Supporting Information

Nuclear oxidation of a major peroxidation DNA adduct, M_1dG, in the Genome

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Supplemental Figures:

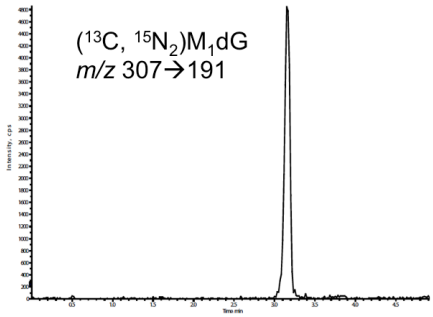
Supplemental Figure 1. (A) Representative LC-MS (SRM) chromatogram of the [^{13}C,$^{15}N_2$]-M_1dG internal standard. **(B)** Representative LC-MS (SRM) chromatogram of the M_1dG channel of the [^{13}C, $^{15}N_2$]-M_1dG internal standard. **(C)**. Representative LC-MS (SRM) chromatogram of the [$^{15}N_5$]-6-oxo-M_1dG internal standard. **(D)**. Representative LC-MS (SRM) chromatogram of the 6-oxo-M_1dG channel of the [$^{15}N_5$]-6-oxo-M_1dG internal standard.

Supplemental Figure 2. Representative CID of the M_1dG peak from a representative sample from RKO cells treated with adenine propenal showing its fragmentation pattern. Fragment at 188.1 corresponds to the loss of deoxyribose. The CID pattern confirms that the peak detected in the cellular samples is M_1dG.

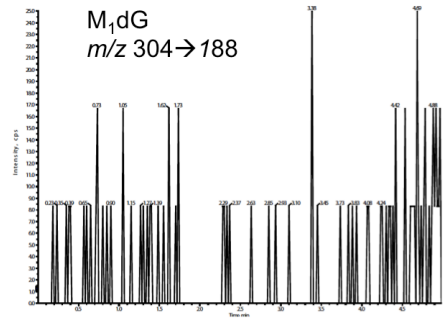
Supplemental Figure 3. Representative CID of the 6-oxo-M_1dG peak from a representative sample from RKO cells treated with adenine propenal showing its fragmentation pattern. Fragment at 204.1 corresponds to the loss of deoxyribose. The CID pattern confirms that the peak detected in the cellular samples is 6-oxo-M_1dG.

Figure S1.

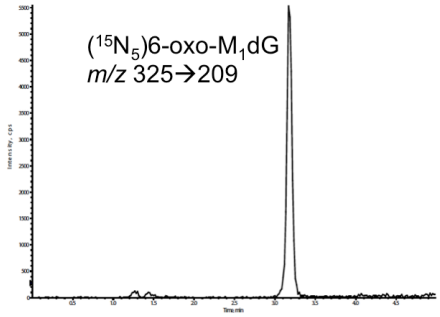
A.



B.



C.



D.

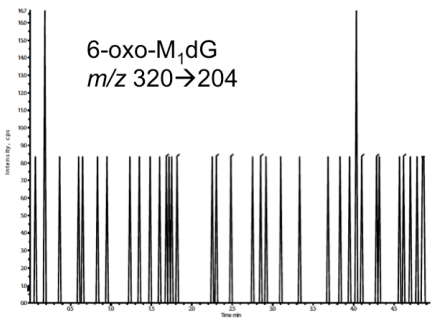


Figure S2.

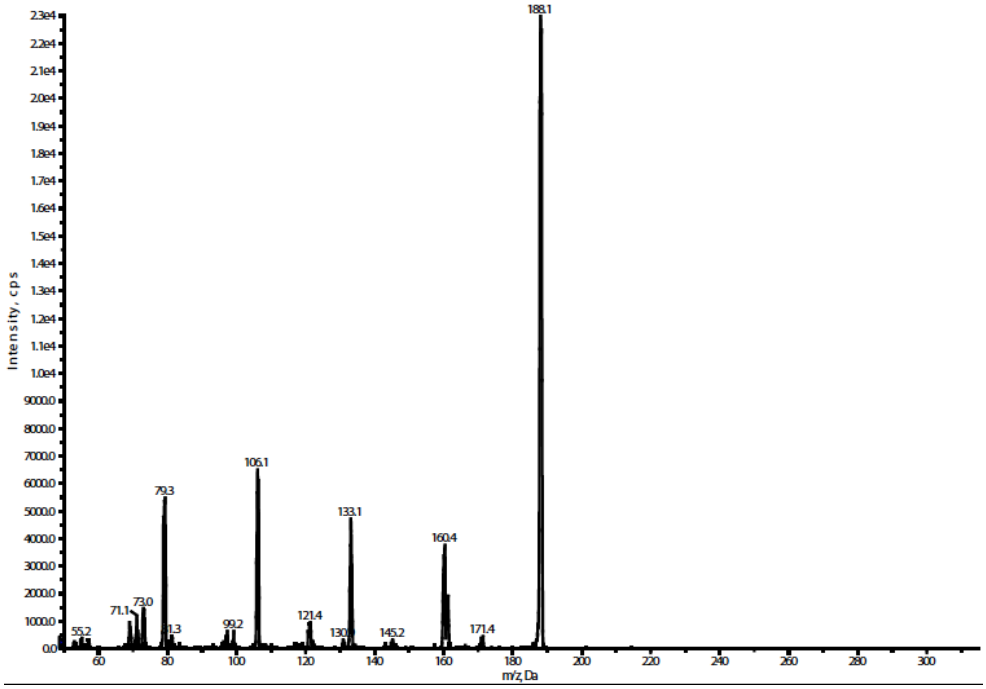
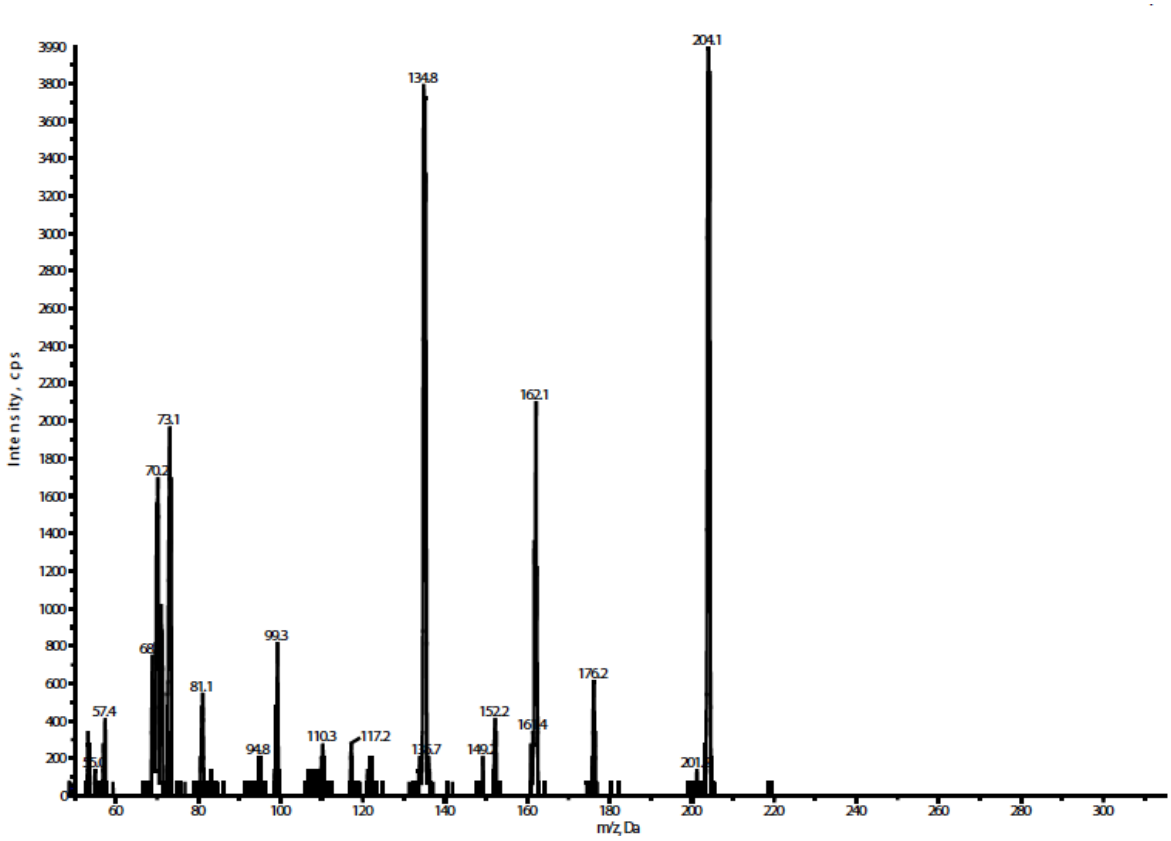

Intensity, cps
m/z, Da
188.1
106.1
79.3
133.1
160.4
73.0
71.1
55.2
99.2
121.4
145.2
171.4

Figure S3.


Intensity, cps
m/z, Da
204.1
134.8
162.1
73.1
70.2
99.3
81.1
176.2
57.4
152.2
110.3
117.2
94.8