## Structure of the 1,N<sup>2</sup>-Etheno-2'-deoxyguanosine Adduct in Duplex DNA at pH 8.6<sup>†</sup> <u>Revised Manuscript</u> <u>Supporting Information</u>

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Running Title: 1,N<sup>2</sup>-ɛdG Adduct in Duplex DNA

<sup>†</sup>We congratulate Professor Lawrence J. Marnett upon the occasion of his sixtieth birthday.

<b>Table S1.</b> Non-exchangeable Proton Chemical Shifts <sup>a</sup> of $1, N^2$ - $\varepsilon$ dG duplex in D <sub>2</sub> O <sup>b</sup> at p	Η
8.6, 7 °C.	

Nucleotide	H8	H6	H1′	H2′	H2"	H3′	H5/H2/CH <sub>3</sub>
$C^1$		7.68	5.78	2.03	2.46	4.75	5.94
G <sup>2</sup>	8.02		5.94	2.72	2.79	5.02	
C <sup>3</sup>		7.44	5.65	2.13	2.44	4.89	5.49
$\mathrm{A}^4$	8.37		6.24	2.68	2.91	5.02	7.22
$T^5$		7.06	5.78	1.74	2.09	4.79	1.50
X6	7.94		5.12	2.34	2.14	4.83	
<b>G</b> <sup>7</sup>	7.84		5.45	2.74	2.77	5.02	
A <sup>8</sup>	8.18		5.92	2.77	2.87	5.08	7.22
A <sup>9</sup>	8.19		6.22	2.60	2.93	5.00	7.75
$T^{10}$		7.18	5.92	2.09	2.50	4.87	1.33
C <sup>11</sup>		7.59	6.06	2.23	2.50	4.85	5.65
C <sup>12</sup>		7.67	6.26	2.28	2.28	4.59	5.77
G <sup>13</sup>	7.88		5.67	2.56	2.74	4.85	
$G^{14}$	7.89		5.69	2.74	2.83	5.06	
A <sup>15</sup>	8.29		6.33	2.74	2.99	5.08	7.89
T <sup>16</sup>		7.23	6.00	2.03	2.60	4.89	1.33
T <sup>17</sup>		7.39	6.10	2.13	2.52	4.98	1.56
C <sup>18</sup>		7.66	6.14	2.36	2.40	4.96	5.77
C <sup>19</sup>		7.63	6.04	1.91	2.25	4.79	5.79
A <sup>20</sup>	8.51		6.31	2.81	3.01	5.06	7.54
$T^{21}$		7.14	5.73	2.07	2.42	4.89	1.42
G <sup>22</sup>	7.91		5.86	2.64	2.70	4.98	

C <sup>23</sup>		7.40	5.77	1.93	2.36	4.85	5.43
G <sup>24</sup>	7.98		6.18	2.38	2.64	4.71	

<sup>a</sup> Values in parts per million. <sup>b</sup>10 mM phosphate buffer in D<sub>2</sub>O, 100 mM NaCl, pH 8.6. Etheno protons H7 and H6 observed at 6.54 and 6.21 ppm, respectively.

Base Pair	$C N^4 H_{nhb}$	C N <sup>4</sup> H) <sub>hb</sub>	G N1H	T N3H
$C^1 \bullet G^{24}$	7.14	8.17		
$G^2 \bullet C^{23}$	6.67	8.48	13.1	
$C^{3} \bullet G^{22}$	6.60	8.40	12.7	
$A^{4\bullet}T^{21}$				13.5
$T^5 \bullet A^{20}$				13.4
X <sup>6</sup> •C <sup>19</sup>	nd	nd	nd	
$G^{7} \bullet C^{18}$	6.91	7.97	12.0	
$A^{8} \bullet T^{17}$				13.8
A <sup>9</sup> •T <sup>16</sup>				13.7
$T^{10} \bullet A^{15}$				13.6
$C^{11} \bullet G^{14}$	6.71	8.40	12.8	
$C^{12} \bullet G^{13}$	7.17	8.33		

**Table S2.** Exchangeable Proton Chemical Shifts<sup>a</sup> of  $1,N^2$ - $\varepsilon$ dG duplex in H<sub>2</sub>O<sup>b</sup> at pH 8.6, 7 °C.

<sup>a</sup> Values in parts per million. <sup>b</sup>10 mM phosphate buffer in H<sub>2</sub>O, 100 mM NaCl, pH 8.6. nh and nhb refer to the hydrogen-bonded and non-hydrogen bonded cytidine amino protons, respectively. nd, not detected. **Table S3.** Distribution of Experimental NOE Restraints Among Nucleotide Units of the 1,*N*<sup>2</sup>-εdG duplex<sup>a</sup> in D<sub>2</sub>O Buffer at pH 8.6.

Nucleotide	Number of Restraints								
Nucleonide	intra-nucleotide	inter-nucleotide <sup>b</sup>	Cross-strand	Total					
C <sup>1</sup>	20	4	0	24					
G <sup>2</sup>	7	13	0	20					
C <sup>3</sup>	9	14	0	24					
$A^4$	9	15	1	25					
T <sup>5</sup>	12	13	0	25					
X <sup>6</sup>	11	6	6	23					
G <sup>7</sup>	7	7	0	14					
A <sup>8</sup>	10	9	0	19					
A <sup>9</sup>	9	15	2	26					
T <sup>10</sup>	10	17	0	27					
C <sup>11</sup>	9	13	1	23					
C <sup>12</sup>	8	6	0	14					
G <sup>13</sup>	16	3	0	19					
G <sup>14</sup>	8	7	0	15					
A <sup>15</sup>	11	15	2	28					
T <sup>16</sup>	10	21	0	31					
T <sup>17</sup>	8	16	1	25					
C <sup>18</sup>	6	8	0	14					
C <sup>19</sup>	11	7	2	20					
A <sup>20</sup>	11	16	1	28					
T <sup>21</sup>	9	14	0	23					
G <sup>22</sup>	7	11	0	18					

 C<sup>23</sup>
 9
 13
 0
 22

 G<sup>24</sup>
 9
 5
 0
 14

<sup>a</sup>Nucleotides  $C^1 \rightarrow C^{12}$  and  $G^{13} \rightarrow G^{24}$  are in the modified and complimentary strand, respectively. <sup>b</sup>The internucleotide NOEs includes both 3'- and 5'-neighbors.

**Figure S1.** NOE cross peak intensities between the base protons and the sugar H1' protons of the attached deoxyribose moieties. **A.** The modified strand. **B.** The complementary strand. Black bars represent intraresidue cross-peaks. Gray bars represent interresidue cross-peaks.



**Figure S2.** The partial charges assigned to the 1,*N*<sup>2</sup>-ɛdG adduct, calculated using the RESP protocol in the program GAUSSIAN98.

