This article was downloaded by:

On: 25 January 2011

Access details: Access Details: Free Access

Publisher Taylor & Francis

Informa Ltd Registered in England and Wales Registered Number: 1072954 Registered office: Mortimer House, 37-

41 Mortimer Street, London W1T 3JH, UK



Nucleosides, Nucleotides and Nucleic Acids

Publication details, including instructions for authors and subscription information: http://www.informaworld.com/smpp/title~content=t713597286

Mismatch Discrimination of Base-Modified Nucleic Acids and their Constituents: Non-Watson-Crick Base Pairing Induced by Tautomerization

Frank Seela^{ab}; Xiaohua Peng^{ab}; Kuiying Xu^{ab}

^a Laboratorium für Organische und Bioorganische Chemie, Universität Osnabrück, Osnabrück, Germany ^b Laboratory of Bioorganic Chemistry and Chemical Biology, Center for Nanotechnology, Münster, Germany

To cite this Article Seela, Frank , Peng, Xiaohua and Xu, Kuiying(2007) 'Mismatch Discrimination of Base-Modified Nucleic Acids and their Constituents: Non-Watson-Crick Base Pairing Induced by Tautomerization', Nucleosides, Nucleotides and Nucleic Acids, 26: 10, 1569 - 1572

To link to this Article: DOI: 10.1080/15257770701547164 URL: http://dx.doi.org/10.1080/15257770701547164

PLEASE SCROLL DOWN FOR ARTICLE

Full terms and conditions of use: http://www.informaworld.com/terms-and-conditions-of-access.pdf

This article may be used for research, teaching and private study purposes. Any substantial or systematic reproduction, re-distribution, re-selling, loan or sub-licensing, systematic supply or distribution in any form to anyone is expressly forbidden.

The publisher does not give any warranty express or implied or make any representation that the contents will be complete or accurate or up to date. The accuracy of any instructions, formulae and drug doses should be independently verified with primary sources. The publisher shall not be liable for any loss, actions, claims, proceedings, demand or costs or damages whatsoever or howsoever caused arising directly or indirectly in connection with or arising out of the use of this material.

Nucleosides, Nucleotides, and Nucleic Acids, 26:1569-1572, 2007

Copyright © Taylor & Francis Group, LLC ISSN: 1525-7770 print / 1532-2335 online DOI: 10.1080/15257770701547164



MISMATCH DISCRIMINATION OF BASE-MODIFIED NUCLEIC ACIDS AND THEIR CONSTITUENTS: NON-WATSON-CRICK BASE PAIRING INDUCED BY TAUTOMERIZATION

Frank Seela, Xiaohua Peng, and Kuiying Xu

Laboratorium für Organische und Bioorganische Chemie, Universität Osnabrück, Osnabrück, Germany and Laboratory of Bioorganic Chemistry and Chemical Biology, Center for Nanotechnology, Münster, Germany

□ The effect of tautomerism on the mismatch discrimination was studied on 7-deazapurine and 8-aza-7-deazapurine analogues of isoguanosine. 7-Halogenated 7-deaza-2'-deoxyisoguanosines show better base pair discrimination than 2'-deoxyisoguanosine due to the more favored keto tautomer formation. 8-Aza-7-deazaisoguanosine and its 7-halogeno derivatives also show higher keto tautomer population than that of isoguanosine, but the 7-halogens do not bias the tautomeric equilibrium significantly as it is observed for the 7-deaza-2'-deoxyisoguanosine derivatives.

Keywords Nucleosides; oligonucleotides; 7-deazapurines; 8-aza-7-deazapurines; base pair discrimination; tautomers

INTRODUCTION

It is well documented that canonical DNA constituents can form stable non-Watson-Crick base pairs. However, the degree of fidelity achieved by the association of Watson-Crick base pairs is not sufficient to maintain genetic integrity. DNA damages are caused by many factors, e.g., oxidation, irradiation and normal metabolic processes inside the cell. Furthermore, the tautomerism of the canonical nucleobases and the formation of wobble base pairs lead to the DNA mutation.^[1] Diagnostic tools have been developed to detect mutation (single nucleotide polymorphisms, SNPs) by hybridization in solution or on polymer surfaces (biochips). Modified nucleosides carrying fluorescent dyes (e.g., 7-deazapurine nucleoside triphosphates) are used in these protocols.^[2] Here, the effect of the tautomerism was studied with 7-deaza-2'-deoxyisoguanosine, 8-aza-7-deazaisoguanosine and their

Financial support by the Bundesministerium für Forschung and Technologie (ChemBiotech, Germany), Idenix Pharmaceuticals (USA), and the Roche Diagnostics GmbH (Germany) is gratefully acknowledged.

Address correspondence to Frank Seela, Laboratorium für Organische und Bioorganische Chemie, Universität Osnabrück, Barbarastra β e 7, 49069 Onsabrück, Germany. E-mail: Frank. Seela@unionsabrück.de

1570 F. Seela et al.

7-halogenated derivatives. Experiments regarding mismatch discrimination are performed.

RESULTS AND DISCUSSION

Base Recognition of 7-Deaza-2'-Deoxyisoguanosine, 8-Aza-7-Deazaisoguanosine, and Their 7-Halogenated Derivatives

2'-Deoxyisoguanosine (iG_d, 1) is one of the most promiscuous nucleosides. The ambiguity of base pairing caused by the tautomerism of iG_d (motifs I and II; Scheme 1) limits its use as an additional "letter in the genetic alphabet."[3] This has been correlated to a particular property of the isoguanine base which adopts the form of an enol tautomer much more easily than the canonical DNA nucleobases $(K_{\text{TAUT}} = [\text{keto}]/[\text{enol}] \approx 10)$. [4] 7-Deaza-2'-deoxyisoguanosine (c^7iG_d , **2a**) shows a more favored keto form population $(K_{\text{TAUT}} \approx 10^3)$ than that of iG_d . [5] However, compound **2a** decreases the duplex stability and still shows base pair ambiguity as is observed for iG_d . We have found that the introduction of 7-halogen substituents greatly increases the keto content with $K_{\text{TAUT}} \approx 10,000$ for 7-chloro-, or 7-bromo-7-deaza-2'-deoxyisoguanosine (2b) (Scheme 1).^[6] As we want to combine the advantages of high duplex stability with better base discrimination, compounds 2a, b were incorporated into oligonucleotides and their base recognition was investigated in both parallel (ps) and antiparallel duplex (aps) DNA.

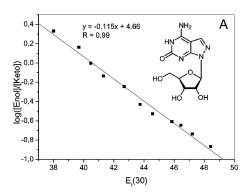
SCHEME 1 Tautomerism of iG_d and **2**, **3** and base pair motifs.

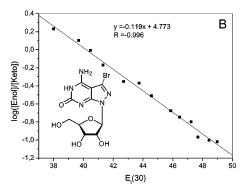
	3'-d(ATCCA X TTATGA)-5' (aps) 5'-d(TAGGT Y AATACT)-3'					5'-d(TiCATAAiCT X XAT)-3' e (ps) 5'-d(AG TATT GA Y CTA)-3'				
$\mathbf{X} \backslash \mathbf{Y}^c$	iC	С	G	T	A	С	G	T	A	iC
iG _d	54	44	44	42 (-12)	32	44	36	33 (-11)	34	40
2a 2b	54 57	43 47	42 45	42 (-12) 40 (- 17)	29 27	44 49	33 36	31 (-13) 31 (- 18)	32 31	37 45

TABLE 1 $T_{\rm m}$ -values of duplexes with mismatches opposite to $iG_{\rm d}$ and $2a,b^{a,b}$

As shown in Table 1, the incorporation of the 7-halogenated derivatives **2b** results in an increase of the $T_{\rm m}$ -values both in *aps* and *ps* DNA duplexes base pairing with iC($\Delta T_{\rm m} = +3.0 \sim 5.0^{\circ}$ C). Furthermore, compound **2b** shows an enhanced base discrimination (in particular against dT) both in *aps* and *ps* DNA indicated from higher $|\Delta T_{m}|$ values ($1 \approx 2a < 2b$). Apparently, a tridentate mispair cannot be formed between the halogenated compound **2b** with dT due to the low content of the enol form ($K_{\rm TAUT} \approx 10^4$), while the base pair formed with 2'-deoxy-5-methylsocytidine (motif IIIa; *aps* DNA) or 2'-deoxycytidine (*ps* DNA) is predominant.

Next, the tautomeric equilibria of the 8-aza-7-deazaisoguanosines $\bf 3a$, $\bf b$ were determined as described. [5,6] Scheme 2 shows the linear relationship of $\log[K_{\rm TAUT}]$ and the $E_{\rm T}(30)^{[9]}$ in the range of 38-50. The $K_{\rm TAUT}$ was calculated by extrapolating this relationship to the $E_{\rm T}(30)$ value of water (63.1). The tautomeric equilibrium constants for compound $\bf 3a$, $\bf b$ were found as 400 for $\bf 3a$, 570 for $\bf 3b$ (Table 2). These values are lower than those of the corresponding 7-deazapurine derivatives $\bf 2a$, $\bf b$ but higher





SCHEME 2 Plot of $\log([\text{enol}]/[\text{keto}])$ versus $E_T(30)$ for compound **3a** (A) and **3b** (B) in mixtures of dioxane ($E_T(30) = 36.0$) and water ($E_T(30) = 63.1$).

 $[^]a$ Measured in 1.0 M NaCl, 0.1 M MgCl₂, and 60 mM Na-cacodylate buffer, pH 7.0, with 5 μ M + 5 μ M single-strand concentration.

^bThe data in parentheses are $\Delta T_{\rm m}$ ($T_{\rm m}^{\rm base\,mismatch} - T_{\rm m}^{\rm base\,match}$).

 $[^]c$ d(iC) = Me iC_d = 5-Methyl-2'-deoxyisocytidine. c X stands for the bases in the row (iG_d, 2a, 2b), Y for the bases in the line (iC, C, G, T, A).

1572 F. Seela et al.

TABLE 2 Tautomeric equilibrium constants for isoguanine derivatives^a

Compound	K_{TAUT}	Compound	K _{TAUT}
2a	1880	3a	400
2b	19700	3b	570

 $^{{}^{}a}K_{\text{TAUT}} = [\text{keto}]/[\text{enol}], \text{ measured in dioxane-water mixtures.}$

than that of isoguanosine (10).^[4] That means compared to **2a, b**, the nucleosides **3a, b** are more enolized; however, compared to isoguanosine, compound **3a, b** enol population is lower. The base pairing properties of 8-aza-7-deaza-2'-deoxyisoguanosine and 7-halogenated derivatives have already been reported.^[8] Studies on the base pair discrimination within oligonucleotides are still in progress.

In summary, 7-halogenated 7-deaza-2'-deoxyisoguanosines, show better base discrimination than 2'-deoxyisoguanosine or 7-deaza-2'-deoxyisoguanosine due to the increased keto population ($K_{\rm TAUT}$ of 20,000 vs. 2000 or 10). In comparison, in 8-aza-7-deazaisoguanosines, 7-halogen substituents do not affect the tautomeric equilibrium significantly.

REFERENCES

- Topal, M.D.; Fresco, J.R. Complementary base pairing and the origin of substitution mutations. Nature 1976, 263, 285–289.
- Prober, J.M.; Trainor, G.L.; Dam, R.J.; Hobbs, F.W.; Robertson, C.W.; Zagursky, R.J.; Cocuzza, A.J.; Jensen, M.A.; Baumeister, K. A system for rapid DNA sequencing with fluorescent chain-terminating dideoxynucleotides. *Science* 1987, 238, 336–341.
- Switzer, C.; Moroney, S.E.; Benner, S.A. Enzymatic incorporation of a new base pair into DNA and RNA. J. Am. Chem. Soc. 1989, 111, 8322–8323.
- Sepiol, J.; Kazimierczuk, Z.; Shugar, D. Tautomerism of isoguanosine and solvent-induced keto-enol equilibrium. Z. Naturforsch. C 1976, 31, 361–370.
- Martinot, T.A.; Benner, S.A. Artificial genetic systems: Exploiting the "aromaticity" formalism to improve the tautomeric ratio for isoguanosine derivatives. J. Org. Chem. 2004, 69, 3972–3975.
- Seela, F.; Peng, X.; Li, H. Base-pairing, tautomerism, and mismatch discrimination of 7-halogenated 7-deaza-2'-deoxyisoguanosine: Oligonucleotide duplexes with parallel and antiparallel chain orientation. *J. Am. Chem. Soc.* 2005, 127, 7739–7751.
- Von Jouanne, J.; Palmer, D.A.; Kelm, H. The use of solvent E_T-values in high pressure kinetics. Bull. Chem. Soc. Jpn. 1978, 51, 463–465.
- Seela, F.; Kröschel, R. The base pairing properties of 8-aza-7-deaza-2'-deoxyisoguanosine and 7halogenated derivatives in oligonucleotide duplex with parallel and antiparallel chain orientation. Nucleic Acids Res. 2003, 31, 7150–7158.