Introduction

The ends of eukaryotic chromosomes are usually capped by telomeres, which protect the chromosomes from damage and degradation (1). Besides, telomeres are essential for genomic integrity and cell viability (2). When the conventional DNA replication process cannot completely synthesize chromosomal ends, proliferating cells that lacking de novo synthesis of telomeric DNA eventually enter a growth arrest state called replicated senescence (3,4). In addition, the synthesis and maintenance of telomeric DNA is dependent on telomerase activity (5), while human telomerase reverse transcriptase (hTERT) catalytic subunit hTERT seemed to be the rate-limiting factor for telomerase activity (6). hTERT has been found in many highly proliferating cells, such as early embryos, reproductive tissue in testis and ovary, stem cells (7). However, most hematopoiesis stem cell and somatic tissues contain undetectable levels of hTERT (8).

Ovarian cancer is one of the most deadly female malignant tumors in the world and is characterized by abnormal proliferation of ovarian cells, with a five-year survival rate is only about 35% (9). Tumor development in ovarian cancer is a complex process involving multiple factors and stages. In addition to genetic factors associated

ShRNA-mediated silencing of hTERT promote apoptosis and senescence in human ovarian cancer cells

Chen Su¹, Yi Luo², Yaying Yang¹, Yongfen Yi¹

¹Department of Pathology, Molecular Medicine and Tumor Center, Chongqing Medical University, Chongqing 400016, China; ²Department of Gynecology and Obstetrics, The First Affiliated Hospital of Chongqing Medical University, Chongqing 400016, China

Contributions: (I) Conception and design: All authors; (II) Administrative support: Y Yi, Y Yang; (III) Provision of study materials: Yi Luo; (IV) Collection and assembly of data: All authors; (V) Data analysis and interpretation: All authors; (VI) Manuscript writing: All authors; (VII) Final approval of manuscript: All authors.

Background: Human telomerase reverse transcriptase (hTERT) is a crucial oncogene and a key factor in cell immortalization. Apart from hematopoiesis stem cell and germ cells, normal cells and tissues almost no expression of this gene. However, the positive expression rate of hTERT in malignant tumors is as high as 80–95%. This study was designed to investigate the mechanism of knockdown of hTERT expression in the regulation of ovarian cancer carcinogenesis in nude mice transplantation model.

Methods: We identified the function of hTERT in the treatment of ovarian cancer by constructing a nude mice transplantation tumor model. Then analyzed the mechanism of hTERT in modulating ovarian cancer cell apoptosis and senescence by immunohistochemically staining.

Results: Our analysis identified hTERT as a novel regulator of ovarian cancer progression and is one of the most significantly upregulated genes in ovarian cancer. Down-regulation of hTERT inhibits proliferation of ovarian cancer cells by promoting apoptosis and the expression of the senescence gene p53 and p21.

Conclusions: hTERT is a key oncogene in ovarian tumorigenesis and metastasis; the downregulation of hTERT significantly inhibit the proliferation of ovarian cancer and have an effect on treatment of ovarian cancer by activating cell apoptosis and senescence pathway.

Keywords: Human telomerase reverse transcriptase (hTERT); ovarian cancer; apoptosis; P53

doi: 10.21037/tcr.2019.03.17

View this article at: http://dx.doi.org/10.21037/tcr.2019.03.17
with carcinogenesis, telomere abnormalities also play a key role in cancer progression (10). hTERT is considered to be a rate-limiting factor for telomerase activity. In addition, hTERT is frequently expressed in the early stages of cancer development and allows cancer clones to bypass mitotic breakdown and replicative senescence (11). Thus, regulation of telomerase activity will have important effects on many developmental processes, including aging, proliferation, differentiation and tumorigenesis. Whether to prevent the progression of ovarian cancer by inactivating the hTERT gene is unclear.

In this study, we interfered with the hTERT gene by using a nude mice model of ovarian cancer transplantation. Then, we used immunohistochemistry staining and electron microscopy to detect the apoptotic phenotype of the cancerous tissues. We also investigated the senescence gene \( p53 \) and \( p21 \) expression in transplanted ovarian cancer tissues. Our experiments provide a theoretical basis and experimental basis for the clinical application of targeting hTERT gene therapy in ovarian cancer.

**Methods**

**Animals**

Four-week female BALB/c nude mice were provided by the Experimental Animal Center of Chongqing Medical University, body weight is about 14.2±2.1 g. The nude mice were cultured in SPF grade animal laboratory. Ovarian cancer cells A2780 and SKOV3 were obtained from ATCC. We cultured the cells at 37 °C, 5% CO\(_2\). After the cell confluence 80–90%, we collected the cells and adjusted the cell concentration to 5×10\(^7\) cells/mL. We subcutaneous inject the 200 μL cells in the back of the mice neck. When the nodule diameter size is up to 0.5 cm, we recorded the size of the subcutaneous nodule and determined the xenografts.

All animal experiments were conducted in accordance with the Guide for the Care and Use of Laboratory Animals and approved by the Animal Experimentation Ethics Committee of the Chongqing Medical University.

**Gene therapy assay by target hTERT**

We constructed the pG2 plasmid containing the hTERT gene based on the molecular clone protocol. The xenograft mice models were divided into three groups, 6 mice per group were separately treated with 30 μg pGCR plus lipofectamine, 200 μL saline and 30 μg pG2 plus lipofectamine. We separately injected the control plasmid and hTERT plasmid and saline into three group mice every two days, and the total number of infections were 7 times. During these two weeks, we observed and recorded the mice's spirit, diet and activates. We also recorded the size of the xenografts.

**Immunohistochemistry**

Immunohistochemistry stain experiments according to the protocol for two-step detection. Paraffin sections of xenografts and other mice organ tissues were cut into 4 μM slices. Then the slides were deparaffinized in xylene and rehydrated using descending concentrations of ethanol. Following, we performed antigen retrieval using 0.01 mol/L of citrate buffer (pH=6.0) in a microwave. Endogenous peroxidase activity was blocked with 3% hydrogen peroxide for 10min at room temperature. After washing with PBS, the sections were incubated with blocked serum for 1h. Tissue sections were then stained with primary monoclonal rabbit antibody against \( p53, p21 \) (Santa Cruz). Tissue slides were incubated with primary antibody overnight at 4 °C. The horseradish peroxidase (HRP)-labeled goat anti-rabbit IgG treated the specimens at room temperature for about 30 min to 1 hour. Positive staining was visualized with DAB. The \( p53, p21 \), hTERT and apoptosis were evaluated by two pathologists blind record the results. The positive staining was counted in five separate specimens and statistical analysis was performed.

**H&E staining**

Xenograft tumors were immersed in 4% neutral formaldehyde and HE staining was completed according to previously described (12). According to the HE staining, Observe the cell phenotype in different groups. The remaining tumor tissue was stored in a −80 °C refrigerator.

**Terminal deoxynucleotidyl transferase mediated dUTP nick end labeling (TUNEL) assay of xenograft tissue sections**

Paraffin-embedded tissue sections were deparaffinized, washed and permeabilized as described above according to the instructions for the in situ cell death assay kit. After treatment with 3% H\(_2\)O\(_2\) treated 10 min at room temperature, we treated the slides with protease K at 37 °C
Translational Cancer Research, Vol 8, No 2 April 2019

569

tcr.amegroups.com

for 10 min. After preparing the TUNEL reaction reagents, add 30 μL TUNEL reaction buffer and incubate at 37 ℃ for 60 min. After washing three times with PBS, add DAB for staining reaction and hematoxylin staining the nuclear, neutral gum sealing pieces and obtained the photography by microscopy.

**Western blot analysis**

The xenograft tissues were lysed by RIPA buffer and protein content was measured using Enhanced BCA Protein Assay kit. The tissue lysates were then separated on a SDS-PAGE gels, transferred to a polyvinylidene fluoride (PVDF) membranes, and blocked with 5% non-fat milk for 2 hr at room temperature. PVDF membranes were incubated with primary antibodies (The following antibodies β-tubulin and hTERT were purchased from Santa Cruz). Then the membranes were incubated with BeyoECL plus, and detected by ChemiDocTM XRS+ imaging system. Data was measured using ImageJ 1.48 u software. All experiments were repeated at least three times with similar results.

**Statistical analysis**

For all quantified data, the mean ± SEM is presented. Statistical significance between the two experimental groups was characterized as an asterisk, and comparisons were made using Student’s t-test. P value<0.05 was considered significant.

**Results**

*Inject hTERT shRNA plasmids significantly decreased hTERT expression in A2780 and SKOV3 induced xenograft tumor*

We constructed the transplantation tumor model and studied the function of hTERT inhibiting tumors. In this study, we constructed the hTERT shRNA plasmid and injected it into xenograft tumors plus lipofectamine. After two weeks, we detected hTERT expression in tumor tissues. As shown in Figure 1A,B, hTERT was significantly decreased in tumors. We repeated the experiment three times and confirmed that hTERT shRNA worked well in xenograft tumors (Figure 1C).

**Knockdown expression of hTERT can significantly inhibit the ovarian tumor growth**

To test whether hTERT plays a role in the inhibition of ovarian cancer, we constructed a nude transplanted mice model by injection of human ovarian cancer cell A2780 or SKOV3 cells. After xenograft formation, we injected the hTERT plasmids for 7 times in two weeks. As shown in Figure 2A, the results showed that the size of transplanted tumor became small after knocking down hTERT expression. Comparing the tumor size at different times, we can see that the tumors was significantly reduced after 6 days of injection of the pG2 plasmid. However, there was no significant change in the pGCR group and the saline group (Figure 2B,C). Moreover, we detected the weight of xenograft tumors and found that the pG2 group which inhibit the hTERT can obviously decreased tumor weight (Figure 2D,E).

*bTERT affect the tumor cell apoptosis and necrosis*

To investigated the role of hTERT in tumor proliferation, we stained xenograft tissue with H&E. As shown in Figure 3A,B, the results showed that knockdown of hTERT...
hTERT downregulated expression significantly inhibits ovarian tumor growth. (A). Decreased expression of hTERT can obviously inhibit A2780 and SKOV3 xenograft tumor in nude mice model. (B,C) The tumor volume changes during the modifying hTERT gene expression in A2780 (B) and SKOV3 (C). (D,E) The ovarian tumor weight changes after down-regulated hTERT in A2780 or SKOV3 induced xenograft tumors (*, P<0.05; **, P<0.01). hTERT, human telomerase reverse transcriptase.

expression significantly increased tumor cell apoptosis and necrosis in A2780 and SKOV3 xenograft tumors. However, in the Saline and pGCR groups, tumor cells were phenotypically normal and well proliferated, the nuclear are clear and presented a large number of pathological split phases. After TUNEL staining, as shown in Figure 3C,D, the percentage of positive TUNEL staining in the pG2 group was as high as 58.47%±4.75%, while in the saline and pGCR control group, the percentage of TUNEL positive was only 2.40%±1.41% and 1.89%±1.65%.Significant increase in the pG2 group in A2780 xenograft tumors. However, in SKOV3 xenograft tumor which lack p53 gene expression also showed that increased TUNEL positive cells, though compared with A2780 pG2 group, it is lower. To investigate the details of cell death, we examined the cell microstructure by transmission electron microscope (Figure 3E) and found that the apoptotic bodies in A2780 pG2 group and the nuclear membrane was pyknotic, it is classic apoptosis characters. However, in SKOV3 xenografts, the cells are necrotic, there are no apoptotic bodies, and the cells are destroyed. Above all, the results indicated that hTERT affects apoptosis dependent on the p53 pathway and induces necrosis in the absence of p53.

bTERT affect senescence by p53 pathway

As we all known, p53 is a famous transcription factor, in addition to participating in the apoptosis pathway, p53 is also involved in the senescence pathway. Whether p53 affects cell aging in this mice model has not yet been known. To investigate this hypothesis, we stained the p53 downstream gene p21 and found that p21 was significantly increased after down-regulated hTERT expression in A2780 xenografts, whereas in p53 defected SKOV3 xenografts, there was no p21 expression changes (Figure 4). In the end, we concluded that the effect of hTERT on aging depends
Figure 3 Knockdown hTERT expression can promote cell necrosis and apoptosis. (A,B) HE staining of xenografted tumor tissues after different treatments; (C,D) TUNEL staining the xenograft tumors; (E) observation of cell phenotype by transmission electron microscopy (*, P<0.05). Magnification: 100x. hTERT, human telomerase reverse transcriptase.
on the p53 pathway.

**Discussion**

Most of the tumor tissues have been identified by hTERT expression, and more than 85% of cancer cells contain genetic aberrations and hTERT overexpression (13). In this study, our data analysis led us to believe that hTERT is also increased in the patients with ovarian cancer. In our mice model studies, we found that hTERT affects ovarian cancer cell apoptosis and aging dependent on the p53 pathway. In addition, down-regulation of hTERT can also induce necrosis of p53-deficient ovarian cancer cells. hTERT can be used for the diagnosis and prognosis for ovarian cancer in the future, or be able to identify the risk of tumor progression and establish the appropriate therapy.

Down-regulation of p53 and necrotic pathways is well known in several cancers (14,15). In addition to the p53 mutation in most cancers, it has also been downregulated in cancer cells. In support of this point, current studies indicate that there is a significant correlation between hTERT and p53 induced apoptosis and senescence. Moreover, hTERT also significantly regulates necrosis in p53-deficient cancer cells. Although previous studies have shown that hTERT is considered to be an important factor in the development of ovarian cancer, and the Pi3K/AKT/mTOR pathway is involved in the carcinogenesis of hTERT regulation (16). Although hTERT expression does not directly cause cancer, it can contribute to the long-life span of cancer cells and subsequently increased tumor formation (17). Although there is no evidence that hTERT affects tumor cell survival depends on the transcription factor p53. In this study, we constructed two mice xenograft tumor mice models and confirmed our hypothesis. Although we do not have more

---

**Figure 4** hTERT regulate transcription factor p53 and its downstream gene p21 expression. (A,B) Immunostaining the xenograft tumor tissue after different treatments using p53 antibody. (C,D) The expression of the p53 downstream gene p21 was analyzed by p21 antibody immunostaining (*, P<0.05). Magnification: 100x. hTERT, human telomerase reverse transcriptase.
evidence to support this mechanism or association between clinicopathological parameters and the individual expression of hTERT and p53. To date, data regarding the prognostic value of hTERT in ovarian cancer are inconclusive. In our study, we found that the therapeutic value of hTERT in affecting ovarian cancer cells survival.

**Conclusions**

We reported that the function of hTERT in ovarian xenograft tumor mice, the first data to document the therapeutic effect of down-regulated hTERT expression. Furthermore, high expression of hTERT in ovarian cancer may suggest that it may also be a potential prognostic marker of ovarian cancer.

**Acknowledgements**

We thank Miss Jing Chen (Chongqing Medical University) who provided purely technical help.

**Footnote**

*Conflicts of Interest:* The authors have no conflicts of interest to declare.

*Ethical Statement:* All animal experiments were conducted in accordance with the Guide for the Care and Use of Laboratory Animals and approved by the Animal Experimentation Ethics Committee of the Chongqing Medical University.

**References**


**Cite this article as:** Su C, Luo Y, Yang Y, Yi Y. ShRNA-mediated silencing of hTERT promote apoptosis and senescence in human ovarian cancer cells. Transl Cancer Res 2019;8(2):567-573. doi: 10.21037/tcr.2019.03.17