

Combined treatment of gemcitabine with indole-3-carbinol or metformin on drug efficacy in pancreatic cancer cell lines: The role of human equilibrative nucleoside transporters

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Abstract

Pancreatic cancer is one of the most lethal carcinomas in the United States. In accord with the American Cancer Society pancreatic cancer is anticipated to move from the third to the second leading cause of deaths in the United States by 2020. Although the standard treatment for advanced pancreatic cancer is gemcitabine (GEM), the response rate is less than 20%. Chemoresistance is a hallmark of this cancer, and modulation of drug transporters expression has been shown to increase cancer drug efficacy. Studies have shown that human equilibrative nucleoside transporters (hENTs) expression patterns may predict GEM treatment efficacy. This study investigated whether or not GEM in combination with metformin (MET) or indole-3-carbinol (I3C) increases cytotoxicity and modulates hENT1 and hENT4. Pancreatic cancer cells from males and females were treated for 24 or 72h with GEM and/or MET or I3C. Cell viability, drug interactions, and protein and mRNA expression levels of hENTs were assessed. Treatment with GEM and/or MET or I3C showed cell line specific reductions in pancreatic cancer cell proliferation, and modulation of hENT1 and hENT4 expression. Response to GEM and MET/I3C may be dependent upon the genetic profile of the tumor and the level of expression of a specific transporter. The sensitivity of GEM could depend on the method of treatment, whether cells were pre-treated with MET or I3C, which studies, including our own, have showed that pre-treatment with I3C increased upregulation of hENT1 expression in pancreatic cancer cell lines.

Keywords: pancreatic cancer; gemcitabine; indole-3-carbinol; metformin; human equilibrative nucleoside transporters

Introduction

Pancreatic cancer is the eleventh most common cancer in the US but represents the third leading cause of cancer deaths in men and women [1]. The prognosis is extremely poor, with a 5-years relative survival rate of only 7%, due to the fact that pancreatic cancer is usually asymptomatic in the early stages and most cases are diagnosed at late stages of the disease [1]. Despite significant improvements in understanding molecular and epigenetic changes of pancreatic cancer, the prognosis and management remain unchanged. Treatment and advances in early detection of pancreatic cancer, therefore, remain of critical importance. Effective treatments or novel therapeutic approaches are currently being investigated in clinical trials [2]. Although gemcitabine (GEM) is the gold standard for advance stages of this disease, improvement is needed to enhance further its efficacy for patient survival.

GEM, a deoxycytidine nucleoside analog, is used in advanced pancreatic cancer patients with locally advanced or metastatic cancer [3]. Pancreatic cancer is a highly resistant cancer, due mainly to up regulation of multidrug resistance genes [4, 5]. In addition, patients with various polymorphisms in these genes can have enhanced or decreased efficacy of drugs for treatment [4, 5]. Patients treated with GEM can also develop resistance to this drug. Therefore, it is important to identify other drug agents

that, when combined with GEM, will increase the efficacy of GEM.

Previously published data from our laboratory demonstrated enhanced efficacy of GEM with the dietary agent indole-3-carbinol (I3C) in pancreatic cancer [6] and further showed that polymorphisms in ABCB1 in pancreatic cancer cell lines affected GEM efficacy [7]. I3C is a common phytochemical found in cruciferous vegetables. Studies have shown that dietary phytochemicals, such as I3C, may be a promising nontoxic chemopreventive agent with multiple anti-tumor activities, including apoptotic, anti-proliferative, and anti-angiogenic activities [8, 9]. Several clinical trials have investigated the use of I3C for cancer

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Received 14 November 2017 Revised 4 January 2018 Accepted 15 January 2018 Published 23 January 2018

Citation: Joseph S, Word B, Lyn-Cook B. Combined treatment of gemcitabine with indole-3-carbinol or metformin on drug efficacy in pancreatic cancer cell lines: The role of human equilibrative nucleoside transporters. J Cancer Res Ther. 2018; 6(2):6-17. DOI: [10.14312/2052-4994.2018-2](https://doi.org/10.14312/2052-4994.2018-2)

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treatment and prevention. In these clinical trials, I3C has been reported to be effective in both the cervix [10] and larynx [11] against precancerous lesions caused by human papillomavirus. In more recent clinical trials, the use of I3C has been investigated in the elimination of cancer-influencing exogenous estrogens in patients with prostate, breast and uterine cancers [8, 12-14].

Studies have shown that the expression of several genes may be predictors of GEM efficacy in pancreatic cancer patients [15, 16]. One of the predictor genes for GEM is human equilibrative nucleoside transporter-1 (hENT1), a transporter responsible for the uptake of GEM in human cells [6, 16]. hENT4, another member of the hENT family, has been identified as one of several drug transporters mediating the renal and intestinal uptake of metformin (MET) [17, 18]. MET is the primary drug used for the treatment of type 2 diabetes mellitus, and has become a drug of interest for the treatment of pancreatic cancer [19-23]. Several studies have suggested that patients taking MET for the treatment of diabetes also have a decreased incidence of pancreatic cancer [24, 25]. MET has also been implicated in its ability to increase sensitivity of pancreatic cancer cells to GEM [24]. Other studies have concluded, however, that there is no association between MET use and cancer risk or prognosis [26, 27].

This study was performed to evaluate the duo-combined drug effects of GEM and/or MET or I3C in pancreatic cancer cells in relation to chemoresistance. Several pancreatic cancer cell lines were examined for cell viability and drug interactions when treated with GEM, MET, and I3C in mono or combination therapy for 24 or 72 h. We further investigated whether GEM in combination treatment with MET or I3C could modulate hENT1 and hENT4 expression.

Materials and methods

Material

I3C, MET, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT), sodium dodecyl sulfate (SDS), hydrochloric acid (HCl), dimethyl sulfoxide (DMSO), bovine serum albumin (BSA), and complete mini protease inhibitor cocktail were purchased from Sigma (St. Louis, MO). GEM was a gift from Eli Lilly (Indianapolis, IN). Fetal bovine serum (FBS), Roswell Park Memorial Institute (RPMI) 1640, Dulbecco's Minimal Eagle's Medium (DMEM), phosphate saline buffer (PBS), glutamine, 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid (HEPES), and sodium pyruvate were purchased from Gibco (Grand Island, NY).

Cell culture

Human pancreatic cancer cell lines from males MIA PaCa2 and PANC1 and females ASPC1, PANC1, and Su.86.86 were obtained from the American Type Culture Collection (ATCC; Manassas, VA). Cell lines were cultured in DMEM (PANC1 and MIA PaCa2) or RPMI 1640 (SU.86.86 and ASPC1) with 10% FBS, 100 U/mL penicillin, and 100 µg/mL streptomycin (Invitrogen, Carlsbad, CA) at 37°C in a humidified atmosphere containing 5% CO₂. Glutamine, HEPES, and sodium pyruvate supplements were added to maintain proper cell growth for SU.86.86 cells.

MTT assay for cell proliferation

Using the MTT assay, a colorimetric assay for measuring

cell metabolic activity, cells were seeded in 96-well plates (1 × 10⁴ cells per well), and were treated with therapeutic agents for 24 or 72 h at 37°C in a humidified atmosphere containing 5% CO₂. At the end of the treatment period, MTT solution (5 mg/mL) was added to each well (10 µL) and incubated for 4 h at 37°C. Subsequently, a SDS/HCl solution (100 µL) was added. Optical density was determined using a BioTek Cytation3 imaging reader (Winooski, VT) at 595nm. Drug concentrations of GEM (500 µM), MET (400 µM), I3C (250 or 500 µM) were used for the assay. These doses were selected based on a dose-range study. The coefficient of drug interaction (CDI) used to determine whether two drugs are synergistic, additive, or antagonist was calculated with the following equation:

$$CDI = \frac{(\text{ratio of cell viability for drug A})(\text{ratio of cell viability for drug B})}{\text{ratio of cell viability for drug combination AB}}$$

RNA extraction and cDNA synthesis

Total RNA were extracted from cells using RNeasy Mini Kit (Qiagen, Valencia, CA) according to manufacturer's instructions. Nucleic acid purities and concentrations were determined using a NanoDrop Spectrophotometer ND1000 (ThermoFisher, Grand Island, NY). All RNA samples had A260/A280 ratios of 1.9 to 2.1. The integrity and quality of the RNA were assessed using a Biorad Experion Automated Electrophoresis Station (Hercules, CA), and samples with RNA integrity number (RIN) values between 8 and 10 were utilized. cDNA was synthesized (reverse transcribed polymerase chain reaction (RT-PCR)) from 0.4 µg of high quality RNA (RIN>8) using the Advantage RT-for-PCR Kit according to the manufacturer's protocol (Clontech Laboratories Inc., Mountain View, CA).

Quantitative real time- polymerase chain reaction (QRT-PCR)

For QRT-PCR, Multiplex master mix solution was obtained from Biorad (Hercules, CA). Primers for SLC 29A1-FAM (hENT1) and β-Actin-VIC were obtained from Life Technologies (Grand Island, NY), and those for SLC29A4 (hENT4) are obtained from Integrated DNA Technologies (Coralville, Iowa). Triplicate experiments were assayed using a Biorad CFX96 C1000 System (Biorad). Delta-Delta Ct (ΔΔC_t) calculations were used to determine the change in the relative level of gene expression. β-Actin functioned as the reference gene.

Western blots

After the indicated treatment time, pancreatic cancer cells were washed with PBS, harvested by scraping and lysed in 100 µL RIPA buffer (Thermo Scientific, Grand Island, NY) supplemented with protease inhibitor. Whole cell protein extracts (30 µg) were electrophoresed on TGX stain-free gel (Biorad), and transferred to polyvinylidene difluoride (low fluorescence PVDF; Biorad) membranes. For determining protein expression level of hENT1 and hENT4, membranes were blocked with 3% bovine serum albumin and incubated overnight at 4°C with primary antibody, anti-SLC29A1 (hENT1: rabbit polyclonal; Origene; Rockville, MD) and anti-SLC29A4 (hENT4: mouse monoclonal, Abcam, Cambridge, MA). Membranes were washed and incubated with horseradish peroxidase (HRP) conjugated secondary antibody, goat anti-rabbit IgG or goat anti-mouse IgG (Thermo Scientific). Immunoreactive bands were detected

by chemiluminescence using ChemiDoc Touch Imaging System (Biorad). The intensity of each band was quantified by dosimetry using Image Lab™ Software (Biorad). Total protein was measured to ensure the protein loading (Supplemental Figure 1).

Statistical analysis

Statistical analyses were performed using GraphPad Prism software Version 6.0 (San Diego, CA). One-way analysis of variance (ANOVA) with Tukey’s method to adjust for multiple comparisons was used to determine statistically significance. A p value of <0.05 was considered to be significant.

Results

MTT assays were performed to investigate the percentage of inhibition of proliferation in pancreatic cancer cell lines when treated with GEM and/or MET or I3C, monotherapy

and combination therapy. Treatment with 500 μM GEM resulted in time-dependent inhibition of cell proliferation in MIA PaCa2, PANC1, ASPC1, and SU.86.86 pancreatic cancer cell lines compared to the untreated control. The inhibition of cell proliferation in MIA PaCa2, PANC1, ASPC1, and SU.86.86 pancreatic cancer cell lines at 24 h was 29.5, 50.2, 16.3, and 5.9%, respectively; and at 72 h was 71.6, 81.0, 40.6, and 76.2%, respectively (Figure 1 and 2). Female cell lines, ASPC1 and SU.86.86, were more resistant to GEM treatment than the male cell lines (MIA PaCa2 and PANC1). Treatment with 400 μM MET resulted in decreased inhibition of proliferation in MIA PaCa2, PANC1, ASPC1, and SU.86.86 pancreatic cancer cell lines (24 h: 36.8, 50.2, 8.2, and 0.8%; and 72 h: 23.7, 42.2, 13.8, and 18.6%, respectively) (Figure 1). In male pancreatic cancer cell lines, MIA PaCa2 and PANC1, there was a slight decrease in inhibition of proliferation as treatment time increased; however, there was an increase in inhibition of proliferation in female cell lines, ASPC1 and SU.86.86, as treatment time increased.

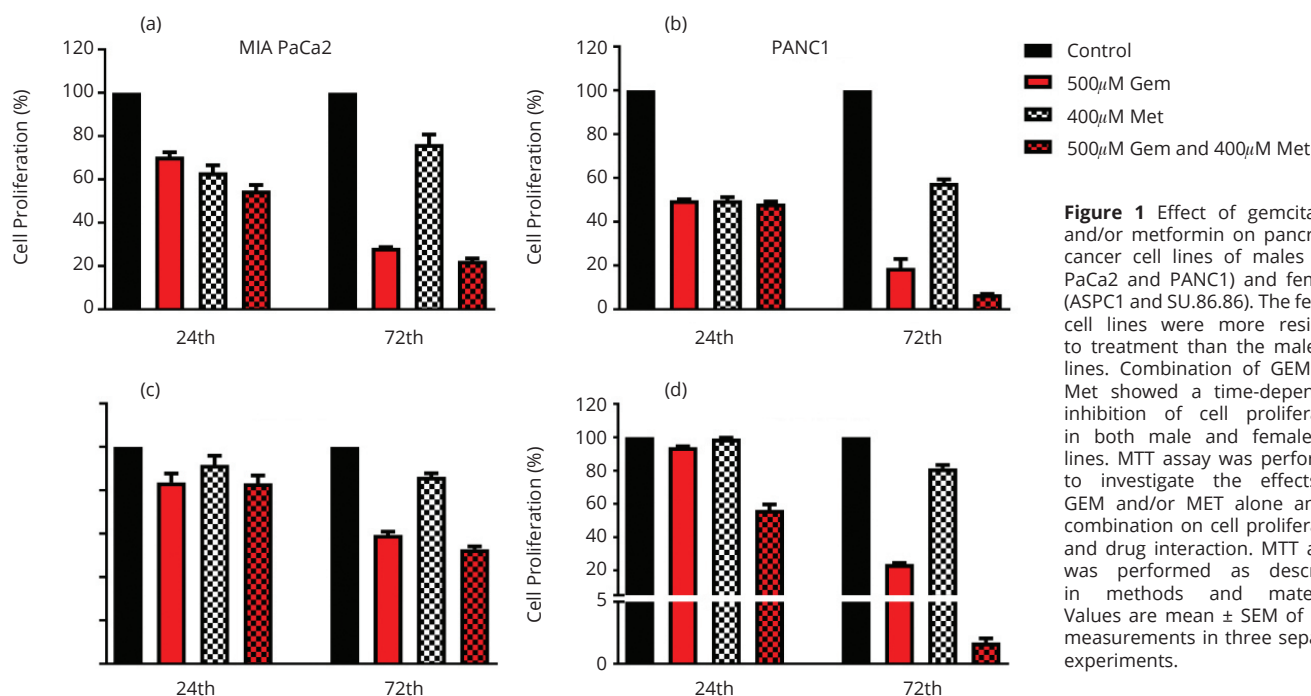


Figure 1 Effect of gemcitabine and/or metformin on pancreatic cancer cell lines of males (MIA PaCa2 and PANC1) and females (ASPC1 and SU.86.86). The female cell lines were more resistant to treatment than the male cell lines. Combination of GEM and Met showed a time-dependent inhibition of cell proliferation in both male and female cell lines. MTT assay was performed to investigate the effects of GEM and/or MET alone and in combination on cell proliferation and drug interaction. MTT assay was performed as described in methods and materials. Values are mean ± SEM of eight measurements in three separate experiments.

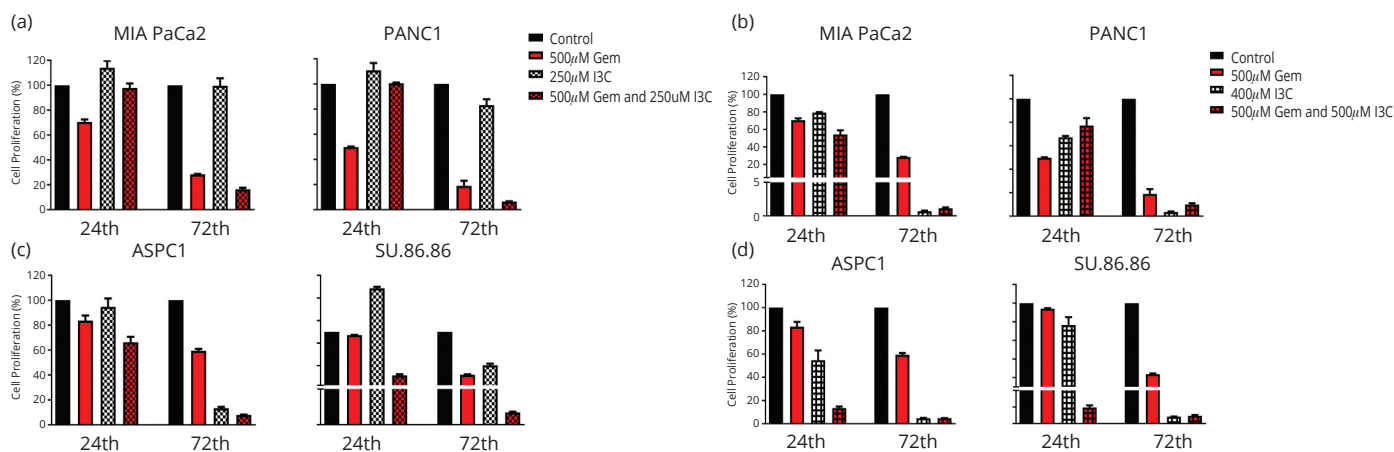


Figure 2 Effect of gemcitabine and/or indole-3-carbinol on pancreatic cancer cell lines of males (MIA PaCa2 and PANC1) and females (ASPC1 and SU.86.86). Combine GEM and I3C induced a higher percentage of inhibition in the female pancreatic cancer cells compared to GEM alone. MTT assay was performed to investigate the effects of GEM and/or I3C alone and in combination on cell proliferation and drug interaction. MTT assay was performed as described in methods and materials. Values are mean ± SEM of eight measurements in three separate experiments.

A combination treatment of 500 μ M GEM and 400 μ M MET resulted in time-dependent inhibition of cell proliferation in MIA PaCa2, PANC1, ASPC1 and SU.86.86 pancreatic cancer cell lines (24 h: 45.2, 51.7, 16.7, and 43.6%; and 72 h: 77.5, 93.0, 47.1, and 98.3%, respectively) (Figure 1). Combination therapy of GEM and MET induced a higher percentage of inhibition of proliferation in pancreatic cancer cells compared to GEM monotherapy at both time points, ranging from 0.4 - 37.7% greater inhibition (Figure 1). However, the summations of cell proliferation inhibition for the individual drugs are higher than the combined drug effect in each of the cell lines with the exception of SU.86.86 (Figure 1). The CDI indicated that treatment with GEM and MET for 24h: ASPC1 and Su.86.86, and 72h: ASPC1, PANC1 and Su.86.86 were synergistic (Table 1). The statistical significance of cell proliferation is found in Table

2. Treatment with I3C resulted in a time- and concentration-dependent increase in inhibition of proliferation of MIA PaCa2, PANC1, ASPC1, and SU.86.86 pancreatic cancer cell lines. After 24 h, the low dose of I3C (250 μ M) increased cell proliferation in MIA PaCa2, PANC1, and SU.86.86 cells lines by 14.1, 10.8, and 77.1%, respectively, demonstrating that these cells were not affected by the concentration of I3C. In ASPC1 cells, the lower dose of I3C inhibited cell proliferation by 5.2% (Figure 2a). After 72h treatment time, low dose of I3C (250 μ M) inhibited cell proliferation in MIA PaCa2, PANC1, ASPC1, and SU.86.86 cells lines by 0.3, 16.8, 86.6, and 89.4%, respectively (Figure 2a). By contrast, the high dose of I3C (500 μ M) resulted in an inhibition of proliferation of 21.0, 32.7, 45.4, and 23.6% at 24h, and cell death 99.3, 96.4, 95.2, and 97.9% at 72h, respectively (Figure 2b).

Table 1 Coefficient of drug interaction (CDI) for combination therapy.

Cell line-treatment time	500 μ M Gem + 250 μ M I3c	500 μ M Gem + 500 μ M I3C	500 μ M Gem + 400 μ M Met
MIAPaCa2-24h (M)	1.2	1.0	1.2
MIAPaCa2-72h	0.6	5.7	1.0
ASPC1-24h (F)	0.8	0.3	0.8
ASPC1-72h	1.0	1.7	0.4
PANC1-24h (M)	1.8	2.3	1.9
PANC1-72h	0.4	14.6	0.6
SU.86.86-24h (F)	0.1	0.1	0.8
Su.86.86-72h	0.2	4.8	0.1

Abbreviations: M: male; F: female; CDI: =1.0 (additive), >1.0 (antagonist), <1.0 (synergist).

Table 2 (a) Statistical analysis of MTT assay. Data represents p-value.

	MIAPaCa2 (Male)	
	24h	72h
Control vs. 400 μ M Met	< 0.0001	0.0002
Control vs. 500 μ M Gem	0.0003	< 0.0001
Control vs. 250 μ M I3C	0.1752	> 0.9999
Control vs. 500 μ M I3C	0.0286	< 0.0001
Control vs. 500 μ M Gem and 400 μ M Met	< 0.0001	< 0.0001
Control vs. 500 μ M Gem and 250 μ M I3C	> 0.9999	< 0.0001
Control vs. 500 μ M Gem and 500 μ M I3C	< 0.0001	< 0.0001
400 μ M Met vs. 400 μ M Met and 500 μ M Gem	0.7663	< 0.0001
500 μ M Gem vs. 500 μ M Gem and 400 μ M Met	0.0961	0.8553
500 μ M Gem vs. 500 μ M Gem and 250 μ M I3C	0.0008	0.1089
500 μ M Gem vs. 500 μ M Gem and 500 μ M I3C	0.0800	< 0.0001
250 μ M I3C vs. 500 μ M I3C	0.0001	< 0.0001
250 μ M I3C vs. 500 μ M Gem and 250 μ M I3C	0.0805	< 0.0001
500 μ M I3C vs. 500 μ M Gem and 500 μ M I3C	0.0071	> 0.9999
500 μ M Gem and 500 μ M I3C vs. 500 μ M Gem and 250 μ M I3C	< 0.0001	0.0192

Table 2 (b) Statistical analysis of MTT assay. Data represents p-value.

<i>Treatment</i>	<i>ASPC1 (Female)</i>	
	<i>24h</i>	<i>72h</i>
Control vs. 400µM Met	0.922	< 0.0001
Control vs. 500µM Gem	0.2255	< 0.0001
Control vs. 250µM I3C	0.9957	< 0.0001
Control vs. 500µM I3C	0.9986	< 0.0001
Control vs. 500µM Gem and 400µM Met	0.0005	< 0.0001
Control vs. 500µM Gem and 250µM I3C	0.2041	< 0.0001
Control vs. 500µM Gem and 500µM I3C	< 0.0001	< 0.0001
400µM Met vs. 400µM Met and 500µM Gem	0.0106	< 0.0001
500µM Gem vs. 500µM Gem and 400µM Met	0.1649	< 0.0001
500µM Gem vs. 500µMGem and 250µM I3C	> 0.9999	0.0017
500µM Gem vs. 500µM Gem and 500µM I3C	0.0015	< 0.0001
250µM I3C vs. 500µM I3C	> 0.9999	> 0.9999
250µM I3C vs. 500µM Gem and 250µM I3C	0.6581	< 0.0001
500µM I3C vs. 500µM Gem and 500µM I3C	< 0.0001	0.0002
500µM Gem and 500µM I3C vs. 500µMGem and 250µM I3C	0.0017	< 0.0001

Table 2 (c) Statistical analysis of MTT assay. Data represents p-value.

<i>Treatment</i>	<i>PANC1 (Male)</i>	
	<i>24h</i>	<i>72h</i>
Control vs. 400µM Met	< 0.0001	< 0.0001
Control vs. 500µM Gem	< 0.0001	< 0.0001
Control vs. 250µM I3C	0.1745	0.0006
Control vs. 500µM I3C	< 0.0001	< 0.0001
Control vs. 500µM Gem and 400µM Met	< 0.0001	< 0.0001
Control vs. 500µM Gem and 250µM I3C	> 0.9999	< 0.0001
Control vs. 500µM Gem and 500µM I3C	0.0012	< 0.0001
400µM Met vs. 400µM Met and 500µM Gem	> 0.9999	< 0.0001
500µM Gem vs. 500µM Gem and 400µM Met	> 0.9999	0.0187
500µM Gem vs. 500µMGem and 250µM I3C	< 0.0001	0.0131
500µM Gem vs. 500µM Gem and 500µM I3C	0.0001	0.1247
250µM I3C vs. 500µM I3C	< 0.0001	< 0.0001
250µM I3C vs. 500µM Gem and 250µM I3C	0.2189	< 0.0001
500µM I3C vs. 500µM Gem and 500µM I3C	0.4859	0.53
500µM Gem and 500µM I3C vs. 500µMGem and 250µM I3C	0.0009	0.9745

Table 2 (d) Statistical analysis of MTT assay. Data represents p-value.

Treatment	SU.86.86 (Female)	
	24h	72h
Control vs. 400µM Met	> 0.9999	< 0.0001
Control vs. 500µM Gem	0.8911	< 0.0001
Control vs. 250µM I3C	< 0.0001	< 0.0001
Control vs. 500µM I3C	< 0.0001	< 0.0001
Control vs. 500µM Gem and 400µM Met	< 0.0001	< 0.0001
Control vs. 500µM Gem and 250µM I3C	0.0123	< 0.0001
Control vs. 500µM Gem and 500µM I3C	< 0.0001	< 0.0001
400µM Met vs. 400µM Met and 500µM Gem	< 0.0001	< 0.0001
500µM Gem vs. 500µM Gem and 400µM Met	< 0.0001	< 0.0001
500µM Gem vs. 500µM Gem and 250µM I3C	0.2215	0.9998
500µM Gem vs. 500µM Gem and 500µM I3C	< 0.0001	< 0.0001
250µM I3C vs. 500µM I3C	0.9996	< 0.0001
250µM I3C vs. 500µM Gem and 250µM I3C	< 0.0001	< 0.0001
500µM I3C vs. 500µM Gem and 500µM I3C	< 0.0001	< 0.0001
500µM Gem and 500µM I3C vs. 500µM Gem and 250µM I3C	< 0.0001	< 0.0001

Combined GEM and I3C therapy (both low and high dose) induced a higher percentage of inhibition of proliferation in female pancreatic cancer cells (ASPC1 and SU.86.86) compared to GEM monotherapy at both time points, ranging from 17.5%-89.2% (Figure 2). However, at 24h treatment time, GEM monotherapy inhibited cell proliferation at a higher rate in MIA PaCa2 compared to combination therapy with low dose of I3C, and in PANC1 compared to combination therapy with either dose of I3C (Figure 2). At 72h, combined therapy of GEM and I3C (both low and high dose) induced a higher percentage of inhibition of proliferation in male pancreatic cancer cells (MIA PaCa2 and PANC1) compared to GEM monotherapy at both time points, ranging from 9.1%-27.2% (Figure 2). The combined drug effects of GEM and I3C are higher than summations of cell proliferation inhibition for the two drugs for the following cell lines (I3C dose and treatment time): APSC1 and SU.86.86 (24h, both I3C doses), and MIA PaCa2 (72h, low dose I3C) (Figure 2).

The CDI indicated that GEM and low dose I3C (250µM) were found to be synergistic in MIA PaCa2 (72h), ASPC1 (24h), PANC1 (72h), and SU.86.86 (24h and 72h); and GEM and high dose I3C (500µM) were found to be synergistic in ASPC1 (24h), and SU.86.86 (24h) (Table 1). The statistical significance of cell proliferation is found in Table 2a-d.

The expression levels of two drug transporter genes, hENT1 and hENT4, were investigated in pancreatic cancer cell lines before and after drug treatment with 500µM GEM and/or 400µM MET or 250µM I3C. On average MET induced lower mRNA expression of hENT1 in pancreatic cancer cells compared to GEM. The highest mRNA expression

of hENT1 induced by MET was found in MIA PaCa2 cells (72h) (Figure 3a). Combined therapy of GEM and MET was found to enhance the hENT1 mRNA expression only in SU.86.86 cells after 72h treatment time by 10% compared to the summation of percent increase induced by the individual drugs (Figure 3a). Protein analysis did not confirm this finding (Figure 3c). 250µM I3C induced hENT1 mRNA expression in all cell lines except ASPC1 in a time-dependent manner (Figure 3b). Indole-3-carbinol induced higher mRNA expression of hENT1 in MIA PaCa2 (24h and 72h), PANC1 (72h) and SU.86.86 (72h) pancreatic cancer cells compared to GEM. Combined therapy of GEM and 250µM I3C was found to enhance the hENT1 mRNA expression only in SU.86.86 cells after 72h treatment time by 40% compared to the summation of percent increase induced by the individual drugs (Figure 3b). Protein analysis confirmed this finding in SU.86.86 cells treated with GEM and 250µM I3C for 72h (2.1-fold, Figure 3c). On average GEM induced higher mRNA expression levels of hENT4 compared to MET (Figure 4b).

Combined treatment of GEM and MET enhanced hENT4 mRNA expression in SU.86.86 pancreatic cancer cell lines after 24h and 72h treatment time by 70% and 80% compared to the summation of percent increase induced by the individual drugs (Figure 4a). Protein analysis confirmed this finding in SU.86.86 cells treated with GEM and MET for 24h and 72h (2.0- and 1.9-fold, respectively; Figure 3c). Combined treatment of GEM and 250µM I3C did not enhance hENT4 mRNA expression in pancreatic cancer cell lines after 24h and 72h treatment time compared to the summation of the individual drugs. The statistical significance of mRNA expression analysis is found in Table 3a and b.

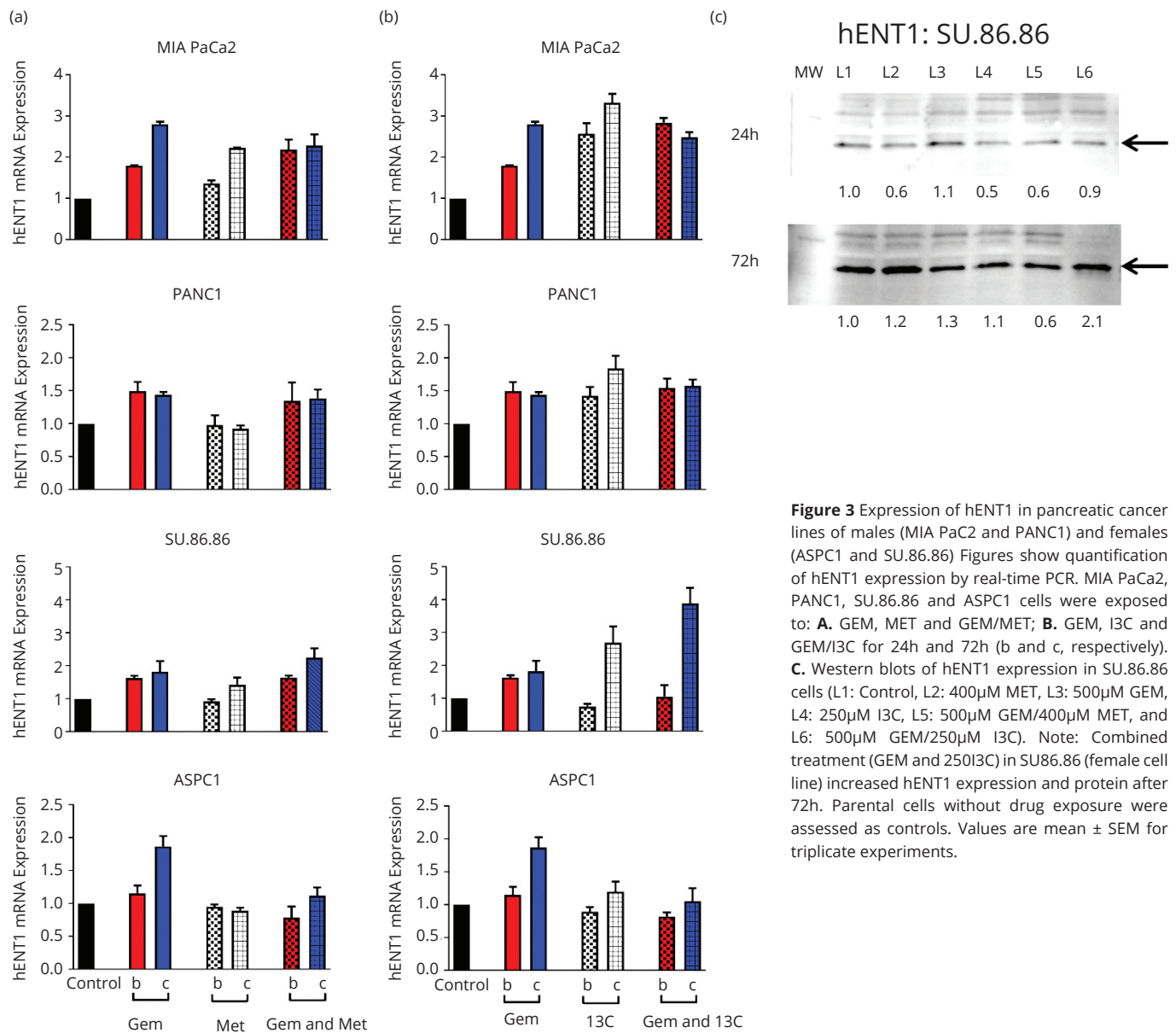


Figure 3 Expression of hENT1 in pancreatic cancer lines of males (MIA PaCa2 and PANC1) and females (ASPC1 and SU.86.86) Figures show quantification of hENT1 expression by real-time PCR. MIA PaCa2, PANC1, SU.86.86 and ASPC1 cells were exposed to: **A.** GEM, MET and GEM/MET; **B.** GEM, I3C and GEM/I3C for 24h and 72h (b and c, respectively). **C.** Western blots of hENT1 expression in SU.86.86 cells (L1: Control, L2: 400µM MET, L3: 500µM GEM, L4: 250µM I3C, L5: 500µM GEM/400µM MET, and L6: 500µM GEM/250µM I3C). Note: Combined treatment (GEM and 250I3C) in SU86.86 (female cell line) increased hENT1 expression and protein after 72h. Parental cells without drug exposure were assessed as controls. Values are mean ± SEM for triplicate experiments.

Table 3 (a) Statistical analysis of hENT1 mRNA expression. Data represents p-value.

		MIA PaCa2 (M)		PANC1 (M)		ASPC1 (F)		Su.86.86 (F)	
		24h	72h	24h	72h	24h	72h	24h	72h
hENT1	Control vs. 400 µM Met	0.4668	0.0058	> 0.9999	0.9857	0.9989	0.9886	0.9989	0.8658
	Control vs. 500µM Gem	0.0386	0.0001	0.3136	0.0393	0.8346	0.0051	0.1006	0.3301
	Control vs. 250µM I3C	0.0003	< 0.0001	0.4596	0.001	0.9592	0.8748	0.8493	0.0238
	Control vs. 500µM Gem + 400µM Met	0.0012	0.0018	0.6528	0.0784	0.5772	0.9843	0.0891	0.0669
	Control vs. 500µM Gem + 250µM I3C	<0.0001	0.0015	0.2265	0.0155	0.7515	0.9995	0.9999	0.0005
	400 µM Met vs. 500µM Gem + 400µM Met	0.017	0.9999	0.5948	0.03	0.7804	0.8025	0.0485	0.3284
	500µM Gem vs. 500µM Gem + 400µM Met	0.5181	0.2763	0.9846	0.9968	0.1152	0.015	>0.9999	0.8642
	500µM Gem vs. 500µM Gem + 250µM I3C	0.0076	0.7948	0.9999	0.9074	0.1864	0.0084	0.1469	0.0067
	250µM I3C vs. 500µM Gem + 250µM I3C	0.828	0.06	0.9934	0.5131	0.9932	0.9642	0.7308	0.1968
	500µM Gem + 400µM Met vs. 500µM Gem + 250µM I3C	0.0647	0.963	0.9446	0.7228	0.9995	0.9992	0.1306	0.0294

Abbreviations: M: male; F: female.

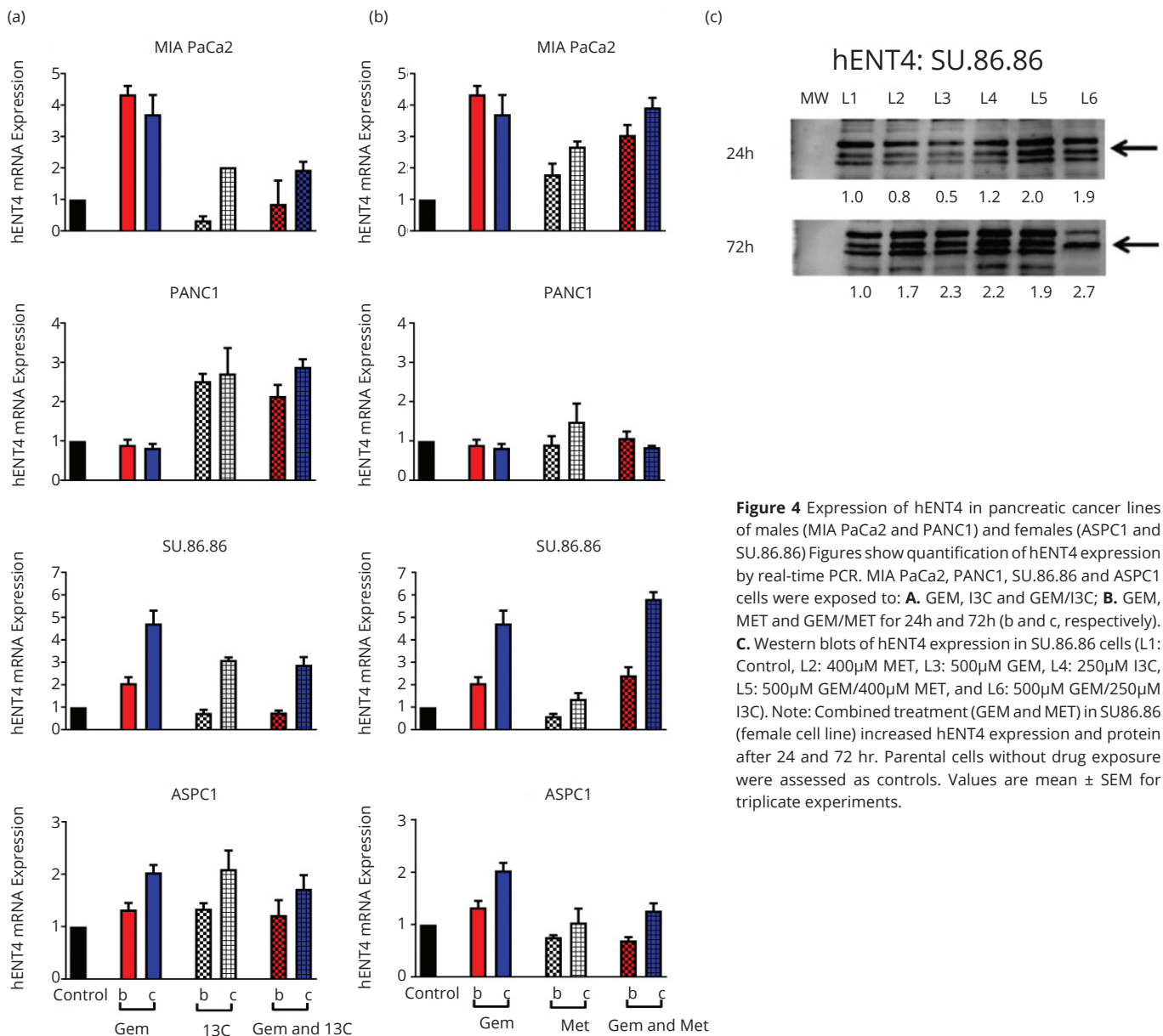


Figure 4 Expression of hENT4 in pancreatic cancer lines of males (MIA PaCa2 and PANC1) and females (ASPC1 and SU.86.86) Figures show quantification of hENT4 expression by real-time PCR. MIA PaCa2, PANC1, SU.86.86 and ASPC1 cells were exposed to: **A.** GEM, I3C and GEM/I3C; **B.** GEM, MET and GEM/MET for 24h and 72h (b and c, respectively). **C.** Western blots of hENT4 expression in SU.86.86 cells (L1: Control, L2: 400µM MET, L3: 500µM GEM, L4: 250µM I3C, L5: 500µM GEM/400µM MET, and L6: 500µM GEM/250µM I3C). Note: Combined treatment (GEM and MET) in SU86.86 (female cell line) increased hENT4 expression and protein after 24 and 72 hr. Parental cells without drug exposure were assessed as controls. Values are mean ± SEM for triplicate experiments.

Table 3 (b) Statistical analysis of hENT4 mRNA expression. Data represents p-value.

		MIA PaCa2 (M)		PANC1 (M)		ASPC1 (F)		Su.86.86 (F)	
		24h	72h	24h	72h	24h	72h	24h	72h
hENT4	Control vs. 400 µM Met	0.6799	0.0297	0.9992	0.7963	0.5714	>0.9999	0.845	0.9623
	Control vs. 500µM Gem	0.0005	0.0009	0.9987	0.9969	0.2757	0.1174	0.054	0.0001
	Control vs. 250µM I3C	0.8146	0.3818	0.0008	0.0236	0.3263	0.0503	0.97	0.0197
	Control vs. 500µM Gem + 400µM Met	0.023	0.0005	0.9995	0.9982	0.343	0.9561	0.0106	<0.0001
	Control vs. 500µM Gem + 250µM I3C	0.9998	0.3561	0.0081	0.0132	0.725	0.296	0.9796	0.0376
	400 µM Met vs. 500µM Gem + 400µM Met	0.2527	0.1383	0.9843	0.5779	0.9968	0.9776	0.0044	<0.0001
	500µM Gem vs. 500µM Gem + 400µM Met	0.2246	0.9962	0.9798	>0.9999	0.0117	0.3396	0.8411	0.2518
	500µM Gem vs. 500µM Gem + 250µM I3C	0.0003	0.0217	0.0043	0.0073	0.983	0.946	0.0346	0.042
	250µM I3C vs. 500µM Gem + 250µM I3C	0.9214	> 0.9999	0.6886	0.999	0.9772	0.8407	>0.9999	0.9984
	500µM Gem + 400µM Met vs. 500µM Gem + 250µM I3C	0.0147	0.0102	0.0136	0.0078	0.0638	0.7228	0.0082	0.002

Abbreviations: M: male; F: female.

Discussion

GEM monotherapy has been the standard of treatment for metastatic pancreatic cancer patients for several years; however, the majority of patients treated with GEM eventually become resistant to the drug. Pancreatic cancer cell lines possess high intrinsic and acquired chemoresistance. Pang et al. showed that polymorphisms in ABCB1 in pancreatic cancer cell lines affected GEM efficacy, where MIA PaCa2 cells exhibit a ABCB12677TT polymorphism, and are more sensitive to GEM than PANC1, SU.86.86 and ASPC1 [7]. Therefore, studies have investigated the use of GEM in combination therapy for the re-sensitization and treatment of pancreatic cancer. Combination therapy of GEM and erlotinib was shown to increase the median survival of patients with metastatic pancreatic cancer by two weeks [28-30]. However, this treatment had substantial side effects, as well as high cost compared to the modest survival benefits [30, 31]. The multidrug combination of leucovorin, fluorouracil, irinotecan, and oxaliplatin also known as folfirinix, was found to increase the median survival of 4.3 months. Despite the improved median of survival, these treatments were found to have significantly higher levels of toxicity [32]. One of the more recent combination therapies is GEM plus nab-paclitaxel. This treatment increased the median survival by 1.87 months, increased the overall survival at 1 and 2 years, and had reasonable adverse effects [33]. Several of these combination therapies are currently being used for patients with good performance status based on their Eastern Cooperative Oncology Group (ECOG) performance status [30]. For metastatic pancreatic cancer patients with poor performance status, GEM (monotherapy) remains the recommended treatment. Therefore, despite advances in chemotherapeutics and the understanding of the biology of pancreatic cancer, there has been limited progress in therapy options for metastatic pancreatic cancer, and the need to evaluate other potential therapeutics continues. In the present study, we evaluated the combined drug effects of GEM and/or MET or I3C in pancreatic cancer cells of males and females. Several pancreatic cancer cell lines were examined for cell viability, drug interaction (summation and synergistic effect), and modulation of critical transporters, hENT1 and hENT4, expression when treated with GEM, MET, I3C in mono or GEM combination therapy for 24h and 72h. The female cell lines, ASPC1 and SU86.86 were found to exhibit synergistic drug behavior when treated with GEM and MET at both time points.

I3C, a compound found in Brassica vegetables such as broccoli, has been found to be well tolerated in high doses in both animals and humans [13, 34], and to be a promising nontoxic chemopreventive agent [9]. While potential use in humans has been shown to be feasible, concerns remain as to its tendency to induce activating enzymes, such as cytochrome P450s, and its promoting activity under uncertain experimental conditions [66]. Under certain experimental conditions, I3C can act as a tumor promoter in rodents [35, 36]. In this study we observed that a low dose of I3C (250 μ M) induced an increase in cell proliferation of pancreatic cancer cells after 24 h of treatment. Despite this concern, several studies have shown that I3C exhibit its anticancer properties in various cancer cells [37-39]. I3C

has inhibited tumorigenesis in breast, liver, lung, cervix, and gastrointestinal tract in different animal models [13, 14, 40-42]. Inhibition of tumorigenesis using I3C occurs via different mechanisms [43, 44]. The current study demonstrated that treatment with I3C resulted in a time- and concentration-dependent increase in inhibition of proliferation of MIA PaCa2, PANC1, ASPC1, and SU.86.86 pancreatic cancer cell lines, where male pancreatic cancer cells, MIA PaCa2 and PANC1, treated with low dose of I3C (250 μ M) exhibited the lowest cytotoxicity levels.

As part of this study, we were also interested in whether members of the hENT family of transporters, namely hENT1 and hENT4, play a role in the increased cytotoxicity of GEM combined therapies. To evaluate the role of hENTs in GEM combined therapies, we compared the current study to a previous study performed by our lab. The most noticeable difference between the previous study and the current study is the method of treatment. In the previous study, cells were initially pre-treated with I3C for 24 h and subsequently treated with both I3C and GEM [6]. Based on the observations from that study, pre-treatment with I3C followed by treatment with I3C and GEM not only significantly decreased pancreatic cancer cell viability by enhancing the efficacy of GEM but required less drug concentration to achieve this outcome [6]. The study also demonstrated I3C enhancement of GEM cytotoxicity in pancreatic cancer through the upregulation of hENT1, which is involved in the transport of GEM [6]. In the current study the pancreatic cancer cells were not pre-treated with I3C before subsequent combination drug therapy treatment. While both studies demonstrated that treatment with I3C and GEM could significantly decrease cell viability in pancreatic cancer cell line compared to GEM-only treatment, there was a difference in their regulation of hENT1. Pancreatic cancer cells treated with GEM and I3C in the current study did not have significantly upregulated hENT1 expression, with the exception of SU.86.86 cells after 72h treatment. GEM and I3C combined therapy also did not upregulate the expression of hENT4. This difference in treatment methods could account for differences in drug responses and hENT expression levels. We postulate that pre-treatment with I3C followed by subsequent combined therapy with GEM better sensitizes the pancreatic cancer cells to GEM therapy through one of several mechanisms. This may occur through several biological mechanisms, such as, the inhibition of cell proliferation through modulation of various proteins involved in induction of pro-apoptotic proteins, inhibition of anti-apoptotic proteins, the inhibition of signaling pathways involved in cell survival, the induction of cytochrome P450, and cell cycle regulation [12, 43]. In breast cancer it has previously been shown that I3C synergizes with tamoxifen to more stringently induce G1 cell cycle arrest through the additive inhibition of CDK2 kinase activity [45]. I3C has also been shown to down regulate of miR-21, which overexpression is linked to chemoresistance in pancreatic cancer [46], reactivate p16^{INK4a} tumor suppression gene through hypomethylation of the promoter [47], and a previous study performed by our lab showed that pre-treating with I3C enhanced the expression of hENT1 [6]. Consequently, pre-treatment of pancreatic cancer cells with I3C may potentially sensitize pancreatic cancer cells through many

of the above biological mechanisms which increase GEM efficacy.

Another therapeutic agent used in combination with GEM is MET. MET is a well-known inexpensive anti-diabetic agent that has been shown to have anticancer properties [48-52]. Studies have illustrated that the mortality rate from cancer was lower with diabetics patients taking MET (pre-treatment with MET) as part of their anti-diabetic medication compared to those not taking MET and to other insulin therapies [22, 53-56]. In this study, we demonstrate that MET monotherapy inhibition of cell proliferation in pancreatic cancer cells was cell line specific. The inhibition of cell proliferation in cancer cells using MET is supported by several studies, where MET has been shown to exert *in vitro* inhibition of proliferation in prostate, ovarian, colorectal and breast cancer cells [56-61], to significantly prevent pancreatic cancer development in hamsters [62], and to target pancreatic cancer stem cells [23]. In recent years, there has been an increase in the number of clinical trials investigating the involvement of MET in cancer. Most of these trials involve MET in combination therapy. In a double-blind, placebo-controlled phase II clinical trial of MET in pancreatic cancer treatment with a survival endpoint, there was no advantage for the addition of MET to erlotinib and GEM [26]. The use of MET and paclitaxel as a second line treatment for advanced pancreatic cancer was found to be poorly tolerated and the primary end point (disease control rate) was not met [63]. Some of the limitations noted in these clinical studies are self-reporting from diabetic patients, misclassification of patients, incomplete medication records, time-related bias studies, and small population size [19, 21, 53, 64, 65]. Therefore, studies interested in whether MET sensitize pancreatic cancer cells to GEM and enhances the capacity of GEM to inhibit invasion and proliferation of pancreatic cancer cells are continuing to be evaluated.

Conclusion

In this study, pancreatic cancer cells treated with GEM and MET demonstrated time-dependent decrease in cell viability. Overall the female cell lines, ASPC1 and SU.86.86, were found to exhibit synergistic drug behavior when treated with GEM and MET at both time points. Overall the efficacy of MET in combination with GEM treatment was cell line specific and may also be dependent on the method of treatment. MET may potentially have a similar drug response to I3C, where the treatment method may influence the cytotoxicity and hENT expression in pancreatic cancer cells. We propose that pre-treatment with MET may enhance cytotoxicity and hENT expression in pancreatic cells; however, further studies are needed to verify this hypothesis. Combination treatment of GEM and MET or I3C has the potential to enhance the ability of GEM and inhibit cell proliferation of pancreatic cancer cells through members of the hENT family of transporters. However, genetic variability in hENT drug transporters is also an important factor in gemcitabine response. In addition, MET targets a number of signaling pathways that play roles in cell proliferation and angiogenesis, such as mTOR. The overall effect of GEM combination, therapy in pancreatic cancer cells may be cell line- and treatment method-specific, where pre-treatment with I3C or MET

better sensitizes pancreatic cancer cells to GEM. To better understand the impact of treatment method, a detailed time-, concentration- and treatment method-dependent matched study will have to be performed to confirm these initial findings.

Competing interests

The authors declare that they have no competing interests.

Funding

This study was funded by the FDA National Center for Toxicological Research.

Acknowledgement

SJ was supported by an appointment to the Postgraduate Research Program at the National Center for Toxicological Research administered by the Oak Ridge Institute for Science Education through an interagency agreement between the U.S. Department of Energy and the U.S. Food and Drug Administration. The views presented in this report do not necessarily reflect those of the US Food and Drug Administration.

Supplementary data

Supplementary data associated with this article can be found, at <http://nobleresearch.org/Doi/10.14312/2052-4994.2018-2>.

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