Suppressed protein expression of the death-associated protein kinase enhances 5-fluorouracil-sensitivity but not etoposide-sensitivity in human endometrial adenocarcinoma cells

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Abstract. Targeted knockdown of the death-associated protein kinase (DAPK) expression in the endometrial adenocarcinoma HHUA cells reportedly induces cell death by enhancing the tumor necrosis factor-related apoptosis-inducing ligand (TRAIL)-mediated apoptosis in an autocrine/paracrine manner. This suggests that endogenous DAPK is a potential candidate for a molecularly targeted anticancer therapy for patients with endometrial adenocarcinoma. To investigate the role of endogenous DAPK in anticancer drug sensitivity, we examined effects on cellular anticancer drug sensitivities of transfections with 5 different specific DAPK small-interfering RNAs (siRNAs) into HHUA cells. DAPK siRNA transfections strongly enhanced 5-fluorouracil (5FU)-sensitivity, but not etoposide-sensitivity, of HHUA cells compared with control siRNA-transfected cells. These results indicate that etoposide-stimulated cell death signals may share or include TRAIL-mediated apoptotic signals, and that 5FU-stimulated cell death signals may be independent from TRAIL-mediated apoptotic signals induced by DAPK siRNA transfections. Moreover, 5FU-combined chemotherapy with DAPK siRNA transfection may show stronger anticancer effects on patients with endometrial adenocarcinoma than does chemotherapy alone.

Introduction

The death-associated protein kinase (DAPK) cDNA was isolated as a positive mediator of apoptosis triggered by IFN-γ from human cervical carcinoma cells (1). Investigations have revealed that DAPK functions as a Ca2+/calmodulin-dependent serine/threonine kinase to regulate cell death or cell survival (1-14). However, DAPK’s physiological functions have not been fully clarified. Loss of DAPK expression has been implicated in tumorigenesis and metastasis (3,4,13), thereby suggesting a crucial role for DAPK in the apoptotic process under pathological conditions. On the other hand, inhibition of DAPK expression in HeLa cells, 3T3 fibroblasts and primary human vascular smooth muscle cells with an antisense DAPK was found to increase apoptosis (6,12).

In a human cervical squamous cell carcinoma cell line ME180, DAPK protein expression is constitutively suppressed but can be strongly induced by treatment with a demethylation agent, 5-aza-2′-deoxycytidine (5-aza-CdR), and a histone deacetylation inhibitor, trichostatin (15). However, in ME180-derived cisplatin (CDDP)-resistant cell lines, DAPK protein expression can not be induced by treatment with 5-aza-CdR and trichostatin. Although DAPK mRNA is expressed in the CDDP-resistant cells as in ME180 parent cells, DAPK mRNA translation is strongly suppressed in CDDP-resistant cells (15). These facts suggest that strong suppression of DAPK protein induction is involved in acquisition of CDDP resistance, and that DAPK protein regulates anticancer drug sensitivity and/or acquired anticancer drug resistance in cancer cells.

We recently reported that targeted knockdown of DAPK protein expression in the HHUA cell line (16), a highly differentiated endometrial adenocarcinoma cell line, induced cell death by enhancing tumor necrosis factor-related apoptosis-inducing ligand (TRAIL)-mediated apoptosis in an autocrine/paracrine manner (17). These findings suggest that endogenous DAPK is a candidate for molecularly targeted anticancer therapy for patients with endometrial adenocarcinoma, and that targeted knockdown of DAPK protein expression may enhance anticancer drug sensitivity in cancer cells. Therefore, we utilized specific DAPK small-interfering RNAs (siRNAs) to knock down endogenous DAPK protein expression in HHUA cells to directly investigate the role of endogenous DAPK in anticancer drug sensitivity.

Materials and methods

Cell line and culture. The HHUA cell line (16) was obtained from Riken Cell Bank (Tsukuba, Japan). Cells were cultured in OPTI-MEM (Invitrogen Corp., Carlsbad, CA) supplemented with 5% fetal bovine serum (FBS) (Equitech Bio Inc., Ingram,
TX), penicillin (100 U/ml), streptomycin (100 U/ml) and Fungizone (0.25 μg/ml; Invitrogen Corp.) in 5% CO₂ and 95% air at 37˚C.

**Transfection of DAPK siRNAs.** Five DAPK siRNA duplexes were designed and synthesized by iGENE Therapeutics Inc. (Tsukuba, Japan). The 5 siRNA sequences are shown in Table I. A negative control siRNA was purchased from Ambion Inc. (Austin, TX). Lipofectamine 2000 (Invitrogen Corp.) was used as the transfection reagent according to the manufacturer’s instructions. For experiments, cells were seeded in 6-well plates (2.5x10⁵ cells/well) or 10-cm dishes (2x10⁶ cells/dish), cultured for 24 h and then transfected with the DAPK siRNAs or control siRNA at a final concentration of 25 or 50 nM. Subsequently, the cells were cultured for 24-48 h for mRNA analyses and 48-72 h for protein assays before being harvested as indicated.

**Anticancer drug-sensitivity assays.** All anticancer drugs were kind gifts from pharmaceutical companies. 5-fluorouracil (5FU) was provided by Kirin-Kyowa-Hakko Co., Ltd. (Tokyo, Japan). Cisplatin (CDDP) was provided by Nihon-Kayaku Co., Ltd. (Tokyo, Japan). Etoposide (VP16) was provided by Bristol-Myers Squibb Japan Co., Ltd. (Tokyo, Japan). The effects of anticancer drugs on cell growth were assayed as follows. Cells in the log phase were detached with 0.25% trypsin/1 mM EDTA (Gibco-BRL) and cultured overnight in 96-well plates (5x10³ cells/well). On day 2, various concentrations of anticancer drugs were added to the cells. On day 4, the numbers of viable cells were evaluated using a cell proliferation assay kit (Dojin, Tokyo, Japan) and expressed as the percentages of viable cells (%) relative to the mean number of viable untreated cells. All experiments were performed in triplicate. Data are expressed as the mean ± SD, and comparative data (n=6) were analyzed by ANOVA.

**Western blotting.** For Western blot analysis, the cells were collected at 48-72 h after transfection with the DAPK siRNAs or control siRNA and lysed in phosphate-buffered saline containing 1% NP-40, 0.1% sodium dodecyl sulfate, Complete protease inhibitor cocktail (Roche Diagnostics, Corp., Indianapolis, IN) and 1 mM phenylmethyl sulfonyl fluoride. Protein concentrations of the cell lysates were assayed as follows. Cells in the log phase were detained with 0.25% trypsin/1 mM EDTA (Gibco-BRL) and cultured overnight in 96-well plates (5x10³ cells/well). On day 2, various concentrations of anticancer drugs were added to the cells. On day 4, the numbers of viable cells were evaluated using a cell proliferation assay kit (Dojin, Tokyo, Japan) and expressed as the percentages of viable cells (%) relative to the mean number of viable untreated cells. All experiments were performed in triplicate. Data are expressed as the mean ± SD, and comparative data (n=6) were analyzed by ANOVA.

<table>
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<th>Number of siRNA</th>
<th>siRNA sequence</th>
<th>Control siRNA</th>
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<td>3'-AU-CCAGUUCGAAUUCCUAGUGUGU-5'</td>
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</table>

(A) DAPK protein expression was strongly suppressed by transfection by each of 5 DAPK siRNAs. (B) Cleavage of PARP in DAPK siRNA-transfected cells.
RNase protection assay (RPA). Total RNA was isolated from the cultured cells using an RNeasy Mini Kit (Qiagen Inc., Valencia, CA) for RNase protection assays. A multiprobe RPA was performed using a RiboQuant™ hcc-2 Multiprobe Template Set and an RPA Kit (BD PharMingen) according to the manufacturer's protocol. Biotin-conjugated probes were prepared using a Non-Rad In Vitro Transcription Kit (BD PharMingen). The probes were combined with 10 μg of total RNA isolated from DAPK siRNA-transfected or control siRNA-transfected HHUA cells. After denaturation at 95°C for 3 min, the mixtures were allowed to hybridize at 56°C for 12-16 h before RNase digestion. RNA hybrids were separated in 4.75% acrylamide/8 M urea denaturing gels, and the protected fragments were transferred to positively charged nylon membranes. The fragments on the membranes were detected using the RPA Kit and the protected fragments were quantified with a Luminocapture apparatus and lane analyzer software (ATTO Corp.). The relative expression of each mRNA species was calculated after normalization by the expression of the housekeeping gene L32.

Statistical analysis. The data were expressed as means ± SD. Comparisons between experimental groups were performed by analysis of variance (ANOVA). If the ANOVA was significant, post-hoc comparisons were conducted using Scheffe’s test. Statistical significance was set at P<0.05.

Results

All 5 DAPK siRNA-transfected HHUA cells showed strong suppression of DAPK protein expression (Fig. 1A). Concurrently, specific downregulation of endogenous DAPK expression by DAPK siRNA transfection induced cleavage of PARP, an apoptosis marker, in the cells (Fig. 1B). No non-specific inhibitory effects of DAPK siRNA were detected on ß-actin protein expression, used as an internal control (Fig. 1A).

Next, anticancer drug-sensitivity tests of the DAPK siRNA-transfected HHUA cells were performed using 3 drugs, 5FU, CDDP and VP16. The anticancer drug-sensitivity curves were compared with those of the control siRNA-transfected
Our recent research showed that DAPK siRNA transfections can significantly enhance 5FU-sensitivity. This is the first report to directly show that suppressed DAPK was apparently suppressed in the cells.

siRNA-transfected cells, although p18 mRNA expression expressions (p130, Rb, p107, p53, p27, p21, p16) in DAPK molecules in the DAPK siRNA-transfected cells. As shown in Fig. 3, only p18 mRNA was significantly suppressed by DAPK knockdown. Although p18, a member of cyclin-dependent kinase inhibitor, is known to function as a positive or negative mediator in different apoptotic signals. Therefore, when DAPK is used for a molecular target in anticancer therapy, it should be first examined whether DAPK molecules play antiapoptotic roles in the target cells.

5FU is a major anticancer drug that is clinically used for patients with various types of cancers. Because 5FU is a well-known time-dependent anticancer drug, we have examined RNA expressions of cell cycle-regulating molecules in the DAPK siRNA-transfected cells by RPA. As shown in Fig. 3, only p18 mRNA was significantly suppressed by DAPK knockdown. Although p18, a member of cyclin-dependent kinase inhibitor, is reported to be a tumor suppressor gene (19-21), further investigations are needed to clarify whether suppressed p18 expression can affect anticancer drug sensitivity of the cells.

HHUA cells. As shown in Fig. 2, all 5 different DAPK siRNA-transfected cells exhibited much higher 5FU-sensitivities than the control siRNA-transfected cells. However, VP16-sensitivities in any DAPK siRNA-transfected cells could not be enhanced at all. As for CDDP-sensitivities of the DAPK siRNA-transfected cells, transfections with 4 of 5 DAPK siRNAs slightly enhanced CDDP-sensitivity while DAPK siRNA 003-transfected cells did not show any enhancement of CDDP-sensitivity compared with control siRNA-transfected cells.

5FU is a well known time-dependent anticancer drug. Therefore, to investigate the mechanisms of enhanced 5FU-sensitivity by DAPK siRNA transfections, RPA was done to examine mRNA expressions of cell cycle-regulating molecules in the DAPK siRNA-transfected cells. As shown in Fig. 3, DAPK siRNA transfections did not affect 7 mRNA expressions (p130, Rb, p107, p53, p27, p21, p16) in DAPK siRNA-transfected cells, although p18 mRNA expression was apparently suppressed in the cells.

Discussion

This is the first report to directly show that suppressed DAPK protein expression can significantly enhance 5FU-sensitivity. Our recent research showed that DAPK siRNA transfections induce cell deaths in human endometrial adenocarcinoma cells, uterine carcinosarcoma cells and uterine leiomyosarcoma cells (18). In a human endometrial adenocarcinoma cell line HHUA, DAPK siRNA transfection induces TRAIL-mediated apoptosis in autocrine/paracrine manner (17). In the present study, we have shown that DAPK siRNA transfections strongly enhance 5FU-sensitivity, but not VP16-sensitivity, in HHUA cells. These results indicate that DAPK can be a convincing target for anticancer therapy, and that 5FU-combined chemotherapy with DAPK siRNA transfection may have a stronger anticancer effect on patients with endometrial adenocarcinoma than the chemotherapy alone.

Our recent study showed that DAPK siRNA transfections into HHUA cells induce TRAIL-mediated apoptosis in cells (17). Since DAPK siRNA transfections did not affect VP16-sensitivity in the cells at all, the VP16-stimulated cell death signals may share or include TRAIL-mediated apoptotic signals. On the other hand, findings that DAPK siRNA transfections strongly enhance 5FU-sensitivity suggest that 5FU-stimulated cell death signals may be independent from the TRAIL-mediated apoptotic signals induced by DAPK siRNA transfections.

As shown in Fig. 2, transfections with 4 out of 5 DAPK siRNA slightly enhanced CDDP-sensitivity of HHUA cells. In a previous study, strong suppression of DAPK protein induction was shown to be involved in acquired CDDP-resistance in ME180 cells (15). Contrary to this report, the present study has showed that CDDP-sensitivity was slightly enhanced in the DAPK siRNA-transfected cells. Taken together, these results present a disagreement between DAPK protein expression and CDDP-sensitivity between in HHUA cells and in ME180 cells. The disagreement may be caused by the cell lineage of the examined cancer cells, because DAPK is known to function as a positive or negative mediator in different apoptotic signals. Therefore, when DAPK is used for a molecular target in anticancer therapy, it should be first examined whether DAPK molecules play antiapoptotic roles in the target cells.

5FU is a major anticancer drug that is clinically used for patients with various types of cancers. Because 5FU is a well-known time-dependent anticancer drug, we have examined RNA expressions of cell cycle-regulating molecules in the DAPK siRNA-transfected cells by RPA. As shown in Fig. 3, only p18 mRNA was significantly suppressed by DAPK knockdown. Although p18, a member of cyclin-dependent kinase inhibitor, is reported to be a tumor suppressor gene (19-21), further investigations are needed to clarify whether suppressed p18 expression can affect anticancer drug sensitivity of the cells.

References


