RESEARCH ARTICLE

siRNA-mediated Inhibition of hTERC Enhances Radiosensitivity of Cervical Cancer

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Abstract

Background: To investigate the influence of telomerase activity, apoptosis, radiosensitivity of cervical cancer after siRNA-mediated knockdown of telomerase RNA and evaluate in vivo growth with gene interference.

Methods: We studied siRNA-targeting-telomerase RNA transfection into the Hela cell line. Expression of hTERC mRNA was detected by RT-PCR and telomerase activity was measured by the TRAP assay. Growth inhibition was determined by MTT assay and radiosensitivity of the cervical cancer cells was examined by colony formation assay. In addition, effects of hTERC inhibition in vivo were studied by injection of siRNA-transfected Hela cells into nude mice.

Results: The hTERC siRNA effectively downregulated the expression of hTERC mRNA and also reduced the telomerase activity to 30% of the untreated control value. The viability of hTERC siRNA transfected Hela cells was reduced by 44.7% after transfection. After radiation treatment, the radiosensitivity of Hela cells with hTERC knockdown was increased. In vivo, the tumors developing from the hTERC siRNA-transfected cells were of reduced size, indicating that the hTERT siRNA also depressed the tumorigenic potential of the Hela cells.

Conclusions: Our results supported the concept of siRNA-mediated inhibition of telomerase mRNA which could inhibit the expression of hTERC and telomerase activity. Furthermore, radiosensitivity was upregulated after knockdown the hTERC in vivo and in vitro.

Keywords: siRNA-mediated inhibition - hTERC - cervical cancer

Introduction

Cervical cancer is one of the most common malignancies among females. Approximately, 250,000 women die each year from cervical cancer and 80% are from developing countries (Parkin et al., 2005; 2006). Ionizing radiation plays a key role in the treatment of cervical cancer but the frequent occurrence of radiation-resistant tumors is a common clinical problem.

Telomeres are nucleoprotein structures at the end of eukaryotic chromosomes that function to maintain genomic stability. In cells which are lack telomerase or with inadequate levels of telomerase activity, telomeres shorten with each cell division. This progressive eventually leads to a situation where cells stop dividing and enter into replicative senescence (Blackbum et al., 2005). As we know, telomeres is detected in approximately 90% of all malignant tumors (Heselmeyer et al., 2003), but not in normal cells. The special charactrictic of telomerase has made it not only was used as a predictive parameter of radiation but also be target of gene therapy which is widely reserched. Telomerase is comprised by telomerase RNA component, which is the template of telomere, the catalytic subunit hTERT, and telomerase-associated protein (TAP). In the past researches the human telomerase reverse transcriptase as the catalytic component of the telomerase activity regulation was highlighted. But more and more recent studies showed that both components are required for telomerase activity and telomere length in vitro (Feng et al., 1995; Cristofari et al., 2006). The telomase RNA is also important for telomere maintenance.

Radiation therapy is an important approach in cervical cancer therapy. But the clinical utility of radiotherapy are limited by characteristic radiation resistant. Increasing the sensitivity of tumor cells to radiotherapy would improve the outcome in patients with cervical cancer. Several studies have shown that there is a close relationship between the telomerase activity and the cellular response to irradiation. Ionizing radiation induces a variety of base alterations and DNA strand breaks, of which the unrepaired double-strand breaks (DSB) are the most lethal lesions leading to cell death. Slijepcevic study highlights the fact that the DNA DSB repair-associated proteins can maintenance of chromosomal integrity by balancing repair activities and telomere maintenance, in which telomerase is also involved (Slijepcevic et al., 2005).

There are several approaches to be used for blocking the telomerase activity such as pharmacological drugs chemical agents, ribozymes and antisense molecules. And RNA interference is an important method of inhibiting the gene express. RNAi is induced by double-stranded, 21- to 23-nucleotide RNAs that induced the sequence-
specific degradation of homologous RNAs. The strong and specific suppression of gene expression by RNAi is currently being evaluated as a potentially useful method for developing gene-silencing therapies for cancer (Tong et al., 2005; Zhang et al., 2006).

In the present study, to clarify more closely possible connections between telomerase and relative intrinsic radioresistance, we designed siRNA targeting telomerase RNA and delivered plasmids siRNA into the human Hela cell line. Those stably transfected Hela cell showed a significant decreased of expression of the telomerase mRNA and enhancement in radiosensitivity.

Materials and Methods

Cell cultures

The Hela cell line was kind gift from the Experiment Central of Second Affiliated Hospital of Harbin Medical University. The Hela cell line was grown in DMEM-HIGH Glu medium (Gibco, Paisley, Scotland, UK) supplemented with 10%calf serum and at 37 °C with 5% in a humidified incubator.

siRNA preparation

The siRNAs sequences for hTERC were designed by a commercial software based on the principle for siRNA. The siRNA-hTERC-I sense sequence is 5’- CGAAGGGTCCTTGGGAAT-3’. The siRNA-hTERC-II sense sequence is 5’- TGTCCTCATGGGTATGA-3’. The siRNA-hTERC-III sense sequence is 5’- GGAGCATGTTGAGGTCTAT-3’. And a negative control siRNA sequence is 5’- TTCTCCGAACGTGTCACGT-3’. The three sequences for hTERC were submitted to BLAST principle for siRNA. The siRNAs sequences for hTERC were designed by a commercial software based on the principle for siRNA. The siRNA-hTERC-I sense sequence is 5’- CGAAGGGTCCTTGGGAAT-3’. The siRNA-hTERC-II sense sequence is 5’- TGTCCTCATGGGTATGA-3’. The siRNA-hTERC-III sense sequence is 5’- GGAGCATGTTGAGGTCTAT-3’. And a negative control siRNA sequence is 5’- TTCTCCGAACGTGTCACGT-3’.

Cell Transfection

The Hela cell in logarithmic growth phase were chosen and then were plated at 2×10^5 in each well of six-well plates. Each well contained DMEM without antibiotics or FBS. After incubation for 24 h, the transfections were performed using Lipofectamine2000 transfection reagent (Invitrogen, USA) according to the manufacture’s guidebook. Transfection efficiency was valued by a fluorescence microscope.

Reverse Transcription and semiquantitative

Approximately 1×10^6 different kinds of cells including untransfected cells, the cells transfected with parental vector and the cells transfected with siRNA vector were harvested, respectively. TRIzol agent was used for total RNA extraction and then synthesized cDNA by an RT-PCR kit. The CDNA was added to a final volume of 20uL PCR reaction. The cycling program was performed as follow:94°C for three minutes; 94 °C for 40 s, 60 °C for 40 s and 72 °C for 90 s (35 cycles); and 72 °C for ten minutes. The primers sequence for hTERC are shown as follows: forward: 5'-CTGGGAGGGGTGTTGACCATT-3’, reverse:5’-CGAAGGGTCCTTGGGAAT-3’, 5'-CTGGGAGGGGTGTTGACCATT-3’. The β-actin was used as internal reference. The primer sequence were as follow: forward: 5’-CGGGAACTGTGCGTGAT-3’, reverse: 5’-CAAAGGGTGGAGGTGGTC-3’. The PCR products were visualized on agarose gel.

MTT assay

Five groups were set up. Cells in each groups plated at 103 in 6 wells of 96-well seeded and then incubated overnight. Each well was added to 20 uL of 5 mg/mL MTT and the absorption at 492 nm was detected on spectrometre. The figures were recorded every 24 h for 5 days.

Telomerase activity assay

Telomerase activity was examined by telomeric repeat amplification protocol (TRAP). In brief, the transfected cells were harvested and adjusted to a density of 2 × 10^5 cells and then were added to 200uL lysate. After incubated on ice for 30 minutes, the mixture was centrifuged at 12000 r/min and 4 °C for 25 minutes. The supernatant was transferred to a 1.5 mL DEPC-treated tube. For the TRAP reaction 2 uL of cell extraction was added to 25uL of reaction mixture and DEPC-treated water was added to a final volume of 50 uL. The extension reaction was performed at 30 °C for 30min and then denatured at 90°C for 3 min. CX Primer was then added. The mixture was denatured at 94 °C for 5min, and underwent 30 Cycles of 94 °C for 45 s, 50 °C for 45 s, and 72 °C for 90 s with a final extension reaction at 72 °C for 5 min. The A value of the product was measured after this reaction.

Clongenic survival assay

The cells were seeded in 60mm dishes for about 10 hours, then cells were treated with different dose (0Gy, 2Gy, 4Gy, 6Gy, 8Gy) of 6MV X-ray radiation by a 23EX accelerator. After cultured for 14 days, the clones were fixed and stained. The clones that more than 50 cells were counted. The plating efficiency and survival fraction were calculated. Sensitizer enhancement ratio was defined as the radiation does required to obtain a specific surviving fraction(SF).

Xenograft mice study

BALB/c nude mice were maintained in pathogen-conditions within the Laboratory Animal Resource Center at the Harbin Medical University. 5×10^3 Hela cells were injected into the left flanks, respectively, of 18 mice of 8-10 weeks. After the tumors reached a mean volume of 50 mm^3 ,the treatment was given. The 30 mice were divided into 6 groups, untreated, IR-alone, hTERC siRNAIII-alone, control siRNA-alone, IR+hTERC siRNAIII, IR+control siRNA. The mice in the IR-alone, IR+hTERC siRNAIII, IR+control siRNA groups were subjected to 6Gy irradiation. After received irradiation, the hTERC siRNAIII and control hTERC were injected in situ. The injection was repeated twice. The volume of the tumor sizes were subsequently measured every three days, the results represented the mean value of three independent
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Figure 1. Representative Photograph Showing the Uptake and Cellular Distribution Pattern of GFP-Conjugated siRNAs in Hela Cells, as Detected by Fluorescence Microscopy (×400)

Figure 2. Levels of Expression of hTERT mRNA in Hela Cells after Transfection with a Plasmid Encoding hTERC siRNA for 48 h. (A) Semiquantitative RT-PCR for hTERC. Expression of β-actin was used as an internal control. The data are representative of 3 independent experiments. (B) Determination of hTERT mRNA levels after the treatments with hTERC siRNA. Data are shown as mean ± SD from three independent experiments (*p<0.05 compared with the control mean values)

Statistics analysis

With SPSS software, the results were presented as means±SD. Statistical analysis was done using one way analysis of variance (ANOVA) for multiple samples and Student’s t-test for comparing paired sample sets. Statistical significance was considered at P value less than 0.05.

Results

siRNA are efficiently internalized.

After transfection for 24 hours, the transfected Hela cells were observed by fluorescence microscopy (Figure 1). The transfection efficiency was more than 65%.

The expression of hTERT mRNA were downregulation in Hela cells

We examined the changes in expression of hTERT after the treatments (Figure 2). Transfection with a plasmid encoding hTERC siRNAII and hTERC siRNAIII resulted in marked downregulation of cognate mRNA in Hela cell. Compared with blank group, the expression of Hela cell transfected with siRNAII and siRNAIII was inhibited which was 59.1% and 44.6% respectively (P<0.05). But hTERT mRNA levels were not altered after transfection with hTERC scrambled siRNA and hTERC siRNAI. The expression of β-actin was used as an internal control for the mRNA analyses. The siRNA-hTERC-III vector had the strongest suppressive effect on hTERC mRNA expression.

Marked reduction in cell viability

We observed that the cell viability had different degrees of reduction in the Hela cell which were treated with the hTERC siRNAII and siRNAIII, while the viability of transfexion with hTERC siRNAI had not change compared with the control group. And the hTERC siRNAIII vector had the strongest suppressive.

The telomerase activity of Hela cells with hTERC siRNA are significant decrease

Telomerase activity was detected by TRAP-PCR kit and the A value was 1.907 ± 0.065, 1.917 ± 0.032, 0.87 ± 0.074, respectively. The ratio of inhibition of telomerase activity in Hela cells exposed to the hTERC siRNAIII was 54.4% (p<0.05). And telomerase activity of Hela cell with hTERT siRNAIII treatment was marked declined.
Many researches suggested that telomerase activity closely correlated with level of hTERT transcription in most malignancy include breast cancer, malignant glioma, meningioma, cervical cancer. There have been reports which demonstrated that RNAi against telomerase reverse transcriptase (hTERT), could successfully inhibit telomerase activity in several cancer cell lines (Mo et al., 2003; Natarajan et al., 2004; Nakamura et al., 2005; Souza et al., 2006; Zhang et al., 2006; Gandellini et al., 2007) and can enhance these radiosensitivity and chemosensitivity (Dong et al., 2009; George et al., 2010; Meng et al., 2012). However, the role of hTERC was not attracted enough attention as other target for telomerase activity regulation. hTERC gene encodes for an RNA unit of telomerase that maintains the length of telomeres through cellular divisions. When this gene is overexpressed, the cell avoid undergoing apoptosis, potentially leading to tumorigenesis. Several studies over years from biochemistry, promoter studies and mouse models has proved that the hTERC has esssential contribution in telomere maintain and telomerase activity (Ren et al., 2003; Caimey et al., 2008). It is well known that a high number of chromosomal alterations have been described related to dysplastic lesions of the uterine cervix. Among them, the most frequent are 3q gains, specifically those involving 3q26 chromosome bands, where human telomerase RNA gene (hTERC) is located (Alamed et al., 2009; Zheng et al., 2010; Gao et al., 2011). In the present study, we designed three siRNAs target at hTERC, after transfecting Hela cells respectively, we detected that the expression of telomerase RNA reduced more or less, and the most powerful siRNA directed against hTERC resulted in more than 70% suppression at the mRNA levels. The decrease in the cell viability and proliferation could be responsible for the downregulation of the hTERC. Furthermore, we noticed that the various siRNAs has different suppression efficiency on mRNA expression. This date show that effectiveness of siRNA inhibition may be highly sequence-dependent.

Cervical cancer is the second most common female cancer worldwide and its main therapeutic method is radiotherapy. Increasing the sensitivity of tumor cells to radiotherapy would improve outcome in patients with cervical cancer. Ionizing radiation can induce many base alteration and the most lethal lesions leading to cell death is the unrepaired double-strand breaks. The effectiveness of ionizing radiation is closely related to the dose dependent capacity of the ionizing radiation to cause DNA damages and the capability of cells to repair the damages. Many recent studies show that the telomerase involved in the progress of the DNA DSB repair. And several studies also supports that the telomerase activity and the telomere status have impacts on the response to ionizing radiation (Aravindan et al., 2011). Current literature indicates that telomerase has a radiation-inducible function lead to resistance of cells to irradiation. Ogawa et al. (1998) reported that the advanced cancers of oral cavity and oropharynx with higher telomerase activity were not sensitive to radiation therapy but the other cancers with lower telomerase activity were sensitive to radiation therapy. Rubio and colleagues (Rubio et al., 2004) reported that high telomerase activity provided...
resistance to irradiation in human fibroblasts with short telomeres, and this resistance was dependent on the ability of telomerase to maintain telomere functionality. In our study the Hela cells transfected by siRNA, the telomerase activity was significantly reduced while the radiosensitivity was upregulation. This is consistent with previous studies. The in vivo tumor growth experiment also demonstrated impaired growth of the tumors formed by the hTERC siRNA transfected Hela cells.

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