

Original Article

The expression profile of Oct4 and Sox2 in the carcinogenesis of oral mucosa

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Abstract: This study is to detect the co-expression of embryonic stem cell-related markers (Oct4 and Sox2) in the carcinogenesis of oral mucosa. The expression profile of these markers was studied by immunohistochemistry assay in rat and human samples. The normal oral mucosa (20 cases) and the transforming oral mucosa (20 cases) were performed in rat samples. The precancerous lesions (20 cases), OSCCs in primary site (116 cases), corresponding epithelial non-cancer tissues adjacent to the OSCC (20 cases) and 46 paired metastatic OSCCs in lymph nodes were performed in human samples. The co-expression of the two markers was defined as both of them are positively detected in the same site of one case under one selected field of microscope. The results indicated that Oct4 and Sox2 were individually detected in normal oral mucosa, but they cannot be co-expressed in the same site of one case. The co-expression of Oct4 and Sox2 (Oct4⁺Sox2⁺) was frequently detected in the transforming oral mucosa of rat (16/20), precancerous lesions of human (12/20) and epithelial non-cancer tissues adjacent to the OSCC (18/20). Also, Oct4⁺Sox2⁺ profile was remarkable noted in the primary sites of OSCCs (38/116). In the 46 paired OSCCs (primary sites with lymph node metastasis), Oct4⁺Sox2⁺ profile (8/46) was less frequently detected than Oct4^{low/+}Sox2^{low/+} (14/46) profile in the metastatic sites. To conclude, this study suggests Oct4 and Sox2 are expressed in normal oral mucosa, premalignant diseases, primary sites of OSCCs and metastasis sites of OSCCs. Oct4⁺Sox2⁺ profile may contribute to the malignant transformation of oral mucosa.

Keywords: Carcinogenesis, Oct4, Sox2

Introduction

Oral squamous cell carcinoma (OSCC) occurs in the lining of upper digestive tract, has a remarkable incidence worldwide [1]. OSCC often occurs with metastasis to adjacent and distant organs, and patients suffer from pain, facial nerve paralysis, pathological fractures, recurrence and frequently die under very distressing circumstances [2]. The majority of OSCC patients often have cervical lymph nodal metastasis when diagnosed even at early stage. Hence, it is critical to understand key molecular mechanisms in the carcinogenesis and spread of OSCC, with a view to design individualized- targeted therapies.

Cancer stem cells (CSCs) have been defined as a unique subpopulation in cancers that possess the ability to initiate neoplasm and sustain tumour self-renewal [3]. Also, self-renewal is

the property of embryonic stem cells (ESCs), thus suggesting common molecules might exist between CSCs and ESCs [4]. Embryonic stem cell markers (Oct4 and Sox2) can be co-expressed in ESCs, are essential for the maintenance of ESCs pluripotency [5]. In addition, somatic cells can be reprogrammed into induced pluripotent stem cells (iPSC) by Oct4, Sox2 in combination with Klf4 (Kruppel-like factor 4) and c-Myc [6]. Seeding iPSC cells in immune-deficient mice can form teratomas [7]. Therefore, the process of reprogramming of iPSC is similar to the generation of cancer stem cells.

Oct4 and Sox2 are expressed in many types of cancer [8, 9]. Knocking down these genes could decrease tumour-sphere formation *in vitro* and inhibit tumour formation in xenograft mouse models [8]. The individual expression of Oct4 and Sox2 are noted in lung adenocarcinoma

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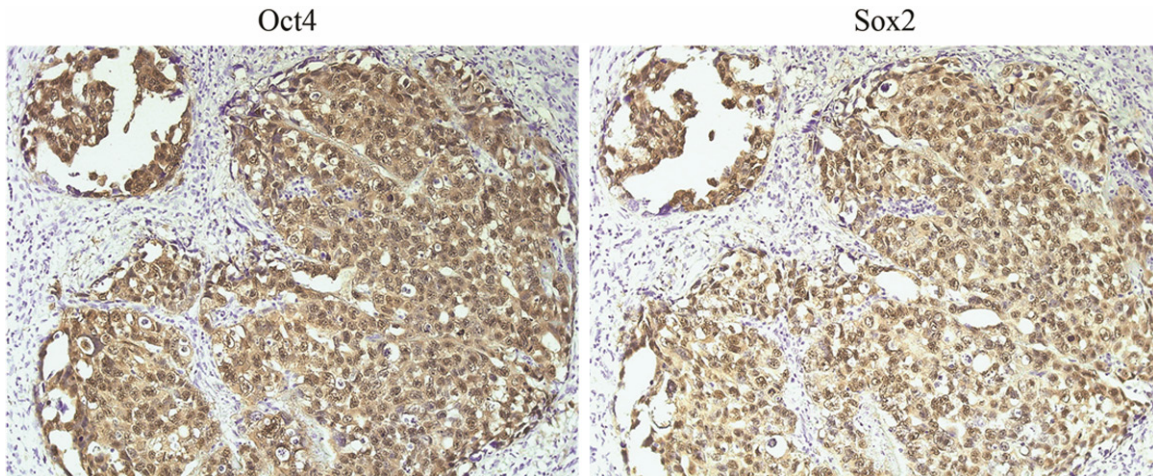


Figure 1. The expression profile of Oct4 and Sox2 in an embryonal carcinoma. Under one selected field of microscope ($\times 200$), Oct4 and Sox2 are co-expressed in cell nuclei (Oct4⁺Sox2⁺).

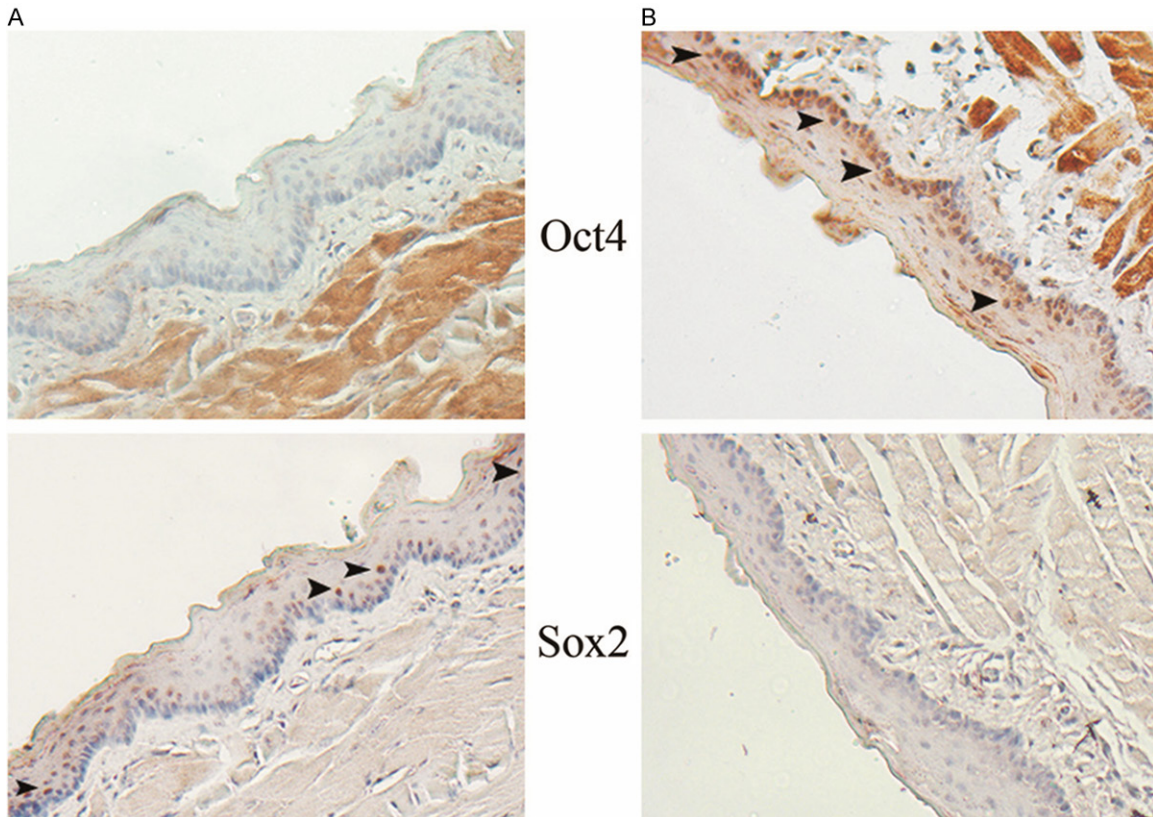


Figure 2. The expression profile of Oct4 and Sox2 in normal oral mucosa of rat samples. Under one selected field of microscope ($\times 160$), Oct4 and Sox2 nuclear positive cells are individually detected in different cases, respectively. The arrow heads show the locations of these positive cells are basal layers.

[10], gliomas [11], gastric carcinoma [12] and OSCCs [9]. Oct4 has two isoforms in human tissues, Oct-4A and Oct-4B [13]. Oct-4A protein, localized to the nuclei of ESCs, is functional for

maintaining the pluripotency of ESCs and promoting tumorigenesis. Oct-4B protein, localized to the cytoplasm of ESCs, is not sufficient to maintain undifferentiated state and activate

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Table 1. Association of expression profile of ESC-related markers in rat samples

	Oct4 positive (%)	Sox2 positive (%)	Both high expression (%)	Both low expression or negative (%)
Normal (20 cases)	16 (80)	5 (25)*	0 (0)#	1 (5)
Experimental (20 cases)	17(85)	16 (80)**	14 (70)##	2 (10)

*VS**, p<0.05; #VS##, p<0.01.

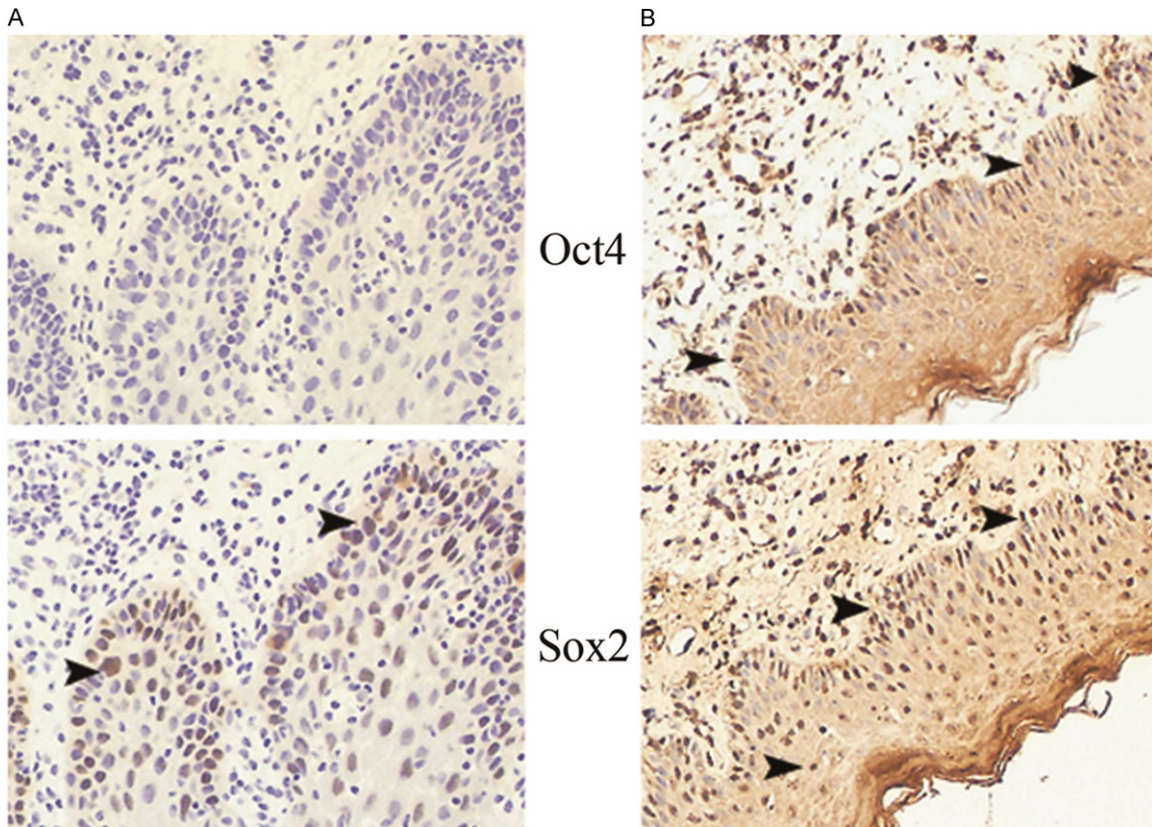


Figure 3. The expression profile of Oct4 and Sox2 in precancerous lesions of human samples ($\times 160$). A: A representative sample of the expression profile of Oct4 and Sox2 in OLK. Under one selected field of microscope, nuclear positive cells of Sox2 are demonstrated (arrow heads). The expression of Oct4 is negative. B: A representative sample of the expression profile of Oct4 and Sox2 in OLP. Under one selected field of microscope, Oct4 and Sox2 double positive cells (Oct4⁺Sox2⁺) are noted (arrow heads).

transcription of Oct4 gene [14]. The two variants of Oct4 often lead to misunderstandings in stem cell research and current knowledge of cancer progression.

At present, the correlations of Oct4 and Sox2 with metastasis features of OSCC patient remains poorly understood. In consideration that cancer stem cells might be responsible for metastasis of cancer, we hypothesized that these molecules (Oct4 and Sox2) should represent cancer stem cells and contribute to the carcinogenesis of oral mucosa. In this study, we aimed to investigate the differential expression

profile of Oct4 and Sox2 in our OSCC cohort as tumour progress in different grades. In addition, the expression patterns of these ESCs proteins in normal oral mucosa, transforming oral mucosa and precancerous lesions of oral mucosa also were examined.

Materials and methods

Patients and specimens

All biopsies of 116 OSCC patients, epithelial non-cancer tissues adjacent to the OSCC (20 cases) and 20 precancerous lesions were col-

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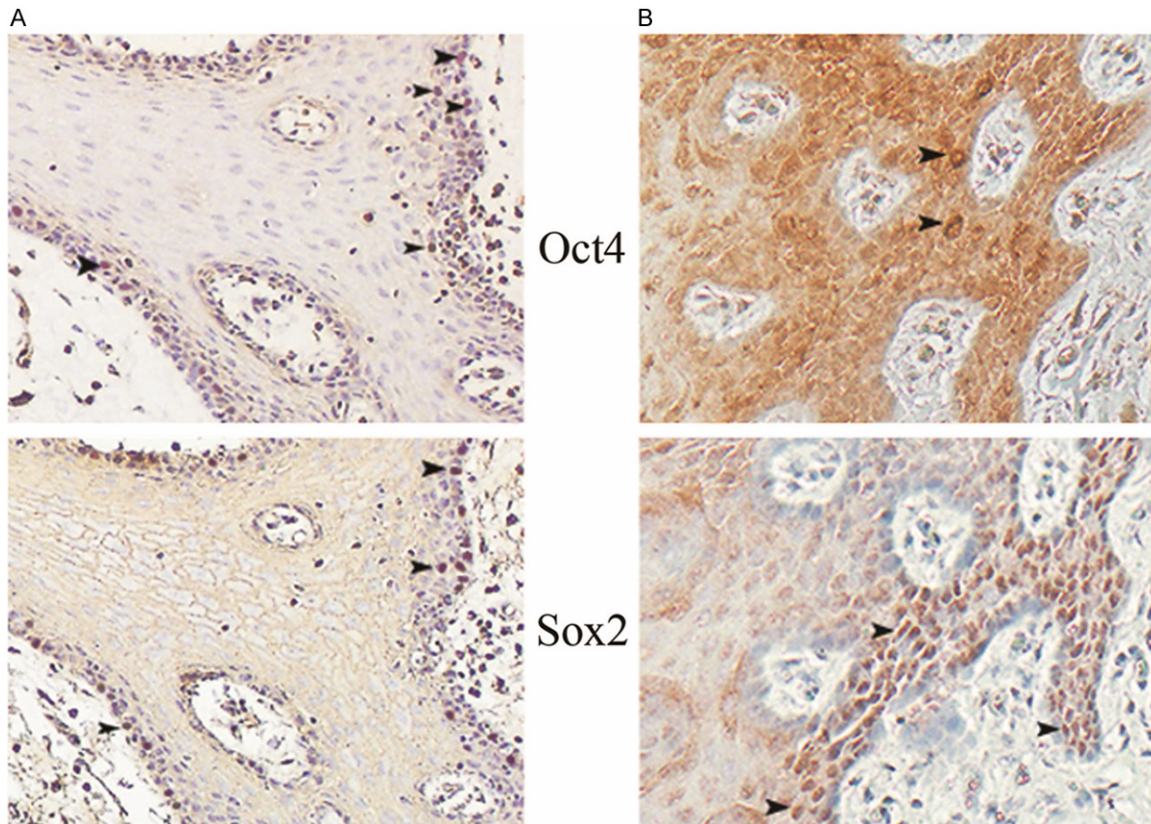


Figure 4. The expression profile of Oct4 and Sox2 in the adjacent non-cancer tissues of human samples and in the transforming tissues of rat samples. The adjacent non-cancer tissues of human samples (A) and the transforming tissues of rat samples (B) show simple hyperplasia in histology. In these two models, Oct4 and Sox2 double nuclear positive cells (Oct4⁺Sox2⁺) are frequently observed under one selected field of microscope (arrow heads $\times 160$).

lected from the Department of Pathology, The First Affiliated Hospital of Zhengzhou University, China, between 2011 and 2012. All patients of OSCC did not undergo preoperative radiotherapy and chemotherapy. The clinical data of patients was reviewed according to the International Union against Cancer rules for Head/Neck Cancer reporting [15]. Written informed consent was obtained from all of patients in this analysis and this study was approved by Zhengzhou University Ethics Committee.

Forty adult BALB/c rats (16-18 g, all males) were obtained from the Laboratory Animal Center of Zhengzhou University (Zhengzhou, China). The Ethical Committee of Experimental Animals of Zhengzhou University approved all procedures. Forty rats were randomly equally divided into experimental group (20 cases) and control group (20 cases). The 0.5% 4-nitroquinoline 1-oxide (4-NQO) in propylene glycol was

applied on dorsal of tongue three times per week in experimental group rats. After 6 weeks, all forty rats were sacrificed. The tongues were cut off and fixed in 10% neutral formalin for immunohistochemistry assay.

Immunohistochemistry assay

All formalin-fixed, paraffin embedded specimens (human and rat) were used in this study. 5 μ m serial paraffin sections were cut off and de-waxed using xylene and re-hydrated through graded alcohols and water. Tissue endogenous peroxidase activity was blocked by incubating sections with 3% H₂O₂. Antigen retrieval was performed by incubation in 10 mM citrate buffer (pH 6.0) in a pressure vessel for 5 min at 120°C. After the slides were cooled, they were rinsed with distilled water and then covered with 0.1% trypsin (Invitrogen, Grand Land, NY, USA) (pH=7.8) in room temperature for 5 min. The slides were then treated with 5% bovine

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Table 2. Clinicopathological factors in 116 patients with OSCCs

	No. of cases
OSCCs in primary site (116 cases)	
Tongue	50
Gingiva	20
Cheek	40
Hard palate	6
Gender	
Male	80
Female	36
Age	
<50 years	46
>50 years	70
Pathological grade	
Well	54
Mod	46
Poor	16
Clinical grade	
I-II	70
III-IV	46
OSCCs in lymph nodes (46 cases)	
N1	20
N2	22
N3	4

serum albumin (Invitrogen) to block non-specific binding. The staining of immunohistochemistry assay was done separately for Oct4 and Sox2. The concentration of each primary antibody was Oct4 (Oct-4A, Cell Signaling Technology, Danvers, MA, USA, 1:50) and Sox2 (Stemgent, Cambridge, MA, USA, 1:30). In each reaction, the primary antibody was incubated at 4°C overnight. Then, secondary antibody (horseradish peroxidase labelled goat-anti mouse or rabbit; Invitrogen) was added at room temperature for 30 min, washed three times with PBS. The sections were then stained with diaminobenzidine (Invitrogen) for 3 min. Positive control was an embryonal carcinoma from ovary. Replacement of the antibodies by normal serum was used as negative control. The immunoreactivity of the two proteins was evaluated by two independent pathologists who had no knowledge of the clinicopathologic factors of the patients. The co-expression of these ESC markers (Oct4 and Sox2) is defined as both of them are positively detected in the same site of one case under one selected field of microscope.

Statistical analysis

Comparisons between groups were performed using the paired Student's t-test. The significance level was taken at $P < 0.05$. The statistical tests were performed with the program, Statistical Package for Social Sciences (SPSS version 14.0, Chicago, IL, USA).

Results

Expression profile and localization of Oct4 and Sox2 in embryonal carcinoma

Three ESC-related markers (Oct4 and Sox2) were nuclear co-expressed in embryonal carcinoma (**Figure 1**). The expression profile and localization of these markers in embryonal carcinoma was set as a standard in the following immunohistochemical study.

Expression profile of Oct4 and Sox2 in rat samples (normal oral mucosa and transforming oral mucosa)

The immunohistochemical results showed that Oct4 and Sox2 could be detected in the basal layer of normal oral mucosa. Sox2 (**Figure 2A**) and Oct4 (**Figure 2B**) exhibited nuclear staining, respectively. Of note, they cannot be co-expressed in the same site of one case. In transforming oral mucosa, simple hyperplasia was exhibited. Sox2 nuclear staining was much more frequently detected than that in normal oral mucosa (**Table 1**). Of note, the nuclear co-expression of Oct4 and Sox2 was observed in these transforming samples.

Expression profile of Oct4 and Sox2 in human samples (precancerous lesions, epithelial non-cancer tissues adjacent to the OSCC and OSCCs)

There were 12 oral leukoplakia (OLK) and 8 oral lichen planus (OLP) involved in this group of precancerous lesions. Oct4 (14/20) and Sox2 (18/20) were positively detected, respectively. In the co-expression study, the Oct4⁺Sox2⁺ profile (12 cases) was predominantly found (**Figure 3B**).

Similar results for ESC-related markers were demonstrated in the epithelial non-cancer tissues adjacent to the OSCCs. Oct4 (nuclear) and Sox2 (nuclear) were positively detected, respec-

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Table 3. Association of expression profile of ESC-related markers in precancerous lesions and OSCCs

	Oct4 positive (%)	Sox2 positive (%)	Both high expression (%)	Both low expression or negative (%)
Precancerous lesions (20 cases)	14 (70)	18 (90)	12 (60)	1 (5)
Primary sites of OSCCs (116 cases)	70 (60)	74 (64)	38 (33) ^a	11 (9) ^{aa}
Paired OSCCs (46 cases)				
Primary sites	26 (57)*	46 (100) [#]	19 (41)	0 (0)
Metastatic sites	22 (48)	30 (65)	8 (17) ^b	14 (30) ^{bb}
Nodal-negative OSCCs (70 cases)	44 (63)**	28 (40) ^{##}	19 (27)	11 (16)

*VS**, p>0.05; #VS##, p<0.05; ^aVS^{aa}, p<0.05; ^bVS^{bb}, p<0.05.

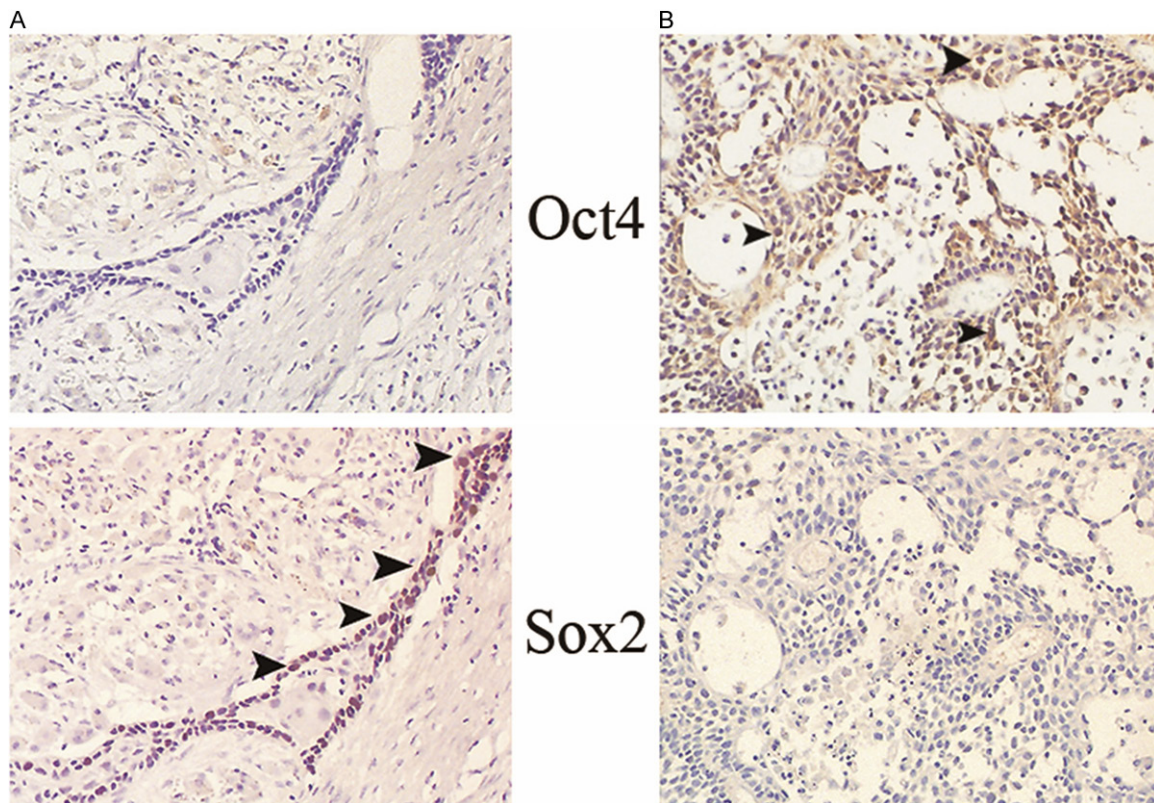


Figure 5. The expression profile of Oct4 and Sox2 in a paired OSCC (well differentiation). Sox2 single positive pattern in the primary site (A), Oct4 single positive pattern in the metastatic site (B) are noted. All the positive cells are demonstrated by arrow heads in the magnification of $\times 160$.

tively. There were 18 cases showing Oct4⁺Sox2⁺ profile in the co-expression study (Figure 4A).

In the OSCC group, there were 116 patients with (46 cases) or without lymph node metastasis (70 cases) involved in this study. According to the TNM classification, 70 cases without lymph node metastasis were in grade I-II (T₁₋₂N₀M₀), 46 cases with lymph node metastasis were in grade III-IV (T₁₋₄N₁₋₃M₀). The clinicopathological details of these OSCC patients were summarized in Table 2. The differential

expression profile of Oct4 and Sox2 were summarized in Table 3. In the primary site of 116 cases, the individual expression of Oct4 (70/116) and Sox2 (74/116) were detected in differential clinical grade of OSCCs. The Oct4⁺Sox2⁺ profile (38/116) was more frequently demonstrated than Oct4^{low/-}Sox2^{low/-} (11/116) profile. In the 46 cases of lymph nodes, the individual expression of Oct4 (22/46) and Sox2 (30/46) also were detected. However, the Oct4^{low/-}Sox2^{low/-} profile (14/46) was more frequently observed than Oct4⁺Sox2⁺

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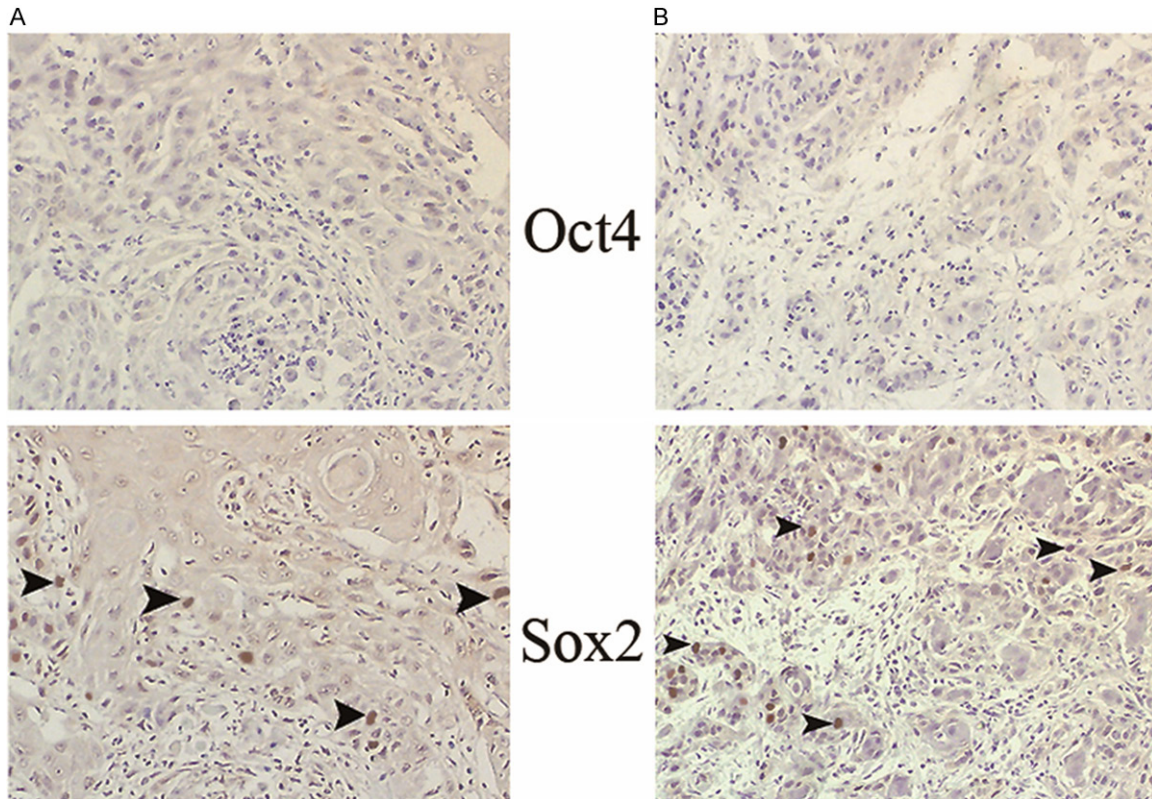


Figure 6. The expression profile of Oct4 and Sox2 in a paired OSCC (moderate differentiation). Sox2 single positive pattern in the primary site (A) and in the metastatic site (B) are noted, respectively. All the positive cells are demonstrated by arrow heads in the magnification of $\times 160$.

profile (8/46) that contrary to the expression pattern in the primary sites. Within the 46 paired OSCCs (primary sites with lymph node metastasis, grade III-IV), Sox2 profile (single expression or co-expression with Oct4) was detected in all the primary sites (46/46) (Figures 5-7). However, the expression profile of Sox2 in the lymph nodes was not always in line with their corresponding primary sites. e.g. Oct4⁺Sox2⁺ double positive profile was in the primary site but Oct4^{low/-}Sox2^{low/-} double negative profile was in the metastasis site (Figure 7).

Discussion

As transcriptional factors, Oct4 and Sox2 can be co-expressed in ESCs, described as a nuclear protein [16]. Similarly, in embryonal carcinoma (positive control), this double positive co-expression was noted in this study. Epidermal stem cells are adult stem cells without pluripotency, locating in basal layers of oral mucosa [17]. In our study, the individual expression of

Oct4 and Sox2 was shown in the epithelial basal layers. Of note, the double positive co-expression profile of these markers cannot be demonstrated in normal oral mucosa. In contrast, the co-expression of Oct4 and Sox2 (Oct4⁺Sox2⁺ profile) was frequently detected in transforming oral mucosa of rat, precancerous lesions of human, epithelial non-cancer tissues adjacent to the OSCC and primary sites of OSCCs.

The importance of the hyperplasia - dysplasia - neoplasia continuum in many epithelial tissues is now well understood [18]. In this study, the clinical cases (precancerous lesions of human, epithelial non-cancer tissues adjacent to the OSCC) and the samples from animal model (transforming oral mucosa of rat) displayed simple hyperplasia. They were in the early stage of carcinogenesis of oral mucosa. The information from them suggested Oct4⁺Sox2⁺ profile should reflect a situation in which carcinomatous changes of epithelial cells were in process. Together with the data from the primary sites of

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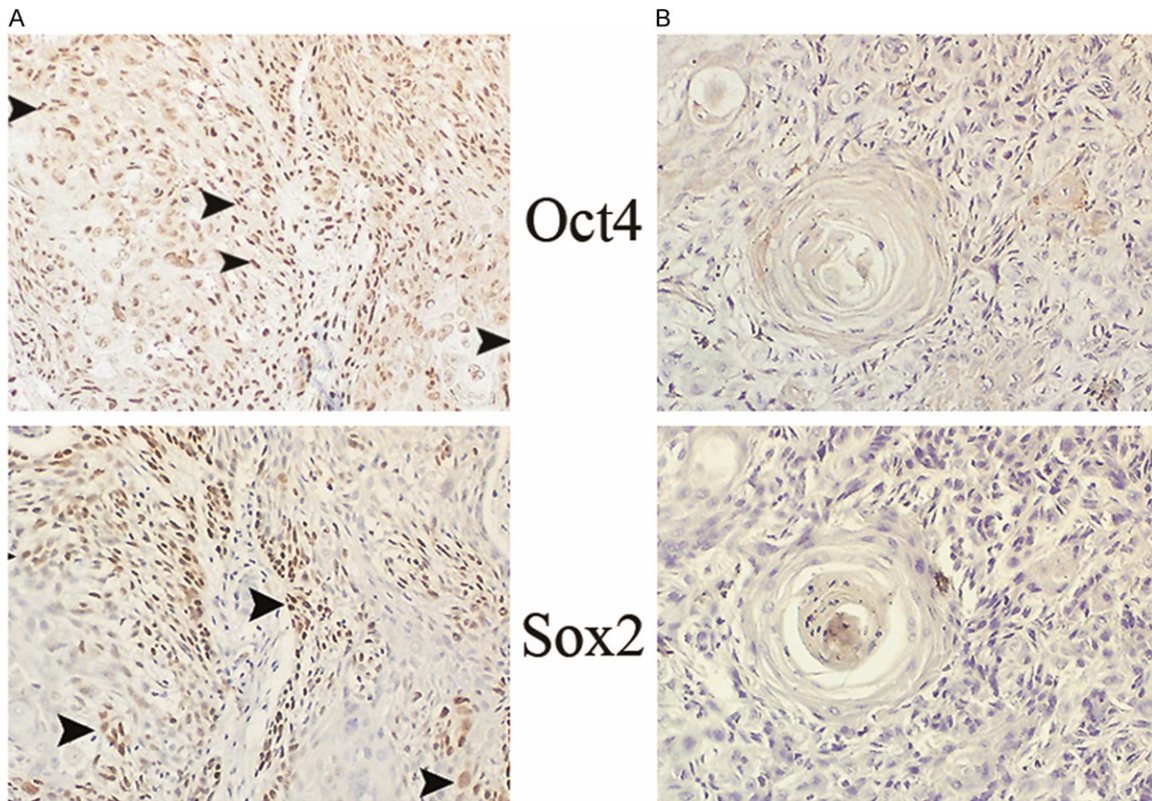


Figure 7. The expression profile of Oct4 and Sox2 in a paired OSCC (poor differentiation). Oct4 and Sox2 double positive pattern in the primary site (A), Oct4 and Sox2 double negative pattern in the metastatic site (B) are noted. All the positive cells are demonstrated by arrow heads in the magnification of $\times 160$.

OSCCs, we proposed that these double positive cells (Oct4⁺Sox2⁺) might be cancer stem cells. Of note, although Oct4⁺Sox2⁺ profile was detected in the metastatic OSCCs as well, Oct4^{low/-}Sox2^{low/-} profile was the more frequently expression pattern in the metastatic sites. Hence, metastatic cancer cells may own the independent evolution profile once the cells fight for survival in a strange environment, while cancer cells in lymph nodes are disseminated from the primary sites of OSCCs.

The correlation of Oct4 expression with cisplatin resistance in OSCC was well studied [19]. While, the relationship between Sox2 expression and tumour behavior of OSCC is under debate. Du et al. [20] showed a significant association of high Sox2 nuclear expression in nodal-negative (pNO) oral tongue squamous cell carcinoma with poor overall and disease-free survival. Michifuri et al. [21] further indicated the poor prognosis accompanying with lymph node metastasis was significant correlated with the high expression of Sox2 in the

primary sites of OSCCs. However, by using of tissue microarray technique and immunohistochemistry assay, Züllig et al. [22] showed high expression levels of Sox2 significantly correlated with negative lymph node status implying good prognosis in OSCC patients. Züllig and the colleagues indicated that the heterogeneity of primary tumors might be one of the reasons for such controversial results [22]. In our study, the Oct4 expression was not correlated with lymph node metastasis (nodal-positive VS nodal-negative, $P > 0.05$). In contrast, 100% of Sox2 expression was detected in the primary sites of 46 nodal-positive OSCCs, while only 40% of Sox2 expression was detected in the primary sites of 70 nodal-negative OSCCs. The Sox2 expression was significantly correlated with lymph node metastasis (nodal-positive VS nodal-negative, $P < 0.05$). The results of us were in line with the findings of Michifuri's group.

In summary, this study suggests that Oct4⁺Sox2⁺ profile should be the biomarker of stem cells which drive epithelial cells to OSCCs. Contrast

to traditional studies, this co-expression profile could narrow the misunderstandings caused by tumour heterogeneity in searching cancer stem cells.

Acknowledgements

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Disclosure of conflict of interest

None.

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