Detection of Endogenous DNA Double-strand Breaks in Oral Squamous Epithelial Lesions by P53-binding Protein 1

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Abstract. Background/Aim: P53-binding protein 1 (53BP1) is one of the DNA damage response (DDR) molecules. This study aimed to assess 53BP1 expression by immunofluorescence (IF) as a biomarker to differentiate between oral squamous epithelial lesions (OSELs). Materials and Methods: We analyzed 129 archival oral biopsy samples, including 18 benign squamous lesions (BSLs), 37 low-grade dysplasias (LGDs), 22 high-grade dysplasias (HGDs), and 52 oral squamous cell carcinomas (OSCCs). 53BP1 and Ki-67 expressions were examined by double IF to assess the type of 53BP1 expression. Results: We found that OSCC exhibited several 53BP1 nuclear foci, particularly high-DNA damage response (HDDR) and large focus (LF)-type, suggesting the presence of endogenous DNA double-strand breaks in the cancer genome, which could disrupt DDR and induce genomic injury. We also found a difference in 53BP1 expression between LGD and HGD, but not between BSL and LGD. Among the Ki-67-positive cells, HDDRand LF-type expressions were higher in OSELs of higher grades. Conclusion: 53BP1 expression can be a valuable biomarker for OSELs to help estimate the grade of oral epithelial dysplasia.

This article is freely accessible online.

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Key Words: 53BP1, genome instability, immunofluorescence, oral cancer, tumor biomarkers.

Oral squamous cell carcinoma (OSCC) incidence is estimated to be more than 500,000 annually (1). It is commonly preceded by oral epithelial dysplasia (OED), and its 5-year survival rate can be improved by early-stage diagnosis (2). The 4th edition of the WHO Classification of Head and Neck Tumours (2017) defines OED as 'a spectrum of architectural and cytological epithelial changes caused by accumulation of genetic changes associated with an increased risk of progression to squamous cell carcinoma (SCC) (3). However, dysplasia grading is poorly reproducible between pathologists. Although OED is traditionally divided into three severity grades, a binary system is also advocated to improve reproducibility. Consensus grading after reviewing by multiple pathologists may enhance diagnostic reliability. The Japanese Society of Oral Oncology recommends a binary system, such as lowgrade and high-grade dysplasia (LGD and HGD), to diagnose OED based on clinical management (4). HGD may include carcinoma in situ, and commonly develop into SCC, hence HGD lesions are usually recommended for active surgical intervention (5). Most LGDs never progress to carcinoma, so a 'wait and see' policy may be adopted for these lesions. Thus, the clinical management of OED is dependent on the histopathological diagnosis by biopsy.

Histological findings of architectural and cytological disturbances in biopsy specimens are critical for OED diagnosis. However, there is lack of consistent evidence to indicate translation of individual features into a dysplasia grade. Thus, the issue of poor consensus is based on the current diagnostic criteria underlying the low reproducibility of OED differential diagnosis and necessitates establishment of an ancillary technique to diagnose OED. P53-binding protein 1 (53BP1) is a DNA damage response (DDR) molecule with a BRCA1 C-terminal domain (6). 53BP1

rapidly forms nuclear foci (NF) with other DDR molecules, like ATM and yH2AX, at the site of DNA double-strand breaks (DSBs) in response to ionizing radiation (7, 8). Our previous dual-color immunofluorescence (IF) analysis revealed frequent co-localization of 53BP1 and vH2AX NF, consistent with DNA DSBs induction in irradiated rat thyroid follicular cells (9). Additionally, we demonstrated the frequent co-localization of 53BP1 and yH2AX NF in untreated human thyroid follicular carcinomas, suggesting occurrence of endogenous DNA DSBs and DDR activation as hallmarks of tumor genome instability (9). Furthermore, we previously demonstrated that the number of 53BP1 NF in uterine cervical cells appeared to increase with cancer progression (10). The distribution of 53BP1 NF was similar to that of punctate human papilloma virus (HPV) signals as determined by in situ hybridization (ISH) and also to the pattern of p16^{INK4a} overexpression, an established surrogate marker for HPV infection in cervical squamous intraepithelial lesions (SILs) (10). Thus, 53BP1 NF are associated with viral infection and replication stress, and the IF analysis of 53BP1 expression can be a useful tool to estimate the grade of SIL during cervical carcinogenesis (10). We hypothesize that the expression of 53BP1 may be an indicator of an endogenous genomic instability that drives tumorigenesis. This study aimed to evaluate type of 53BP1 expression as an indicator of genomic instability in oral squamous epithelial lesion.

Materials and Methods

Patients and histological evaluation. We found total 158 biopsy cases of oral lesions which were diagnosed as mild (n=62)/ moderate (n=23)/ severe (n=12) dysplasia and SCC (n=61) at the Nagasaki University Hospital between 2016 and 2017. Pathological diagnosis was confirmed by two certified pathologists (S.M. and M.N.) according to the 4th WHO Classification of Head and Neck Tumours (Figure 1) (3). OED dysplasia grading was performed using the binary system (LGD or HGD). All samples were formalin-fixed and paraffin-embedded tissues that were prepared for IF, immunohistochemistry (IHC), and fluorescence in situ hybridization (FISH) studies. This study was performed retrospectively in accordance with the tenets of the Declaration of Helsinki. The Ethics Committee of Nagasaki University approved the study (approval date: August 27, 2020; #15062617-4) and waived the need for informed consent. All methods were carried out in accordance with relevant guidelines and regulations. Patient profiles were anonymized by coding and collectively summarized with the obtained data as the final dataset. Patients were able to opt out of the study by following the instructions provided on the institute website.

IF analysis of 53BP1 expression. 53BP1 expression was examined with Ki-67 expression by double IF analysis to assess the extent and integrity of DDR. After antigen retrieval via microwave treatment in Target Retrieval Solution, Citrate pH 6 (Agilent Technologies, Santa Clara, CA, USA), deparaffinized sections were preincubated

with Dako Protein Block, Serum-Free (DakoCytomation, Glostrup, Denmark), then incubated with polyclonal rabbit anti-53BP1 antibody (1:1,000; Bethyl Laboratories, Montgomery, TX, USA), and monoclonal mouse anti-Ki-67 antibody (clone MIB-1; 1:50; DakoCytomation). The samples were then incubated with Alexa Fluor® 488-conjugated goat anti-rabbit and Alexa Fluor® 546conjugated goat anti-mouse antibodies (1:2,000; Invitrogen, Carlsbad, CA, USA), and mounted using VECTASHIELD® HardSet™ Mounting Medium with DAPI (Vector Laboratories, Burlingame, CA, USA). The stained sections (minimum 25-30 slices per field) were photographed at 1,000-fold magnification using the Z-stack function on a High Standard All-in-One Fluorescence Microscope (Biorevo BZ-X710; KEYENCE, Osaka, Japan), which enabled the delineation of all 53BP1 NF in the nucleus. The 53BP1 signals were measured using the image analysis software provided with the Biorevo BZ-X710 microscope. 53BP1 immunoreactivity type was classified as: i) stable (no staining), ii) low DDR (LDDR; 1 or 2 discrete NF, <1.0 µm), iii) high DDR (HDDR; 3 or more discrete NF, <1.0 µm, or diffuse nuclear staining), or iv) large foci (LF; discrete NF, ≥1.0 µm). Each type was categorized as either Ki-67 negative or positive, yielding eight categories (Figure 2). The percentage of nuclei with each type of 53BP1 immunoreactivity was calculated in at least five consecutive fields along the basement membrane of each squamous lesion.

IHC for p53 and p16^{INK4a} expression. Deparaffinized 4-µm sections were pre-treated by heating in a microwave in a pH 9.0 phosphate buffer for 20 min, then incubated with anti-p53 monoclonal antibody (1:200; Leica Biosystems, Nussloch, Germany) or mouse anti-p16INK4a monoclonal antibody (1:2; Roche Diagnostics, Basel, Switzerland) for 15 min at 37°C. After incubation for 8 min with the primary reagent of the Bond polymer system (Leica Biosystems), the samples were reacted with the polymer reagent (peroxidase-labelled polymer-conjugated anti-mouse polyclonal antibody and anti-rabbit polyclonal antibody) for 8 min. All samples were then incubated with 3,3,-diaminobenzidine hydrogen peroxide for 10 min and counterstained with hematoxylin. The squamous epithelial lesions were considered positive for 53BP1 if nuclear staining from the basal side was ≥50%. The level of p16INK4a immunoreactivity was evaluated according to the following criteria: no staining, focal nuclear and cytoplasmic staining, or diffuse and intense nuclear and cytoplasmic staining.

Dual-color FISH analysis of the 9p21 locus. To evaluate the association between p16INK4a immunoreactivity and loss of the p16INK4a gene, dual-color FISH with Vysis LSI CDKN2A SpectrumOrange/CEP9 SpectrumGreen Probes (Abbott, Chicago, IL, USA) and a Histra FISH pre-treatment kit (Jokoh, Tokyo, Japan) was carried out according to the manufacturers' instructions. FISH analysis was performed for 5 BSL cases, which included 2 samples negative and 3 samples diffusely positive for p16INK4a by IHC, and 10 SCC cases, including 5 samples each of negative and positive p16^{INK4a} by IHC. Briefly, after deparaffinization and microwave treatment in pre-treatment solution for 20 min at 94°C, 4-µm tissue sections were pre-digested with 0.06% protease for 45 min at 39°C. The tissue sections were denatured with FISH probes at 74°C for 2 min, and subsequently incubated at 37°C for 16 h. The sections were washed in 50% formamide and 2×standard saline citrate, followed by counterstaining with 4',6-diamidino-2-phenylindole. FISH signals were automatically analyzed by a fluorescence microscope

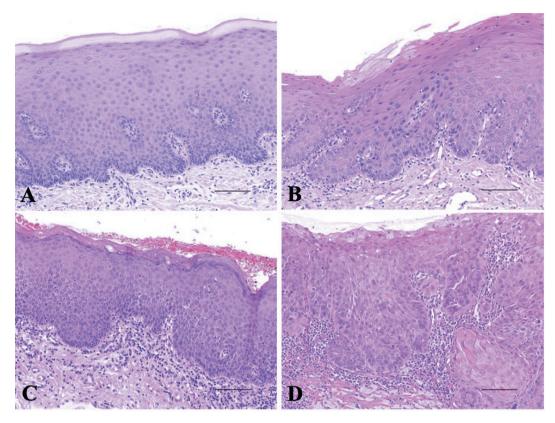


Figure 1. Representative histological images of oral epithelial lesions. (A) Benign squamous lesion. (B) Low-grade dysplasia. (C) High-grade dysplasia. (D) Oral squamous cell carcinoma. The scale bars indicate 100 µm.

(OLYMPUS BX63, Olympus, Tokyo, Japan) with the image analysis software Bioview DUET™ IFU (Bioview, Rehovot, Israel).

Statistical analysis. We assessed associations between histological type of an oral squamous epithelial lesion, type of 53BP1 expression, and level of p16^{INK4a} and p53 immunoreactivity with the Jonckheere–Terpstra, Cochran–Armitage, and *t*-tests. A logistic regression model and receiver operating characteristic (ROC) curve were used to evaluate the significance of 53BP1 expression using IF as a diagnostic test to differentiate HGD and OSCC. The closest distance from the top left corner (point (0,1)) was used to identify the optimal cut-off value on the ROC curve. The PHREG procedure in SAS 8.2 software (SAS Institute, Cary, NC, USA) was used for this calculation. All tests were two-tailed, and a *p*-value<0.05 was considered as statistically significant.

Results

Clinicopathological profiles of patients. The clinicopathological profiles are summarized in Table I. Total 129 (81.6%) out of 158 cases were available in this study, while other 29 (18.4%) cases were excluded from the study because of discordant diagnoses (n=16), poor staining quality in IF analysis (n=10), or lost their squamous lesion (n=3). Histologically, our samples

included 18 benign squamous lesion (BSL) (average age: 68.7 years), 37 LGD (average age: 70.9 years), 22 HGD (average age: 69.0 years), and 52 OSCC (average age: 72.3 years) samples. Representative lesion images are shown in Figure 1.

53BP1 expression in oral squamous epithelial lesions. The results of IF analyses for 53BP1 expression in the oral squamous epithelial lesion are presented in Table II, and representative images are depicted in Figure 2. The histological type of oral squamous epithelial lesion was significantly associated with the 53BP1 expression type (p<0.0001). In BSL samples, 74.7% of nuclei had stable expression, and only 2.6% of nuclei had HDDR- or LF-type expression. Similarly, 82.8% of nuclei in OED samples had stable expression, and only 2.7% had HDDR- or LF-type expression. In OSCC samples, 48.7% of nuclei had stable expression, and 31.4% of nuclei showed HDDR- or LF-type expression. The ratio of nuclei with 53BP1 NF amongst the counted nuclei was significantly higher in OSCC samples than in other lesions (p<0.0001), whereas no significant difference was observed between BSL, and LGD and HGD OED samples. If 6.8 % was adopted as the cut-off value

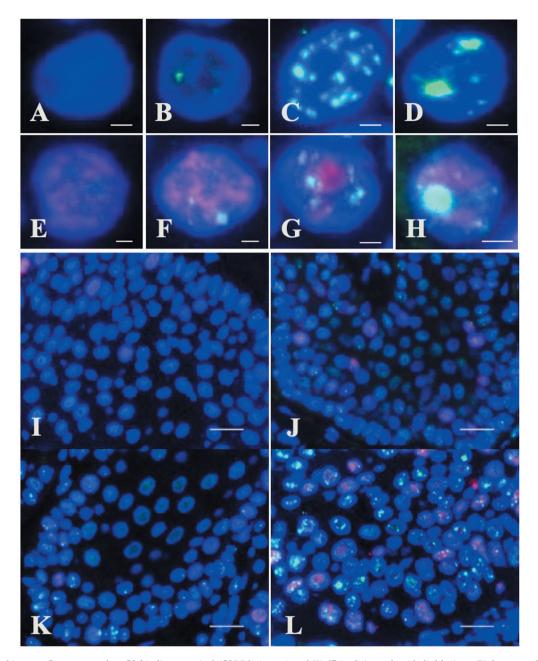


Figure 2. Dual immunofluorescence for p53-binding protein 1 (53BP1) (green) and Ki-67 (red) in oral epithelial lesions. Eight types of p53-binding protein 1 (53BP1) expression were observed by immunofluorescence using Ki-67 expression (A-H). (A and E): Stable type, no or faint nuclear 53BP1 staining. (B and F): Low DNA damage response (DDR)-type, characterized by 1 or 2 discrete 53BP1 nuclear foci. (C and G): High DDR-type, with 3 or more discrete 53BP1 nuclear foci. (D and H): Large discrete nuclear foci measuring ≥1µm. The scale bars indicate 2µm. (I): Benign squamous lesion. (J): Low-grade dysplasia. (K): High-grade dysplasia. (L): Oral squamous cell carcinoma. The scale bars indicate 20 µm.

[area under the curve (AUC): 0.862, distance: 0.247] for the diagnosis of HGD or OSCC, the sensitivity and specificity were 77.0% and 91.9%, respectively. If 15.1% was adopted as the cut-off value (AUC: 0.983, distance: 0.108) to diagnose OSCC, the sensitivity and specificity were 94.2% and 91.0%, respectively.

Among Ki-67–positive cells, HDDR- and LF-type 53BP1 expression were observed more frequently in samples with higher histological grades: 0.3%, 0.6%, and 8.0% of nuclei in BSL, OED, and OSCC samples, respectively. If 1.1% was adopted as the cut-off value (AUC: 0.878, distance: 0.26) to diagnose HGD or OSCC, the sensitivity and specificity were

Table I. Clinicopathologic profiles of patients included in the study.

	n	M/F	Age	Sites				Clinical diagnoses						
			(range)	Т	В	G	Р	0	Lichen	Leukoplakia	Erosion/ ulcer	Neoplasia	Carcinoma	0
BSL	18	9/9	68.7 (43-86)	1 (5.6%)	4 (22.2%)	9 (50%)	3 (16.7%)	1 (5.6%)	1 (5.6%)	4 (22.2%)	1 (5.6%)	3 (16.7%)	6 (33.3%)	3 (16.7%)
OED	59	33/26	70.2 (35-92)	25 (42.4%)	14 (23.7%)	17 (28.8%)	2 (3.4%)	1 (1.7%)	3 (5.1%)	44 (74.6%)	3 (5.1%)	4 (6.8%)	5 (8.5%)	0
LGD	37	21/16	70.9 (47-92)	14 (37.8%)	7 (18.9%)	14 (37.8%)	2 (5.4%)	0	3 (8.1%)	27 (73.0%)	1 (2.7%)	4 (10.8%)	2 (5.4%)	0
HGD	22	12/10	69.0 (35-83)	11 (50.0%)	7 (31.8%)	3 (13.6%)	0	1 (4.5%)	0	17 (77.2%)	2 (9.1%)	0	3 (13.6%)	0
OSCC	52	25/27	72.3 (29-99)	14 (26.9%)	11 (21.2%)	18 (34.6%)	5 (9.6%)	4 (7.7%)	2 (3.8%)	4 (7.7%)	2 (3.8%)	7 (13.5%)	34 (65.4%)	3 (5.8%)

M: Male; F: female; T: tongue; B: buccal mucosa; G: gingiva; P: palatine; O: others; BSL: benign squamous lesion; OED: oral epithelial dysplasia; LGD: low-grade dysplasia; HGD: high-grade dysplasia; OSCC: oral squamous cell carcinoma.

Table II. Types of p53-binding protein 1 (53BP1) expression in oral squamous epithelial lesions.

	Counted nuclei		• 1	P1 expression ted nuclei		Ki-67-positive nuclei	Type of 53BP1 expression in Ki-67-positive nuclei			
		Stable	LDDR	HDDR	LF		Stable	LDDR	HDDR	LF
BSL	10,121	7,558 (74.7%)	2,303 (22.8%)	244 (2.4%)	16 (0.2%)	1,512 (14.9%)	1,224 (12.1%)	257 (2.5%)	31 (0.3%)	0 (0.0%)
OED	37,434	31,013 (82.8%)	5,379 (14.4%)	875 (2.3%)	167 (0.4%)	4.241 (11.3%)	3,271 (8.7%)	761 (2.0%)	185 (0.5%)	25 (0.1%)
LGD	20,312	16,490 (81.2%)	3,317 (16.3%)	456 (2.2%)	49 (0.2%)	2,011 (9.9%)	1,562 (7.7%)	389 (1.9%)	58 (0.3%)	(0.0%)
HGD	17,122	14,523 (84.8%)	2,062 (12.0%)	419 (2.4%)	118 (0.7%)	2,231 (13.0%)	1,709 (10.0%)	372 (2.2%)	127 (0.7%)	23 (0.1%)
OSCC	32,283	15,731 (48.7%)	6,422 (19.9%)	9,270 (28.7%) 0001*	860 (2.7%)	7,192 (22.3%)	3,297 (10.2%)	1,316 (4.1%)	2,472 (7.7%) 0001*	107 (0.3%)

LDDR: Low DNA damage response; HDDR: high DNA damage response; LF: large foci; BSL: benign squamous lesion; OED: oral epithelial dysplasia; LGD: low-grade dysplasia; HGD: high-grade dysplasia; OSCC: oral squamous cell carcinoma.*by Jonckheere-Tepstra test.

75.7% and 91.9%, respectively. Finally, if 2.6% was adopted as the cut-off value (AUC: 0.931, distance: 0.108) to diagnose OSCC, the sensitivity and specificity were 94.2% and 91.0%, respectively.

Immunoreactivity of p53 and p16^{INK4a} expression in oral squamous epithelial lesions. The IHC results for p53 and p16^{INK4a} expression in oral squamous epithelial lesions are presented in Table III, and representative images are depicted in Figure 3. The percentage of cases with p53 immunoreactivity was significantly higher in samples of higher histological grade: 5.6%, 6.8%, and 28.8% in BSL, OED, and OSCC cases, respectively. Conversely, p16^{INK4a} immunoreactivity was significantly lower in samples of

Table III. Immunohistochemical results for p53 and p16^{INK4a} expression in oral squamous epithelial lesions.

	n	p53 positive	p16 ^{INK4a} positive			
			Diffuse	Focal		
BSL	18	1 (5.6%)	6 (33.3%)	10 (55.6%)		
OED	59	4 (6.8%)	10 (16.9%)	31 (52.5%)		
LGD	37	1 (2.7%)	10 (27.0%)	20 (54.1%)		
HGD	22	3 (13.6%)	0	11 (50.0%)		
OSCC	52	15 (28.8%)	10 (19.2%)	11 (21.2%)		
		p=0.0009*	p=0.5223*	p=0.0004*		

BSL: Benign squamous lesion; OED: oral epithelial dysplasia; LGD: low-grade dysplasia; HGD: high-grade dysplasia; OSCC: oral squamous cell carcinoma. *By Cochran-Armitage test.

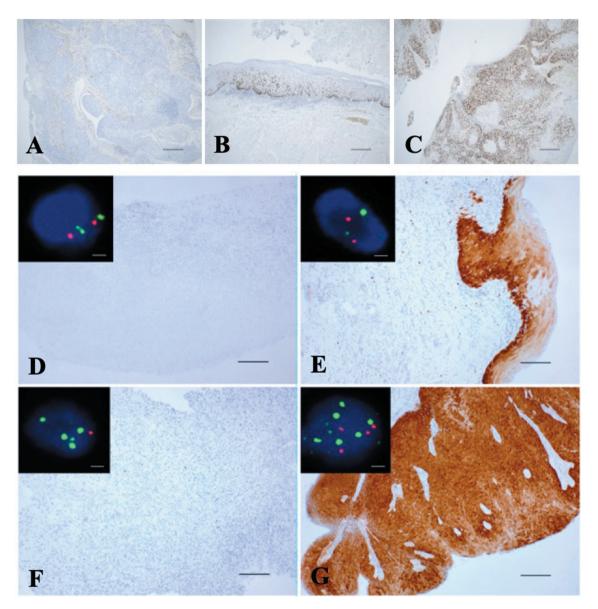


Figure 3. Immunohistochemistry analysis for p53 expression in oral squamous epithelial lesions (A-C), and comparison between immunohistochemical p16^{INK4a} expression and presence of the 9q21 locus (orange) by dual-color fluorescence in situ hybridization (FISH) using the CEP9 probe (green) in oral squamous epithelial lesions (D-G). (A): Benign squamous lesion. (B): Oral epithelial dysplasia. (C): Oral squamous cell carcinoma. (D and E): Two cases of benign squamous lesions had a wild type 9p21 locus (orange/green=1) regardless of p16^{INK4a} immunoreactivity. (F and G): Two cases of oral squamous cell carcinoma exhibited loss of the 9p21 locus (orange/green <0.5) regardless of p16^{INK4a} immunoreactivity. The scale bars indicate 400 µm for immunohistochemical p53 and p16^{INK4a} expressions and 2 µm for FISH analysis.

higher histological grade: 88.9%, 69.4%, and 40.3% in BSL, OED, and OSCC cases, respectively.

FISH analysis of the 9p21 locus in oral squamous epithelial lesions. Representative FISH images for the 9p21 locus in oral squamous epithelial lesions are depicted along with p16^{INK4a} immunoreactivity in Figure 3 to facilitate comparison in the

respective cases. The average number of nuclei exhibiting loss of the 9p21 locus (9p21 [orange]-to-CEP9 [green] signal ratio <0.5 is shown in Table IV. The results are shown in comparison to p16^{INK4a} immunoreactivity in BSL and OSCC samples. Loss of the 9p21 locus was significantly more common in OSCC samples than BSL samples, however it was not associated with p16^{INK4a} immunoreactivity.

Table IV. Comparison between p16^{INK4a} immunoreactivity and chromosome 9q21 alteration by dual-color FISH for in oral squamous epithelial lesions.

Туре	n	p16 ^{INK4a}	Average ratio of nuclei with loss of 9p21 per case by FISH
BSL	2	-	49.5%*↑
	3	2+	53.0%*↑
OSCC	5	-	84.6%↑
	5	2+	81.2%

FISH: Fluorescence *in situ* hybridization; BSL: benign squamous lesion; OSCC: oral squamous cell carcinoma. $\uparrow p < 0.0001$, *p = 0.6914, **p = 0.2010 by *t*-test.

Discussion

We observed a higher frequency of 53BP1 NF in HGD and OSCC samples than in BSL and LGD, and in the OSCC compared to the OED samples, in biopsied oral squamous epithelial lesions. These findings suggest that endogenous DNA DSBs activate DDR in higher-grade lesions, contributing to histological disturbance and malignant transformation. We hypothesized that the co-localization of 53BP1 NF and Ki-67 expression in the nucleus is an abnormal DDR phenotype, because DDR is normally activated during cell-cycle arrest (11, 12). Our results reveal frequent co-localization of 53BP1 NF and Ki-67 expression in OSCC cell nuclei, suggesting perturbed DDR in cancer cells. Impairment of the DDR machinery is frequently seen in several human cancers, and it induces genomic instability, a hallmark of malignancy. Thus, we believe that analyzing 53BP1 expression by IF can be useful for estimation of the level of genomic instability and malignant potential of human tumors. Indeed, our previous reports have demonstrated that the type of 53BP1 expression is significantly associated with the malignant potency of several tumors (13, 14).

Regarding oral epithelial lesions, other authors suggested that a high level of 53BP1 immunoreactivity was associated with a risk of oral lichen planus neoplastic transformation (15). Furthermore, in a Fanconi anemia (FA) patient suffering relapsing OSCC, the defective DNA repair was detected in tissue specimens of skin, oral mucosa and tumor with an excess of unrepaired DSBs in the proliferating zone by analyzing DDR proteins, including 53BP1 (16). FA is molecular pathologically characterized by cellular hypersensitivity to DNA damaging agents that induce DNA interstrand crosslinking, impairing strand separation and unwinding, and ultimately hindering replication and transcription, therefore, is associated with an increased predisposition to not only hematologic but also nonhematologic tumors, particularly SSC of the aerodigestive and anogenital tract (17). Because this

study also showed that the 53BP1 NF analysis of irradiated blood lymphocytes allowed to identify patients with an impaired DSB repair capacity, defective function of 53BP1 molecule may play a significant pathological role in malignant transformation of OED. Interestingly, a recent evidence suggested that poly (adenosine diphosphate–ribose) polymerase (PARP) inhibitor olaparib, which causes an accumulation of DNA damage and tumor-cell death by preventing DDR machinery, potentiates anticancer drug cytotoxicity via 53BP1 in esophageal SCC cells (18). Thus, 53BP1 can be an attractive candidate of molecular-targeted therapy for SCC in aerodigestive tract.

Histopathological OED grading by biopsy is the gold standard for determining a management approach (19-21). The degree of dysplasia is reportedly an independent marker for the prediction of malignant transformation, as HGD samples are associated with a 2.78-fold higher risk of progression to cancer than LGD samples (22). Importantly, our results indicate that IF-based analysis of 53BP1 expression is an outstanding diagnostic test to distinguish OSCC from OED or BSL, and HGD from LGD or BSL, according to a statistic textbook (23). However, the clinical application of this test is limited by the need for a fluorescence microscope and training to evaluate the staining results, which requires an understanding of the morphological features.

No single biomarker can stratify the cancer risk associated with OED. We found that p53 immunopositivity was significantly higher in OSCC than in OED and BSL samples, however, only 28.8% of OSCC cases were positive, indicating that impairment of p53 tumor suppressor function is not the main pathway in oral carcinogenesis. IHC revealed a stepwise decrease in p16^{INK4a} expression with oral carcinogenesis progression. Immunohistochemical p16^{INK4a} expression is a surrogate marker of HPV infection, and p16^{INK4a} IHC is an auxiliary technique for diagnosis of uterine cervical SIL. We inferred that 53BP1 expression pattern is a valuable biomarker for identifying the grade of cervical lesions caused by the integration of HR-HPV into the host genome (10). However, in oral lesions, up to one-third of OSCC cases are p16INK4apositive, and viral DNA is detectable by polymerase chain reaction in about 28% cases (24, 25). ISH, a more sensitive method of detecting HR-HPV, revealed very few positive cases (1-10%) (26, 27), suggesting a low incidence of HPVdriven OSCC. Therefore, p16^{INK4a} immunoreactivity is not concordant with HPV infection or dysplasia grade in the oral cavity, unlike in the uterine cervix. In addition, the current study demonstrates that reduced p16INK4a levels are not associated with loss of the 9p21 locus in oral lesions, which is a critical early event during carcinogenesis (28, 29). Thus, p16^{INK4a} expression level is not a marker of the malignant transformation of oral lesions. Expression of p16^{INK4a} in BSL represents a response to growth stress caused by various

stimuli, including inflammatory factors (30), suggesting that molecular cell-cycle regulation mechanisms are intact in benign lesions. In this study, 53BP1 NF were observed in 25.4% of BSL cases, although HDDR- and LF-type 53BP1 were rare. Similarly, our previous study on the esophagus revealed more 53BP1 NF in non-neoplastic epithelium than in neoplastic lesions. OSCC is typically associated with risk factors such as tobacco use, alcohol consumption, and betel quid chewing (31, 32). As the normal oral and esophageal squamous epitheliums are continually exposed to genotoxic agents in food, beverages, and tobacco, DDR-type 53BP1 immunoreactivity in the non-neoplastic epithelium may represent a minor genotoxic injury induced by external factors.

We propose that 53BP1 expression can be a valuable biomarker to estimate the endogenously occurring DNA DSBs or altered DDR in oral squamous epithelial lesion. The type of 53BP1 expression was associated with grade of OED, thereby influencing clinical management. Further studies are required to confirm the utility of IF analysis of 53BP1 expression as an auxiliary histologic technique to diagnose OED.

Conflicts of Interest

The Authors declare no competing interests.

Authors' Contributions

T.I., K.M., and M.N. designed and performed experiments, and wrote the article; K.T., N.U., C.O., Y.M., and Y.A. gave technical support and conceptual advice; H.K. performed statistical analyses. H.K., S.M., and M.N. confirmed pathological diagnoses. M.N. reviewed and revised the manuscript.

Acknowledgements

This work was supported in part by the Atomic Bomb Disease Institute (Nagasaki University) and is a Joint Research by Hiroshima University, Nagasaki University, and Fukushima Medical University Research Base for Radiation Accidents and Medical Science. We wish to thank Editage (www.editage.jp) for English language editing.

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Received June 27, 2021 Revised July 24, 2021 Accepted August 24, 2021