

FREE RADICAL DNA ADDUCT 8-OH-DEOXYGUANOSINE AFFECTS ACTIVITY OF *Hpa* II AND *Msp* I RESTRICTION ENDONUCLEASES

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8-OH-deoxyguanosine can diminish the ability of the restriction endonucleases *Hpa* II and *Msp* I to cleave DNA. The exact position of the adduct within the recognition site appears to determine the extent of the effect.

KEY WORDS: Reactive oxygen species, 8-OH-deoxyguanosine, 8-hydroxyl-2'-deoxyguanosine, 8-hydroxyguanine, restriction endonuclease.

Reactive oxygen species produce a wide array of damage to cells at the molecular level¹⁻⁶, and have been suggested to be a significant factor both in carcinogenesis and the aging process⁷⁻¹³. 8-OH-deoxyguanosine (8-OH-dG) is one of the more common adducts resulting from both exogenous and endogenous oxidative damage to DNA¹⁴⁻²⁰, and has been shown to affect the specificity of both prokaryotic¹⁸ and human DNA methyltransferases (*Carcinogenesis*, in press). In this study, the presence of 8-OH-dG at the 5' and especially the 3' guanine of a 5'-CCGG-3' *Hpa* II/*Msp* I recognition site reduces the ability of these enzymes to cleave the DNA. Inhibition of cleavage by 8-OH-dG has been previously observed for the enzymes *Nhe* I²¹ and *Apa* LI²².

Synthetic oligonucleotides containing the *Hpa* II/*Msp* I recognition site were employed as substrates in restriction assays (see Figure 1). As previously described¹⁸, oligonucleotides containing 8-OH-dG at specific positions were kindly provided by Dr. Francis Johnson²³. Two of these oligonucleotides also contained 5-methylcytosine at specific sites. Other oligonucleotides were synthesized with either native sequences or sequences containing one or more 5-methylcytosines.

Oligonucleotides were annealed in pairs and approximately 40 pmoles (~500 ng) of each pair was digested for 24 hours at room temperature to avoid the possibility of strand denaturation at higher temperatures. Oligonucleotides were cut with either *Hpa* II (New England Biolabs, 90 units) or *Msp* I (New England Biolabs, 100 units) according to the manufacturer's buffer conditions, or not digested (control). Based upon the unit definition of each enzyme, the number of enzyme recognition sites per oligonucleotides pair (one) and the amount of oligonucleotide digested, these amounts of enzyme represent more than twenty times that required to theoretically cleave all of the sites. Aliquots from each digest were 5'-end labeled using γ -³²P-ATP (Amersham Corporation) and polynucleotide kinase (United States Biochemicals), and

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- A5'-GTACC^MC^{OH}GGTGACACACC-3'
- B5'-GTACC^MCG^{OH}GTGACACACC-3'
- C5'-GTACCC^{OH}GGTGACACACC-3'
- D5'-GTACCCG^{OH}GTGACACACC-3'
- U5'-GTACCCGGTGACACACC-3'
- U'5'-TGGTGTGTCACCGGGTA-3'
- M5'-CTCTGAAGTACC^MCGGTGACACACCA-3'
- M'5'-TGGTGTGTCAC^MCGGGTA-3'
- M25'-CTCTGAAGTAC^MC^MCGGTGACACACCA-3'
- M2'5'-TGGTGTGTCAC^MC^MCGGGTA-3'

FIGURE 1 Oligonucleotide Sequences

electrophoresed through 20% polyacrylamide gels. Gels were subsequently wrapped and exposed to film at room temperature for approximately 10–20 minutes. Densitometry of films for quantification of digestion was performed using an LKB Ultrosan XL Laser Densitometer. Each digestion assay was repeated 2–3 times to confirm the accuracy of the results. The results of the restriction assays and the sequence motifs of the recognition sites are presented in Table I.

TABLE I
Effect of 8-OH-dG on *Hpa* II and *Msp* I DNA Cleavage

Oligo Pair	Motif	<i>Hpa</i> II Cleavage (%)	<i>Msp</i> I Cleavage (%)
U/U'	CCGG	100	100
	GGCC		
U/M'	CCGG	0	100
	GGMC		
M/M'	CMGG	0	100
	GGMC		
M2/M2'	MMGG	0	0
	GGMM		
A/U'	CMXG	0	100
	GGCC		
B/U'	CMGX	0	100
	GGCC		
C/U'	CCXG	68	71
	GGCC		
D/U'	CCGX	0	100
	GGCC		
C/M'	CCXG	0	100
	GGMC		
D/M'	CCGX	0	46
	GGMC		

M = 5-methylcytosine X = 8-OH-deoxyguanosine

Hpa II is a methylation sensitive restriction endonuclease which recognizes 5'-CCGG-3' in double-stranded DNA, but will not cleave the DNA if either or both of the cytosines is 5-methylcytosine; *Msp* I is an isoschizomer of *Hpa* II which will not cleave the sequence if the 5' cytosine is methylated, but will cleave whether or not the 3' cytosine is methylated. In addition, *Msp* I can cleave hemimethylated DNA, whereas *Hpa* II cleaves it poorly or not at all²⁴. As can be seen in Table I, both enzymes exhibit expected activity with respect to native, hemimethylated, and fully methylated sequences (U/U', U/M', M/M' and M2/M2'). The enzymes also demonstrate predicted activity with respect to methylation status in A/U', B/U', and C/M'. However, while *Msp* I cleaves D/U', it has reduced ability to cleave C/U' and D/M'; *Hpa* II does not cleave D/M' (as expected based on methylation status) nor D/U', and has reduced ability to cleave C/U'.

These results indicate that the presence of 8-OH-dG as the 3' guanosine residue on one strand of the 5'-CCGG-3' recognition site completely inhibits normal activity of the *Hpa* II endonuclease (D/U'), but exerts a lesser affect on the activity when in the 5' guanosine position (C/U'). *Msp* I endonuclease activity is diminished when both the 3' guanosine is 8-OH-dG and the 3' cytosine of the complementary strand is 5-methylcytosine (D/M'), and to a lesser extent when the only modification is a 5' 8-OH-dG (C/U'). *Msp* I activity is unaffected by the presence of a 5' 8-OH-dG on one strand and a 3' 5-methylcytosine on the complementary strand (C/M').

The effect of 8-OH-dG on cleavage by these enzymes may be due to a decrease in the ability of the enzymes to recognize and bind to the recognition sequence, similar to the phenomenon we observed previously in studies examining the effect of 8-OH-dG on methylation by *Hpa* II methyltransferase¹⁸. Alternatively, the enzymes may bind to the recognition sequence but be unable to cleave the DNA. Further studies to determine the exact mechanism of cleavage inhibition are necessary.

That oxidative damage to DNA can produce alterations in the ability of restriction endonucleases to cleave DNA at their recognition sites is consistent with our earlier observations that such damage can affect enzymatic methylation of DNA¹⁸. DNA adduct formation via damage by reactive oxygen species may influence other DNA-protein interactions as well.

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