

ANTI TUMOR ACTIVITY OF PCDNA3.1-TUBEROUS a COMPLEX-2 SENSE IN A HUMAN ORAL TONGUE CANCER (SP-C1) CELL LINE

(AKTIVITAS ANTI TUMOR PCDNA3.1-TUBEROUS SCLEROTIC COMPLEX-2 SENSE
PADA SEL KANKER LIDAH MANUSIA (SP-C1))

Supriatno

Department of Oral Medicine
Faculty of Dentistry, Gadjah Mada University,
Jl. Denta Sekip Utara, Yogyakarta, 55281, Indonesia
E-mail: pridentagama_oncolog@yahoo.com

Abstract

Tuberous sclerosis complex (TSC) is a hamartomatous disease with defects in tuberlin (TSC2) that can be autosomal dominant inheritance or spontaneous mutation. Mutation of the TSC-2 gene encoding tuberlin on chromosome 16p13.3 increases to the clinical disorder of tuberous sclerosis characterized by the development of hamartomas. In the present study, the antitumor activity of pcDNA3.1-TSC2 sense in an oral tongue cancer *Supri's clone-1* (SP-C1) was investigated. An expression vector containing sense-oriented rat TSC-2 pcDNA with pcDNA3.1 and transfected to cell (SP-C1) to regulate the expression of TSC-2 gene in each transfectant was constructed. MTT assay was performed to examine the SP-C1 cell growth suppression transfected by pcDNA3.1-TSC2. Caspase-3 and -9 were conducted to observe the induction of cell apoptosis. Western blotting analysis was carried out to determine the protein level of TSC-2, p27^{Kip1} and α -tubulin. The results showed that, overexpression of TSC-2 exerted the growth inhibitory effect of SP-C1 cell and markedly increased apoptosis via caspase-3 and -9 pathways ($P=0.001$). Sense-oriented SP-C1-cDNA3.1-TSC-2 cancer cells have a high expression of p27^{Kip1}. In conclusion, pcDNA3.1-TSC-2 sense increased the antitumor activity of oral tongue cancer SP-C1 cell through p27^{Kip1} induction.

Key words: TSC-2, apoptosis, oral tongue cancer SP-C1, cell growth, p27^{Kip1}

Abstrak

Tuberous sclerosis complex (TSC) merupakan suatu penyakit hamartoma dengan kelainan pada tuberlin (TSC-2) akibat mutasi spontan atau warisan secara autosom dominan. Mutasi gen TSC-2 pada kromosom 16p13.3 meningkatkan kelainan tuberous sclerosis secara klinik dengan karakterisasi berkembangnya hamartoma. Pada penelitian ini dilakukan uji aktivitas antitumor pcDNA3.1-TSC-2 sense pada sel kanker lidah manusia *Supri's clone-1* (SP-C1). Konstruksi ekspresi vektor yang mengandung sense rat TSC-2 pcDNA dengan pcDNA3.1 ditransfeksikan ke sel SP-C1 untuk meregulasi ekspresi gen TSC-2 pada setiap transfeksi. Uji MTT dilakukan untuk menganalisis hambatan pertumbuhan sel SP-C1 yang ditransfeksi dengan pcDNA3.1-TSC-2. Caspase-3 dan -9 digunakan untuk mengobservasi induksi apoptosis sel. Analisis Western blotting dipakai untuk menentukan tingkat protein TSC-2, p27^{Kip1} dan α -tubulin. Hasil penelitian, peningkatan ekspresi TSC-2 menunjukkan efek hambatan pertumbuhan sel SP-C1 dan peningkatan apoptosis melalui jalur caspase-3 dan -9 ($P=0,001$). Sel kanker SP-C1-pc DNA 3.1-TSC-2 mempunyai ekspresi protein p27^{Kip1} yang tinggi. Kesimpulan, pc DNA 3.1-TSC-2 sense meningkatkan aktivitas antitumor sel kanker lidah SP-C1 melalui induksi p27^{Kip1}.

Kata kunci: TSC-2, apoptosis, kanker lidah, pertumbuhan sel, p27^{Kip1}

INTRODUCTION

Cancers of the oral cavity present a major health problem, as indicated by their high incidence in many parts of the world. In Southeast Asia

countries, oral cancers are the most common form of cancer and constitute about a third of all cancers.¹

Oral cancers are characterized by a high degree of local invasion and a high rate of metastasis to the cervical lymph nodes. It frequently shows local

recurrence after initial treatment, probably due to micro invasion and/or micro metastasis of the tumor cells at the primary site.² Treatment of oral cancer is conventionally a combination of surgery, radiotherapy, and chemotherapy. However, the overall survival rates have not improved significantly in the last two decades.³ Also, the prognosis has not changed during the past 10 years.² This highlights the necessity for continued efforts to improve the treatment modalities using TSC2 gene.

Tuberous sclerotic complex (TSC) is known as an autosomal dominant tumor suppressor gene disorder affecting 1 in 6000 live births. TSC presents variably in multiple organs, including the brain, eye, skin, kidney, heart and oral.^{4,5} Linkage analysis of TSC resulted in the identification of two distinct genetic loci on chromosome 9 and 16.⁶ These genes are hamartin/TSC-1 (9q34) and tuberin/TSC-2 (16q13).⁷ Germ-line TSC-1 and TSC-2 mutation appear to be inactivating and loss of heterozygosity at either the TSC-1 or TSC-2 region occurs in TSC tumors.^{8,9} Indicating that TSC-1 and TSC-2 are tumor suppressor genes. TSC-1 and TSC-2 follow the classic retinoblastoma tumor suppressor gene models and appear to function as negative growth regulators.¹⁰ TSC-1 and TSC-2 as tumor suppressor genes and when mutated, they give rise to abnormal cell proliferation and growth.¹¹ The protein product of TSC-2 gene, tuberin, is expressed in variety of different cell types.¹² TSC-2 has a 190 kDa protein that contains a Rap1 GTPase-activating protein (GAP)-related domain and a coiled-coil domain believed to mediate its interaction with hamartin (TSC-1) and a carboxyl terminal GTPase.¹³ Recently, TSC-2 may function as a tumor suppressor by induction of p27^{Kip1} protein.⁵ Interestingly, overexpression of TSC-2 results in reduced cell proliferation *in vitro*¹⁴ and increased amounts of the cell cycle regulator p27^{Kip1} in rat fibroblast.¹⁰ Therefore, TSC-2 is essential for p27^{Kip1} to regulate the cell cycle because tuberin can retain p27^{Kip1} protein in nuclei of cancer cells.¹⁰ In addition, TSC-2 may be closely associated with p27^{Kip1} to exert the function as tumor suppressor gene.

The purpose of the study was to investigate the antitumor activity of pcDNA3.1-TSC-2 sense in oral tongue cancer cell (SP-C1 cell) through induction of p27^{Kip1}.

MATERIALS AND METHODS

SP-C1 cells were isolated from a cervical lymph-node metastasis of an oral squamous cell carcinoma patient in our laboratory.¹⁵ Cells were maintained in Dulbecco's modified eagle medium (DMEM, Sig-

ma, St Louis, MO, USA) supplemented with 10% fetal calf serum (FCS, Moregate BioTech, Bulimba, Australia), 100 µg/ml streptomycin, and 100 units/ml penicillin.

The mammalian expression vectors pcDNA3.1-TSC-2 containing sense-oriented rat TSC-2 cDNA was constructed as follows: pcDNA3.1 (+) (Invitrogen, Carlsbad, CA) is mammalian expression vectors containing a CMV promoter. pcDNA3.1(+) was digested with Xba 1 (Takara Biomedicals, Kusatsu, Japan) and Xho 1 (Takara Biomedicals) and dephosphorylated by calf intestinal alkaline phosphatase (Roche Diagnostics, Mannheim, Germany). The rat TSC-2/pBluescript including the rat TSC-2 cDNA fragment (5.4 kbp Xba 1 and Xho 1 fragment) was obtained as a generous gift from Dr. Koji Harada (Second Department of Oral Maxillofacial Surgery, Tokushima University, Japan). This fragment was ligated to the prepared cloning site of pcDNA3.1 (+) by T4 DNA ligase (Takara Biomedicals). The direction of the ligated fragment was confirmed by sequencing analysis. Furthermore, cells (5x10⁵ cells/dish) were seeded in 100 mm culture dishes (Falcon, USA) in DMEM supplemented with 10% FCS. Twenty-four hours later, the cells were transfected with 5 µg of pcDNA3.1-TSC-2 or pcDNA3.1 without inserted (empty vector or neo) by using the Superfect reagent (Qiagen, Hilden, Germany). The cells were incubated for 48 hours in DMEM containing 10% FCS. Then, trypsinized and seeded at 1:5 ratio in 100 mm culture dishes in DMEM containing 10% FCS. Forty-eight hours later, the cells were switched to a selective medium containing Geneticin (800 mg/ml G418). After 14 days of culture in the selective medium, ten representative G418-resistant clones were isolated and expanded in a 24-well cluster dish (Falcon).

SP-C1-transfected cells were seeded on 96-well plates (Falcon, NJ, USA) at 2 x 10⁴ cells per well in DMEM containing 10% FCS. Cells were incubated for 24 and 48 hours. After incubation, the number of cells was quantitated by BioRad microplate reader machine (Sigma-Aldrich, USA) with 540 nm wavelength. MTT solution contains 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (Sigma-Aldrich) was used in examination.¹⁷

Cell lysates were prepared from SP-C1 transfected cells from a Falcon tissue culture. Briefly, samples containing equal amounts of protein (50 µg) were electrophoresed on a SDS-polyacrylamide gel and transferred to a nitrocellulose filter (PVDF membrane: Bio Rad, Hercules, CA, USA). The filters were blocked in TBS containing 5% nonfat milk powder at 37°C for 1 hour and then incubated with a

1: 500 dilution of the monoclonal antibody against Tuberin (TSC-2) protein (N-19, rabbit polyclonal antibody, Santa Cruz Biotechnology, CA, USA), and p27 protein (clone 1B4, monoclonal antibody, Novocastra Laboratories, New Castle, UK), as the primary antibody and an Amersham ECL kit (Amersham Pharmacia Biotech). Anti- α -tubulin monoclonal antibody (Zymed laboratories, San Fransisco, CA, USA) was used for normalization of Western blot analysis.

Induction of apoptosis was measured using the colorimetric assay kit (caspase-3 and caspase-9; BioVision Research Product, CA, USA) according to the manufacturer's directions. Briefly, equal amounts of tissue extracts prepared from SP-C1 transfected with pcDNA3.1-TSC2 or pcDNA3.1-neo were incubated with the substrate (DVED-pNA or LEHD-pNA) in the assay buffer for 2 hours at 37°C. Absorbance was measured at 405 nm using a microplate reader (BioRad, USA). Each determination was performed in triplicate.

Statistical analysis was performed with a Stat Works Program for Macintosh computers (Cricket Software, Philadelphia, PA, USA). Data was analyzed by Post Hoc test (Student's *t*-test) with significant level 95%.

RESULTS

Relative cell number was evaluated by comparing the absorbent of each cell using MTT assay on 24 and 48 hours. Cell number of SP-C1-pcDNA3.1-TSC-2 was significantly decreased compared with that of SP-C1-pcDNA3.1-neo (untreated control) on 24 and 48 hours ($p=0.001$) (Figure 1).

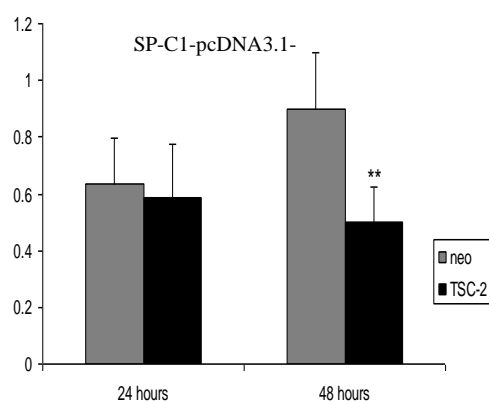


Figure 1. Relative cell number of SP-C1-pcDNA3.1-TSC-2 and SP-C1-pcDNA3.1-neo at 24 and 48 hours by MTT assay (**, $p < 0.001$)

Western blotting analysis revealed the up-regulation of TSC2 and p27^{Kip1} were detected in sense-TSC-2 transfectants when compared with that

in control cells which were transfected with pcDNA3.1-neo. The expression of α -tubulin as an internal control was approximately the same in all of the tumors (Figure 2).

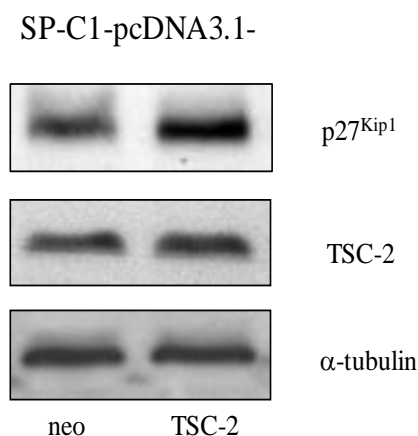


Figure 2. Expressions of TSC-2, p27^{Kip1} and -tubulin in SP-C1-pcDNA3.1-TSC-2 and SP-C1-pcDNA 3.1-neo by Western blotting analyzed

The activity of caspase-3 and -9 in sense-TSC-2 transfectants or control cells was examined using colorimetric assay. As seen in Figure 3, TSC-2 transfectants was markedly increased the proteolytic activities of caspase-3 (2.4 fold) and -9 (1.8 fold) as compared with that of control cells ($p=0.000$) (Figure 3).

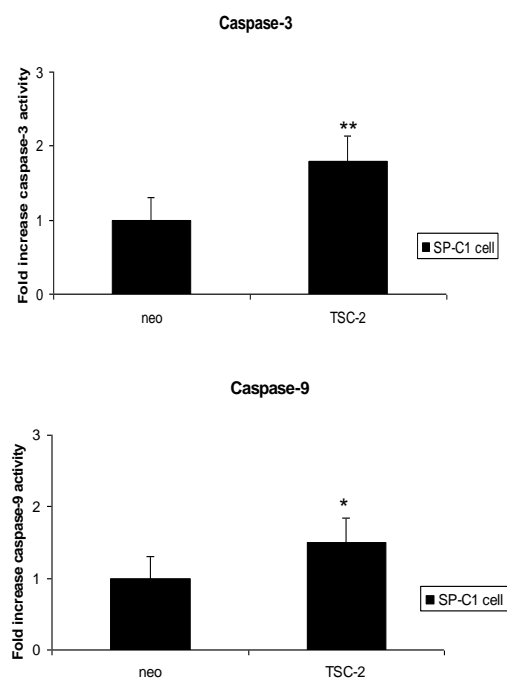


Figure 3. Activation of caspase-3 and -9 in SP-C1-pcDNA3.1-TSC-2 and SP-C1-pcDNA 3.1-neo at 48 hours incubated (**, $p < 0.001$ and *, $p < 0.05$)

DISCUSSION

Treatment strategy using gene transfection technique on human head and neck cancer included oral tongue cancer has a main attention of clinicians or oncologists in the world and in the last decade. TSC-2 is used as a tumor suppressor gene and widely used in suppression of TSC-2 syndrome (hamartoma). The autosomal dominant disease tuberous sclerosis complex (TSC) is caused by mutation in either TSC-1 on chromosome 9q34 encoding hamartin, or TSC-2 on chromosome 16p13.3 encoding tuberlin. Mutations of either the TSC-1 or the TSC-2 gene result in the syndrome of tuberous sclerosis that affects multiple organs with the development of hamartomas including cortical tuber of the brain, angiofibroma of the skin, rhabdomyoma of the heart and angiomyolipoma of the kidney.¹⁹ TSC-2 seemed to play an important role in exerting the function of p27^{Kip1} protein as a cell cycle regulator by retaining p27^{Kip1} protein in nuclei of cancer cells.¹⁰ However, the true biochemical functions of TSC-2 as a tumor suppressor and the underlying mechanisms responsible for pathogenesis of TSC-related hamartomas have not been clarified yet.

In the present study overexpression of TSC-2 exerted the antitumor effect on human oral tongue cancer cells whether they have high expression of p27^{Kip1} protein. These finding may suggest that TSC-2 can regulate the p27^{Kip1} protein at up-stream of p27^{Kip1}. It means that TSC-2 may be more important as a prognostic factor in oral SCC rather than p27^{Kip1}. Next results revealed that a significant suppression of tumor growth of sense transfectant (SP-C1-TSC-2 cells) *in vitro*. These findings may suggest that up-regulation of TSC-2 in tumor cells will be safe for the body and TSC-2 can be used as molecular target for gene therapy if the precise function of TSC-2 can be clarified. Furthermore, apoptosis induction was detected in SP-C1-pcDNA3.1-TSC2 sense through high level of caspase-3 and-9 compared with that of SP-C1-pcDNA3.1-neo. It is suggested that apoptosis induction of SP-C1 cancer cell can arise via extrinsic and intrinsic pathways. Azuma *et al.*²⁰ reported that high level of proteolytic activity caspase-3 showed a high cleavage of protein in receptor-induced apoptosis (extrinsic pathway), and induction of caspase-9 level revealed a high proteolytic activity in chemical-induced apoptosis (intrinsic pathway).

In conclusion, TSC-2 may exert the antitumor effect on human oral tongue cancer (SP-C1) cell through p27^{Kip1} protein induction.

ACKNOWLEDGMENTS

I gratefully acknowledge Prof. Mitsunobu Sato DDS., Ph.D and Dr. Koji Harada DDS., Ph.D, Department of Oral Maxillofacial Surgery and Oncology, School of Dentistry, Tokushima University, for providing some materials and discussion.

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