\text{NH}_2$ ).  $^{13}\text{C}$  nmr (DMSO): 90.57, 116.27, and 149.90 ppm ( $\text{C}=\text{C}$  and  $\text{C}\equiv\text{N}$ ); 135.22 ppm (C-2); and 159.10 ppm (C-amide).

4[5]-aminoimidazole-2,5[4]-dicarboxamide (4). The residual 0.1 M HCl solution from the above extraction procedure was evaporated to dryness *in vacuo* and redissolved in  $\text{H}_2\text{O}$  (7 ml) and then applied to a Sephadex G-15 column. Thirteen fractions were collected (Fig. 2).

Fractions 7–9 were rechromatographed on an Aminex A-25 exclusion column; compound 4, as the major peak in these fractions, was collected (A-25: 13.50 min) and was desalted by chromatography on Sephadex G-15. Mass:  $m/e$  169( $\text{M}^+$ ), 152(100), 136, 124, 109, 107, 81, 55, and 54. IR:  $3320$  and  $1670\ \text{cm}^{-1}$  (amide);  $1590\ \text{cm}^{-1}$ . UV: 306, 251 nm (0.1 M HCl); 312, 253 nm (0.1 M  $\text{NH}_3$ ); 305, 253 nm ( $\text{H}_2\text{O}$ ).  $^1\text{H}$  nmr:  $\delta = 8.75$  and  $8.17$  ppm (both asymmetric peaks);  $\delta = 6.97$  ppm (amino  $\text{NH}_2$ ).

4[5]-N-(aminomethylidene)aminoimidazole-2,5[4]-dicarboxamide (5). The isolation of compound 5 was performed in the same way as described for compound

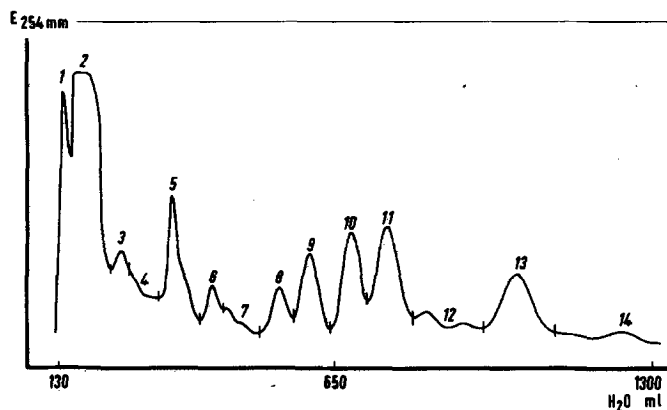


FIG. 2. Sephadex G-15 fractionation of the residual 0.1 M solution (see text).

TABLE I  
HIGH-RESOLUTION MASS SPECTROMETRY OF COMPOUND 3

Mass	C	H	N	O	
$M^+ 151.0494$	5	5	5	1	
134.0239	5	2	4	1	$M^+ - 17 [-NH_3]$
106.0282	4	2	4	0	$[-C=O]$

4, but with fractions 9 and 10 of the initial Sephadex G-15 separation (A-25: 24.50 min). Mass:  $m/e$  196( $M^+$ , 100), 179, 153, 151, 134, 124, 108, 81, 55, and 54. IR: 3400 and 1650  $\text{cm}^{-1}$  (amide); 1600  $\text{cm}^{-1}$ . UV: 307 nm (0.1  $M$  HCl); 310 nm (0.1  $M$   $\text{NH}_3$ ); 302 nm ( $\text{H}_2\text{O}$ ).

## RESULTS

The identification of compound 3 was based on the combined interpretation of the spectra (mass, nuclear magnetic resonance, and infrared spectra). Table 1 presents the molecular formulas for the molecular-ion peak and two successive fragment peaks as determined by high-resolution mass spectrometry.

Infrared spectra supported the structure proposed and showed a broad absorption peak at 3300  $\text{cm}^{-1}$  and a sharp band at 1665  $\text{cm}^{-1}$  which indicated an amide functional group. Another very important peak present was at 2215  $\text{cm}^{-1}$  which suggested a nitrile group attached to a conjugated system. The  $^1\text{H}$  nmr spectrum showed three main signals in the ratio of 1 : 1 : 2. No methine proton was present in

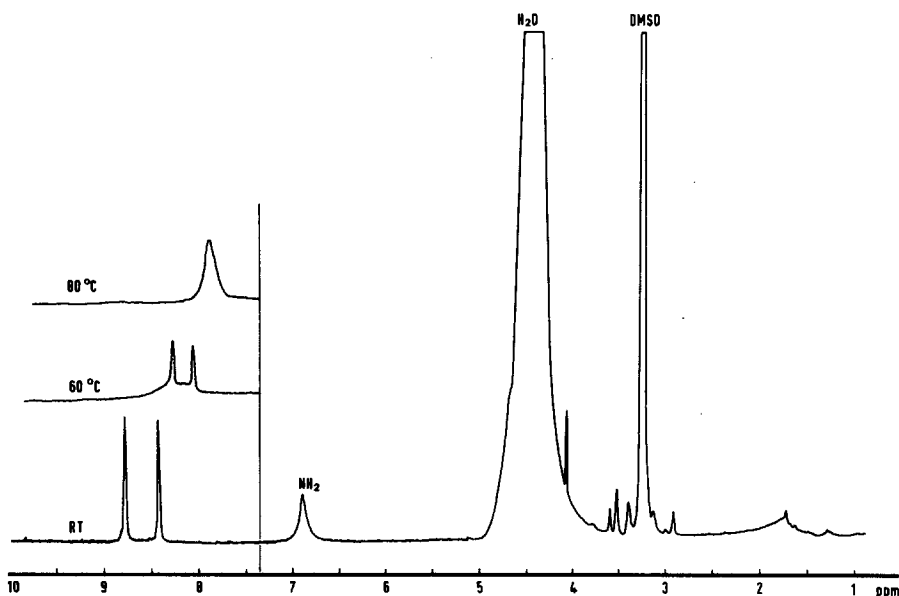


FIG. 3.  $^1\text{H}$  nmr spectra of compound 3 at different temperatures.

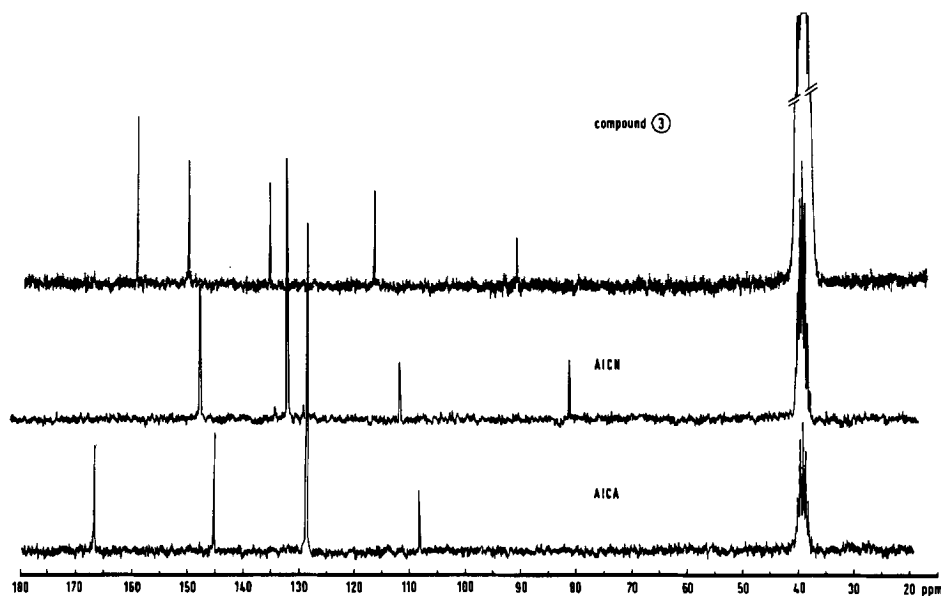


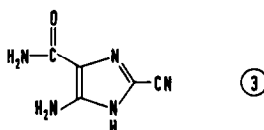
FIG. 4.  $^{13}\text{C}$  nmr spectra of compound 3, AICA, and AICN in DMSO.

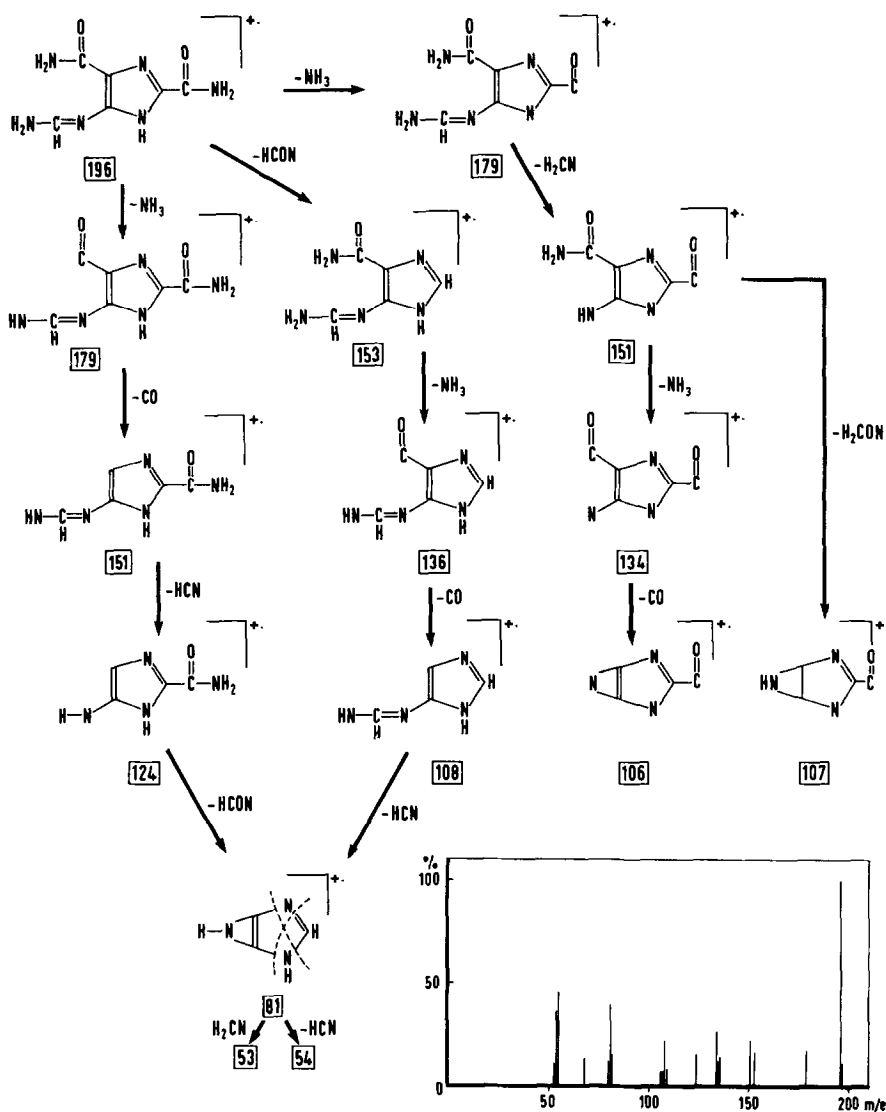
the molecule, because of the facile proton exchange with  $\text{D}_2\text{O}$  indicating only NH or OH functions. When the spectrum was recorded at elevated temperatures ( $80^\circ\text{C}$ ), two of the three signals moved to higher field and showed overlapping, which is a strong indication of two protons belonging to the same functional group (Fig. 3).

Correlation of the  $^{13}\text{C}$  nmr spectrum of isolated compound 3 with those of 4[5]-aminoimidazole-5[4]-carbonitrile (AICN) and 4[5]-aminoimidazole-5[4]-carboxamide (AICA) gave more information (Fig. 4).

The chemical shifts in AICA were 129.8 ppm for the C2 and 146.0 and 108.26 ppm, respectively, for the  $\text{C}=\text{C}$ . The amide absorption occurred at 165.3 ppm (Sadler  $^{13}\text{C}$  nmr, No. 1631c,  $\text{AICA} \cdot \text{HCl}$ ; for the free AICA the amide absorption was at 161.3 ppm). AICN showed almost the same absorption pattern, but without the signal at 165.3 ppm and with new signals at 111 and 80.8 ppm for the nitrile function and the ring-carbon atom attached to the nitrile. The spectrum of compound 3 closely resembled the combined spectra of AICA and AICN. All the imidazole-ring absorption peaks were present as well as an amide function and a nitrile. The substitution of the C-2 position of the imidazole ring can be seen by the decrease in intensity of the absorption at 132 ppm compared to the corresponding peaks in AICN and AICA (absence of a Nuclear Overhauser Effect).

The spectral data suggested a structural formula for the isolated compound 3 of 4[5]-amino-5[4]-carboxamide-2-cyanoimidazole:



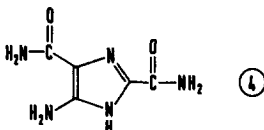


SCHEME 1

The nitrile function at the 4 or 5 position could be excluded by mass fragmentation data. No  $\text{M}^+ - \text{HCN}$  fragment was produced as in the fragmentation of AICN. Another explanation of the positions of the functional groups in compound 3, is the easy fragmentation of  $\text{NH}_3$  by a proton donation of the amino to the amide function in a six-membered ring rearrangement (Mc Lafferty rearrangement).

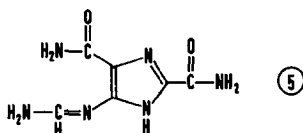
The mass spectrometry of compound 4 showed a  $m/e$  value of 169 as molecular-ion peak, which was established as  $\text{C}_5\text{H}_7\text{N}_5\text{O}_2$  by high resolution (169.0590). The base peak (100%) was  $m/e$  152 (152.0335,  $\text{C}_5\text{H}_4\text{N}_4\text{O}_2$ ). The  $^1\text{H}$  nmr spectrum of the isolated compound 4 resembled the spectrum of 3. Once again two signals were

present at room temperature at 8.75 and 8.17 ppm ( $\text{NH}_2$  amide functional group(s)) and there was also a signal at 6.97 ppm from the amino group, but in the ratio 1:1:1. In the IR spectrum bands at  $3220$  and  $1670\text{ cm}^{-1}$  suggested amino and amido functions whereas the band associated with a conjugated nitrile was absent. These combined spectral data, together with the knowledge that the molecule is a hydrolysis product of compound **3**, supported the structure **4**, viz., 4[5]-aminoimidazole-2,5[4]-dicarboxamide:



The third product from the HCN oligomerization was identified only on the basis of similarity to the compounds **3** and **4** and by mass spectral fragmentation (Scheme 1).

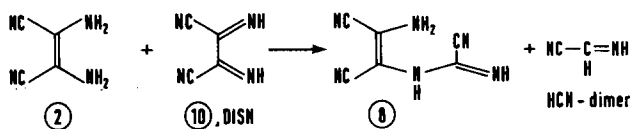
The IR spectrum showed the same kind of absorption bands as for compound **4**,  $3400$  and  $1650\text{ cm}^{-1}$  for an amide function and  $1600\text{ cm}^{-1}$  for the imidazole ring. High-resolution mass spectrometry gave for the molecular-ion peak ( $m/e = 196.0708$ ) a formula of  $\text{C}_6\text{H}_8\text{N}_6\text{O}_2$ . These data suggest the probable structure of 4[5]-*N*-(aminomethylidene)aminoimidazole-2,5[4]-dicarboxamide (**5**):



## DISCUSSION

Prior to the present work, little was known concerning the intermediate compounds formed in the HCN oligomerization beyond the stage of the tetramer. Some attempts at structural analysis have been undertaken (15). However, in the very complex mixture of products formed by the HCN oligomerization it is very difficult—perhaps impossible—to indicate the total range of functional groups present. We previously reported the isolation of the first direct precursor of adenine in HCN-oligomerization mixtures—later confirmed to be adenine-8-carboxamide (**6**) ((8) and unpublished results)—and proposed a mechanism of formation. With the further isolation and identification of compounds **3**, **4**, and **5** we now can refine and extend the proposed mechanism for the formation of adenine (Fig. 5).

An important step in this mechanism is the reaction of the HCN tetramer DAMN (**2**) with a cyanoimino derivative (**7**) leading to the formation of compound **8**. Diiminosuccinonitrile (DISN, compound **10**), an oxidation product of DAMN which can be formed in the presence or absence of oxygen (16, 17), could account for compound **7**:



However, attempts to find support for this reaction were unsuccessful. No higher yields of compound 6 were obtained after DISN was added prior to the oligomerization. A second possibility is the reaction of two molecules of DAMN, one of

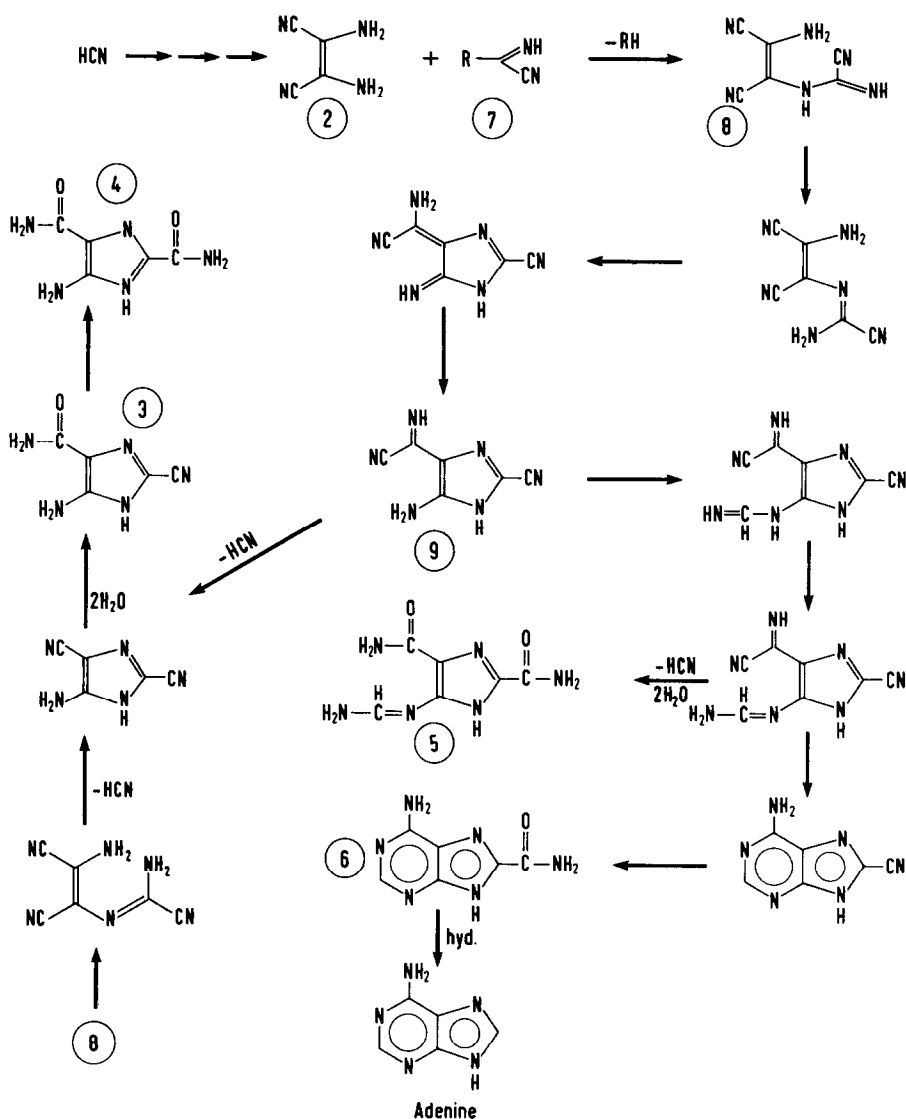


FIG. 5. Suggested pathways for the formation of adenine and other identified products of the HCN oligomerization.



### SCHEME 2

mediate in the route to adenine and is, in fact, the only direct precursor of this purine yet to be identified in dilute, oligomerizing solutions of HCN.

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