

Lysophosphatidic Acid Activates Telomerase in Ovarian Cancer Cells Through Hypoxia–Inducible Factor– 1α and the PI3K Pathway

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ABSTRACT

Telomerase is reactivated in over 90% of tumors and plays critical roles in tumor progression. The mechanisms by which telomerase is upregulated in cancer cells are poorly understood. Here we showed that a bioactive lipid, lysophophatidic acid (LPA), up-regulated the expression of human telomerase reverse transcriptase (hTERT) and telomerase activity in serous ovarian adenocarcinoma cell lines SKOV3, A2780, and HEY, but not in OCC1, a clear cell ovarian cancer cell line. This cell type specific effect of LPA on telomerase regulation may reflect distinctive genetic backgrounds in different histological subtype of ovarian cancer cells. Our data further suggest that the phosphatidylinositol 3-phosphate kinase (PI3K) pathway and hypoxia-inducible factor-1 α (HIF-1 α) are likely to be involved in LPA-induced hTERT expression. Targeting human telomerase by LPA is potentially involved in its role of promoting tumor progression. J. Cell. Biochem. 105: 1194–1201, 2008. © 2008 Wiley-Liss, Inc.

KEY WORDS: LPA; hTERT; TELOMERASE; HIF-1α; PI3K/Akt

elomerase is an RNA-dependent DNA polymerase that synthesizes telomeric DNA and thus counteracts the "endreplication" problem occurring during semi-conservative DNA replication [Shay and Wright, 2005]. Telomerase comprises two essential components: the RNA component (hTR) that serves as a template for telomeric DNA synthesis and the catalytic protein with the human telomerase reverse transcriptase activity (hTERT) that adds the telomeric repeats onto the end of chromosome [Collins, 2006; Stewart and Weinberg, 2006]. The telomerase activity is turned off during embryonic development by transcriptional repression of *hTERT*, which is undetectable in most normal human somatic cells but present in over 90% of cancerous cells [Shay and Bacchetti, 1997]. Telomerase is required for the long-term proliferation potential of human stem cells and cancer cells. Earlier studies have demonstrated that ectopic expression of hTERT in normal human cells leads to extension of life-span and immortalization of many cell types [Bodnar et al., 1998; Harley, 2002]. Inhibition of telomerase in telomerase positive cancer cells induces

cell death [Hahn et al., 1999a; Herbert et al., 1999; Zhang et al., 1999]. Moreover, it has been shown that expression of telomerase was required for the oncogenic transformation of several normal cell types [Hahn et al., 1999b; Drayton et al., 2003; Zongaro et al., 2005; Mizumoto et al., 2006]. Together, these observations support the fundamental roles of telomerase in tumorigenesis. The molecular mechanisms controlling telomerase up-regulation during cellular immortalization and tumorgenesis are beginning to be elucidated [Cong et al., 2002; Dong et al., 2005; Dwyer et al., 2007].

In ovarian tissues, telomerase is absent in normal ovarian surface epithelium and premalignant lesions, but up-regulated in over 95% of ovarian carcinomas [Shay and Bacchetti, 1997]. Concurrently, lysophophatidic acid (LPA), a bioactive lipid growth factor for ovarian cancer, is elevated in both plasma and ascites of ovarian cancer patients and promotes cancer cell growth, migration, invasion, and tumor metastasis [Xu et al., 1998, 2003; Ren et al., 2006]. LPA exhibits pleiotrophic biological activities via the interaction with specific G protein-coupled receptors, LPA₁/Edg-2,

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LPA₂/Edg-4, and LPA₃/Edg-7 [Moolenaar et al., 2004]. It has been shown that LPA promotes ovarian cancer cell proliferation by directly increasing the level of cyclin D1 [Hu et al., 2003], upregulating VEGF, IL-6, and IL-8 expression, which facilitate tumor angiogenesis [Schwartz et al., 2001; Sivashanmugam et al., 2004; Chou et al., 2005; Sako et al., 2006]. In addition, LPA regulates matrix metalloproteinase and urokinase plasminogen activator activity in ovarian cancer cells, which contributes directly to cell invasion and may be involved in ovarian cancer metastasis [Pustilnik et al., 1999; Fishman et al., 2001].

A number of factors and pathways have been implicated in the regulation of hTERT expression and telomerase activity [Cong et al., 2002]. However, the regulation of telomerase expression and activity under pathological conditions is poorly understood. HIF-1 is a heterodimeric transcriptional factor composed of the basic helixloop-helix-PAS protein HIF-1a and the arylhydrocarbon receptor nuclear translocator (ARNT or HIF-1β), play a critical role in tumor progression and angiogenesis by activation of hypoxia-inducible genes expression. HIF-1 α is highly regulated, whereas HIF-1 β is constitutively expressed [Semenza, 2003]. It has been reported that the minimum promoter of hTERT contains two HIF-1 responsive elements (HRE) required for *hTERT* up-regulation by HIF-1 α in hypoxia condition [Nishi et al., 2004]. However, the mechanism of telomerase regulation by HIF-1 α has not been fully elucidated, especially, the role of HIF-1 α in activation of telomerase during tumorigenesis is not known.

Telomerase and LPA have been implicated in tumorigenesis, particularly in ovarian cancer [Xu et al., 1995; Fang et al., 2002; Sengupta et al., 2007], with both of them were elevated in ovarian cancer cells, when compared to normal ovarian surface epithelium cells. More recently, Bermudez et al. [2007] have shown that LPA up-regulates telomerase activity via a VEGF-dependent pathway and identified Sp-1 binding sites in hTERT promoter region for LPA's activity. In this study, we have investigated the regulation of telomerase expression by LPA in additional ovarian cancer cell lines and elucidated the new signaling molecules (phosphatidylinositol 3-phosphate kinase–PI3K and HIF-1 α) involved in LPA-induced up-regulation of *hTERT* expression.

MATERIALS AND METHODS

REAGENTS AND ANTIBODIES

LPA was purchased from Avanti Polar Lipids (Alabaster, AL). LY294002, PD98059, and rapamycin were purchased from Calbiochem (San Diego, CA). Resveratrol was purchased from Sigma-Aldrich (St. Louis, MO). β -actin monoclonal antibody was from Cell Signaling (Beverly, CA) and HIF-1 α monoclonal antibody was from BD (San Diego, CA). Total Akt and Phospho-Akt antibodies were obtained from (Cell Signaling). LipofectAMINE 2000 was obtained from Invitrogen (Carlsbad, CA).

CELL CULTURE

Ovarian cancer cell lines were maintained in RPMI 1640 (Hyclone, Logan) supplemented with 10% fetal bovine serum (Hyclone). For LPA treatment, cells were treated with 40 μ M LPA dissolved in phosphate-buffered saline (PBS) or treated with PBS as control. For

pharmacologic inhibitor treatment, cells were pretreated with inhibitors 10 mM LY294002, 100 μ M PD98059, 10 mM rapamycin or 50 μ M resveratrol dissolved in DMSO for 2 h prior to LPA treatment.

PLASMIDS

p3996del, p1009del, and p342 are *hTERT* promoter-luciferase reporter plasmids in which various lengths of DNA upstream of the initiating ATG codon of hTERT were cloned into pGL2-Enhancer (Promega, Madison, WI). Reporter plasmids p342mt1, p342mt2, and p342mt contain one or two mutations in the two putative binding sites of HIF-1 α at nt -241 and -33, in which CACGTGGC were changed into TTTGTGGC [Nishi et al., 2004]. Site-directed mutagenesis was performed using the plasmid p342 with the following primer pairs: p342mt1, forward, 5'-CAG GAC CGC GCT TCC TTT GTG GCG GAG GGA CTG and reverse, 5'-CAG TCC CTC CGC CAC AAA GGA AGC GCG GTC CTG; p342mt2, forward, 5'-CTG CGT CCT GCT GCG TTT GTG GGA AGC CCT GGC and reverse, 5'-GCC AGG GCT TCC CAC AAA CGC AGC AGG ACG CAG. The mutated sequences in the p342 were confirmed by DNA sequencing and then subcloned into KpnI and BglII sites of pGL2 enhancer plasmid to generate p342mt1, p342mt2, and p342mt.

ASSAY OF TELOMERASE ACTIVITY

Telomerase activity was assayed using *TeloTAGGG* Telomerase PCR–ELISA^{PLUS} kit (Roche, Indianapolis, IN) according to the manufacturer's protocol. To perform the telomerase PCR–ELISA within a linear range, all of cell extracts equivalent to 1,000 cells were used. Absorbance values were measured using the Tecan Genios Microplate Reader and telomerase activity was expressed as relative telomerase activity against negative control.

RT-PCR

Total RNA was collected using TRIzol reagent (Invitrogen). 2 μ g of DNase I (Ambion, Austin) treated RNAs were reverse transcribed to single-strand cDNA using M-MLV Reverse Transcriptase (Promega) with 1 μ g oligo(dT). Primer sequence of *GAPDH*, *hTERT*, and *HIF-1* α are the following: *GAPDH* forward primer 5-CGGAGTCAACGG-ATTTGGTCGTAT-3, *GAPDH* reverse primer 5-TGCTAAGCAGTTG-GTGGTGCAGGA-3; *hTERT* forward primer 5-AGAACGTTCCGC-AGAGAAAA-3, *hTERT* reverse primer 5-ATGTACGGCTGGA-GGTCTGT-3; *HIF-1* α forward primer 5-TCCATGTGACCATGAGG-AAA-3, *HIF-1* α reverse primer 5-TCCATGTGACCATGAGGAAA-3.

WESTERN BLOT AND IMMUNOFLUERSCENCE STAINING ANALYSIS

Cell lysates were prepared in RIPA lysis buffer (50 mM Tris–HCl, pH 8.0, 150 mM NaCl, 1% NP-40, 0.5% Sodium Deoxycholate, 0.1% SDS). Protein concentrations of cell lysates were determined using BCA assay kit in accordance to the manufacturer's instructions. One hundred micrograms of protein extracts were resolved on 12% SDS–PAGE and transferred to PVDF membranes (Bio–Rad, Hercules, CA). Immunoblotting was performed using HIF-1 α antibodies (1:500), serine 473-phospho-Akt antibody (1:1,000), and total Akt antibody (1:1,000). β -actin (1:5,000, Sigma–Aldrich) was used as loading control. For immunofluorescence staining analysis, cells were grown on coverslips and treated with ±LPA for indicated time

periods. The coverslips were fixed with 4% cold ultrapure formaldehyde (Merck, Germany) and permeabilized with 0.5% Triton X-100 (Amresco) for 5 min. Then, the coverslips were blocked in 0.6% goat serum in PBS for 1 h and incubated with primary HIF-1 α antibody (1:20) and secondary fluorescent antibodies (1:200) for 1 h at 37 °C respectively. After washed with PBS, coverslips were counterstained with 0.1 µg/ml 4,6-diamino-2-phenylindole (DAPI, Roche, Switzerland). Images are obtained with a laser scanning confocal microscope.

LUCIFERASE REPORTER ASSAY

HEY cells (6 × 10⁴) were plated in 24-well culture dishes and transiently co-transfected with 0.7 μg of luciferase reporter and 0.1 μg of pCMV-β-gal plasmid using LipofectamineTM 2000 reagent (Invitrogen). Cultures were replenished with medium containing 10% serum at 4 h post-transfection. Prior to collection, HEY cells were treated with or without LPA for 24 h. Luciferase activity was measured at 48 h post-transfection using the Luciferase Assay System (Promega) according to the manufacturers' instructions. Transcriptional activity was expressed as relative luciferase activity after normalization with β-galactosidase activity.

STATISTICAL ANALYSIS

Data were subjected to the Student's *t*-test for determination of statistical significance. P < 0.05 was considered significant.

RESULTS

LPA UP-REGULATED hTERT EXPRESSION IN OVARIAN CANCER CELLS

To further understand the role of LPA in regulating hTERT expression in more commonly used ovarian cancer cell lines, we tested the effect of LPA on *hTERT* mRNA levels in four ovarian cancer cell lines using RT-PCR. We found that LPA significantly up-regulated *hTERT* mRNA in ovarian cancer cell lines SKOV3, A2780, and HEY (all serous adnocarcinoma cell lines), but not in OCC1, a clear cell ovarian cancer cell line (Fig. 1A). LPA signaling is mainly mediated by its specific G-protein-coupled receptors [Moolenaar et al., 2004], we have found that all these four ovarian cancer cells lines expressed LPA receptors LPA₁, LPA₂, and LPA₃ (data not shown), suggesting that factor(s) other than LPA receptors is responsible for the lack of *hTERT* mRNA up-regulation in OCC1 cells.

The time course of LPA-induced telomerase expression was further examined in HEY, a representative ovarian serous adenocarcinoma cell line. We observed that LPA induced *hTERT* RNA expression, but not telomerase RNA (*hTR*), in a time dependent manner. Induction of *hTERT* transcription reached the highest level at 24 h and decreased after 36 h of LPA treatment (Fig. 1B). Correlated with the induction of *hTERT* mRNA level, telomerase activity was increased to maximum at 24 h and decreased after 36 h of LPA treatment (Fig. 1C).



Fig. 1. LPA induced *hTERT* mRNA expression and telomerase activity. A: LPA up-regulated *hTERT* transcription in different ovarian cancer cell lines. The cells were treated with or without LPA for 24 h. Total RNA was isolated and RT-PCR was conducted using specific *hTERT* primers. B: LPA up-regulated *hTERT* transcription in time-dependent manner. The HEY cells (1×10^6) were treated with LPA for the indicated time periods. Total RNA was isolated, and RT-PCR was conducted using specific *hTERT* primers. C: Telomerase activity was measured in cultured HEY cells treated with LPA for the indicated time periods. Results are the mean \pm SD (standard deviation) from three independent experiments in triplicates. Columns, mean; bars, SD; * denoted statistical significance, P < 0.05.

ACTIVATION OF hTERT PROMOTER ACTIVITY BY LPA REQUIRED HIF-1 α

In an effort to understand the mechanisms by which LPA induced *hTERT* expression, we characterized the effect of LPA on the *hTERT* promoter activity in ovarian cancer cells. We observed that LPA was able to stimulate the luciferase activity using *hTERT* promoters containing the full-length 3996 bp, the 1009 bp, or the 342 bp sequences upstream from ATG (Fig. 2A,B), suggesting that the 342 bp sequence contains the responsive element(s) for LPA-induced *hTERT* up-regulation. It has been reported that the minimum promoter of *hTERT* up-regulation by HIF-1 α under hypoxia conditions [Nishi et al., 2004], and LPA induces VEGF expression through activation of HIF-1 α in ovarian cancer cells [Lee et al., 2006]. We tested whether LPA induced *hTERT* transcription via HIF-1 α . There are two hypoxia-responsive elements (HREs) in the 342 promoter region. We mutated each of these sites separately or in





combination [Nishi et al., 2004]. We observed that mutations in each or both of these HRE sites abolished the induction of *hTERT* promoter activity by LPA (Fig. 2C,D), suggesting that both of these sites are required for LPA's effect. We next examined the effects of LPA on HIF-1 α expression. HEY cells were treated by LPA for different times as indicated in Figure 3 and expression of HIF-1 α was examined by RT-PCR and Western blot analyses. Interestingly, LPA did not alter the mRNA levels of *HIF-1\alpha* (Fig. 3A), but upregulated HIF-1 α protein levels from 30 min to 12 h (Fig. 3B), suggesting post-transcriptional regulation is involved. In addition, parallel to increased HIF-1 α protein expression, we observed that HIF-1 α was translocated to nucleus 4 h post of LPA treatment (Fig. 3C). Together, these observations suggest that HIF1 α is likely to be required for LPA's transcription activity on *hTERT*.

It has been shown that resveratrol (3,5,4'-trihydroxystilbene) produced by plants inhibited HIF-1 α expression by promoting its degradation [Park et al., 2007]. To further confirm the role of HIF-1 α in LPA-induced *hTERT* up-regulation, we examined the effects of HIF-1 α inhibitor resveratrol on LPA-induced *hTERT* up-regulation. The ovarian cancer cell lines SKOV3, A2780, HEY, and OCC1 were pretreated by resveratrol for 2 h, followed by LPA treatment. As shown in Figure 3D,E, LPA significantly increased the levels of HIF-1 α protein, and resveratrol suppressed HIF-1 α accumulation induced by LPA in the three serous ovarian cancer cell lines SKOV3, A2780, and HEY. Concomitantly, *hTERT* mRNA levels

have increased by the treatment of LPA, and this induction was suppressed by the HIF-1 α inhibitor resveratrol. However, no significant changes in either HIF-1 α protein levels or *hTERT* expression in respond to LPA have been detected in OCC1 cells, a clear cell ovarian cancer cell line. The differences of LPA responsiveness in these cell lines may be related to the distinctive cellular contexts in different histological subtypes of ovarian cancer cells [Schwartz et al., 2001].

PI3K WAS INVOLVED IN LPA INDUCED hTERT EXPRESSION

Both PI3K and mitogen-activated protein kinase (MAPK) signaling pathway have been implicated in the regulation of HIF-1 α expression [Zhou et al., 2004]. The PI3K pathway regulates HIF-1 α protein level via mammalian target of rapamycin (mTOR)dependent and -independent mechanisms [Pore et al., 2006]. To investigate the signaling pathway mediated HIF-1 α expression by LPA in HEY ovarian cancer cells, we examined the effects of PI3K, MAPK, and mTOR inhibitors on LPA induced *hTERT* expression. HEY cells were pretreated by different inhibitors and *hTERT* expression was monitored by RT-PCR. As shown in Figure 4A, LPA-induced *hTERT* expression was completely attenuated by the PI3K inhibitor LY294002, but only slightly affected by the MAPK inhibitors PD98059 and the mTOR inhibitor rapamycin, suggesting that PI3K, but not MAPK or mTOR, was mainly involved in LPA induced *hTERT* expression in HEY cells.



Fig. 3. LPA induced HIF-1 α over-expression and nuclear translocation. A: effect of LPA on *HIF-1* α transcription. HEY cells were treated by LPA for indicated time periods. Total RNA was isolated and *HIF-1* α transcription was analyzed by RT-PCR with *HIF-1* α specific primers. B: effect of LPA on HIF-1 α protein expression. HEY cells were incubated in the presence of LPA for the indicated time periods. HIF-1 α protein level was determined by immunoblotting with anti-HIF-1 α antibody. C: nuclear translocation of HIF-1 α induced by LPA. The cells seeded in slides were treated with or without LPA (40 μ M) for indicated time periods and cellular distribution of HIF-1 α was analyzed by immunofluorescence staining with anti-HIF-1 α antibody. D: resveratrol inhibited LPA-induced HIF-1 α expression. Cells were incubated with resveratrol (50 μ M) or PBS as control for 2 h prior to LPA treatment. Cells were collected after 4 h of LPA treatment and cell lysates were subjected to Western blotting analysis with the HIF-1 α antibody, or with β -actin antibody for loading control. E: resveratrol blocked LPA-induced *hTERT* expression. Cells were quantified related to β -actin or *GAPDH* respectively by densitometric analysis (band intensity) in B,D,E, representative of three experiments with similar results. Columns, mean; bars, SD.

Akt is known as an important downstream target for PI3K and commonly activated in human cancer. There were a number of studies indicated that PI3K/Akt pathway involved in the regulation of HIF-1 α [Pore et al., 2006]. We characterized the phosphorylation of Akt S473 in response to LPA. Consistent with previous report

[Baudhuin et al., 2002], we found that LPA induced Akt S473 phosphorylation (LPA 40 μ M and 30 min) in HEY ovarian cancer cells. Phosphorylation of Akt S473 was persistent and then decreased to the basal level 4 h after LPA treatment (Fig. 4B). The PI3K inhibitor LY294002 efficiently blocked Akt S473



Fig. 4. PI3K pathway was involved in LPA induced HIF-1a over-expression and hTERT transactivation. A: effect of pharmacologic inhibitors on LPAinduced hTERT transcription. Cells were pretreated for 2 h with vehicle or inhibitors and then exposed to LPA for 24 h. hTERT and GAPDH expression were analyzed by RT-PCR, GAPDH was used as a loading control. B: effect of LPA on Akt phosphorylation. HEY cells were incubated by LPA for indicated time periods. Serine 473-phospho-Akt and total-Akt protein levels were analyzed by Western blotting with specific antibodies. C: effect of pharmacologic inhibitors on LPA induced Akt phosphorylation. Cells were pretreated for 2 h with vehicle or inhibitors and then exposed to LPA for 30 min, Serine 473-phospho-Akt and total-Akt protein levels were analyzed by Western blotting with specific antibodies. D: effect of pharmacologic inhibitors on LPA induced HIF-1a protein over-expression. Cells were pretreated for 2 h with vehicle or inhibitors and then exposed to LPA for 4 h, HIF-1 α protein expression was analyzed by Western blotting with HIF-1 a specific antibodies. Representative of three experiments with similar results.

phosphorylation and HIF-1 α accumulation in response to LPA (Fig. 4C,D). Interestingly, although rapamycin did not affect *hTERT* expression level (Fig. 4A), it did moderately reduce pS473-Akt and HIF-1 α levels (Fig. 4 C,D), suggesting that elevated HIF-1 α may not always correlate with *hTERT* expression.

Collectively, our results suggested that LPA enhanced HIF-1 α expression and nuclear translocation through the PI3K pathway, thereby induced *hTERT* expression and telomerase activity in ovarian cancer cells.

DISCUSSIONS

The importance of telomerase in tumorigenesis has been fully recognized [Shay and Bacchetti, 1997; Hahn et al., 1999b; Shay and

Wright, 2005]. Thus, to understand how telomerase is up-regulated in cancer cells becomes very critical for developing new therapeutic strategies for cancer treatment. Similarly, the role of LPA in ovarian cancer has been extensively studied and its therapeutic potential has been realized [Xu et al., 1998, 2003; Mills and Moolenaar, 2003]. To link these two important molecules in ovarian cancer is very important and significant. We have provided convincing data to show that LPA regulates expression and activity of telomerase in multiple ovarian cancer cells lines, suggesting telomerase is an important molecule though which LPA exerts its oncogenic effects.

Accumulated information suggests that telomerase activity can be regulated at multiple levels by different factors in different cellular context [Cong et al., 2002]. However, the molecular mechanisms and pathways governing telomerase expression and activity are still elusive. Bermudez et al. [2007] has shown in ovarian carcinoma cell lines PA-1, SW626, and a telomerase-negative, nontumorigenic SV40 large-T antigen transfected human ovarian surface epitithelial (IOSE) cell line, that LPA regulates telomerase via a transcriptional factor Sp1 and the ERK1/2 pathway. In this report, we have shown that LPA up-regulated hTERT expression and telomerase activity in HEY, SKOV3, and A2780 cells through activation of HIF-1 α and PI3K. MAPK signaling pathway may not be involved in HEY cells, since the inhibitor of MAPK did not have an effect on LPA-induced hTERT expression. These results suggest LPA may employ distinct signaling pathways in different ovarian cancer cell lines. Since PI3K is an upstream activator of the MAPK signaling pathway in many different ovarian cancer cells, targeting PI3K may be more effective to reach the therapeutic effects in different ovarian cancer.

Ovarian cancer is a highly lethal type of gynecological disease. Based on the type of epithelial differentiation and morphological criteria, there are four major subtypes including serous, mucinous, endometrioid, clear cell. Serous tumors are the most common subtype representing about one-half of all ovarian cancer, the endometrioid adenocarcinomas account for 20-25% of ovarian cancer, whereas clear cell and mucinous adenocarcinomas are less common, which account for less than 10% of ovarian cancer cases [Schwartz et al., 2001; Soslow, 2008]. Recent studies indicate that the different histological types of ovarian cancer likely represent distinct disease entities [Feeley and Wells, 2001]. In support with this notion, it has been shown that clear cell ovarian cancer has a distinctive pattern of gene expression, which distinguishes it from other type of ovarian cancer [Schwartz et al., 2001]. In this report, we showed that LPA induced hTERT expression in ovarian serous adenocarcinomas cell lines SKOV3, A2870, and HEY, but not in OCC1, a clear cell ovarian cancer cell line. Consistently, we showed that LPA increased HIF-1 α expression, this induction was blocked by HIF-1 α inhibitor Resveratrol, which subsequently suppressed LPA-induced hTERT up-expression in ovarian serous adenocarcinomas cell lines SKOV3, A2870, and HEY. Although the effects of resveratrol on hTERT expression and cell proliferation remains to be investigated in ovarian cancer cells, resveratrol efficiently inhibited HIF-1a expression and subsequently blocked hTERT up-regulation induced by LPA, suggesting its potential therapeutic values for ovarian cancer treatment. Interestingly, we found that LPA has no effect on the expression of HIF-1 α and hTERT in the clear cell

ovarian cancer cell line OCC1. This different effect of LPA on telomerase regulation may be associated with distinctive genetic backgrounds in different histological subtype of ovarian cancer cells. Recently, Pectasides et al. [2006] have reviewed the clinical feature of patients with ovarian clear cell carcinomas, they have found that the recurrences are more frequent and survival rates are lower in patients with clear cell carcinomas than those with advanced serous subtype of tumors, suggesting that the intrinsic chemoresistance may be associated with this subtype and novel treatment approaches should be adopted in clinical management for patients with ovarian clear cell carcinomas. Therefore the classification of histological subtype of ovarian carcinomas may have important biological and therapeutic implications.

It is of interesting to note that the *hTERT* promoter activity induced by LPA was decreased to the same degree when either one or two of the HIF-1 α binding sites were mutated, suggesting that both of these sites are required for the activity and the two sites may have a cooperative effect. This is consistent with the report showing that transactivation of *hTERT* promoter by HIF-1 α was abolished when one or two of HIF-1 binding sites were altered [Nishi et al., 2004]. In addition, we observed in the *hTERT* promoter reporter assay that the minimum promoter p342 has greater promoter activity than the fulllength promoter p3996del or p1009del, suggesting that a negative regulatory element(s) may exist upstream of the minimum promoter.

We have shown that LPA not only increased expression of HIF-1 α , but also its nuclear translocation. HIF-1 α is mainly regulated post-transcriptionally through the von Hippel-Lindau protein (VHL) mediated ubiquitin-proteasome degradation [Ke and Costa, 2006]. We have shown that LPA increases the levels of HIF-1a protein rather rapidly (which 30 min), but not the RNA level. These data are fully consistent with post-transcriptional regulation of HIF-1a. HIF-1 α is critical transcription factor for a number of important cancer-related genes in various cancer cells [Bárdos and Ashcroft, 2005]. Thus, its nuclear translocation is very important to its function. We have shown that LPA-induced cell migration and invasion are significantly enhanced under hypoxia conditions, where HIF-1 α plays an important role [Kim et al., 2006]. The current work has added an another link between LPA and HIF-1α, suggesting that increased cellular responses to LPA under hypoxia conditions may be related to LPA's ability to up-regulate expression of HIF-1 α and increase its nuclear translocation. While this remains to be tested directly, a network among LPA-PI3K-HIF-1a -telemerase has been emerging through our work, which may represents targets for ovarian cancer therapy.

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