Guanine Oxidation, and Intra-Pair Proton Transfer and Non-Statistical Dissociation of Guanine·Cytosine Base Pair

Jianbo Liu*, Wenchao Lu, and Yan Sun Department of Chemistry and Biochemistry, Queens College and the Graduate Center of CUNY 65-30 Kissena Blvd., Queens, NY 11367, USA

I. Guanine oxidation mechanism and *the questions we want to address*

Guanine nucleobase is the exclusive DNA target



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The only evidence was from photooxidation of an 8-methyl

Formation of transient endoperoxide?

(major at pH < 7)

- **Concerted** [4+2] cycloaddition across **Q2** 4,8-bond OR stepwise addition initialized by 8-peroxide? R Grüber, A Monari, E Dumont, *JPC A*, 2014, 118, 11612
- An endothermic activation barrier **Q3** for oxidation?

E Dumont, R Grüber, E Bignon, C Morell, Y Moreau,





substituted model guanosine compound at -78°C C. Sheu and C. Foote, JACS, 1993, 115, 10446

A Monari, J-L Ravanat, Nucl. Acids Res., 2016, 44, 56





Top frame shows that isolated intermediate/products were short-lived and decomposed within the TOF in mass spectrometer. Bottom frame shows relaxation of products via water

C A dynamics trajectory shows the formation of 5,8-OO- $[G - H]^-$ in collision of $[G - H]^-$

D Evolution of gas-phase products to end product spiroiminodihydantoin (SP) in solution.

 $E^{-1}O_2$ oxidation of guanine nucleoside (A model study)

Oxidation of protonated [9MG + H]^+ is similar to free guanine: reaction is mediated by a [4 + 2] cycloaddition leading to formation of 5,8-endoperoxide.

Oxidation of deprotonated [9MG – H]⁻ is different and proceeds stepwise, starting with a terminal peroxide that subsequently evolves to a 4,8endoperoxide.

2D-PES calculated

Intra-pair proton transfer and *non-statistical* kinetics of deprotonated [Guanine-Cytosine – H]⁻ pair

- Deprotonation of base pairs is one consequence of ionizing radiation interacting with cells, occuring after dissociative electron attachment.
- Deprotonation of a nucleobase prompts intra-pair proton transfer \rightarrow Shift of actual deprotonation site & Structural perturbation

conventional structure

proton-transferred structure

PT equilibrium constants (K_{PT}), ratio of dissociation rate constants (k_{diss}), and RRKM vs. Exp product branching

E _{col} (eV)	K _{PT}	<i>k</i> _{diss} (conventional structure)/ <i>k</i> _{diss} (proton-transferred structure)	Ratio of [G – H] ⁻ / [C – H] ⁻	
			RRKM	EXP
7HG·[C – H] ⁻ (population 62%)				
2.0	2.28	9.65	22.0	—
2.5	2.03	3.64	7.39	—
3.0	1.86	2.54	4.72	-
9HG·[C – H] ⁻ (population 38%)				
2.0	1.54	0.28	0.43	—
2.5	1.44	0.26	0.37	_
3.0	1.38	0.27	0.37	_
	po	opulation-weighted average for [G·C	– H] ⁻	
2.0	-	-	13.8	48.3
2.5	-	-	4.6	36.7
3.0	-	-	3.1	22.9
9MG·[C − H] ⁻				
2.0	1.27	0.10	0.13	8.1
2.5	1.22	0.17	0.21	5.9
3.0	1.18	0.22	0.26	5.9

Origin for non-statistical base-pair dissociation (?)

2.5

E_{ool} (eV)

1) TS re-crossing for intra-pair PT was observed in trajectories TS geometries are twisted that bring about a dynamic bottleneck along the constrained configuration.

E_{col} (eV)

E_{col} (eV)

2) Location of a critical configuration that separates base pair and dissociation products Base pair dissociation has no reverse barrier and the location of the dividing surface depends on energy. We are trying a rigorous way to locate the critical configuration in reaction.

3) RRKM assumes that IVR renders the distribution of energy a random one. But **CID** may produce short time non-statistical fragmentation.

On-going Work

Trajectories are being propagated starting from both activated $G \cdot [C - H1]^{-1}$ and its proton-transferred tautomer, and at the proton-transfer activation barrier.

W. Lu and J. Liu, *PCCP*, 2016, 18, 32222.

Implications for oxidation of guanosine Acidic and basic media may lead to different oxidation chemistries of guanosine, starting from initial stage. W. Lu, H Teng, and J. Liu, PCCP, 2016, 18, 15223.

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