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Increased oxidative DNA damage, 8-hydroxydeoxyguanosine, in human pterygium

Abstract

Purpose Chronic exposure to ultraviolet (UV) light is a widely accepted aetiological factor in the development of pterygium. UV radiation may induce production of reactive oxygen species via photosensitized oxidation, thus causing oxidative damage. This study was conducted to test the hypothesis that oxidative damage to DNA is increased in pterygium. Methods Immunohistochemical analysis employing a monoclonal antibody specific for 8-hydroxy-2'-deoxyguanosine (8-OHdG), a ubiquitous maker of oxidative stress, was performed in three patients with primary pterygium. The levels of 8-OHdG in DNA isolated from the other 29 pterygium specimens and their adjacent normal conjunctival tissues were determined using enzyme-linked immunosorbent assay (ELISA). Results Immunohistochemistry of 8-OHdG showed a distinct pattern of more extensive and intense staining in the nuclei of pterygium tissue compared with that in their adjacent normal conjunctiva. ELISA also revealed that the average level of 8-OHdG in the pterygium tissues was 4.7-fold higher than that of the corresponding normal conjunctiva (P<0.001).

Conclusions The increased levels of 8-OHdG in the pterygium tissues indicate that oxidative stress could play a role in the development of pterygium. These findings provide new information to better understand the pathogenesis of pterygium and are useful in the prevention and treatment of this disease.

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Introduction

Pterygium, a wing-shaped (pterygos in Greek is wing) fibrovascular growth of conjunctiva extending into the superficial cornea, may cause irritation, redness, and tearing. As the pterygium develops, it may alter the shape of the cornea to cause astigmatism, or grow over the central cornea to threaten the vision. Several factors have been proposed to play a role in the pathogenesis of pterygium, such as antiapoptosis, proinflammatory cytokines, growth factors, immunological responses, viral infections, extracellular matrix remodelling, and genetic factors.¹ Although the detailed molecular mechanism remains unclear, it has been widely accepted that chronic ultraviolet (UV) exposure is a major aetiological factor, which is supported by epidemiological evidence,^{2,3} the ray tracing model,⁴ and the similar histopathological features in pterygium and UV-damaged skin.5,6 The noxious effects of UV radiation are either caused by direct UV radiation or by indirect stress via reactive oxygen species (ROS).7 Excess ROS can overwhelm the antioxidant system and cause oxidative damage to a variety of biological molecules such as lipids, proteins, and DNA. Lu et al⁸ had demonstrated ROS-induced lipid peroxidation in the pterygium tissue. However, very limited information is available regarding the oxidative DNA damage in pterygium.

In a previous study, we showed that patients with Ser326Cys polymorphism of 8-oxoguanine glycosylase I (hOGG1) were more susceptible to pterygium formation.⁹ hOGG1 is a key enzyme responsible for the base excision repair of 8-hydroxy-2'-deoxyguanosine (8-OHdG). ROS produced by UVB and UVA radiation through photosensitized oxidation can target DNA base guanine, giving rise to 8-OHdG in DNA molecules.¹⁰ 8-OHdG has been shown to be a sensitive and stable biomarker for evaluating the degree of oxidative DNA damage.¹¹ Based on these findings, we further hypothesized that oxidative damage to DNA is increased in pterygium tissues. In this study, immunohistochemical staining using a monoclonal antibody to 8-OHdG was performed to compare the distribution of 8-OHdG in pterygium tissues and their corresponding normal conjunctiva. In addition, we provide novel data to further demonstrate that the average level of 8-OHdG in pterygium tissues is higher than that of normal conjunctiva by enzyme-linked immunosorbent assay (ELISA).

Materials and methods

Tissue sampling

Excised primary pterygia and their corresponding adjacent normal conjunctival tissues located under lower eyelid margin from 32 eyes of 32 patients were obtained at surgery from the Taipei Veterans General Hospital, Taiwan. Diagnosis of pterygium was established according to the clinical finding of a wing-shaped fold of conjunctiva and fibrovascular tissue that had invaded the adjacent cornea for which there was no alternative explanation (eg, trauma). All tissue specimens were collected and immediately stored in cryovials with nitrogen substitution and kept in a refrigerator at -70°C until use. Immunohistochemical analysis of 8-OHdG was conducted in the first three patients, and the 8-OHdG levels were measured by ELISA in the other 29 patients. Informed consent was obtained from each subject and the experimental protocol was approved by the Institutional Review Board of Taipei Veterans General Hospital and performed in accordance with the tenets of the Declaration of Helsinki.

Immunohistochemical analysis

The avidin–biotin–peroxidase complex method was first used to determine the content of 8-OHdG.¹² The pterygium tissues were fixed in Bouin's solution for 2 h, sequentially immersed in 50 and 70% ethanol, and followed by dehydration. Specimens were then embedded in paraffin, sectioned at 4 μ m, and mounted on glass slides coated with poly-L-lysine. After being heated in a microwave for 10 min to enhance epitope expression, the specimens were treated with normal rabbit serum (Dako, Kyoto, Japan; diluted 1:75) to block nonspecific binding and then incubated with monoclonal antibody N45.1 (Japan Institute for the Control of Aging, Fukuroi, Japan; 2.5 μ g/ml), followed by addition of biotin-labelled rabbit anti-mouse IgG serum (Dako, Kyoto, Japan; diluted 1:300). Endogenous peroxidase

was blocked by treating the specimens with 3% hydrogen peroxide in methanol for 30 min. After washing with 50 mM Tris-HCl buffer (pH 8.0), the antigen–antibody complexes in specimens were visualized using a streptavidin–horseradish peroxidase detection system (Dako, Kyoto, Japan) with diaminobenzidine as chromogen.

Quantitative analysis of 8-OHdG

The DNA in each specimen was obtained by using the DNA Extractor Kit (Wako Pure Chemical Industries Inc., Osaka, Japan) according to the manufacturer's instructions. To eliminate the production of iatrogenic 8-OHdG, the container had nitrogen substituted through the entire procedure. After extraction, the DNA was dissolved in 60 μ l DNase-free water and stored at -70° C until use. Of the DNA solution, $10 \,\mu$ l was used for measurement of DNA concentration using the PicoGreen dsDNA Quantitation Kit (Molecular Probes, Eugene, OR, USA).

The rest of the extracted DNA solution was digested with nuclease P1 (Roche, Mannheim, Germany) in a solution of 200 mM sodium acetate (pH 5.3), followed by treatment with alkaline phosphatase (Roche, Mannheim, Germany) in 1 M Tris-HCl buffer (pH 7.4). To remove enzymes and other macromolecules, the hydrolysate was filtrated through the Ultrafree Centrifugal Filter C3LGC (Millipore, Billerica, MA, USA) at 10 000 g for 10 min at 4°C.

The content of 8-OHdG in each sample was determined by using the highly sensitive 8-OHdG ELISA kit (Japan Institute for the Control of Aging, Fukuroi, Japan) according to the manufacturer's instructions. The detection range was 0.125–10 ng/ml. ELISA method is more simple and easier than high-performance liquid chromatography with electrochemical detection (HPLC-ECD) with respect to the equipment, analysis time, running cost, and sample volume. The results of the ELISA test correlated well with the data obtained from measurement by HPLC-ECD.¹³

Statistical analysis

Data were analysed using an SPSS statistical package, version 10.0 (SPSS Inc., Chicago, IL, USA). Comparison of 8-OHdG contents between pterygium and normal conjunctival tissue was performed by means of the Wilcoxon signed-rank test. The comparison of the 8-OHdG content in pterygium between different genders was performed by the Wilcoxon rank-sum test. The correlation coefficient between age and the 8-OHdG content in pterygium was calculated. P < 0.05 was considered to be of statistical significance.

Results

Immunohistochemical detection of 8-OHdG

Immunohistochemical analysis of 8-OHdG was conducted in the first three pterygium cases and reproducible results were obtained. Representative results of one case are shown in Figure 1. Strong immunoreactivity was observed in the whole layer of pterygium epithelium and scantly distributed in the subepithelial stroma (Figure 1a). There was an intense and coarse granular staining pattern in the nuclei of pterygium tissue. In contrast, the immunoreactivity in



Figure 1 Immunohistochemical staining for 8-OHdG in pterygium tissue and normal conjunctiva. (a) Strong immunoreactivity was observed in the whole layer of pterygium epithelium and scantly distributed in the subepithelial stroma (arrowhead), revealing an intense and coarse granular staining pattern in the nuclei (arrow). (b) In contrast, the immunoreactivity in normal conjunctival epithelium and stromal layer was weak, showing a faint and small spherical staining pattern near the nuclear membrane (arrow). Some strong staining nuclei were occasionally seen in the superficial layer of normal conjunctival epithelium (arrowhead). Scale bars in (a) and (b) represent $20 \,\mu$ m.

normal conjunctival epithelium and stromal layer was weak, and showed a faint and small spherical staining pattern near the nuclear membrane (Figure 1b).

Determination of 8-OHdG content

The demographic information and the 8-OHdG content in the pterygium tissues of the 29 patients are summarized in Table 1. The mean value of 8-OHdG content was 53.93 ± 8.17 ng of 8-OHdG per mg DNA in pterygium tissues and 11.40 ± 0.80 ng of 8-OHdG per mg DNA in the adjacent normal conjunctiva (Figure 2). The average level of 8-OHdG in pterygium tissues was 4.7-fold higher than that in the normal conjunctiva (P < 0.001). However, the 8-OHdG content in the pterygium tissue was not related to the sex (P = 0.753) or age (P = 0.817) of the patient.

Discussion

In the current study, we demonstrated abundant immunoreactivity of 8-OHdG in pterygium tissues and

Table 1 Demography and 8-OHdG content in pterygiumtissues of 29 patients

Case	Sex	Age	8-OHdG content (ng/mg DNA)	
			Pterygium	Normal conjunctiva
1	F	54	189.47	12.71
2	М	70	130.45	10.01
3	М	69	30.98	11.82
4	F	65	38.18	7.87
5	Μ	71	26.81	5.78
6	М	70	99.69	11.11
7	Μ	70	41.62	12.59
8	М	68	15.93	10.07
9	Μ	82	64.04	6.19
10	Μ	72	159.49	15.11
11	F	68	52.41	10.33
12	F	70	101.62	11.82
13	F	73	23.31	6.78
14	Μ	73	17.13	20.11
15	М	80	51.62	12.59
16	F	44	39.68	8.43
17	М	83	25.82	20.83
18	F	66	27.32	8.56
19	F	57	29.21	9.07
20	Μ	65	42.98	8.74
21	Μ	67	77.78	10.07
22	Μ	70	19.45	8.83
23	F	65	25.32	9.72
24	F	49	17.64	18.43
25	М	70	28.53	13.25
26	М	74	18.34	22.24
27	Μ	78	56.22	11.20
28	Μ	85	75.93	6.07
29	F	78	36.88	10.21



Figure 2 The mean 8-OHdG content in DNA from the pterygium tissues is higher than that of the adjacent normal conjunctiva.

significantly higher levels of 8-OHdG in the DNA of pterygium tissues as compared to its adjacent normal conjunctiva. This finding supports the idea that oxidative damage to DNA may play a role in the development of pterygium.

The pathogenesis of pterygium is not clearly understood. Recent studies have provided evidence indicating that pterygium has both degenerative and proliferative characteristics. The degenerative nature of pterygium revealed from light and electron microscopic examinations showed elastotic degeneration of collagen fibres, activated fibroblasts, and inflammatory infiltrate that resembles the actinic degenerative changes seen in the skin exposed to chronic UV radiation.^{5,6,14} On the other hand, the epithelium of pterygium may display squamous metaplasia.¹⁵ In addition, adjunctive alkylating antineoplastic agent or radiotherapy is usually required to prevent the high recurrent rate in the treatment of pterygium.^{16,17} Furthermore, the presence of oncogenic viruses,18 loss of heterozygosity,19 and abnormal expression of p53 in pterygium tissues²⁰ suggest possible neoplastic nature of pterygium. Oxidative stress induced by either environmental agents or endogenous biochemical reactions has been implicated in the pathogenesis of a variety of degenerative and proliferative diseases.^{21–23} As a reliable biomarker of oxidative DNA damage, higher levels of 8-OHdG have been observed in the lungs of cigarette smokers,²⁴ the liver of patients with chronic hepatitis,²⁵ the breast tissue of benign and malignant tumors,²⁶ the

trabecular meshwork of glaucoma patients,²⁷ the renalcell carcinoma,²⁸ the urine of patients with dystrophinopathy,²⁹ the leucocytes of patients with Leber's hereditary optic neuropathy³⁰ and patients on chronic hemodialysis,³¹ and heart mitochondrial DNA of patients with atrial fibrillation.³² To our knowledge, this is the first study to explore the increase of 8-OHdG content in pterygium tissue.

Apart from the direct DNA damage, oxidative stress has been shown to regulate gene expression by modulating transcription factors, which then induce the expression of various proinflammatory cytokines, angiogenic, and fibrogenic factors.³³ Several reports pointed out the presence of these cytokines and growth factors and/or their receptors in pterygium tissues; in addition, their presence could also be induced by UV irradiation.^{34–37} These evidences suggest that oxidative stress and oxidative damage, especially caused by UV irradiation, may contribute to the pathogenesis of pterygium.

The 8-OHdG frequently mispairs with adenine during DNA replication, which leads to G-C to T-A transversion, ultimately gives rise to gene mutation.³⁸ A number of defense and repair mechanisms have evolved to minimize its accumulation within the genome. Primary defense mechanisms include antioxidants and free radical-scavenging enzymes. Once 8-OHdG is formed, the hOGG1 enzyme, a DNA glycosylase/apurinic lyase, is expressed and involved in the base excision repair of 8-OHdG. Previous studies have revealed a common polymorphism in the hOGG1 gene of functional difference. It has been proved by using an Escherichia coli complementation assay that the DNA repair activity of hOGG1-Cys³²⁶ is lower than that of hOGG1-Ser³²⁶.³⁹ This genetic polymorphism of hOGG1 is associated with the risk of many kinds of neoplasia and high degree of oxidative DNA damage in patients suffering from chronic oxidative stress.^{40,41} Our previous study has shown that individuals who carried the homozygous Cys/Cys genotype were at a 2.2-fold higher risk to develop pterygium. Besides, the patients with Cys/Cys genotype were younger than those with other two genotypes combined (Cys/Ser and Ser/Ser) in the pterygium group.⁹ It is suggested that subjects with the Cys326 allele may have more accumulative oxidative damages and thus are more prone to develop pterygium. Our current findings further proved that there was significant higher oxidative DNA damage in pterygium tissue. However, further studies are warranted to provide more information about the role of hOGG1 and 8-OHdG in the pathogenesis of pterygium.

In conclusion, our study demonstrated increased oxidative damage to DNA in human pterygium tissues. These findings shed new light to better understanding of 830

the pathogenesis of pterygium and may be useful in the prevention and treatment of this disease.

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