



Computational studies of guanine oxidation

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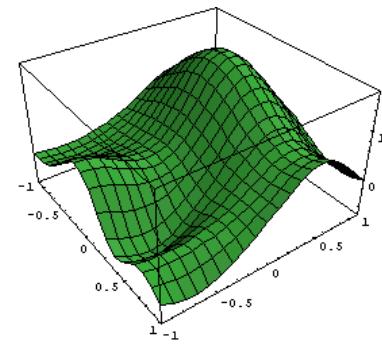
Dr. Jia Zhou

Prof. Richard Lord

Prof. Peng Tao

Prof. Barbara Munk

Dr. Jason Sonk

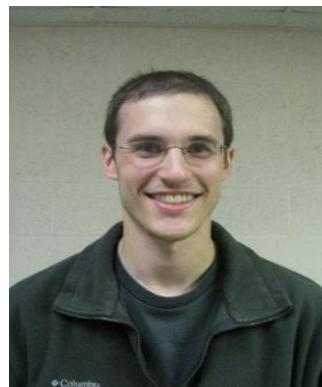




Computational studies of guanine oxidation



Bishnu Thapa



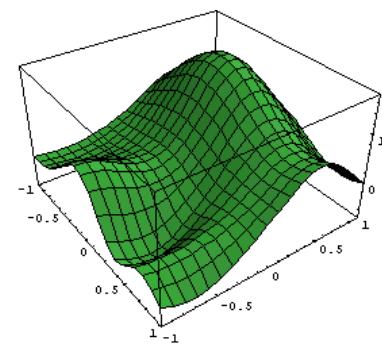
Brian Psciuk



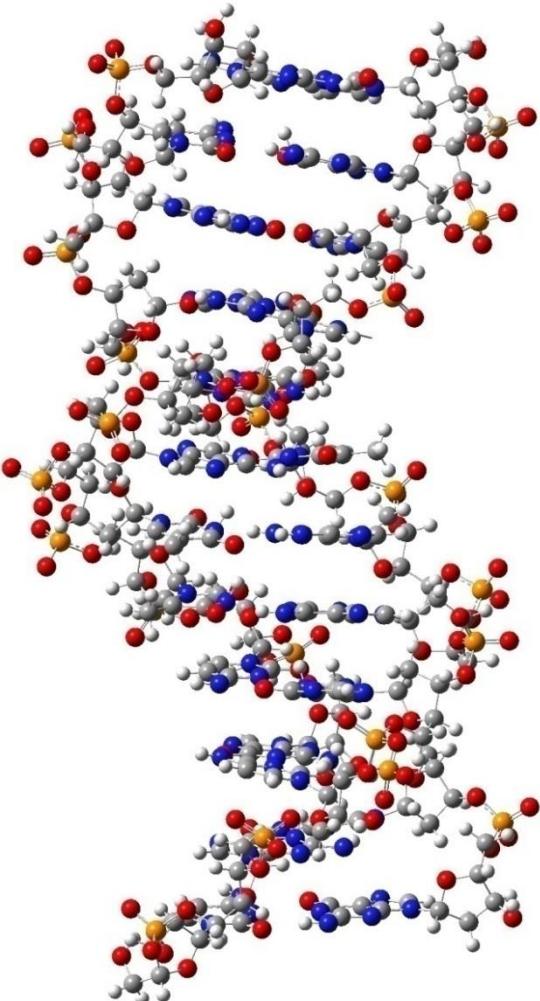
Dr. Barbara Munk



Prof. Cynthia Burrows

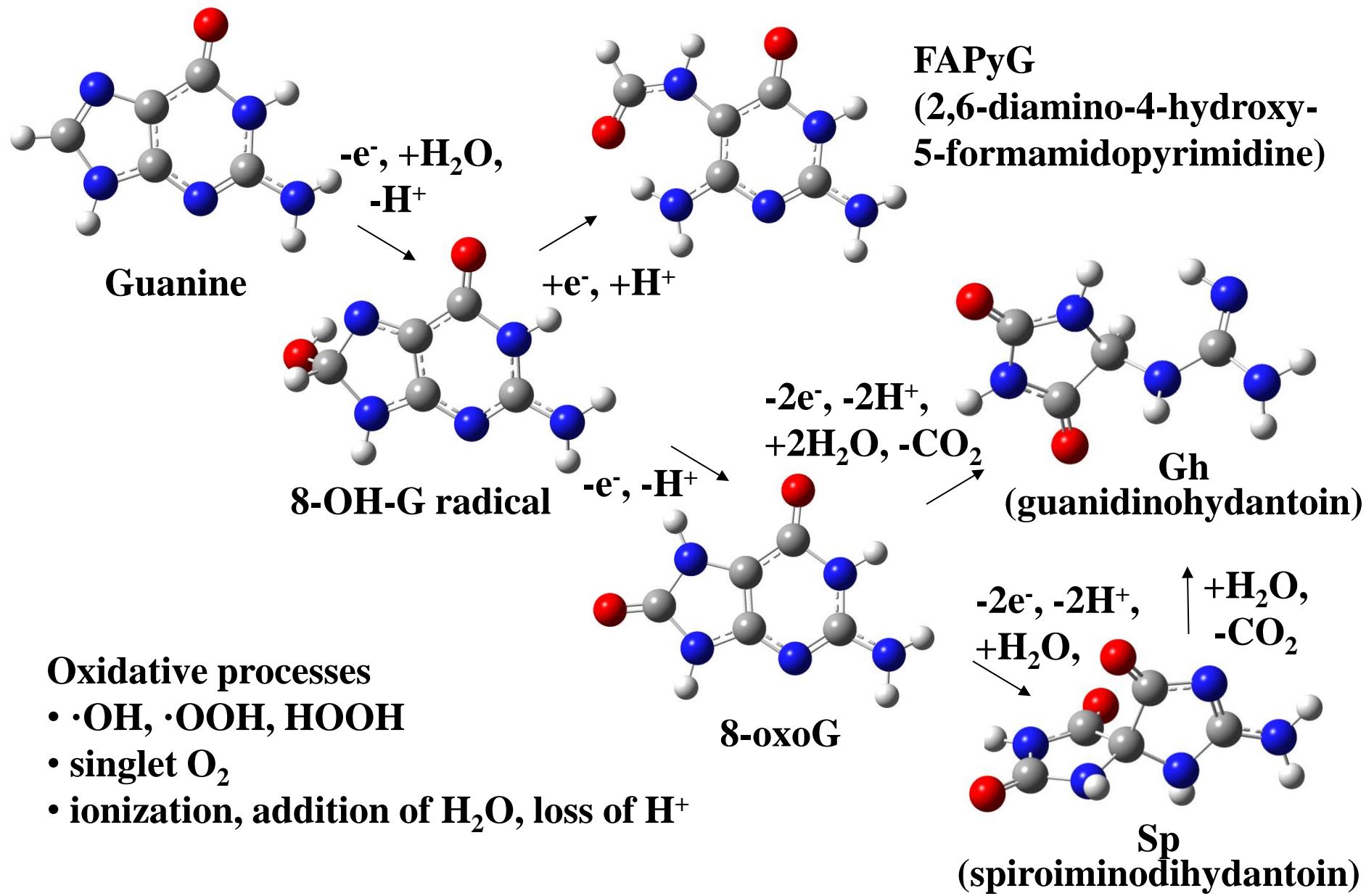


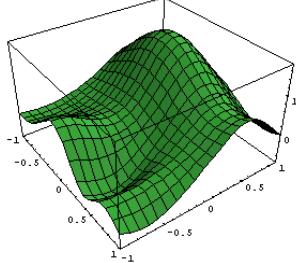
Oxidative Damage to DNA



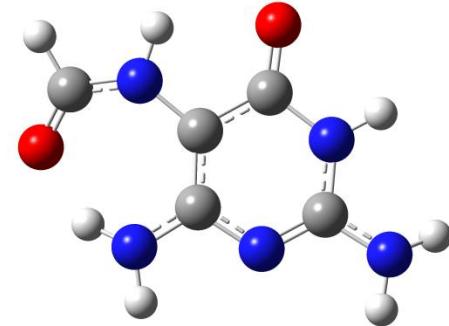
- Studied experimentally by a large number of research groups
 - Burrows et al., *Chem. Rev.*, **1998**, *98*, 1109.
 - Pratviel et al., *Chem.-Eur. J.*, **2006**, *12*, 6018.
 - Gimisis et al., *Eur. J. Org. Chem.*, **2006**, 1351.
 - Delaney et al., *Chem. Res. Toxicol.*, **2008**, *21*, 232.
 - and numerous others
- Caused by a variety of reactive species
 - ◆ O_2^- , ONO_2^- , H_2O_2 , $\cdot\text{OH}$, Cr(VI), etc.
- Can result in the formation of
 - ◆ DNA strand breaks
 - ◆ Protein-DNA crosslinks
 - ◆ Nucleobase mutations
 - ◆ Nucleobase deletions
- Guanine is the most easily oxidized of the nucleobases

Guanine Oxidation Products

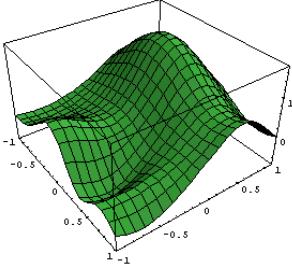




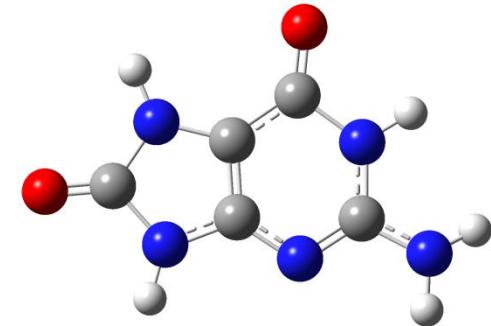
Formamidopyrimidines (FAPyG)



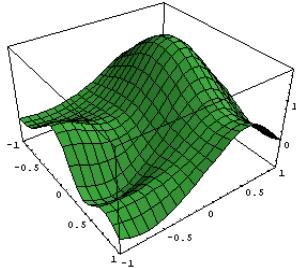
- Produced in vivo with both oxidizing ($\cdot\text{OH}$) and alkylating agents (cyclophosphamide)
- Elevated levels observed in the cortical region of the brain of patients with dementia
- Elevated levels observed in patients with acute lymphoblastic leukaemia, breast cancer and skin cancer. Strong polymerase arresting capability vs 8-oxoG¹
- In vitro
 - ◆ low level of G→T and G→C mutations¹
 - ◆ repaired by *E. coli* DNA glycosylase MutM²



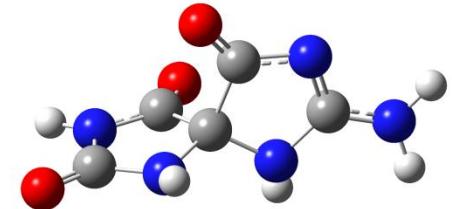
8-Oxoguanine (8-oxoG)



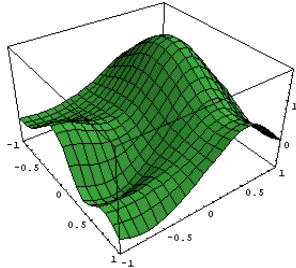
- Most common in vivo (3-4 lesions per 10^7 bases)¹
- Generated at the rate of ca 2000 lesions per cell per day
- Elevated levels observed in patients with neurological disorders such as Parkinson's disease, Alzheimer's disease, and amyotrophic lateral sclerosis (ALS)
- Elevated levels observed in patients with acute lymphoblastic leukemia, and a variety of cancers of the breast, colon, kidney, lung and skin.
- In vitro:
 - ◆ nucleobase bypass (88%) and G→T mutations (7%)²
 - ◆ repaired by *E coli* DNA glycosylases MutM (8-oxoG:C,T,G), Nei (8-oxoG:A) and MutY (8-oxoG:A) and human hOGG1^{2,3}



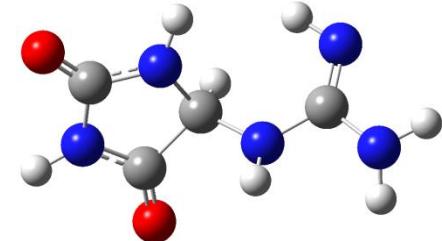
Spiroiminodihydantoin (Sp)



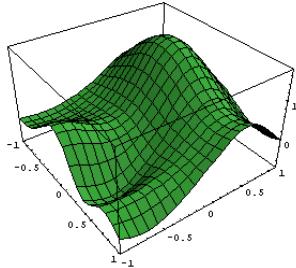
- Produced in vivo following chromate exposure, a known human lung carcinogen¹
- In vitro
 - ◆ nucleobase bypass of 9%
 - ◆ higher mutation frequency than 8-oxoG
 - ◆ G→T (27-41%) and G→C (57-72%) mutations²
 - ◆ Strong block to DNA strand replication²
 - ◆ repaired by *E. coli* Nei (Sp:C,G,A), Nth (Sp:C,G) and MutM (Sp:C,G)²
 - ◆ Not repaired by human hOGG1²



Guanidinohydantoin (Gh)



- Has not yet been isolated from human tissue or fluids
- In vitro data
 - ◆ Nucleobase bypass of 75-80%
 - ◆ Higher mutation rate than 8-oxoG
 - ◆ G→T (1.4%) and G→C (98%) mutations²
 - ◆ Strong block to DNA strand replication²
 - ◆ Repaired by *E coli* Nei (Sp:C,G,A), Nth (Sp:C,G) and MutM (Sp:C,G)
 - ◆ Not repaired by human hOGG1 ²



Methodology

Geometry Optimization:

- (a) B3LYP/6-31+G(d,p)
- (b) IEF-PCM//B3LYP/6-31+G(d,p) (for acyl migrations)

Single points in solution: IEF-PCM/B3LYP/aug-cc-pVTZ

Relative energies in solution:

$$\begin{aligned} & \Delta E_{\text{gas}}(\text{B3LYP/aug-cc-pVTZ//B3LYP/6-31+G(d,p)}) \\ & + \Delta ZPE(\text{B3LYP/6-31+G(d,p)}) \\ & + \Delta \Delta G_{\text{sol}}(\text{IEF-PCM/B3LYP/aug-cc-pVTZ//B3LYP/6-31+G(d,p)}) \end{aligned}$$

Tautomerizations assisted by one or more waters

Cystosine/cytosine cation as proton donor / acceptor

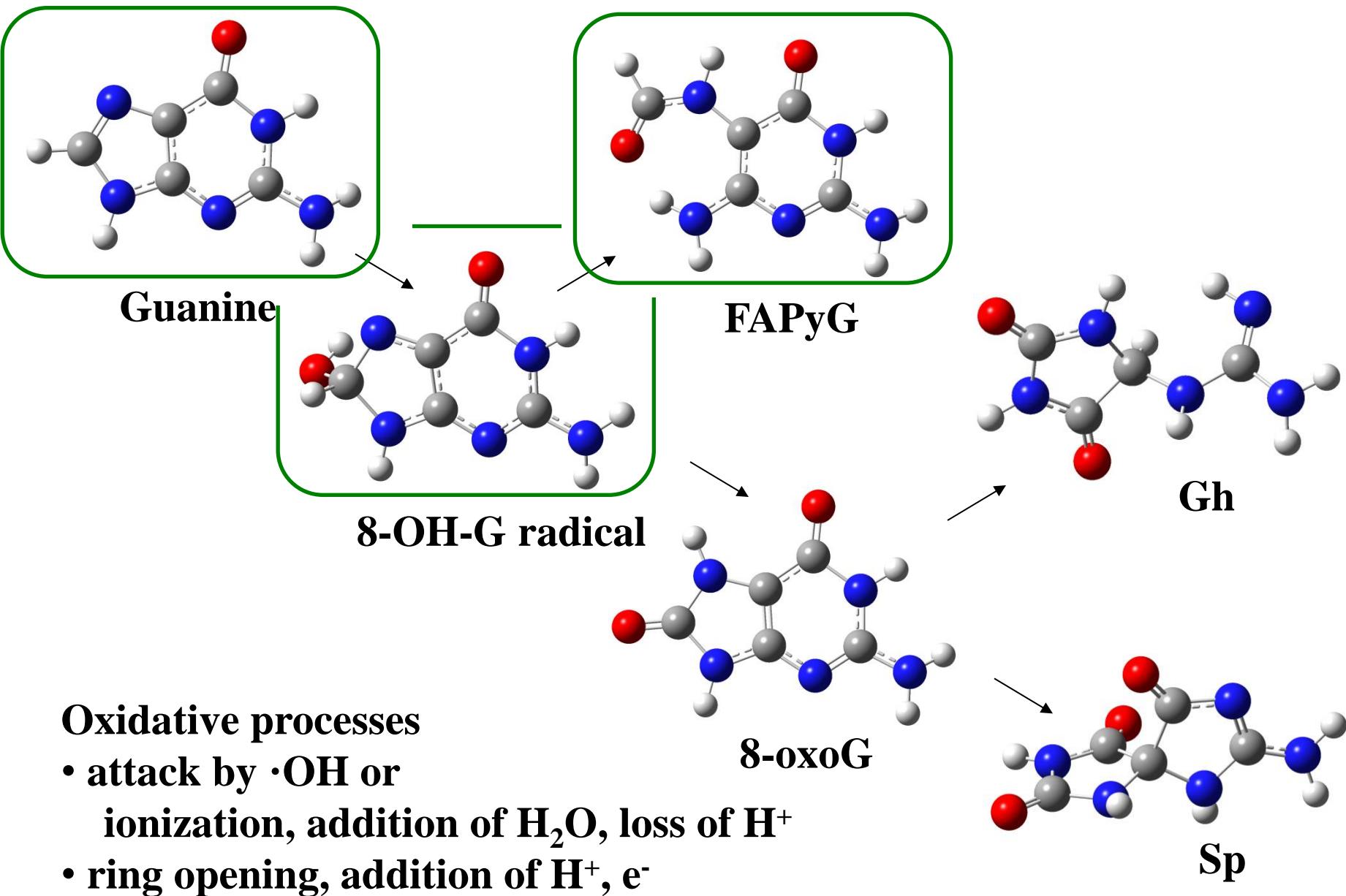
Guanine/guanine radical cation as electron donor / acceptor

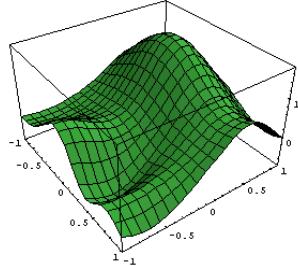
Munk, Burrows, Schlegel, *Chem. Res. Tox.* 2007, 20, 432-444.

Munk, Burrows, Schlegel., *J. Am. Chem. Soc.* 2008, 130, 5245.

Ye, Munk, Miller, Cogbill, Burrows, and Schlegel, *Chem. Res. Toxicol.* 2009, DOI:10.1021/tx800402y

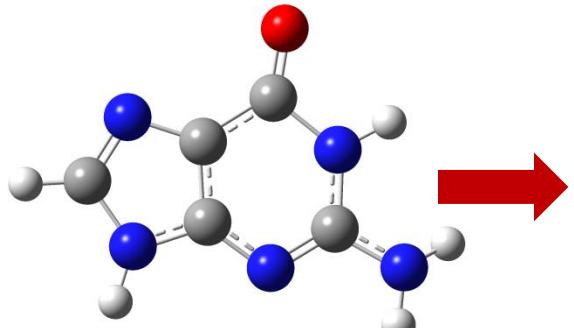
Guanine to 8-OH-G and FAPyG



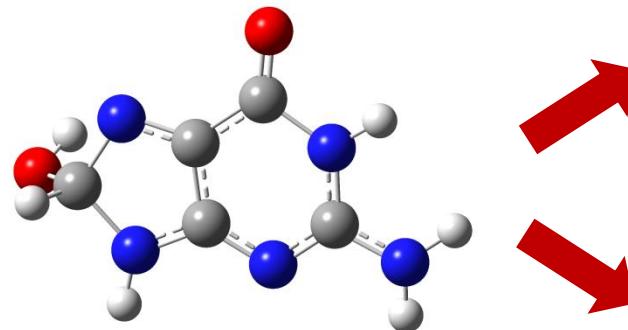


Experimental Studies of the Oxidation Mechanism

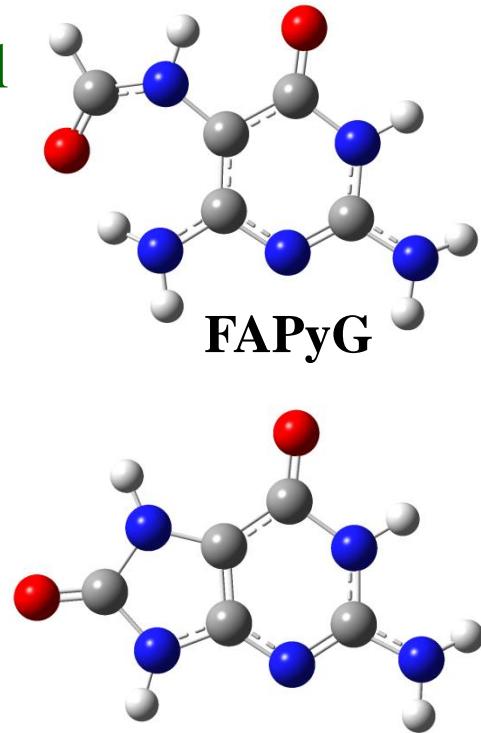
- Several research groups note that both 8-oxoG and FAPyG are probably formed *via* OH radical attack at C8 to form the same intermediate.



Guanine



8-OH-G radical



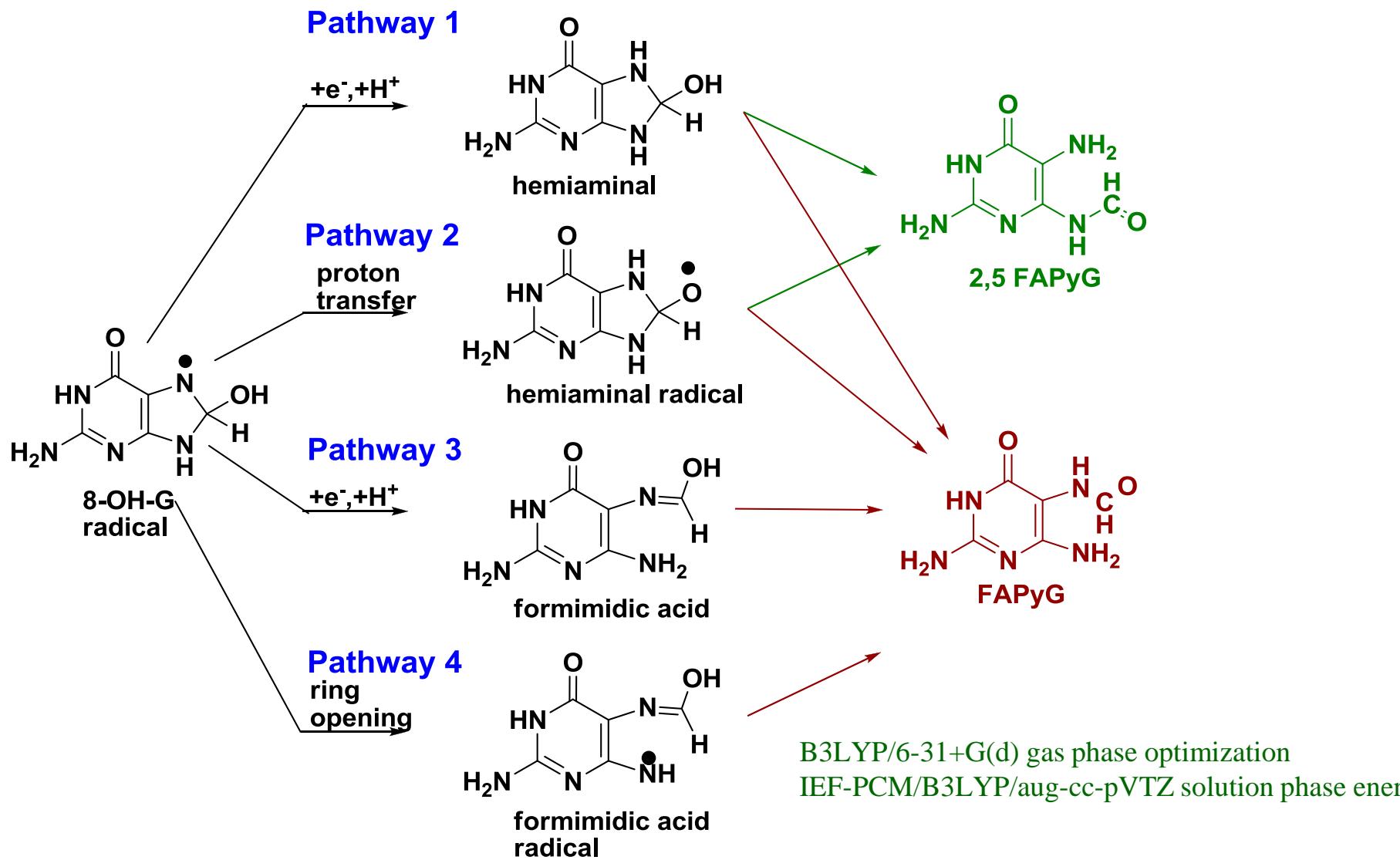
8-oxoG

Steenken, S. *Chem. Rev.* 1989, 89, 503. Breen et al. *Free Radical Biol. Med.* 1995, 18
Burrows et al. *Chem. Rev.* 1998, 98, 1109. Candeias et al. *Chem. Eur. J.* 2000, 6, 475.

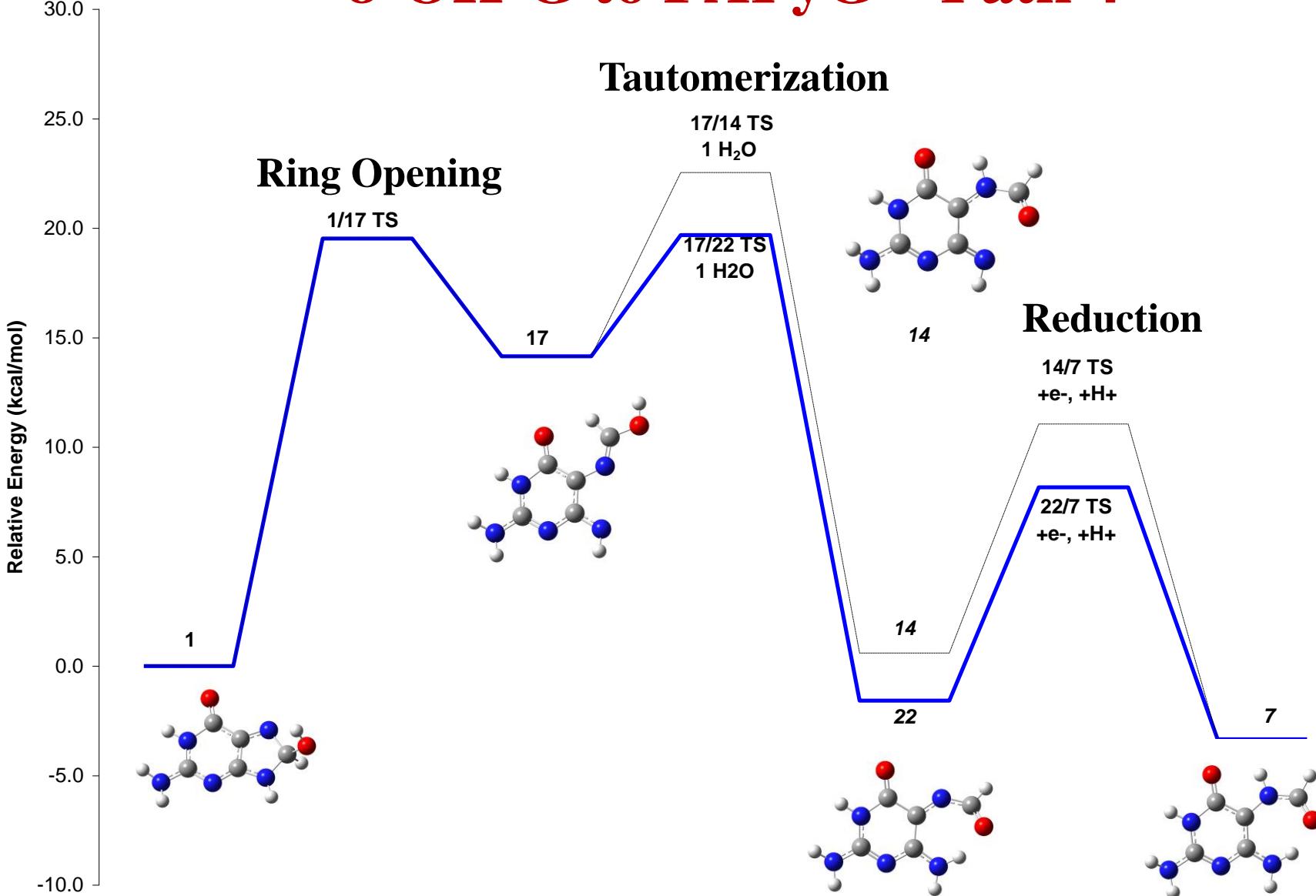
Evans et al. *Mutat. Res.-Rev. Mutat. Res.* 2004, 567, 1. Kalam et al. *Nucleic Acids Res.* 2006, 34, 2305. Birincioglu et al. *J. Am. Chem. Soc.* 2003, 125, 11607. Neeley et al. *Chem. Res. Toxicol.* 2006, 19, 491.

Ober, M., et al.; *J. Am. Chem. Soc.* 2005, 127, 18143-18149. Jiang, Y.L., et al.; *J. Org. Chem.* 2005, 70, 141-149.

Formation of FAPyG from 8-OH-G[•]

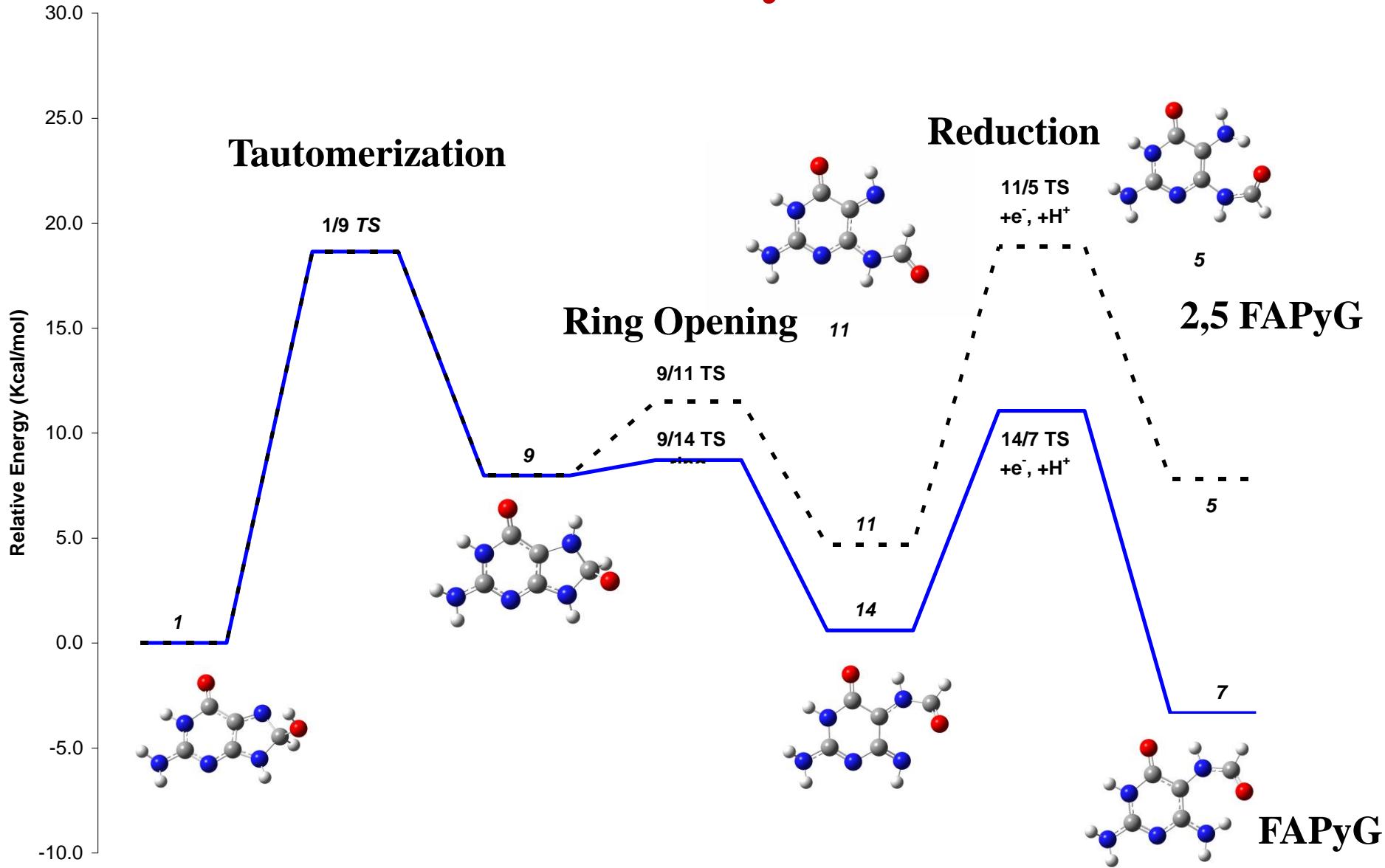


8-OH-G to FAPyG – Path 4

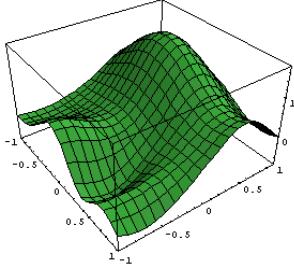


All relative energies were calculated at IEF-PCM/B3LYP/aug-cc-pVTZ//B3LYP/6-31+G(d,p).

8-OH-G to FAPyG – Path 2



All relative energies were calculated at IEF-PCM/B3LYP/aug-cc-pVTZ//B3LYP/6-31+G(d,p).

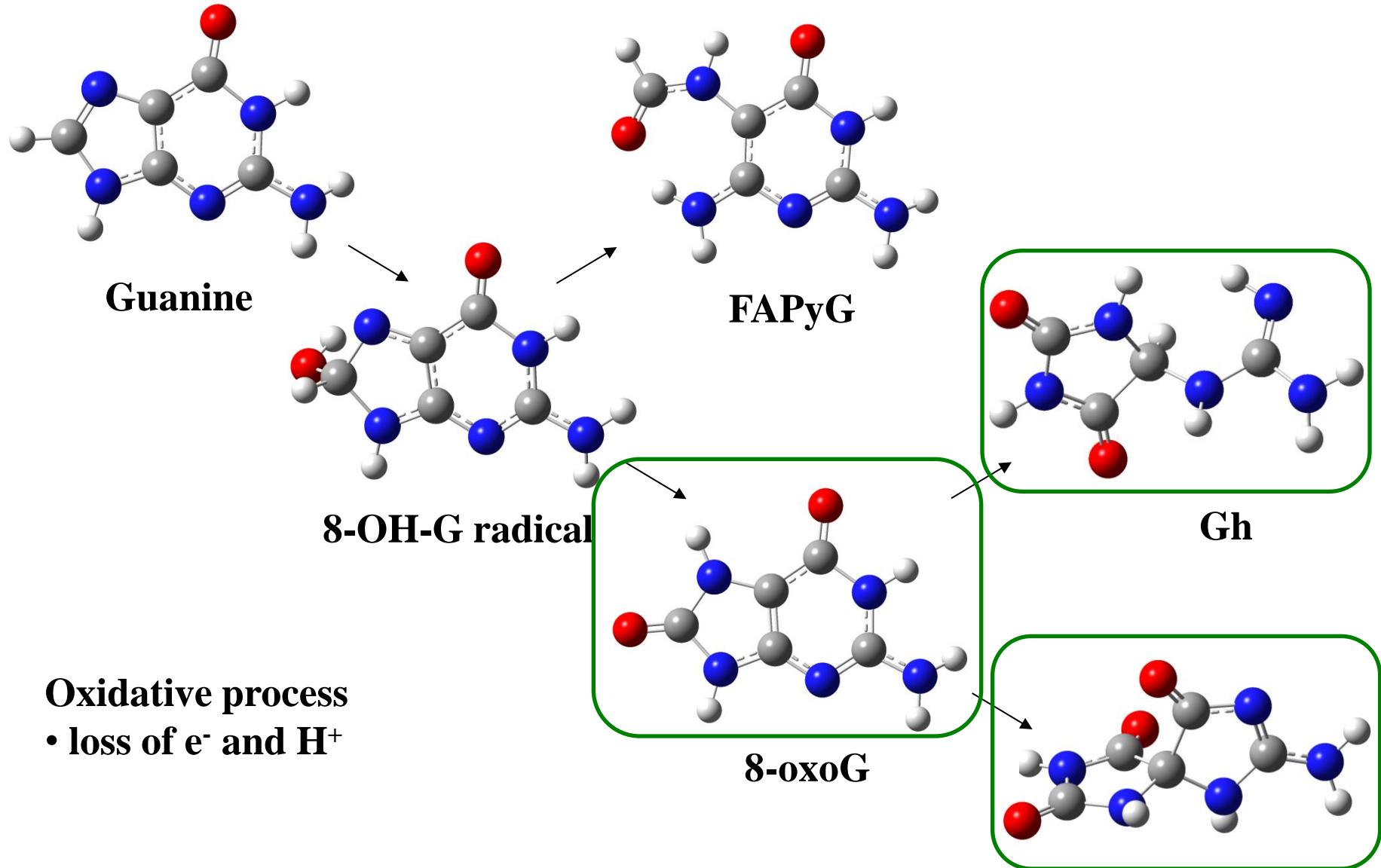


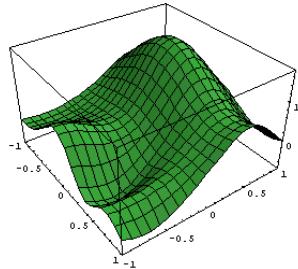
Results

G to FAPyG:

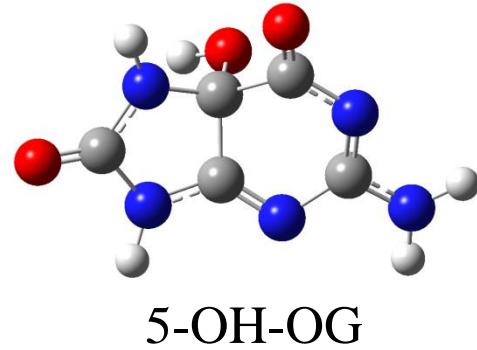
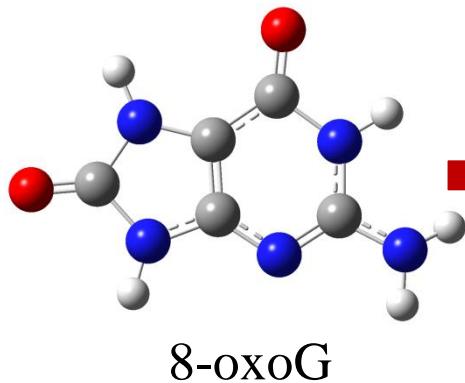
- Of the four reaction pathways studied for formation of FAPyG, Pathway 2, via the 8-oxyguanine radical appears to be the kinetically favored pathway.
- 2,5 FAPyG, a thermodynamically less stable intermediate, may also be formed and IR and NMR calculations suggest that its presence should be differentiable experimentally.

8-oxoG to Gh and Sp

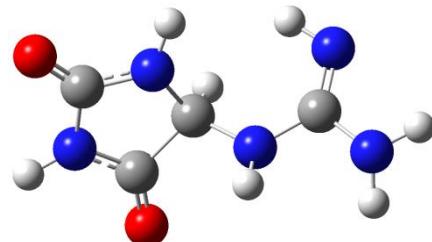
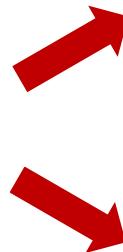




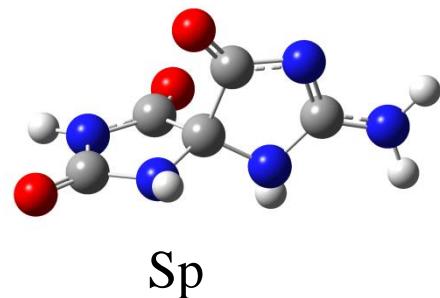
Experimental Studies of 8-Oxoguanine Oxidation



ds DNA
ss DNA, $T < 4^\circ\text{C}$
Nucleosides, pH < 6



ss DNA, $T > 20^\circ\text{C}$
Nucleosides, pH > 7



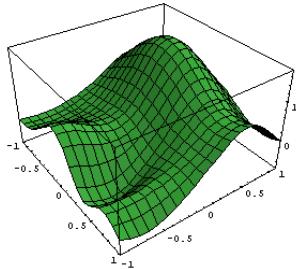
Luo et al., *Org. Lett.*, 2000, 2, 613; *Org. Lett.*, 2001, 3, 280; *Chem. Res. Toxicol.*, 2001, 14, 927.

Kornyushyna et al., *Biochemistry*, 2002, 41, 15304.

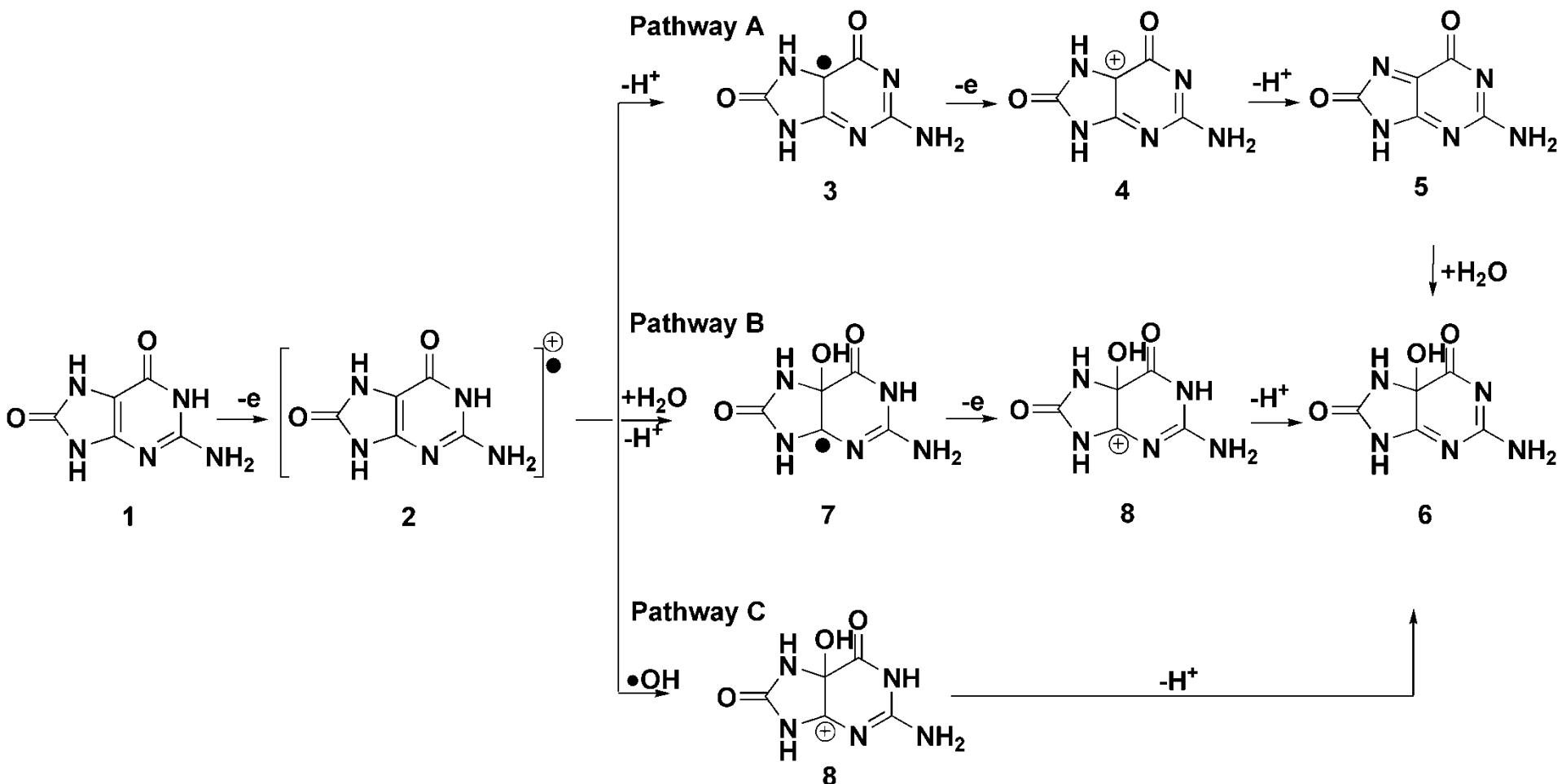
Ye et al., *J. Am. Chem. Soc.*, 2003, 125, 13926.

McCallum et al., *J. Am. Chem. Soc.* 2004, 126, 16777.

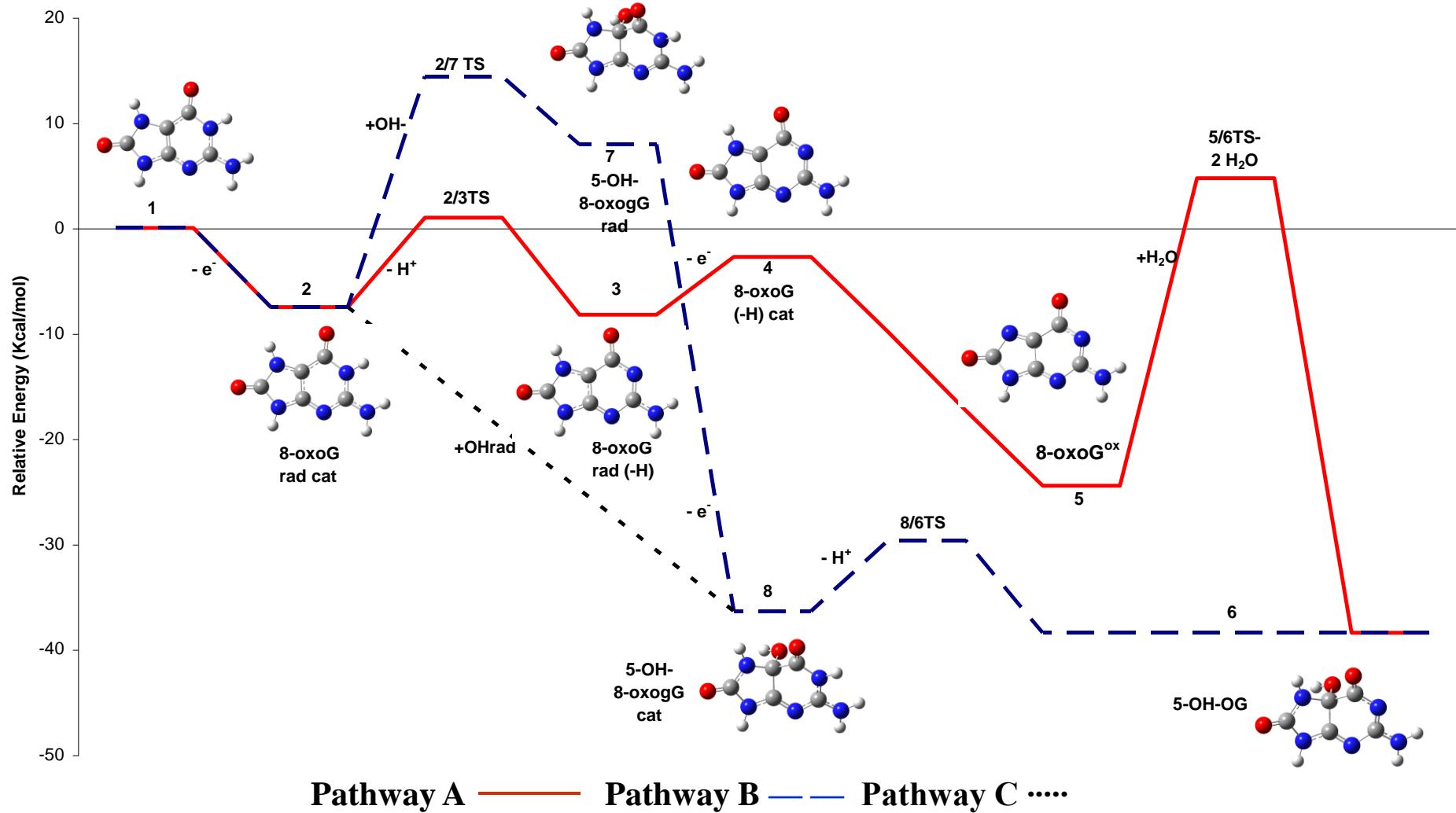
Jia et al., *Biochemistry*, 2005, 44, 13342; *Biochemistry*, 2005, 44, 6043.



8-oxoG to 5-OH-OG Pathways

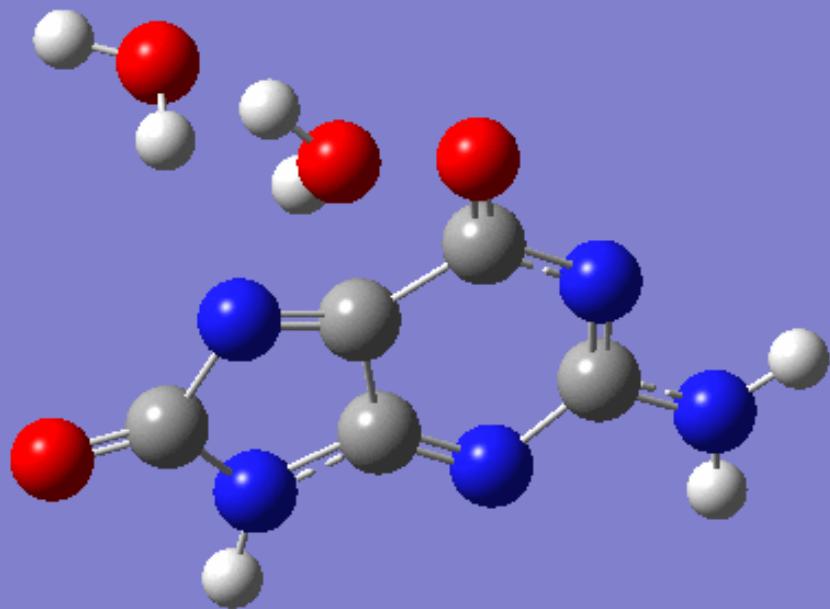


8-oxoG to 5-OH-OG

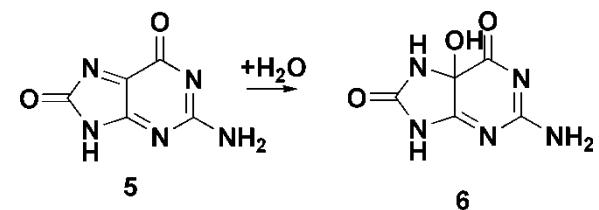


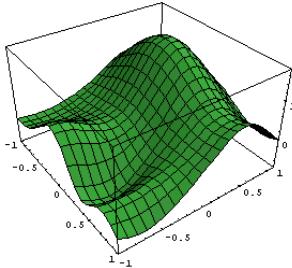
Pathway A ————— Pathway B ————— Pathway C

All relative energies were calculated at IEF-PCM/B3LYP/aug-cc-pVTZ//B3LYP/6-31+G(d,p).



Addition of water to imine



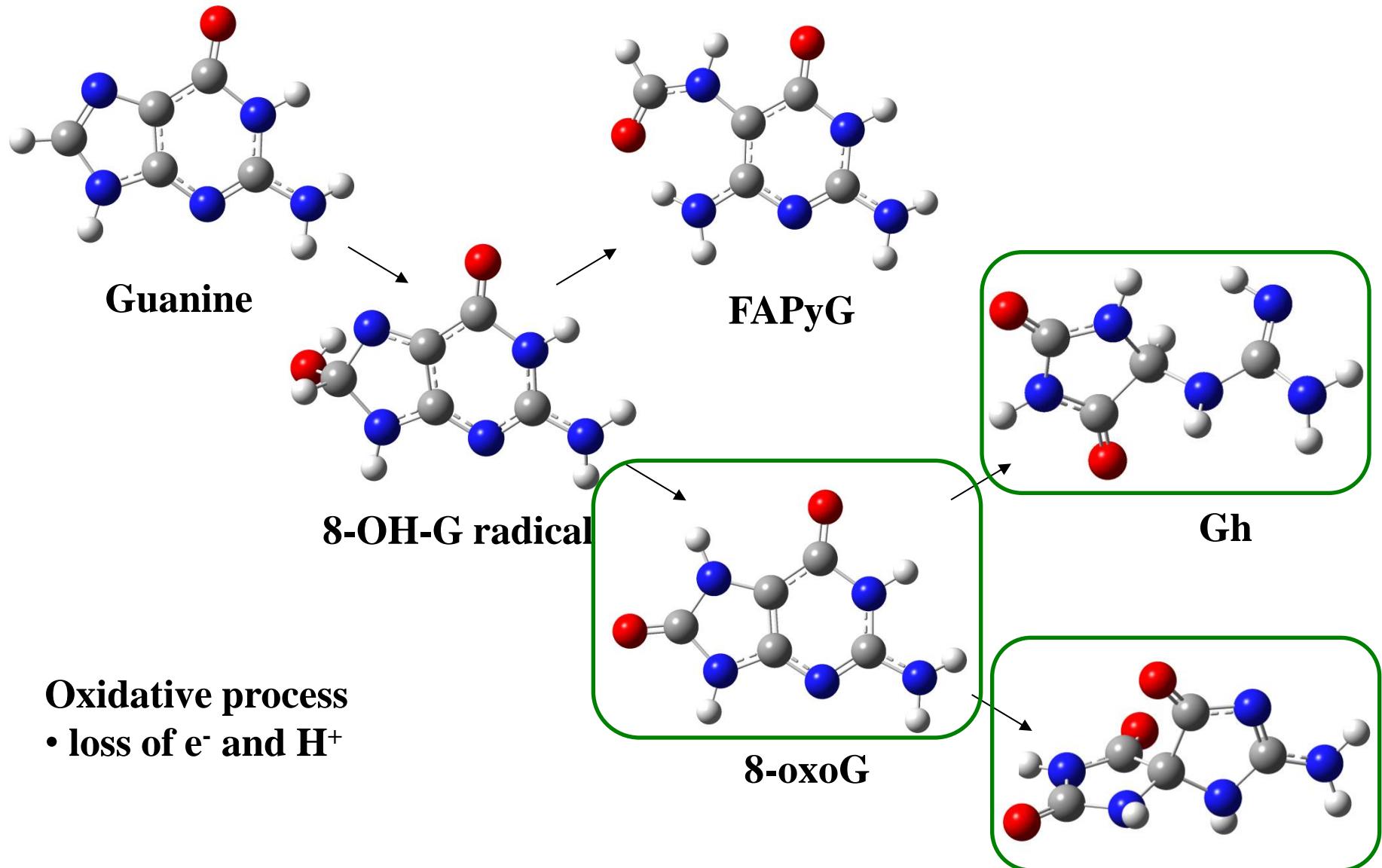


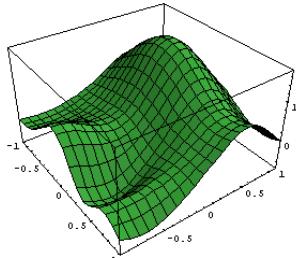
Results

8-oxoG to 5-OH-OG:

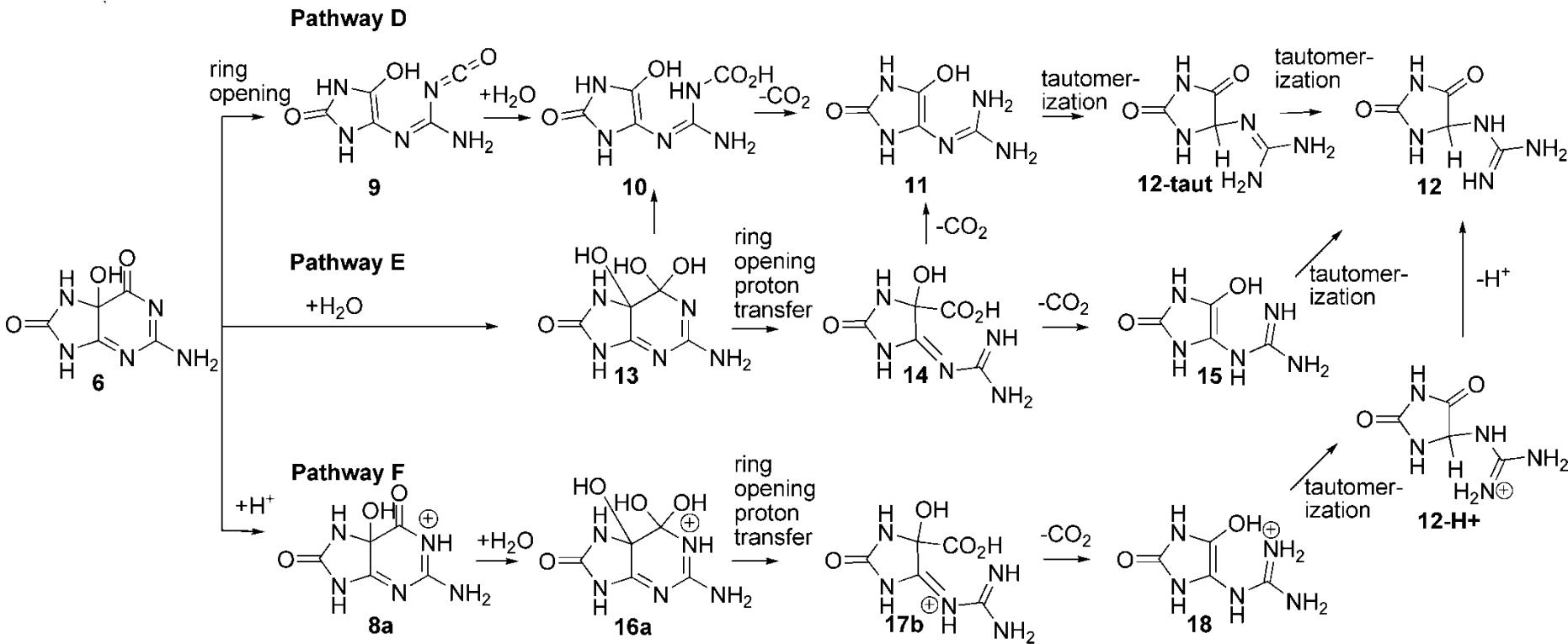
- All three pathways are energetically feasible mechanisms.
- Pathway A may be kinetically favored over Pathway B. Both Pathways A and B depend on the local concentration of oxidizing agent.
- Pathway C, addition of $\cdot\text{OH}$, has lower barriers than either A or B, but will be in competition with other biochemical reactions for the reactive oxygen species, $\cdot\text{OH}$, and may play a smaller role in the production of 5-OH-OG.

5-OH-OG to Gh and Sp

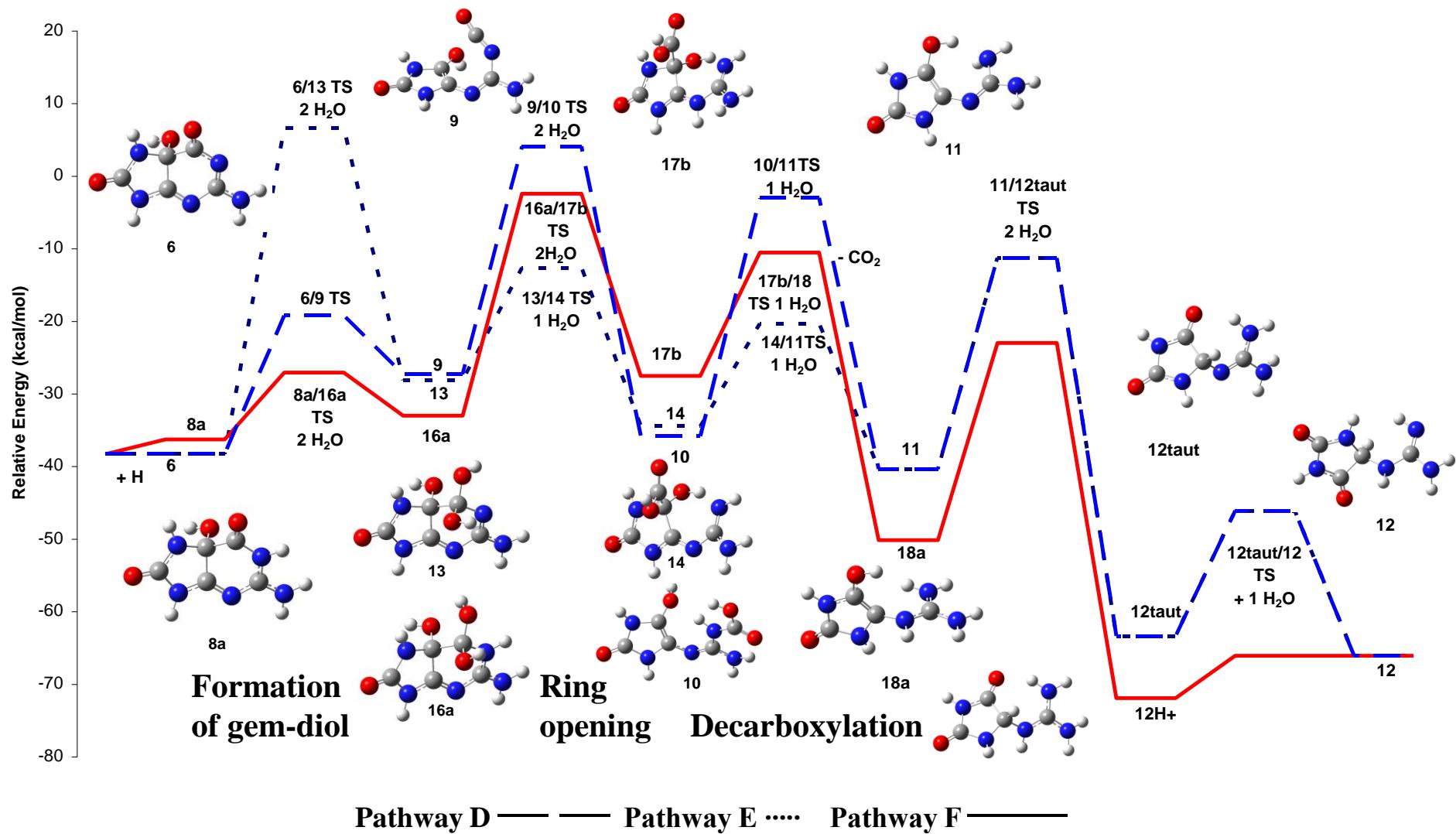




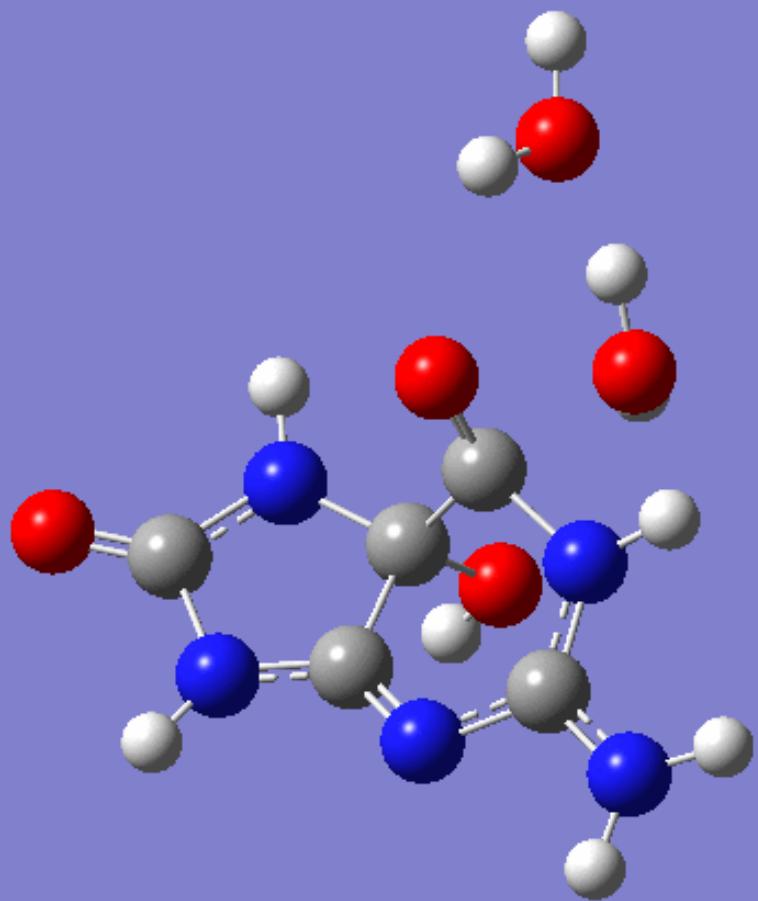
5-OH-OG to Gh Pathways



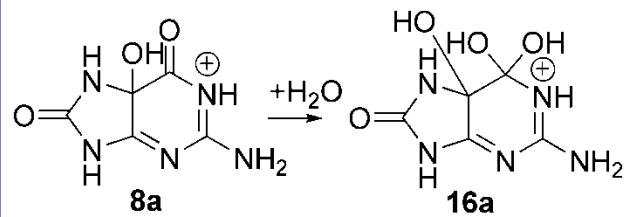
5-OH-OG to Gh

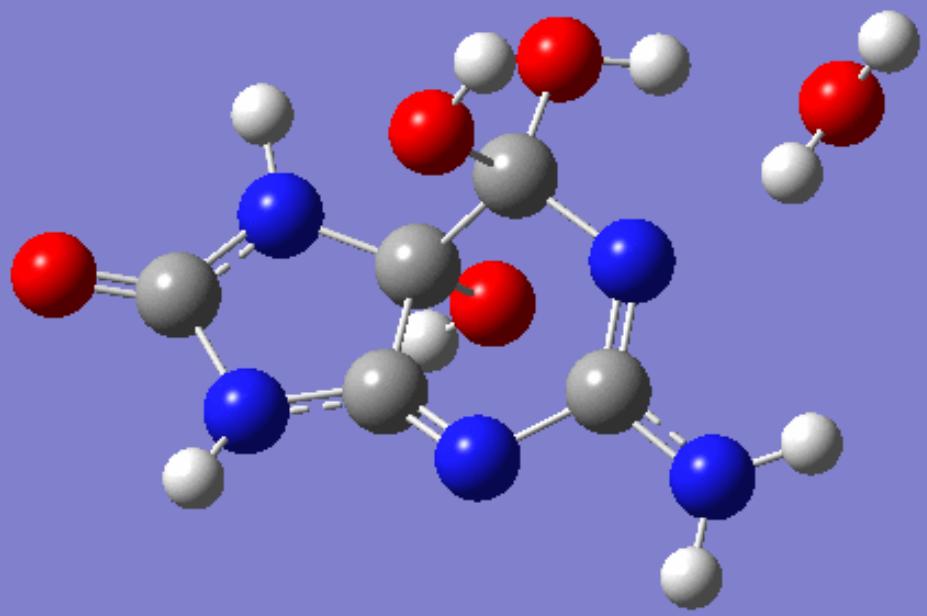


All relative energies were calculated at IEF-PCM/B3LYP/aug-cc-pVTZ//B3LYP/6-31+G(d,p).

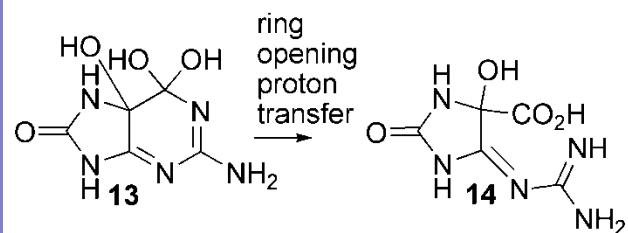


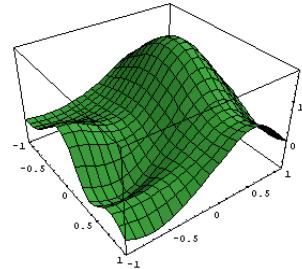
Formation of gem diol



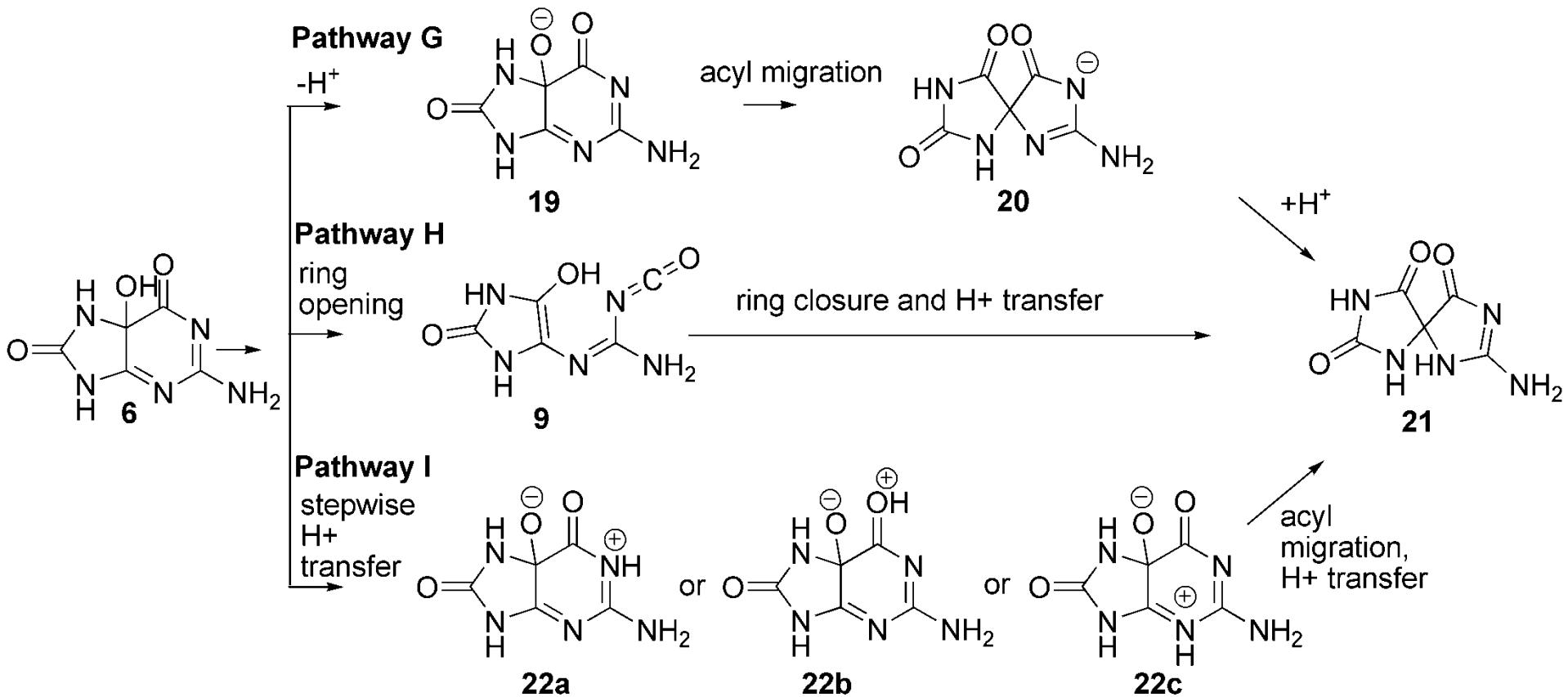


Cleavage of C6-N1 bond

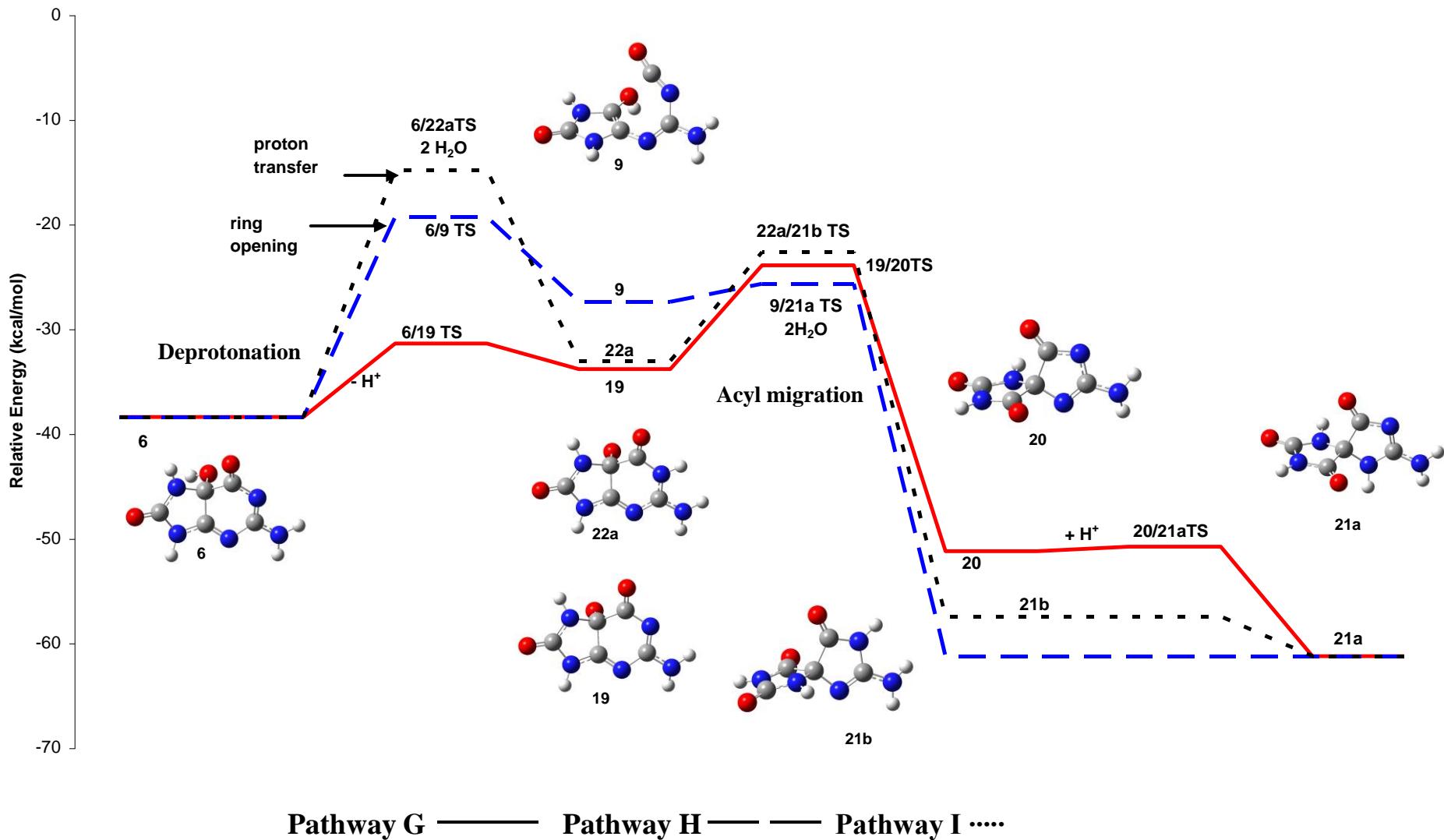




5-OH-OG to Sp Pathways

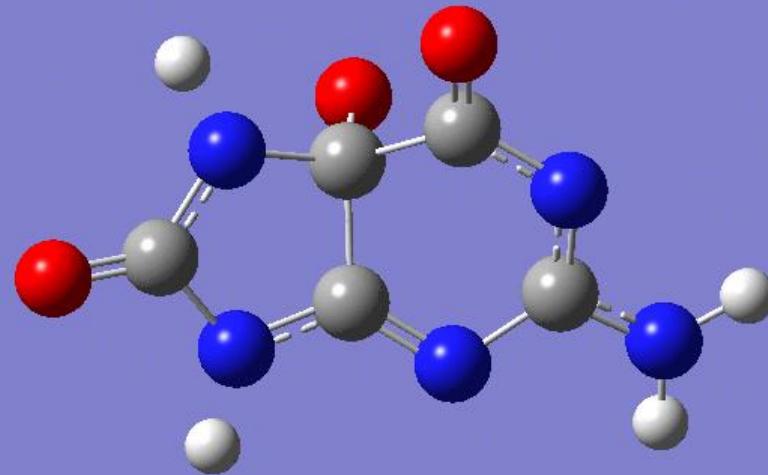
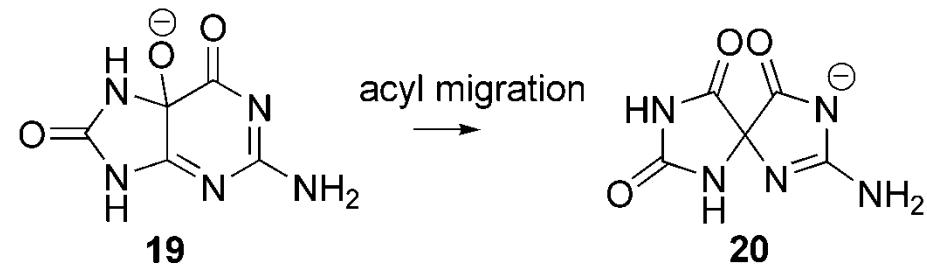


5-OH-OG to Sp



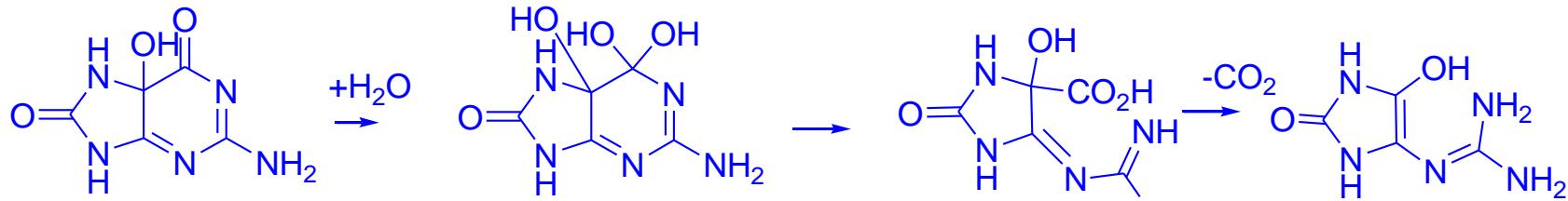
All relative energies were calculated at IEF-PCM/B3LYP/aug-cc-pVTZ//B3LYP/6-31+G(d,p).

Acyl Migration

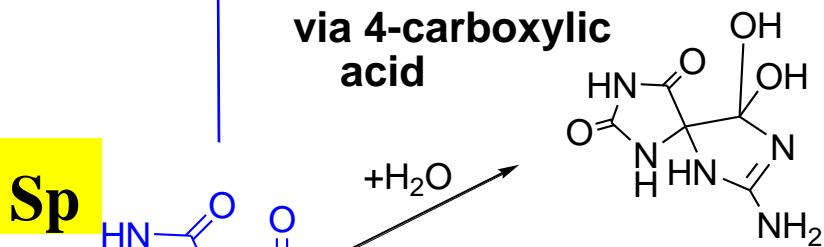


Sp to Gh Pathways

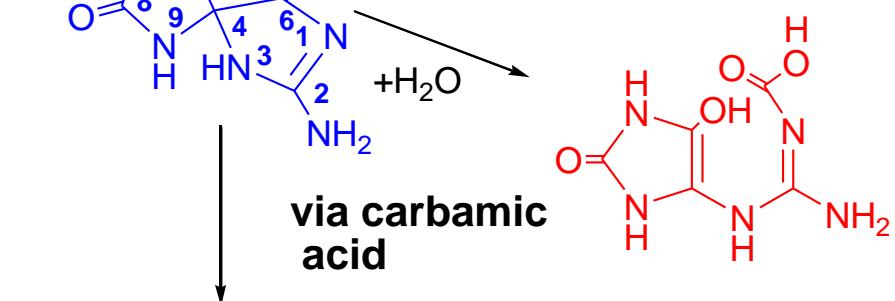
via 5-OH-OG



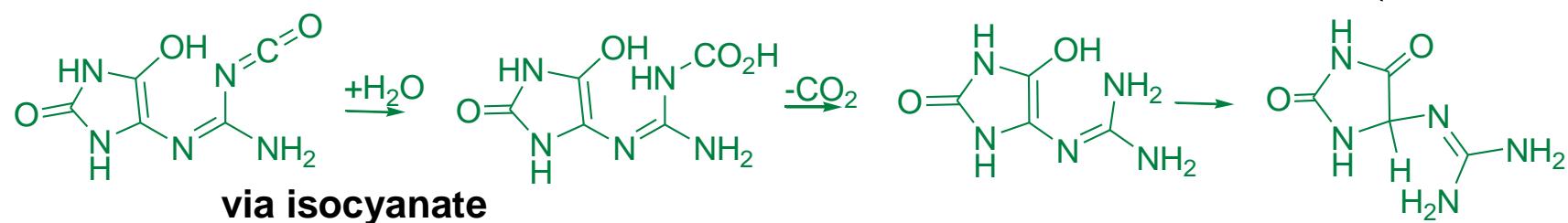
via 4-carboxylic acid



via carbamic acid

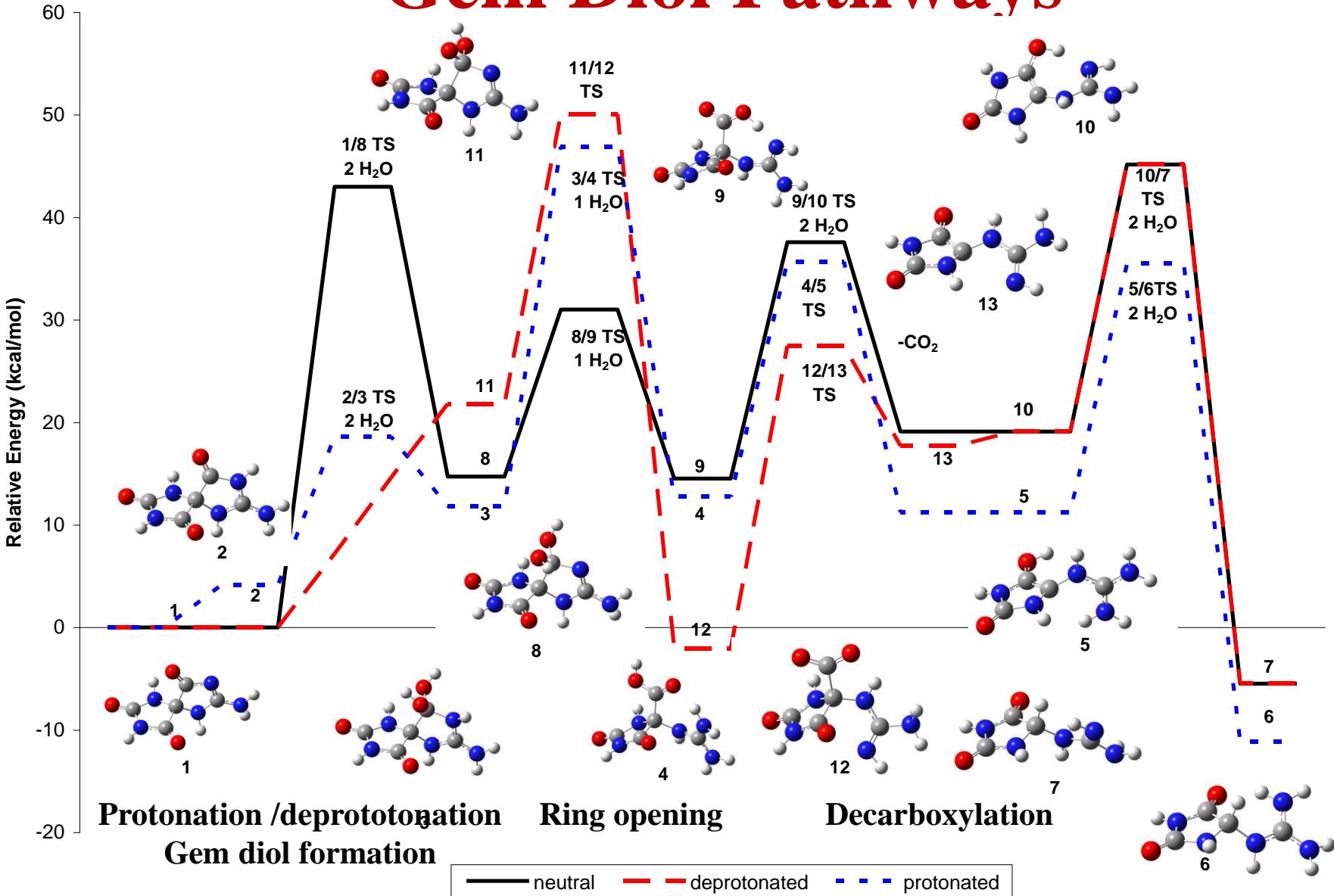


via isocyanate

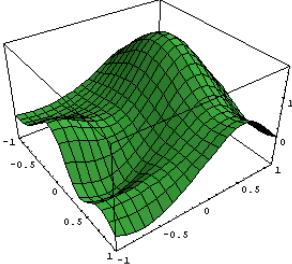


Gh

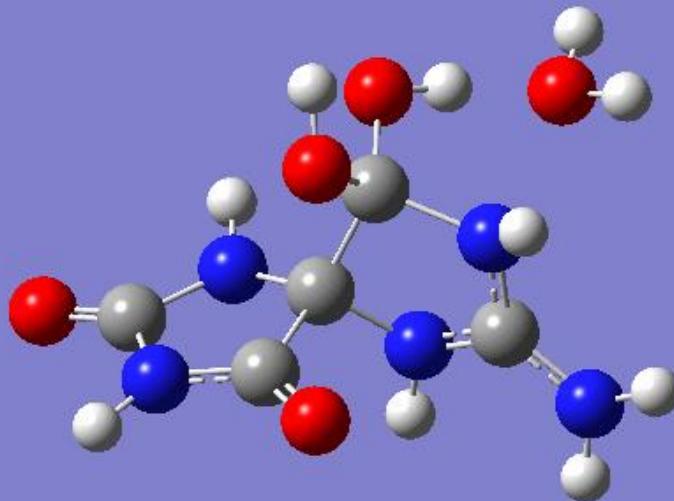
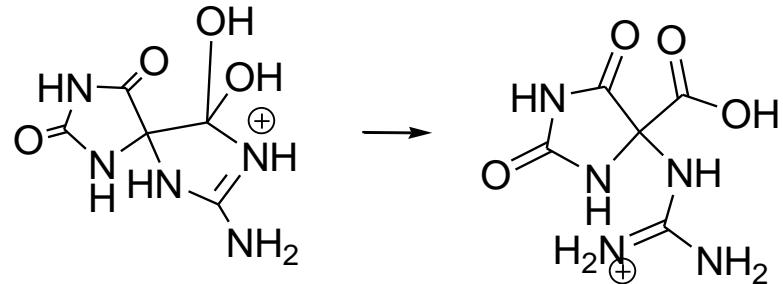
Gem Diol Pathways

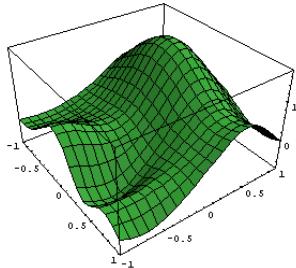


All relative energies were calculated at IEF-PCM/B3LYP/aug-cc-pVTZ//B3LYP/6-31+G(d,p).

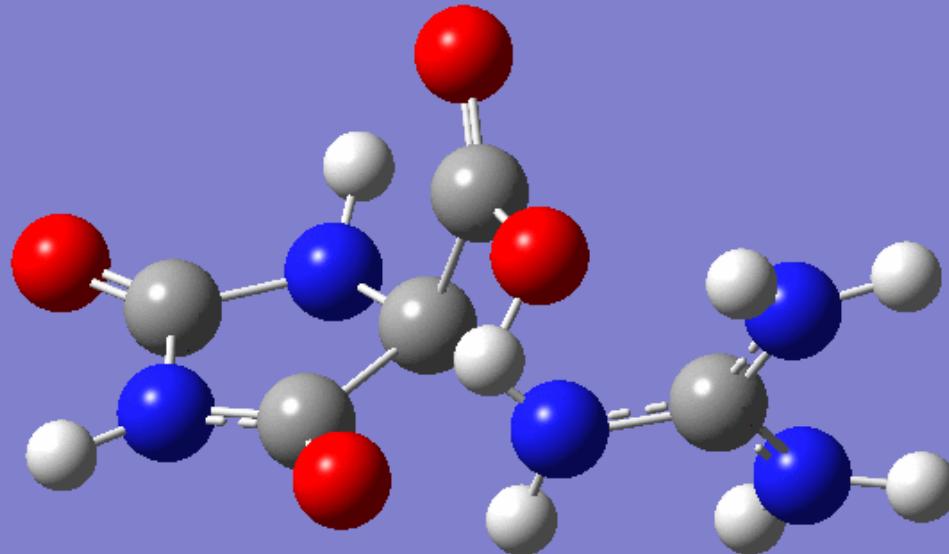
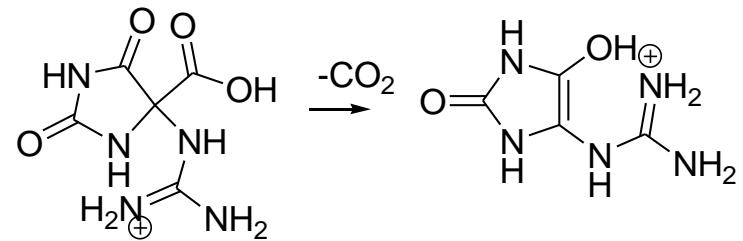


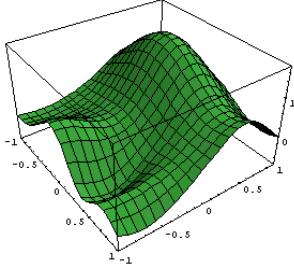
Ring Opening of Protonated Diol





Decarboxylation to form Gh Enol

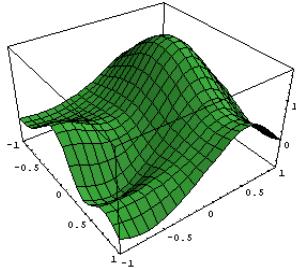




Results

Sp to Gh:

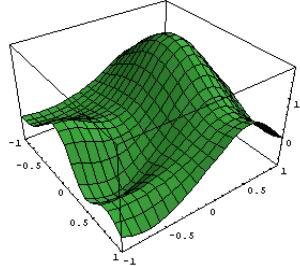
- Most likely pathway is via a Sp C6 gem diol.
- Protonation of Sp at the N1 position is predicted to facilitate formation of the gem diol.
- Consistent with experiment, formation of Gh from Sp is predicted to be favored at low pH.



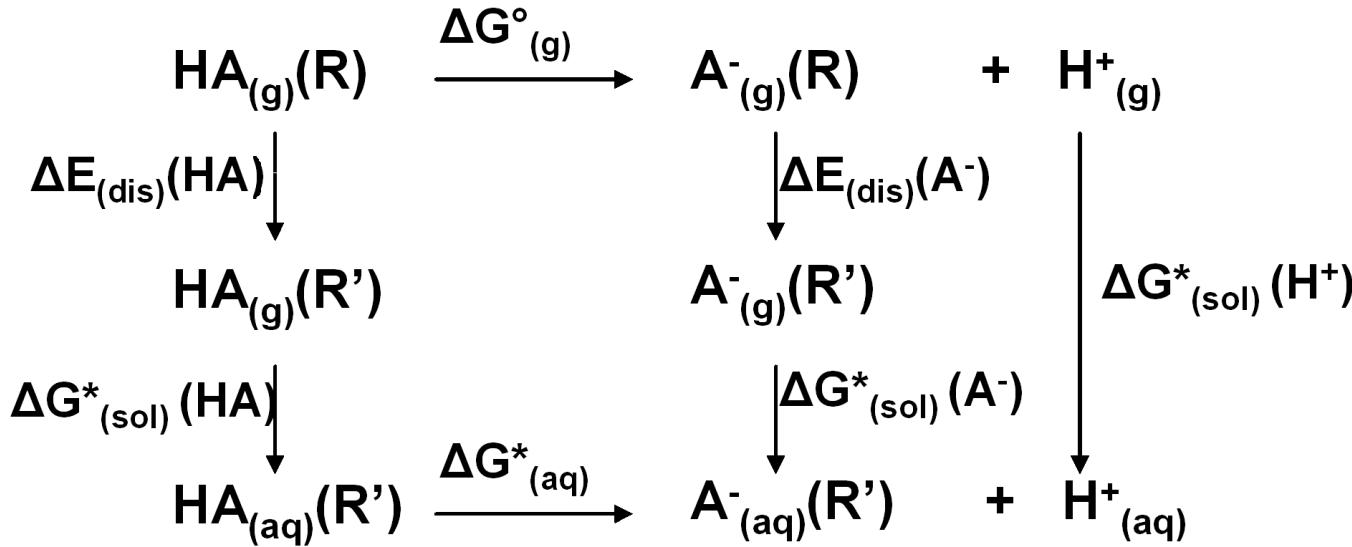
pK_a and Redox Potentials

- Oxidation of guanine occurs in water at physiological pH
- Reaction pathways and barrier heights depend on the protonation / deprotonation of intermediates
- Agents that oxidize guanine can also oxidize other intermediates
- Need a protocol to estimate the pK_a's and redox potentials along the guanine oxidation pathway

Calculation of pK_a's



$$pK_a = \frac{1}{2.303 RT} \Delta G_{(aq)}$$



$$\Delta G^\circ_{(g)} = G^\circ_{(g)}(A^-_R) + G^\circ_{(g)}(H^+) - G^\circ_{(g)}(HA_R)$$

$$\Delta G^*_{solv}(H^+) = G^*_{(aq)}(H^+) - G^*_{(g)}(H^+)$$

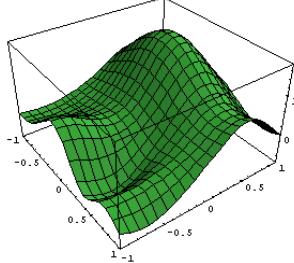
$$\Delta G^*_{(\text{aq})} = G^*_{(\text{aq})}(A^-_{R'}) + G^*_{(\text{aq})}(H^+) - G^*_{(\text{aq})}(HA_{R'})$$

$$\Delta G^*_{solv}(A^-) = G^*_{(aq)}(A^-_{R'}) - G^*_{(g)}(A^-_{R'})$$

$$E_{dis}(A^-) = G^\circ_{(g)}(A^-_{R'}) - G^\circ_{(g)}(A^-_R)$$

$$\Delta G^*_{solv}(HA) = G^*_{(\text{aq})}(HA_{R'}) - G^*_{(g)}(HA_R)$$

$$E_{dis}(HA) = G^\circ_{(g)}(HA_{R'}) - G^\circ_{(g)}(HA_R)$$

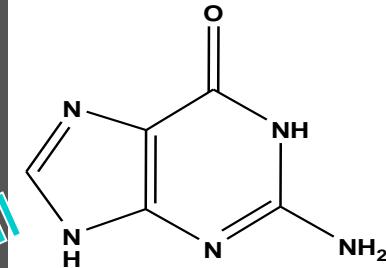
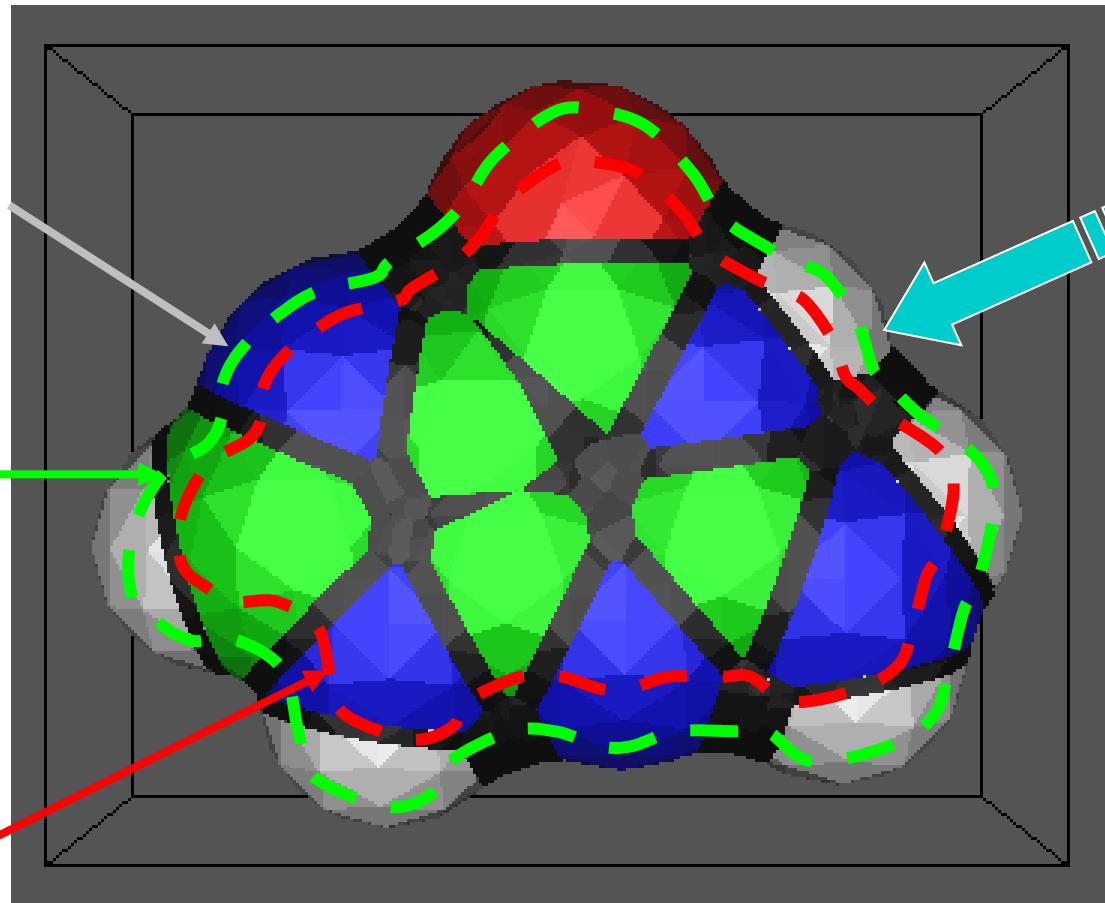


Polarizable Continuum Model for Solvation

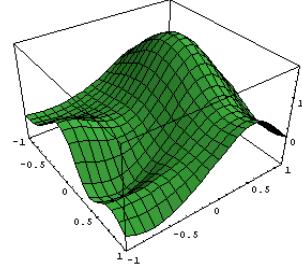
STANDARD
CAVITY FOR
NEUTRAL

CATION

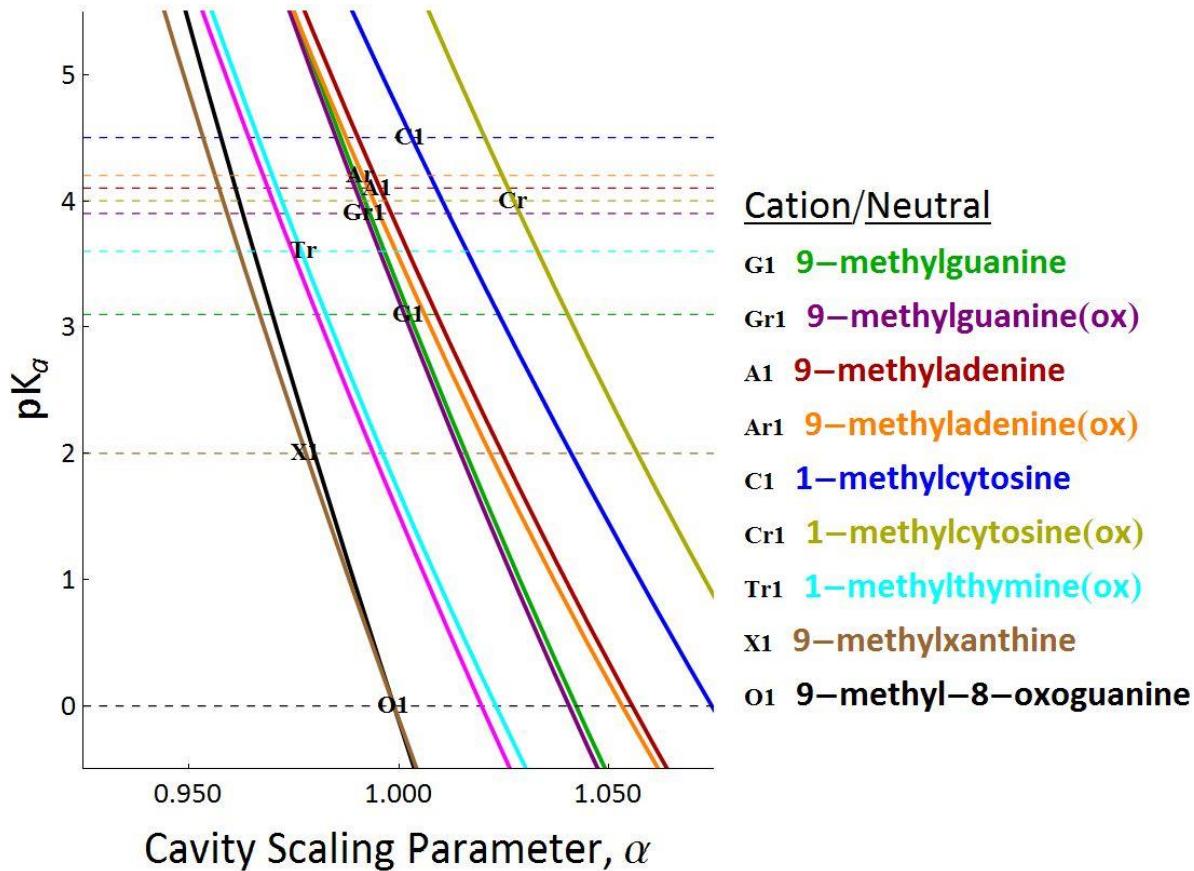
ANION



$$\Delta G_{(aq)} = \Delta G_{el} + \Delta G_{disp} + \Delta G_{rep} + \Delta G_{cav}$$



Cation Cavity Scaling



B3LYP

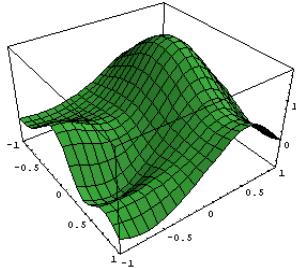
cation $\alpha = 1.000 \pm 0.015$

anion $\alpha = 0.900 \pm 0.050$

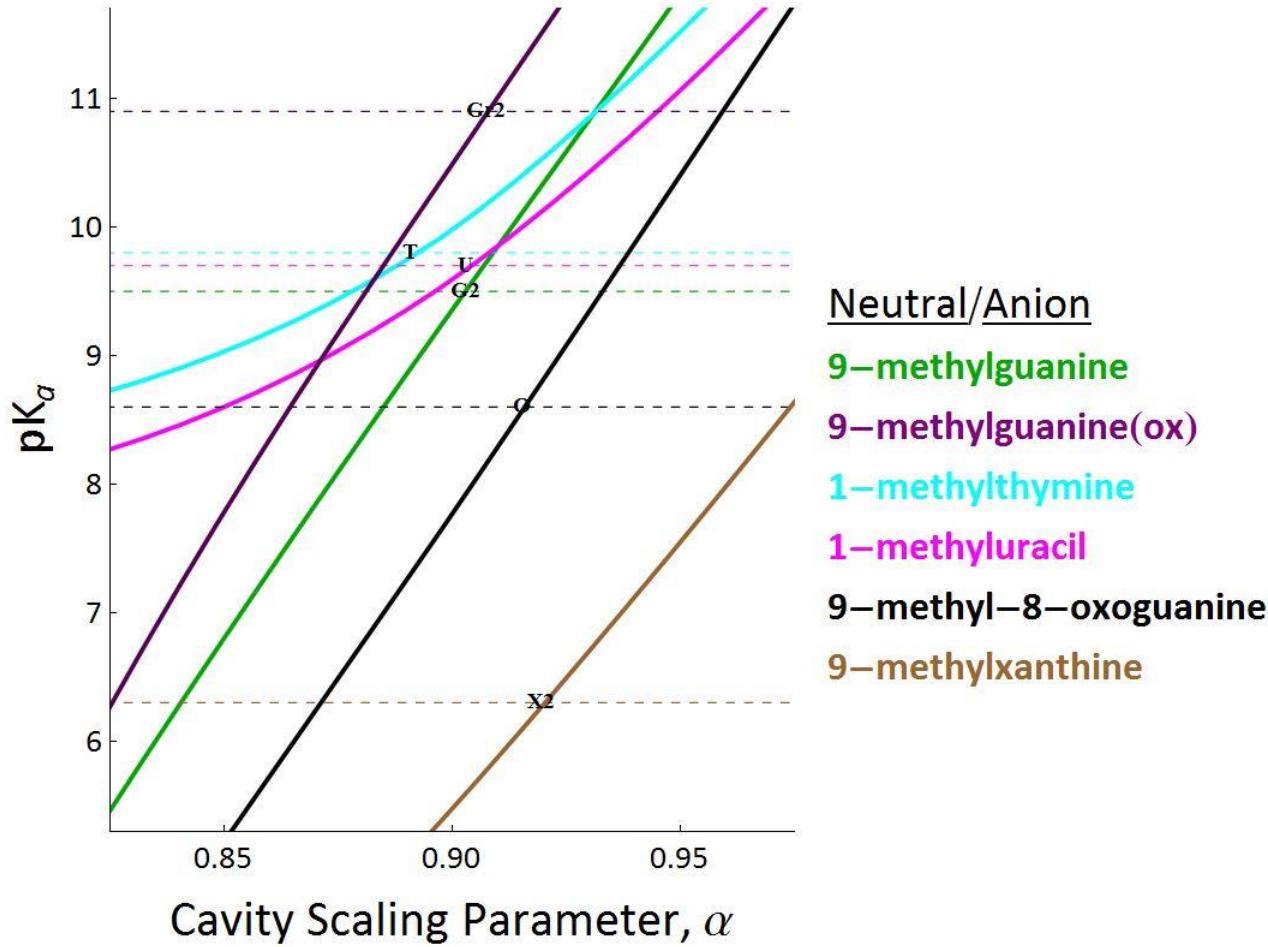
CBS-QB3

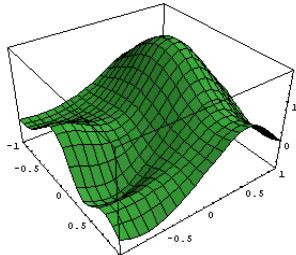
cation $\alpha = 0.975 \pm 0.014$

anion $\alpha = 0.925 \pm 0.058$

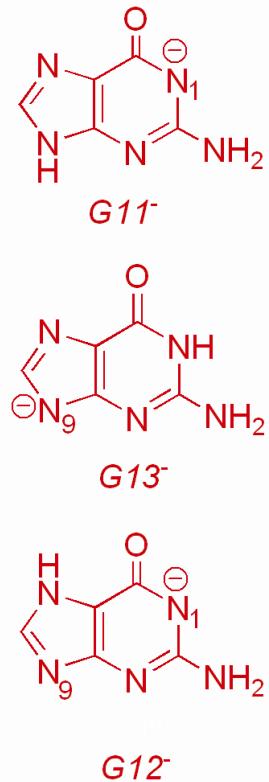
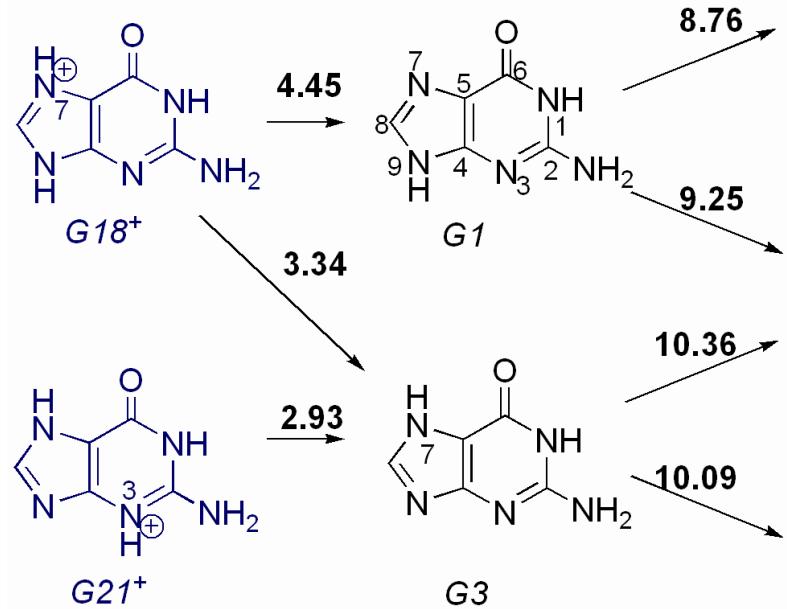


Anion Cavity Scaling





Guanine pK_a's



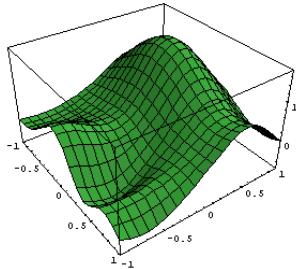
MOL	$\Delta G_{(g)}$ (kcal/mol)	$\Delta G_{(aq)}$ (kcal/mol)	ΔG_{solv} (kcal/mol)
$G18^+$	0.0	0.0	-64.8
$G21^+$	4.3	0.6	-68.9
$G1$	0.3	1.5	-18.5
$G3$	0.0	0.0	-19.0
$G11^-$	2.8	0.0	-75.4
$G12^-$	0.0	0.3	-72.0
$G13^-$	0.8	0.7	-73.1

$$pK_{a1}(\text{calc}) = 3.45$$

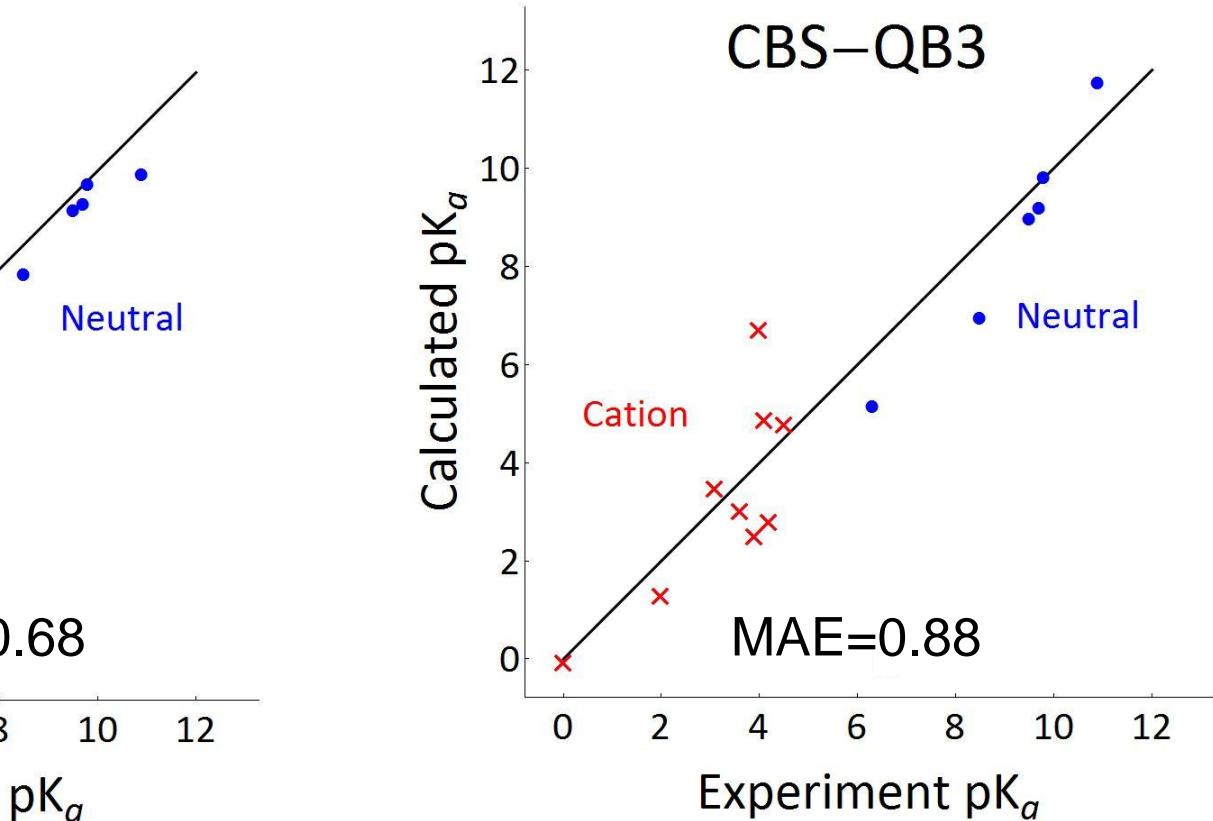
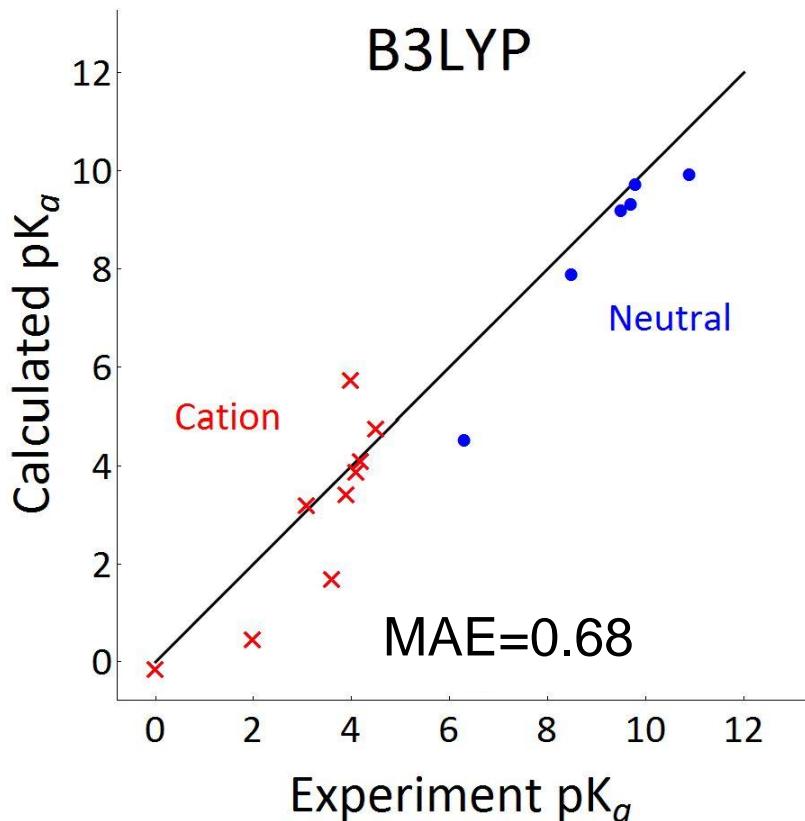
$$pK_{a1}(\text{exp}) = 3.2-3.3$$

$$pK_{a2}(\text{calc}) = 9.6$$

$$pK_{a2}(\text{exp}) = 9.2-9.6$$



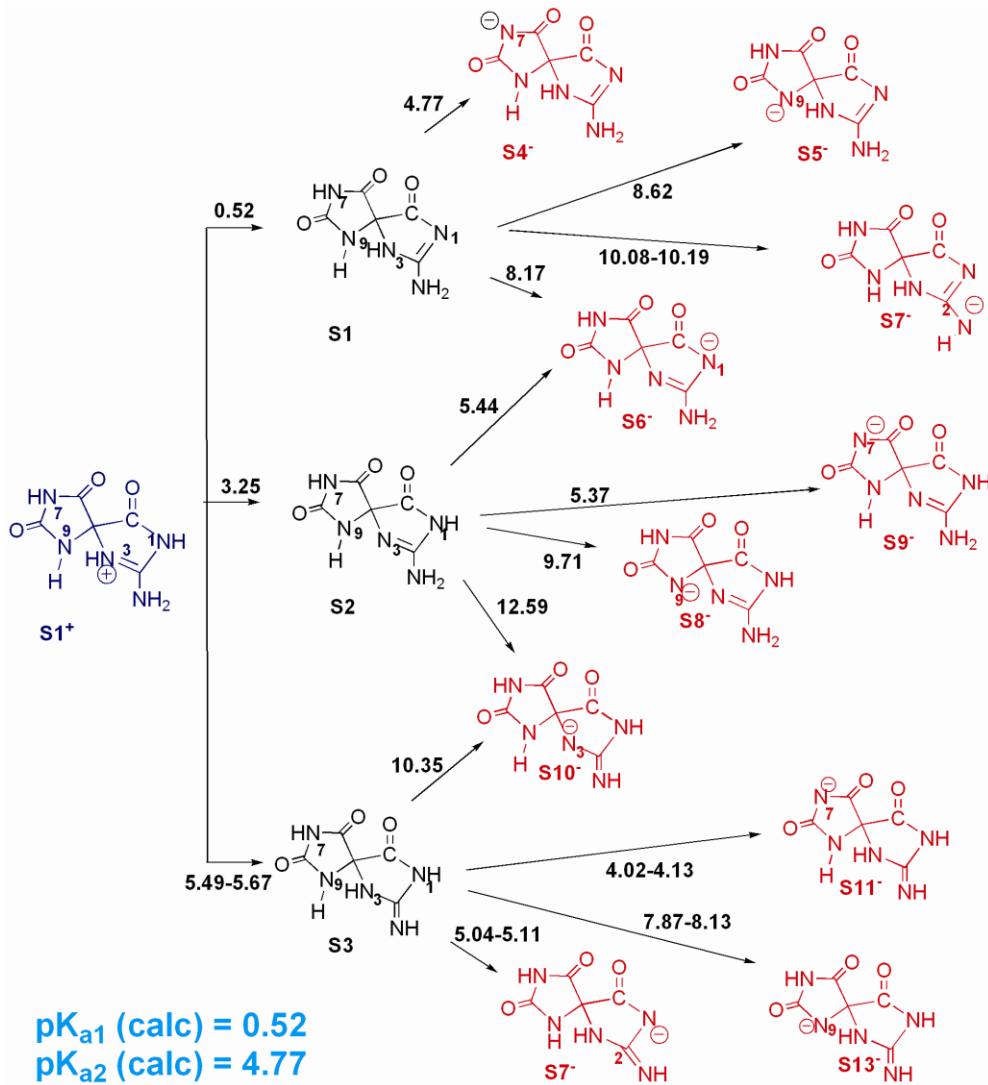
Nucleobase pK_a's



Solvent cavity scaling factors

B3LYP: 1.00 for cations, 0.90 for anions
 CBS–QB3: 0.975 for cations, 0.925 for anions
 1 pK_a = 1.36 kcal/mol = 5.69 kJ/mol

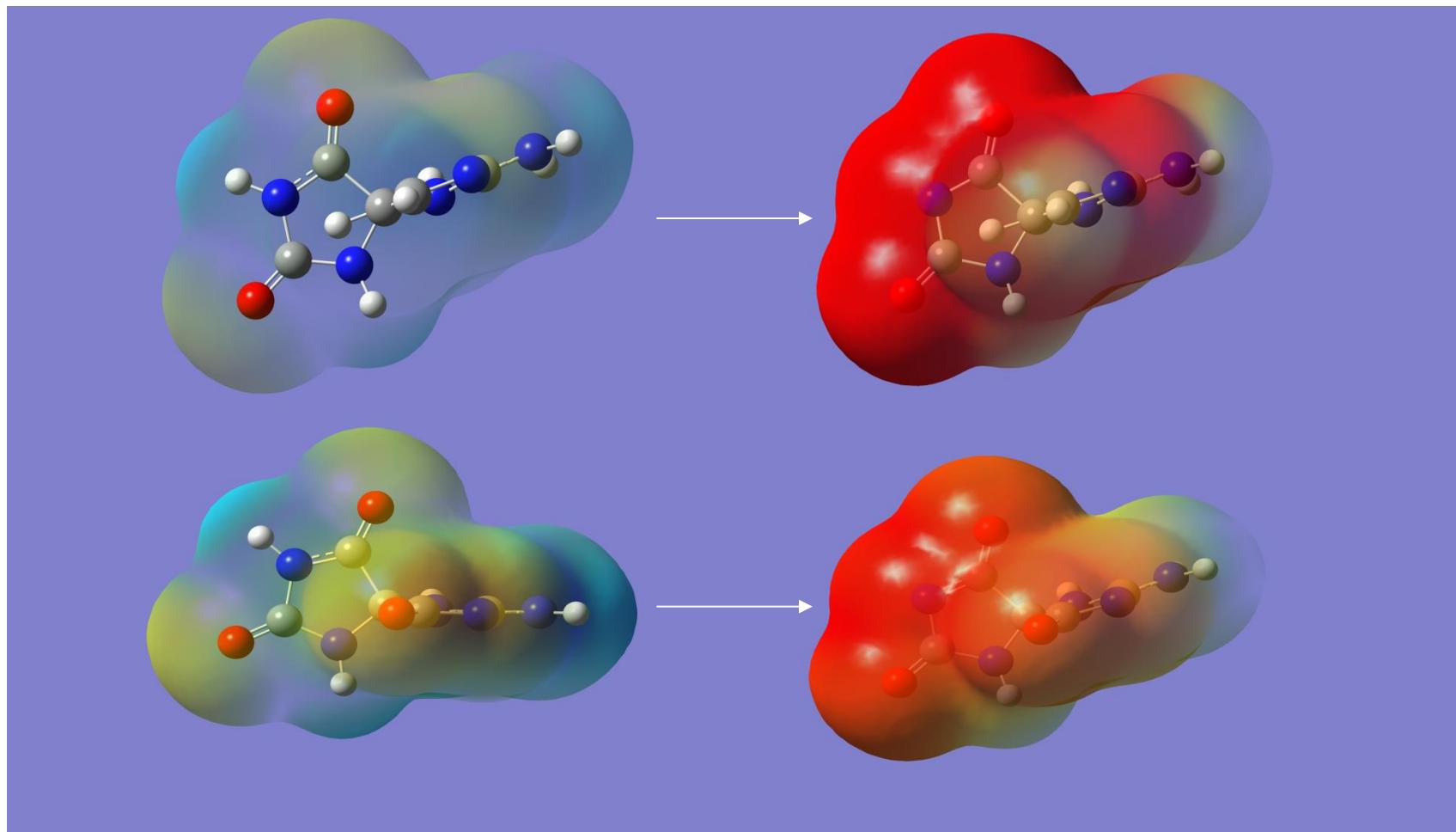
Spiroiminodihydantoin (Sp)



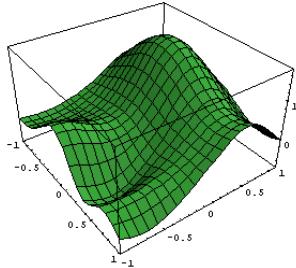
MOL	$\Delta G_{(g)}$ (kcal/mol)	$\Delta G_{(aq)}$ (kcal/mol)	ΔG_{solv} (kcal/mol)
S1⁺	0.0	0.0	-79.2
S1	0.0	0.0	-26.2
S2	1.2	3.7	-22.6
S3	1.3, 2.0	6.8, 7.0	-19.4,-19.6
S4⁻	5.2	0.0	-83.0
S5⁻	11.8	5.3	-83.0
S6⁻	0.9	4.6	-70.9
S7⁻	4.0, 0.0	7.3, 7.4	-72.1,-67.4
S8⁻	18.2	10.5	-83.8
S9⁻	12.0	4.5	-84.3
S10⁻	6.4	14.4	-66.3
S11⁻	1.7, 3.3	5.9, 6.0	-71.2,-72.5
S13⁻	6.9, 8.5	11.4, 11.3	-70.1,-71.6

A total of 1 cation, 21 neutral and 13 anionic tautomers were examined. These results are for the major tautomers in solution.

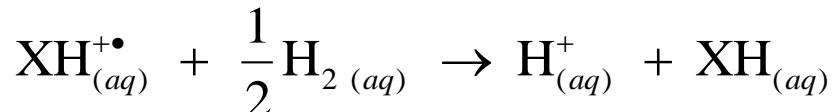
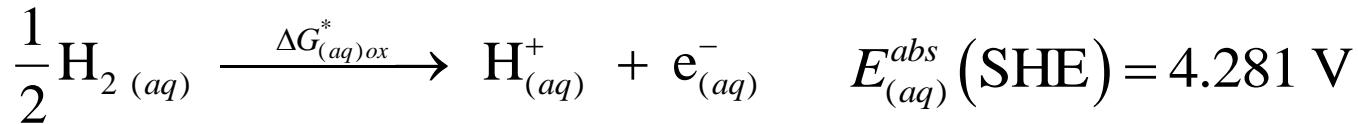
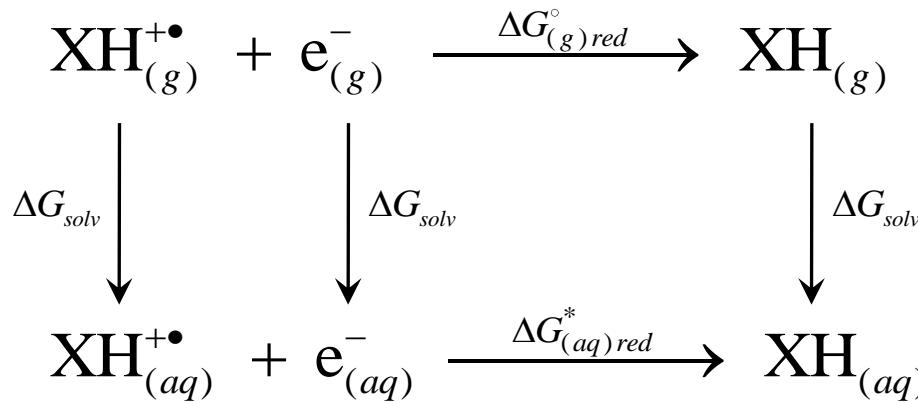
Anomalously low pK_a of Sp



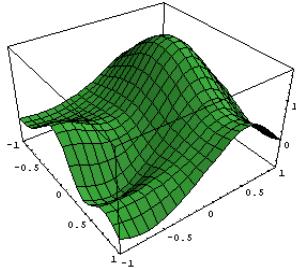
	ΔG_g	$\Delta\Delta G_{\text{solv}}$	ΔE_{dis}	Calc pK _a
Isomer S1 with C6=CH ₂	328.26	-56.84	1.58	6.6
Isomer S1 with C6=O (Sp)	325.26	-56.9	2.11	4.8
ΔpK_a contribution	+2.2	+0.0	-0.4	+1.8



Calculation of Redox Potentials



$$E_{red}^\circ \left(\text{XH}^{+\bullet}/\text{XH} \right) = \frac{-\Delta G_{(aq)red}^*}{nF} - E_{(aq)}^{abs}(\text{SHE})$$



Redox Potentials in Acetonitrile

$E^\circ \left(\text{XH}^{+\bullet} / \text{XH} \right)$

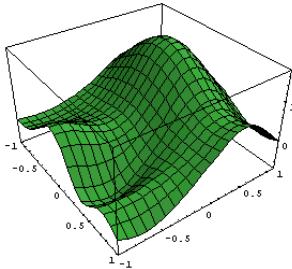
Seidel et al.

One-electron oxidation potentials for DNA nucleosides

Cyclic voltammetry measurements in acetonitrile solution against
saturated calomel electrode ($E_{\text{abs}} = 4.429$ V)

	<i>Seidel et al.</i> <i>Deoxyribose</i>	<i>CBS-QB3</i> <i>Methyl</i>	<i>B3LYP</i> <i>Methyl</i>	<i>B3LYP</i> <i>Ribose</i>
Guanine	1.25	1.14	0.96	1.04
Adenine	1.72	1.58	1.44	1.41
Cytosine	1.90	1.58	1.50	1.57
Thymine	1.87	1.63	1.51	1.60
Uracil	>2.15	1.91	1.81	1.88
<i>MAE to Exp.</i>		0.21	0.33	0.28

(in Volts)



Relative $E^\circ \left(\text{XH}^{+\bullet} / \text{XH} \right)$ in Acetonitrile

	<i>Seidel et al.</i> ¹ <i>Deoxyribose</i>	<i>CBS-QB3</i> ² <i>Methyl</i>	<i>B3LYP</i> ² <i>Methyl</i>	<i>B3LYP</i> ² <i>Ribose</i>
Guanine	-0.47	-0.44	-0.47	-0.37
Adenine	0.00	0.00	0.00	0.00
Cytosine	0.18	0.00	0.06	0.16
Thymine	0.15	0.05	0.07	0.19
Uracil	0.43	0.34	0.37	0.47
Xanthine	-	-0.11	0.03	0.07
8-Oxoguanine	-	-0.77	-0.79	-0.73

(in Volts)

Relative Mean Absolute Errors

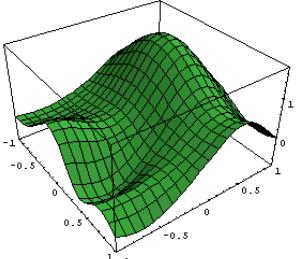
0.10 V CBS-QB3 to Exp.

0.07 V B3LYP to Exp.

0.05 V CBS-QB3 to B3LYP

¹ Seidel, C. A. M.; Schulz, A. and Sauer, M. H. M.; *J. Phys. Chem.* **1996**, *100*, 5541

² Pisciuk, B. T.; Lord, R. L.; Munk, B. H.; Schlegel, H. B.; *J. Chem. Theory Comput.* **2012**, *8*, 5107



Relative Potentials in Aqueous Solution

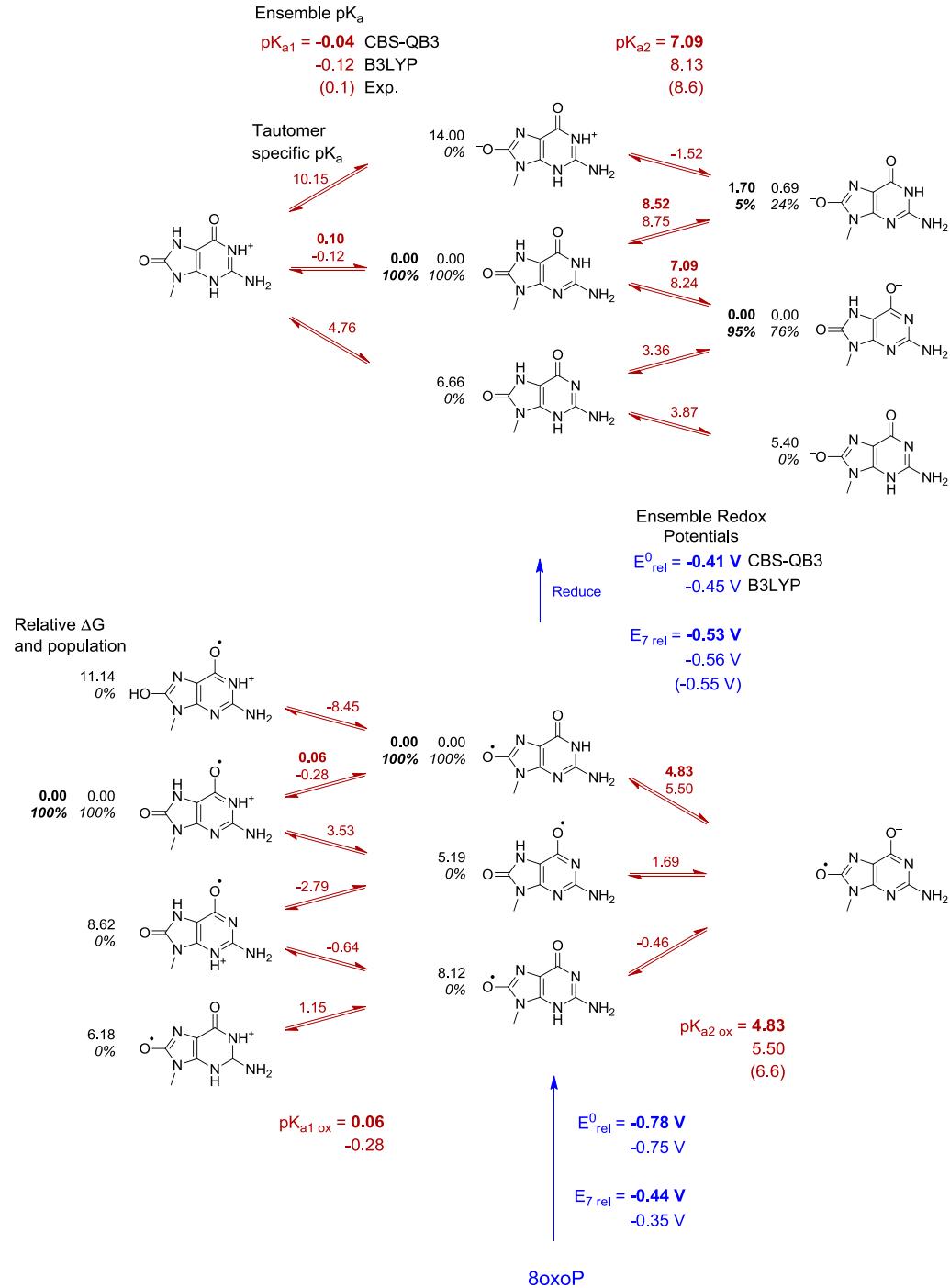
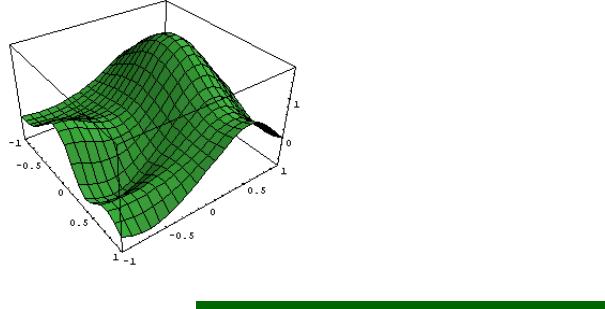
$E_7(X^\bullet, H^+ / XH)$

	<i>Steenken et al.</i>	<i>Fukuzumi et al.</i>	<i>CBS-QB3</i>	<i>B3LYP</i>	<i>B3LYP</i>	<i>Li et al.</i>
	<i>Ribose</i>	<i>Nucleotide</i>	<i>Methyl</i>	<i>Methyl</i>	<i>Ribose</i>	<i>Base</i>
Guanine	-0.13	-0.11	-0.40	-0.42	-0.32	-0.28
Adenine	0.00	0.00	0.00	0.00	0.00	0.00
Cytosine	~ 0.18	0.19	0.31	0.29	0.42	0.38
Thymine	~ 0.28	0.14	0.17	0.07	0.26	0.04
Uracil	-	-	0.51	0.34	0.43	0.24
Xanthine	-	-	-0.35	-0.40	-	-
8-Oxoguanine	-0.68	-	-0.93	-0.98	-	-0.51

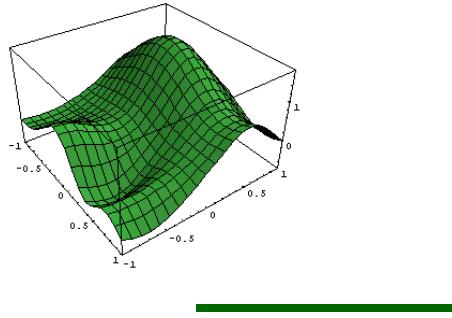
(in Volts)

8oxoGuanine

pK_a 's
and
redox
potentials

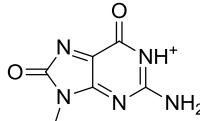


8oxoPurine pK_a 's and redox potentials



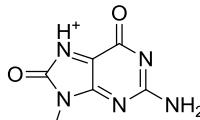
Relative ΔG
and population

0.00 0.00
100% 100%

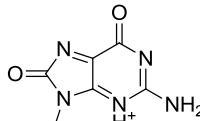


Ensemble pK_a
 $pK_{a1} = 1.21$ CBS-QB3
 -0.04 B3LYP
 Exp.

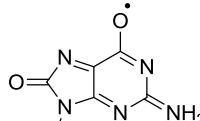
17.36 15.17
0% 0%



10.23 10.15
0% 0%



Tautomer specific pK_a



-8.59
-1.92

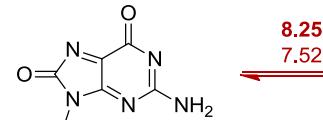
$pK_{a1\ ox} = -8.59$
-1.92

8oxoG_{OX}

$E^0_{\text{rel}} = -0.78 \text{ V}$
-0.75 V

$E_7^{\text{rel}} = -0.44 \text{ V}$
-0.35 V

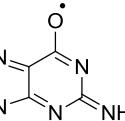
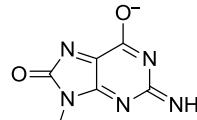
$pK_{a2} = 8.25$
7.52



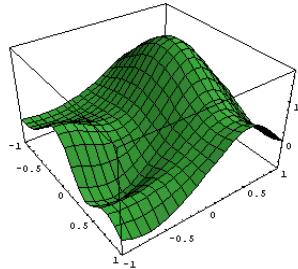
Ensemble Redox
Potentials

$E^0_{\text{rel}} = +1.07 \text{ V}$ CBS-QB3
+1.13 V B3LYP

$E_7^{\text{rel}} = +1.07 \text{ V}$
+1.13 V

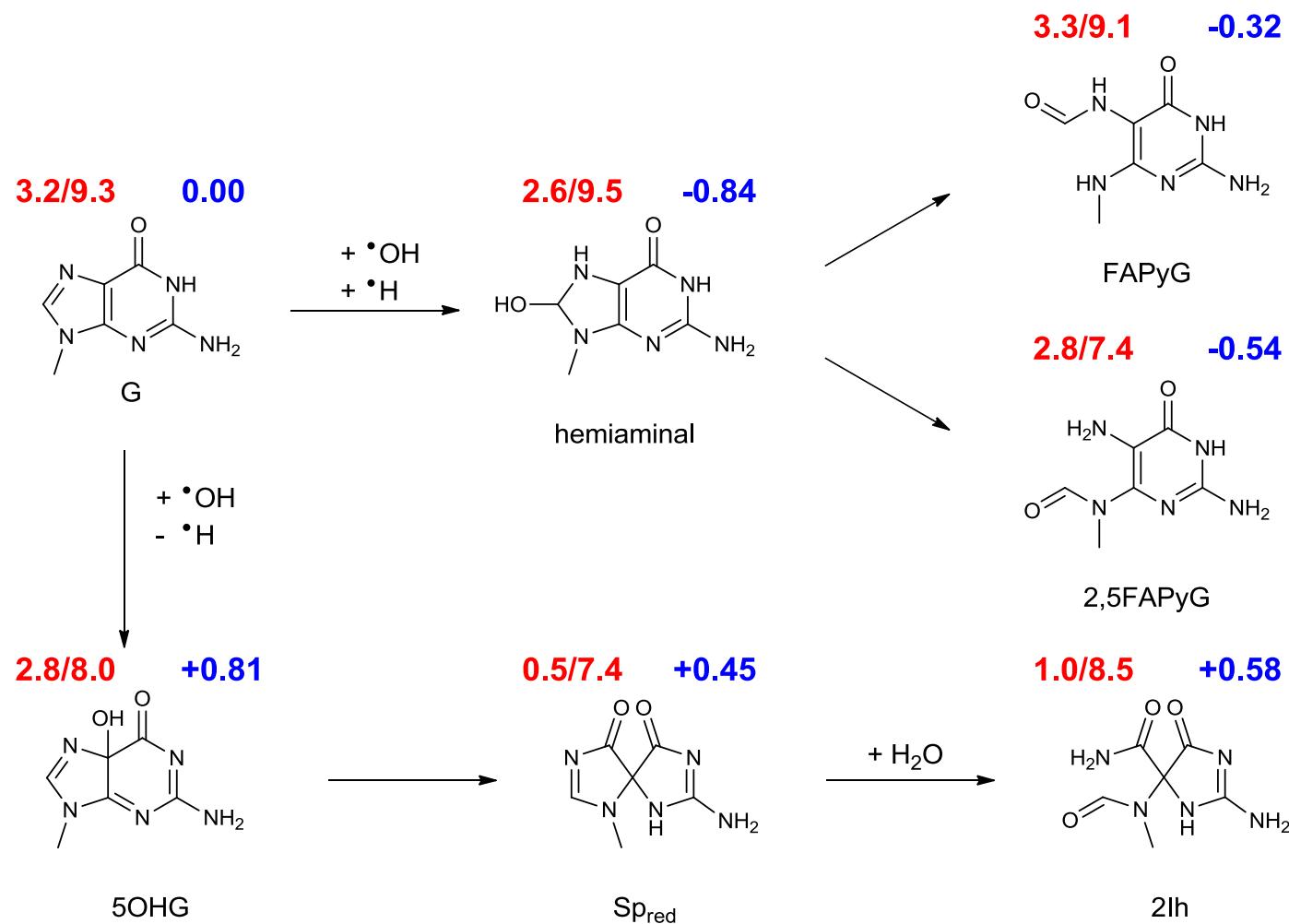


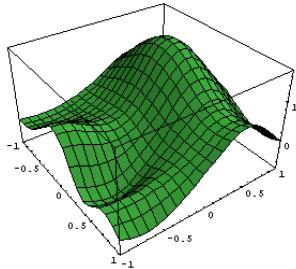
Reduce



Guanine Oxidation to FAPyG and Ih

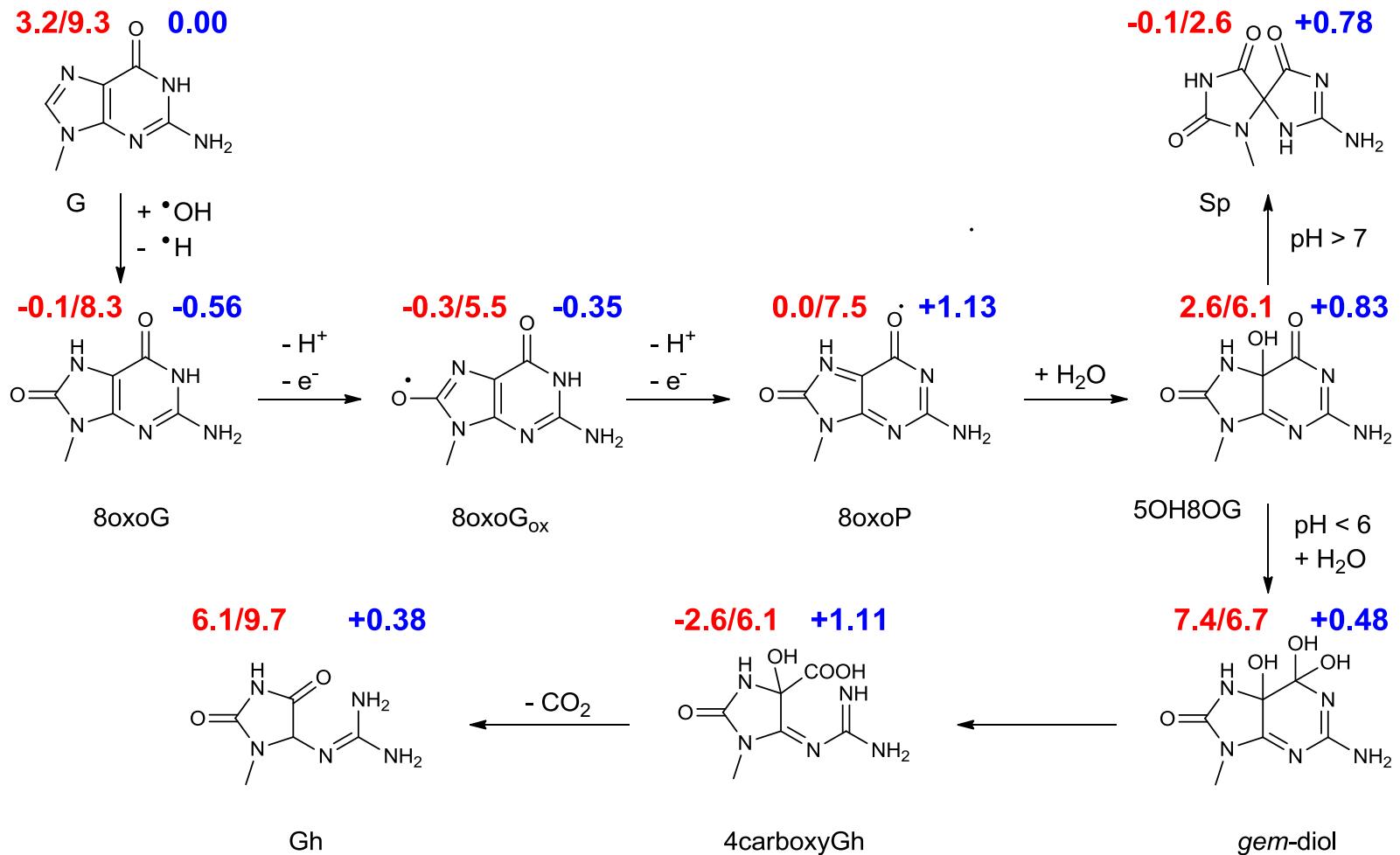
pK_{a1}/pK_{a2} and E_7 relative to G

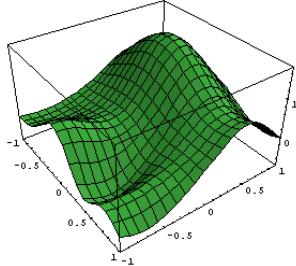




Guanine Oxidation to Sp and Gh

pK_{a1}/pK_{a2} and E_7 relative to G





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Current Research Group

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