

**Structure of the 1,*N*<sup>2</sup>-Ethenodeoxyguanosine Adduct Opposite Cytosine in Duplex DNA: Hoogsteen Base Pairing at pH 5.2<sup>†</sup>**

Supporting Information

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Running Title: The 1,*N*<sup>2</sup>-εdG Adduct in Duplex DNA at pH 5.2

Keywords: Etheno DNA Adducts, DNA Adduct Conformation

<sup>†</sup>This paper is dedicated to the memory of Professor Bea Singer 1922-2005.

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Figure S3	NOESY data showing cross peaks between the C <sup>19</sup> hydrogen bonded amino proton and flanking T <sup>5</sup> imino proton (peak a), and between C <sup>19</sup> non-hydrogen bonded amino proton and flanking T <sup>5</sup> imino proton (peak b).
Figure S4	Chemical shift differences observed for 1, <i>N</i> <sup>2</sup> -εdG-modified duplex as compared to the corresponding unmodified duplex at pH 5.2. $\Delta\delta = [\delta_{\text{unmodified oligodeoxynucleotide}} - \delta_{\text{modified oligodeoxynucleotide}}]$ (ppm).

**Table S1.** Non-exchangeable Proton Chemical Shifts<sup>a</sup> of the Unmodified Duplex<sup>b</sup> at 7 °C.

Nucleotide	H8	H6	H1'	H2'	H2''	H3'	H5/H2/CH <sub>3</sub>
C <sup>1</sup>		7.66	5.72	1.97	2.40	4.69	5.88
G <sup>2</sup>	7.97		5.90	2.66	2.74	4.97	
C <sup>3</sup>		7.40	5.59	2.11	2.42	4.85	5.43
A <sup>4</sup>	8.33		6.21	2.64	2.89	5.00	7.61
T <sup>5</sup>		7.03	5.58	1.76	2.15	4.77	1.39
G <sup>6</sup>	7.74		5.38	2.56	2.60	4.91	
G <sup>7</sup>	7.71		5.47	2.54	2.68	4.97	
A <sup>8</sup>	8.07		5.95	2.66	2.89	5.02	7.18
A <sup>9</sup>	8.09		6.12	2.50	2.89	4.95	7.65
T <sup>10</sup>		7.10	5.88	2.00	2.46	4.83	1.23
C <sup>11</sup>		7.55	6.02	2.17	2.46	4.81	5.59
C <sup>12</sup>		7.64	6.21	2.25	2.25	4.53	5.71
G <sup>13</sup>	7.81		5.60	2.48	2.66	4.79	
G <sup>14</sup>	7.85		5.62	2.68	2.77	5.00	
A <sup>15</sup>	8.25		6.29	2.70	2.97	5.02	7.83
T <sup>16</sup>		7.20	5.96	2.01	2.58	4.83	1.31
T <sup>17</sup>		7.39	6.10	2.11	2.58	4.87	1.52
C <sup>18</sup>		7.51	5.92	2.09	2.44	4.83	5.55
C <sup>19</sup>		7.49	5.26	2.11	2.36	4.83	5.57
A <sup>20</sup>	8.35		6.23	2.70	2.93	5.00	7.64
T <sup>21</sup>		7.11	5.66	1.95	2.33	4.81	1.44
G <sup>22</sup>	7.85		5.78	2.60	2.62	4.79	

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C <sup>23</sup>	7.33	5.73	1.98	2.33	4.79	5.38
G <sup>24</sup>	7.92	6.11	2.32	2.32	4.65	

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<sup>a</sup> Values in parts per million. <sup>b</sup> 10 mM NaH<sub>2</sub>PO<sub>4</sub>, 100 mM NaCl, and 5 μM Na<sub>2</sub>EDTA (pH 5.2).

**Table S2.** Non-exchangeable Proton Chemical Shifts<sup>a</sup> of the 1,*N*<sup>2</sup>- $\epsilon$ dG-modified Duplex<sup>b</sup> at 7 °C.

Nucleotide	H8/H2 <sup>c</sup>	H6	H1'	H2'	H2''	H3'	H5/H2/CH <sub>3</sub>
C <sup>1</sup>		7.65	5.73	1.97	2.40	4.69	5.88
G <sup>2</sup>	7.97		5.89	2.66	2.74	4.97	
C <sup>3</sup>		7.39	5.57	2.07	2.40	4.83	5.41
A <sup>4</sup>	8.27		6.12	2.54	2.70	4.95	7.57
T <sup>5</sup>		6.55	5.63	d	2.42	4.71	1.29
X <sup>6</sup>	7.13		5.71	2.04	d	4.97	
G <sup>7</sup>	7.80		d	2.64	2.64	5.12	
A <sup>8</sup>	8.05		5.94	2.64	2.85	5.02	7.17
A <sup>9</sup>	8.10		6.14	2.52	2.87	4.97	7.65
T <sup>10</sup>		7.11	5.86	2.00	2.46	4.81	1.25
C <sup>11</sup>		7.56	6.02	2.17	2.44	4.81	5.59
C <sup>12</sup>		7.66	6.20	2.25	2.25	4.53	5.73
G <sup>13</sup>	7.80		5.59	2.48	2.64	4.79	
G <sup>14</sup>	7.84		5.59	2.68	2.77	5.00	
A <sup>15</sup>	8.24		6.27	2.70	2.95	5.02	7.83
T <sup>16</sup>		7.20	5.94	2.01	2.56	4.83	1.29
T <sup>17</sup>		7.40	6.08	2.17	2.56	4.87	1.50
C <sup>18</sup>		7.65	6.00	2.25	2.46	4.85	5.67
C <sup>19</sup>		7.56	5.59	1.52	2.33	4.81	5.81
A <sup>20</sup>	8.54		6.22	2.79	2.93	5.00	7.78
T <sup>21</sup>		7.14	5.69	1.95	2.34	4.81	1.43
G <sup>22</sup>	7.85		5.79	2.60	2.62	4.79	

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C <sup>23</sup>	7.33	5.71	1.89	2.33	4.79	5.38
G <sup>24</sup>	7.92	6.12	2.31	2.31	4.65	

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<sup>a</sup> Values in parts per million. <sup>b</sup> 10 mM NaH<sub>2</sub>PO<sub>4</sub>, 100 mM NaCl, and 5 μM Na<sub>2</sub>EDTA (pH 5.2). <sup>c</sup> Imidazole proton of X<sup>6</sup>. Etheno protons H7 and H6 observed at 7.33 ppm. <sup>d</sup> not detected

**Table S3.** Exchangeable Proton Chemical Shifts<sup>a</sup> of the 1,*N*<sup>2</sup>-εdG-modified duplex<sup>b</sup> at 7 °C (pH 5.2).

Base Pair	C( <i>N</i> <sup>4</sup> H) <sub>nhb</sub>	C( <i>N</i> <sup>4</sup> H) <sub>hb</sub>	G( <i>N</i> 1H)	T( <i>N</i> 3H)
C <sup>1</sup> •G <sup>24</sup>	7.16	8.14		
G <sup>2</sup> •C <sup>23</sup>	6.61	8.41	13.03	
C <sup>3</sup> •G <sup>22</sup>	6.51	8.31	12.61	
A <sup>4</sup> •T <sup>21</sup>				13.54
T <sup>5</sup> •A <sup>20</sup>				13.34
X <sup>6</sup> •C <sup>19</sup>	8.95	10.22		
G <sup>7</sup> •C <sup>18</sup>	7.19	8.14	12.34	
A <sup>8</sup> •T <sup>17</sup>				13.61
A <sup>9</sup> •T <sup>16</sup>				13.73
T <sup>10</sup> •A <sup>15</sup>				13.56
C <sup>11</sup> •G <sup>14</sup>	6.85	8.36	12.76	
C <sup>12</sup> •G <sup>13</sup>	7.24	8.31		

<sup>a</sup> Values in parts per million. <sup>b</sup> 10 mM NaH<sub>2</sub>PO<sub>4</sub>, 100 mM NaCl, and 5 μM Na<sub>2</sub>EDTA (pH 5.2). nh and nhb refer to the hydrogen-bonded and non-hydrogen bonded cytidine amino protons, respectively.

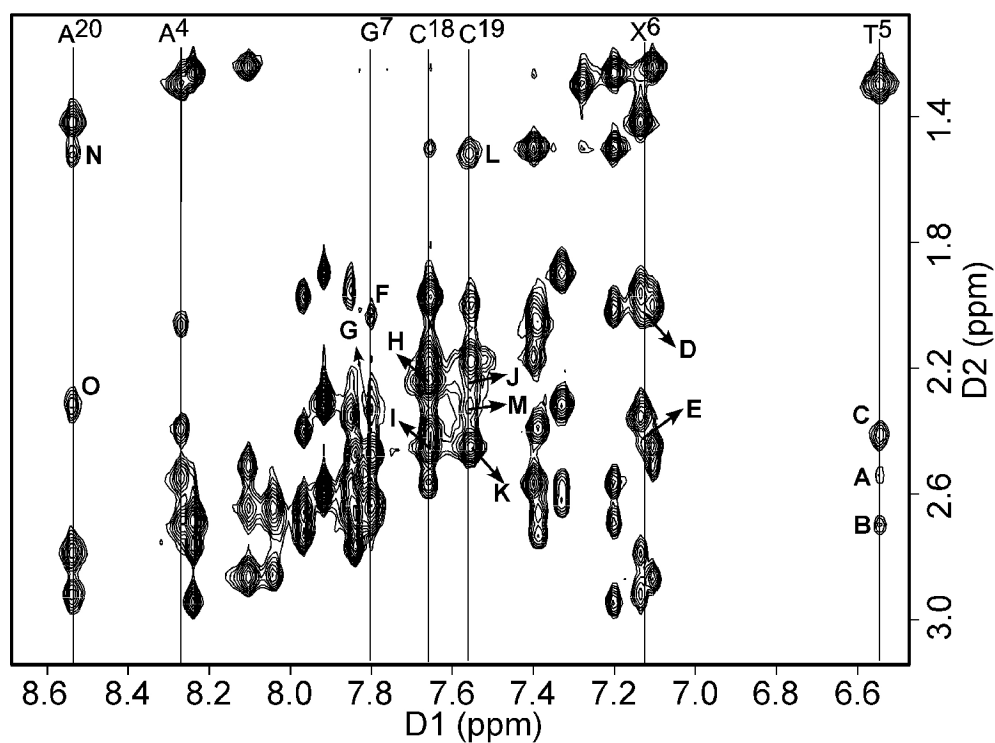
**Table S4.** Exchangeable Proton Chemical Shifts<sup>a</sup> of the Unmodified Duplex<sup>b</sup> at 7 °C (pH 5.2).

Base Pair	C(N <sup>4</sup> H) <sub>nbb</sub>	C(N <sup>4</sup> H) <sub>hb</sub>	G(N1H)	T(N3H)
C <sup>1</sup> •G <sup>24</sup>	7.14	8.15		
G <sup>2</sup> •C <sup>23</sup>	6.63	8.39	13.04	
C <sup>3</sup> •G <sup>22</sup>	6.51	8.35	12.65	
A <sup>4</sup> •T <sup>21</sup>				13.53
T <sup>5</sup> •A <sup>20</sup>				13.61
G <sup>6</sup> •C <sup>19</sup>	6.82	8.47	12.69	
G <sup>7</sup> •C <sup>18</sup>	6.67	8.23	12.51	
A <sup>8</sup> •T <sup>17</sup>				13.80
A <sup>9</sup> •T <sup>16</sup>				13.76
T <sup>10</sup> •A <sup>15</sup>				13.61
C <sup>11</sup> •G <sup>14</sup>	6.86	8.39	12.79	
C <sup>12</sup> •G <sup>13</sup>	7.21	8.33		

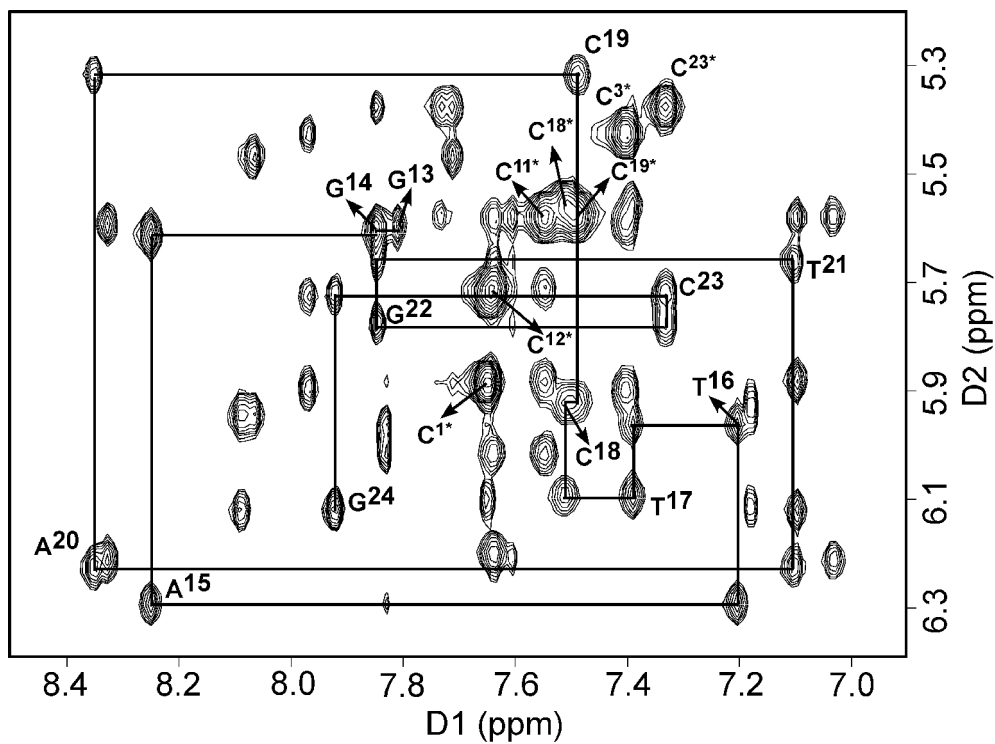
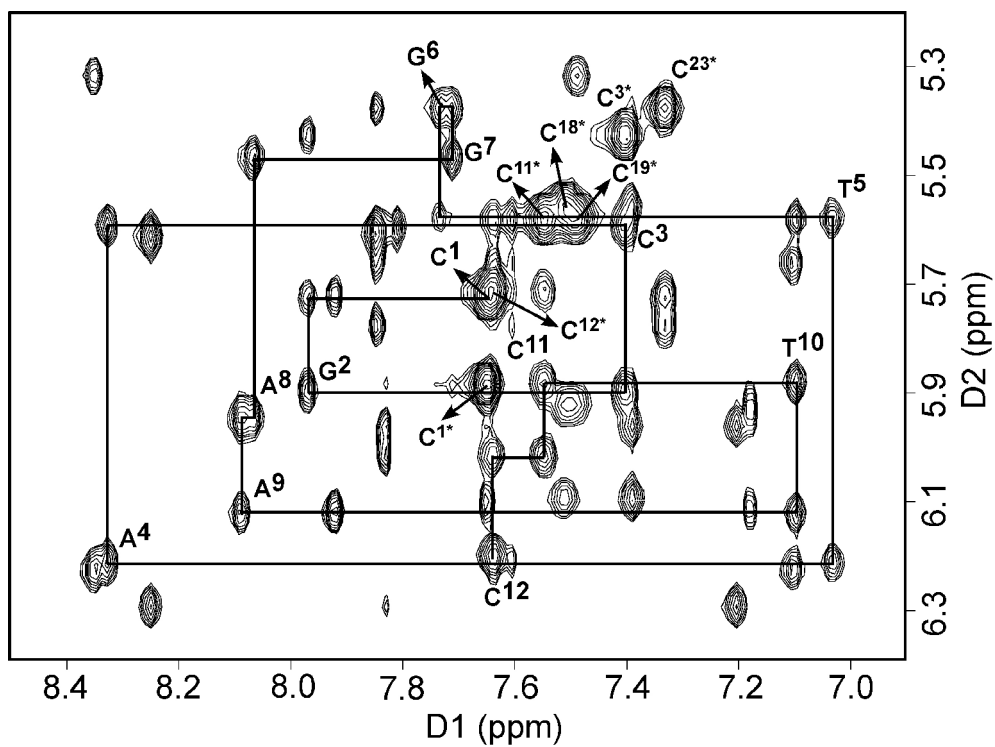
<sup>a</sup> Values in parts per million. <sup>b</sup> 10 mM NaH<sub>2</sub>PO<sub>4</sub>, 100 mM NaCl, and 5 μM Na<sub>2</sub>EDTA (pH 5.2). nh and nbb refer to the hydrogen-bonded and non-hydrogen bonded cytidine amino protons, respectively.



**Figure S1.**  $^1\text{H}$  NOESY spectrum showing cross peaks establishing connectivity between the base protons and the deoxyribose H2' and H2'' protons. Labeled peaks A, T<sup>5</sup> H6→A<sup>4</sup> H2' (weak); B, T<sup>5</sup> H6→A<sup>4</sup> H2'' (weak); C, T<sup>5</sup> H6→T<sup>5</sup> H2''; D, X<sup>6</sup> H2→X<sup>6</sup> H2' (weak, overlapped); E, X<sup>6</sup> H2→T<sup>5</sup> H2'' (weak, overlapped); F, G<sup>7</sup> H8→X<sup>6</sup> H2' (medium); G, G<sup>7</sup> H8→X<sup>6</sup> H2''(medium); H, C<sup>18</sup> H6→C<sup>18</sup> H2' (overlapped); I, C<sup>18</sup> H6→C<sup>18</sup> H2'' (overlapped); J, C<sup>19</sup> H6→C<sup>18</sup> H2' (very weak); K, C<sup>19</sup> H6→C<sup>18</sup> H2'' (overlapped); L, C<sup>19</sup> H6→C<sup>19</sup> H2'; M, C<sup>19</sup> H6→C<sup>19</sup> H2'' (medium); N, A<sup>20</sup> H8→C<sup>19</sup> H2'; O, A<sup>20</sup> H8→C<sup>19</sup> H2''. The spectrum was recorded at 7 °C at 250 ms mixing time (pH 5.2).



**Figure S2.**  $^1\text{H}$  NOESY spectrum showing sequential NOE connectivity from anomeric H1' to 3'-neighbor aromatic protons for the unmodified oligodeoxynucleotide duplex. The spectra were recorded at 7 °C, at 250 ms mixing time (pH 5.2). **A.** Nucleotides  $\text{C}^1 \rightarrow \text{C}^{12}$ . **B.** Nucleotides  $\text{G}^{13} \rightarrow \text{G}^{24}$ . Asterisks indicate cytosine H5-H6 cross peaks.



**Figure S3.** NOESY data (processed with lower threshold value than in Figure 3) showing cross peaks between the C<sup>19</sup> hydrogen bonded amino proton and flanking T<sup>5</sup> imino proton (peak a), and between C<sup>19</sup> non-hydrogen bonded amino proton and flanking T<sup>5</sup> imino proton (peak b). The NOESY spectrum was recorded in 10 mM phosphate buffer (9:1 H<sub>2</sub>O:D<sub>2</sub>O) at 7 °C (pH 5.2). Mixing time was 250 ms.

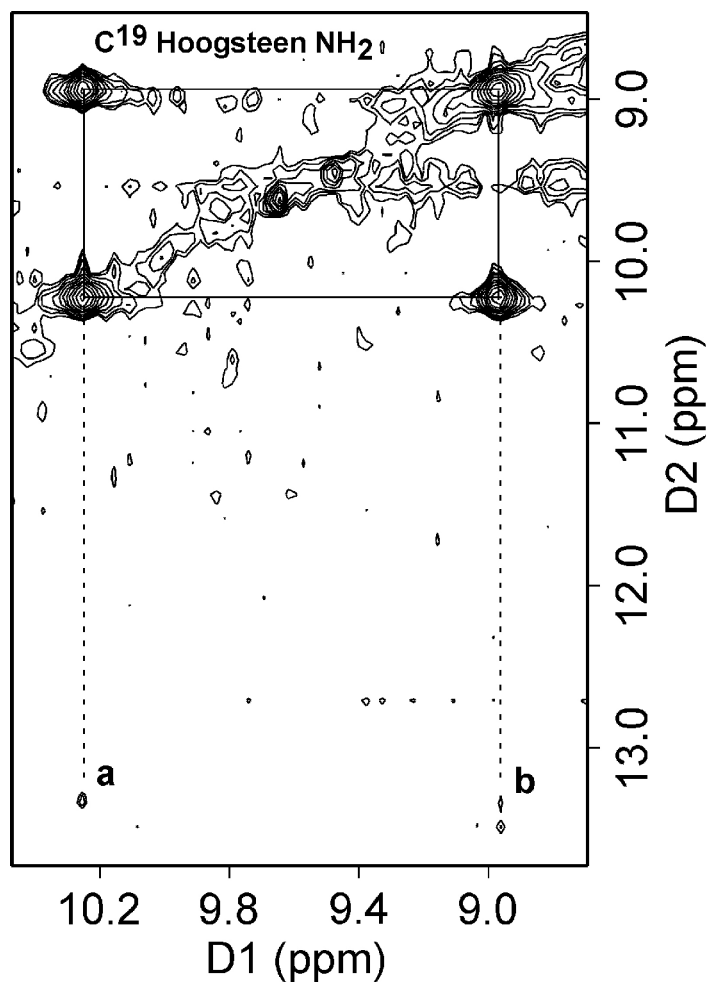


Figure S4 Chemical shift differences observed for 1,*N*<sup>2</sup>-εdG-modified duplex as compared to the corresponding unmodified duplex at pH 5.2.  $\Delta\delta = [\delta_{\text{unmodified oligodeoxynucleotide}} - \delta_{\text{modified oligodeoxynucleotide}}]$  (ppm).

