

Published on Web 02/07/2004

5-Cyanoamino-4-imidazolecarboxamide and Nitrosative Guanine Deamination: Experimental Evidence for Pyrimidine Ring-Opening during Deamination

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Nitrosative guanine deamination via guanine diazonium ion 1 to xanthine was discovered by Strecker in 1861¹ before Kossel identified guanine as a DNA base in 1884.² A century later this chemistry became a central topic in toxicology,3 and it is now well understood that endogenous nitrosation⁴ occurs and that dG-to-dG and dG-to-dA cross-links also are formed.5 Guanine deamination chemistry became interesting once again with the 1996 discovery by Suzuki, et al. that deamination may form oxanine.⁶⁻⁸ We proposed a reaction mechanism that involves dediazoniation of 1 with concomitant pyrimidine ring-opening9 and deprotonation10 leading to intermediate 2^{11} (Scheme 1). It is our hypothesis that the highly reactive dihydroimidazole 2 adds water to form the cyanoamine 3 which then cyclizes to oxanine 5. The cyanoimine 2 also may add water to cyanoimine instead of the ketene moiety, and the resulting ketene might then ring-close to xanthine. Our published theoretical predictions are now being tested experimentally, and as a first step, we targeted the amide 4^{12} for one practical and one conceptual reason (vide infra).

Here, we report the synthesis of **4a** and its recyclization to guanine **6a** and isoguanine **12a**. Importantly, we were able to purify **4a** so that we can firmly establish that **4a** is the precursor for the formations of **6a** and **12a**. We employed the R-group $CH_2-O-CH_2-CH_2-OH$ as a sugar model. This (2-hydroxyethoxy)methyl group occurs in acyclovir, and we indicate this R-group by adding "**a**" to the compound numbers.

The synthesis of **4a** is outlined in Scheme 2. Compound **16** was synthesized with the efficient method by Clausen¹³ via alkylation of **13**, condensation of **14** with benzoyl isothiocyanate, and hydrolysis of **15**. The methylation of **16** in basic aqueous solution gave low yields¹⁴ even though this procedure works well for the related ester.¹⁵ We found the methylation of **16** in the mixed solvent 5/1 acetone/methanol to give **17** in good yield.

Cyanoamine 4a was obtained by methylthioether elimination from 17a. Yamazaki, Okutsu, and Yamada¹⁶ studied the basecatalyzed methylthioether elimination from 17 (R = H, 17h) leading to guanine 6 and isoguanine 12 via cyanoamine 4 (R = H, 4h). The unstable cyanoamine 4h was not isolated and was characterized only by its IR spectrum (ν (C=N) = 2190 cm⁻¹). We have now succeeded in the preparation of the cyanoamine 4a and the isolation of this reactive intermediate. Reactions to form 4a were quenched in liquid nitrogen after 3 min, and after HPLC separation and lyophilization, 4a was stored at -18 °C to prevent cyclization. The cyanoamine 4a was characterized by mass spectrometry (MS, MS/ MS, HRMS), by IR spectroscopy (ν (C=N) = 2138.6 cm⁻¹), and by ¹H and ¹³C NMR spectroscopy. The ¹H NMR spectrum (Figure 1) demonstrates the purity of the synthetic 4a. The cyanoamine 4a may exist in equilibrium with the carbodiimide. Ab initio studies of the parent cyanamide and carbodiimide show a preference of about 7 kcal/mol for the cyanamide,¹⁷ and our IR data also show that the observed tautomer is the cyanoamine 4a. The practical



reason for the choice of 4 is now clear. While 3 cannot be synthesized starting with the acid analogue of 13, with the experience of the synthesis of 4 it appears possible to prepare 3 starting with an ester of 13.

The cyclization of **4a** was studied at room temperature in 0.2M K_2HPO_4/KH_2PO_4 buffer solution (pH = 6.0, 7.0, 8.0, 9.0). Figure 2 shows the HPLC chromatograms recorded at pH = 7.0. There are two products, the guanine **6a** and the isoguanine **12a**, and they were identified by comparison to authentic samples. **6a** is commercially available, and **12a** was synthesized with an adaptation



Figure 1. ¹H NMR spectrum (250 MHz, D₂O, 0.05 N NaOD at 4 °C) of 4a shows signals at δ 7.0 (s, 1H, C₂-H), 5.0 (s, 2H, N1-CH₂), 3.1-3.3 (m, 4H, -CH₂-CH₂-) and establishes the purity of 4a.



Figure 2. HPLC chromatograms of the cyclization of cyanoamine 4a in buffer solution at pH 7 show formation of 6a (aGua) and 12a (aIGua). Final yields are 74.2% 12a and 5.8% 6a with 4.3% 4a remaining.

of the synthesis by Chern, et al.¹⁸ The formation of **6a** from **4a** is a nucleophilic addition of the weakly nucleophilic amide-amino group to the cyanoamine, and this reaction most likely is acidcatalyzed. The formation of 12a is less obvious and does not involve the hydrolysis of 4 with subsequent condensation of the urea-amino group with the amide. Instead, Yamazaki, et al.¹⁶ proposed a mechanism (see SI for details) that involves initial cyclization of the conjugate base of 4. This mechanism accounts for the fact that the amide-O of 4 becomes the carbonyl-O of 12.19 We found that 12a forms faster at lower pH, and we propose a mechanism (see SI) for its formation that does not require the initial deprotonation and that retains all other essential features to proceed via 8 and 10 (Scheme 1). This mechanism involves electrocyclic cyclization, tautomerization, and electrocyclic ring-opening to achieve the O-transfer, and this mechanism also might operate (in addition) at higher pH values.

We have achieved the synthesis of the pure cyanoamine 4a. With the availability of pure 4a, we were able to study its cyclization chemistry. The results of this study have several important implications for nitrosative guanine deamination. (1) The demonstrated formation of 6a from 4a is the model reaction for the



formation of oxanine 5 from 3. (2) Interestingly, the formation of 12a and the explanation of its formation suggests that there is a second pathway to oxanine via an electrocyclic reaction (3 to 7, carbonyl-O as nucleophile), and for X = O, oxanine is the final product of that path because it is known that xanthine 11 does not form from oxanine 7 via rearrangement.⁶ (3) Finally, the conceptual reason for the choice of 4 is as follows: Any potentially formed 2 in nitrosative DNA deamination might add to the amino group of a proximate DNA base, and this addition would lead to Nsubstituted derivatives of 4, as exemplified in Scheme 3. We have synthesized the classical G-to-G cross-link and its structure-isomer 19 by addition of 4a to guanosine.20 The reaction of 4a to 6a strongly suggests that 18 would cyclize to 19 and provide a second path to this cross-link.

Acknowledgment. We are grateful for support by the National Institutes of Health (GM61027).

Supporting Information Available: Details for the syntheses of 17, 4a, and 12a; ¹H and ¹³C NMR spectra of 17, 4a, and 12a; the IR spectrum of 4a; and HPLC chromatograms of the pH-dependent cyclization of 4a (PDF). This material is available free of charge via the Internet at http://pubs.acs.org.

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JA0389523