Targeting the oncogenic protein beta-catenin to enhance chemotherapy outcome against solid human cancers

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Abstract

Background: Beta-catenin is a multifunctional oncogenic protein that contributes fundamentally to cell development and biology. Elevation in expression and activity of β-catenin has been implicated in many cancers and associated with poor prognosis. Beta-catenin is degraded in the cytoplasm by glycogen synthase kinase 3 beta (GSK-3β) through phosphorylation. Cell growth and proliferation is associated with β-catenin translocation from the cytoplasm into the nucleus. This laboratory was the first to demonstrate that selenium-containing compounds can enhance the efficacy and cytotoxicity of anticancer drugs in several preclinical xenograft models. These data provided the basis to identify mechanism of selenium action focusing on β-catenin as a target. This study was designed to: (1) determine whether pharmacological doses of methylseleninic acid (MSeA) have inhibitory effects on the level and the oncogenic activity of β-catenin, (2) investigate the kinetics and the mechanism of β-catenin inhibition, and (3) confirm that inhibition of β-catenin would lead to enhanced cytotoxicity of standard chemotherapeutic drugs.

Results: In six human cancer cell lines, the inhibition of total and nuclear expression of β-catenin by MSeA was dose and time dependent. The involvement of GSK-3β in the degradation of β-catenin was cell type dependent (GSK-3β-dependent in HT-29, whereas GSK-3β-independent in HCT-8). However, the pronounced inhibition of β-catenin by MSeA was independent of various drug treatments and was not reversed after combination therapy. Knockout of β-catenin by ShRNA and its inhibition by MSeA yielded similar enhancement of cytotoxicity of anticancer drugs. Collectively, the generated data demonstrate that β-catenin is a target of MSeA and its inhibition resulted in enhanced cytotoxicity of chemotherapeutic drugs.

Conclusions: This study demonstrates that β-catenin, a molecule associated with drug resistance, is a target of selenium and its inhibition is associated with increased multiple drugs cytotoxicity in various human cancers. Further, degradation of β-catenin by GSK-3β is not a general mechanism but is cell type dependent.

Background

Beta-catenin protein is a vital component of the canonical Wnt/β-catenin signaling pathway, which is described as an oncogenic cause in many human cancers [1]. In head and neck squamous cell carcinomas (HNSCC), over expression of the Wnt/β-catenin signaling pathway increases cell survival and invasion [2]. The higher β-catenin expression in HNSCC patients, the more advanced stage [3] and poor prognosis are observed [4]. Mutations in the gene that encodes β-catenin (CTNNB1) [5] and elevated nuclear β-catenin [6] were implicated in prostate cancers (CaP).

Over 90% of colorectal cancers (CRC) demonstrate a deregulated Wnt/β-catenin signaling pathway [7]. Published studies suggest that unregulated β-catenin, overlapping with adenomatous polyposis coli (APC) mutation, is associated with the initiation of CRC [8-10]. Beta-catenin is expressed in the cytoplasm and the nucleus. The cytoplasm β-catenin, as a component of adherens junctions (AJs) [11], is an essential element of...
cell-to-cell adhesion and stability. The level of cytoplasmic β-catenin is controlled by the activity of a destruction complex that consists of axin, glycogen synthase kinase 3β (GSK-3β) and APC [12-15]. In the absence of Wnt signaling, the complex is assembled and GSK-3β phosphorylates and consequently degrades cytoplasmic β-catenin [14,15]. However, GSK-3β is inactivated in cancer cells by phosphorylation at serine 9, a similar mechanism of GSK-3β inhibition by lithium [16,17]. In the presence of Wnt signaling, β-catenin destruction complex is disassembled by removing axin [18,19] resulting in β-catenin accumulation in the cytoplasm. The accumulated cytoplasmic β-catenin hence enters the nucleus to initiate its oncogenic function. The nuclear β-catenin has an important function in many human malignancies [1] by stimulating cell growth and proliferation. The nuclear β-catenin affects TCF/LEF family transcription factors [20,21] and consequently activates oncogenes such as cyclin D1 [22,23], Myc [24] and many other downstream targets. The nuclear accumulation of β-catenin is a critical step in the activation process of the canonical Wnt signaling pathway and is associated with poor prognosis in cancer patients [25].

In addition to its role in cell growth and adhesion, activated canonical Wnt/β-catenin signaling pathway is linked to cancer stem cells [26,27] that contribute to tumor bulk, recurrence and resistance to chemotherapy. Accordingly, β-catenin inhibitors in combination with standard systemic therapies hold great promise to improve treatment’s efficacy and outcome.

The response rates of combination regimen of irinotecan and 5-fluorouracil/leucovorin (5-FU/LV) is 39% in metastatic CRC [28]. Treatment with oxaliplatin and 5-FU/LV has improved the response rate to 50.7% in CRC [29]. Treatment with docetaxel and prednisone against metastatic CaP resulted in a median survival of 19.2 months [30]. Docetaxel in combination with cisplatin and 5-FU against inoperable advanced HN SSCC resulted in a median progression free survival of 11 months [31]. Although the relative survival in advanced solid tumors is improved by using systemic therapy, the current chemotherapy cure rates are limited. Thus, the development of new regimens is greatly needed to achieve a better clinical outcome.

In our preclinical models, selenium-containing compounds enhanced the efficacy of multiple chemotherapeutic agents (CPT-11 or docetaxel) against various types of cancers (colorectal, head and neck and prostate) [32,33]. In mice bearing human colorectal cancer xenografts (HCT-8), combination treatment of MSC and irinotecan resulted in complete tumor regression (100% CR) that was not observed with each drug alone (30% CR) [32]. Sequential combination of MSeA and docetaxel resulted in synergy enhancing docetaxel-induced cell death in CaP [33]. Multiple mechanisms of the synergy between selenium and other chemotherapeutic agents are proposed. Selenium (Se) is an essential element that possesses antioxidant properties in a form of selenoproteins protecting cells from harmful free radicals [34-36]. The effect of selenium on β-catenin has yet to be investigated.

This study is designed mainly to determine if β-catenin is a target of MSeA in CRC, HNSCC and CaP cancers; and to evaluate the role of GSK-3β in the degradation of β-catenin and if such an effect is associated with enhanced cytotoxicity of anticancer drugs.

**Results**

**Inhibition of β-catenin by MSeA is concentration, time and tumor type dependent**

To evaluate the effect of MSeA on the expression of β-catenin, various tumor cell types were treated with multiple time and doses of MSeA. In all treated cells, MSeA decreased the expression of β-catenin in dose and time dependent manners (Figure 1). The data in Figure 1 indicate that the down regulation of β-catenin is MSeA concentration dependent. In CRC cells (HCT-8 and HT-29), 24 h treatment with 5 μM resulted in completely depletion of β-catenin. In contrast, in HNSCC cells (FaDu and A253), the decrease in β-catenin levels in FaDu cells was achieved with lower concentrations of MSeA (0.5 μM) than in A253 (5 μM) (Figure 1A). In the androgen-independent CaP cells (PC3 and C42), inhibition of β-catenin by MSeA required a high concentration (5 μM) (Figure 1A). The kinetics of β-catenin inhibition by MSeA appears to be tumor type dependent, early in HCT-8, C42, HT-29 and PC3, and late event in FaDu and A253 (Figure 1B).

**MSeA inhibits β-catenin nuclear expression**

To determine whether MSeA down regulates the activity of β-catenin, nuclear and cytoplasmic extracts of colorectal cancer cells were tested for the level of β-catenin before and after MSeA treatment. The data in Figure 2A indicate that β-catenin is predominantly expressed in the nucleus of untreated CRC cells (HCT-8 and HT-29) indicating activation of β-catenin. Treatments with MSeA resulted in inhibition of nuclear expression of β-catenin (Figure 2B). These data suggest that MSeA down regulates β-catenin activity through inhibiting its nuclear expression.

**Inhibition of β-catenin is due to enhanced degradation**

To determine whether the observed down regulation of β-catenin by MSeA results from inhibition of its synthesis or from increased degradation, cells were treated with MSeA and cycloheximide (an inhibitor of de novo protein synthesis [37]) alone and in combination.
**Figure 1** MSeA effect on intracellular expression of β-catenin. Six human cancer cell lines including 2 colorectal (HT-29 and HCT-8), 2 head and neck (FaDu and A253) and 2 prostate (PC3 and C42) were treated with various doses of MSeA (panel 1A) and various times (panel 1B). Western blot analyses show that MSeA inhibits intracellular level of β-catenin in dose and time dependent manners.
Treatments for up to 30 minutes did not affect the expression level of β-catenin (Figure 3A). However, β-catenin is down regulated after 24 h treatment with MSeA alone and in combination with cycloheximide (Figure 3B). These data suggest that the inhibition of β-catenin by MSeA is the result of increased degradation.

The role of GSK-3β in β-catenin degradation is cell type dependent

To determine the mechanism of β-catenin degradation by MSeA, the role of GSK-3β was evaluated in HT-29 and HCT-8 (Figure 4). Treatments with MSeA had no significant effect on the level of total GSK-3β (in HT-29 or HCT-8) and on the level of phosphorylated GSK-3β in HCT-8. In contrast, phosphorylated GSK-3β was significantly decreased in HT-29 (Figure 4A).

To evaluate further the role of GSK-3β, cells were treated with lithium chloride (LiCl) alone (GSK-3β inhibitor [38]) and in combination with MSeA. Treatment with LiCl increased the level of phosphorylated GSK-3β in both cell lines indicating inhibition of GSK-3β (Figure 4B). Combination of MSeA with various doses of LiCl resulted in reversing the down regulation of β-catenin in HT-29 cells but not HCT-8 by MSeA (Figure 4B). Data in Figure 4 demonstrated that the inhibition of β-catenin by MSeA is GSK-3β phosphorylation dependent in HT-29 but independent in HCT-8.

The effect of MSeA in combination therapy on the level and activity of β-catenin

To determine whether MSeA in combination with a chemotherapy agent would affect β-catenin expression and activity, cells were treated with MSeA ± chemotherapeutic agent and analyzed for total and nuclear β-catenin expression. The data in Figure 5A indicate that MSeA inhibited β-catenin expression in all cell lines but neither 7-Ethyl-10-Hydroxycamptothecin (SN-38, the active metabolite of irinotecan) nor docetaxel alone had an effect on the β-catenin levels. Adding SN-38 or docetaxel to MSeA did not interfere with selenium inhibition of β-catenin. The combination of MSeA/SN-38 resulted in even more observed inhibition of total β-catenin when compared with MSeA alone in HT-29, HCT-8 and FaDu cells. In CaP cells, MSeA alone and in combination with docetaxel have similar inhibitory effect on the expression of β-catenin.
β-catenin (Figure 5A). To determine whether the inhibition of β-catenin by MSeA in combination with SN-38 is due to inhibition of the nuclear expression, nuclear extracts of CRC cells were treated with MSeA alone and in combination with SN-38 and evaluated for the level of β-catenin. The combination treatment of MSeA/SN-38 resulted in down regulation of the nuclear expression of β-catenin in HT-29 and HCT-8 cells (Figure 5B). These results indicate that the activity of β-catenin is decreased after the combination therapy.
The inhibition of β-catenin expression by ShRNA or MSeA is associated with enhancement of drug-induced growth inhibition of tumor cell

To determine whether inhibition of β-catenin can be correlated with enhanced efficacy of chemotherapy, β-catenin in tumor cells was knocked down by specific ShRNA. Two β-catenin ShRNA transfectant clones (HCT-8RH7 and HCT-8RF4) that demonstrated inhibition of β-catenin expression when compared with scrambled control (HCT-8SC) and wild type (HCT-8WT) were selected for further testing (Figure 6A).

To evaluate the effect of silencing β-catenin on cell growth, HCT-8WT, HCT-8SC and HCT-8R (HCT-8RH7 and HCT-8RF4) transfectants were treated with various classes of chemotherapeutic agents. Treatments with 0.5 μM of SN-38 were more effective (p < 0.05) against HCT-8R (~50% cell growth inhibition) when compared with all other groups (15% cell growth inhibition).

Figure 5 Combination treatment effect on the expression of β-catenin. Cancer cells (panel 5A) and nuclear extraction of colorectal cancer cells (panel 5B) were treated with MSeA, SN-38 and docetaxel alone or in combination. Expression of β-catenin was determined using western blots. Combination treatments of MSeA/SN-38 continued to decrease expression of β-catenin when compared with all other groups. Combination treatments of MSeA/docetaxel did not reverse the inhibitory effect of MSeA alone on expression of β-catenin (panel 5A). The active form (nuclear expression) of β-catenin was inhibited by the combination treatment of MSeA/SN-38 when compared with other group (panel 5B).
Figure 6 The effect of various drug treatments on cell growth and proliferation in β-catenin knockout cells. Beta-catenin was silenced in colorectal cancer cells (HCT-8, panel 6A). HCT-8 wild type cells (HCT-8WT), scramble controls (HCT-8SC) and 2 β-catenin knockout clones (HCT-8RH1 and HCT-8RF4) were tested. Expression of β-catenin was significantly lower in HCT-8RH1 and HCT-8RF4 when compared with all other groups (panel 6A). Cancer cell growth and proliferation were determined in HCT-8WT, HCT-8SC, HCT-8RH7 and HCT-8RF4 after treatment with various doses of SN-38, docetaxel, paclitaxel, oxaliplatin, 5-FU and topotecan using SRB assay. HCT-8RH7 and HCT-8RF4 cells were significantly more sensitive to various drug treatments when compared with HCT-8WT and HCT-8SC (panel 6B). * denotes a p value of less than 0.05 when compared with all other groups.
inhibition in HCT-8SC or HCT-8WT, Figure 6B). Other
doses of SN-38 resulted in similar patterns of inhibition
cell growth (Figure 6B). In similar fashion, treatments
with various doses of docetaxel, paclitaxel, oxaliplatin, 5-
FU and topotecan significantly enhanced efficacy against
cell growth of HCT-8RH7 and HCT-8RF4 when com-
pared with all other groups of HCT-8SC or HCT-8WT
(Figure 6B).

To confirm further that inhibition of β-catenin is asso-
ciated with enhanced cytotoxicity of anticancer drugs,
tumor cells were treated with MSeA alone and in com-
bination with SN-38 and the results were correlated
with the levels of β-catenin (table 1). The data in table 1
demonstrated a relationship between enhanced cytotoxi-
city of SN-38 and inhibition of β-catenin by MSeA or
ShRNA. Thus, these data support the initial hypothesis
that inhibition of β-catenin by MSeA is a critical deter-
minant of drug response.

Discussion
Beta-catenin oncogenic protein is widely expressed in
many human malignancies [1] including HNSCC [2-4],
CaP [5,6] and CRC [8-10]. Beta-catenin is involved in
cell growth [22-24], adhesion [11] and stemness [26,27].
Beta-catenin is found in multiple cellular locations
including intracellular membrane, cytoplasm and
nucleus. The nuclear accumulation of β-catenin indi-
cates the activation of its oncogenic form that stimulates
transcription factors and genes [22-24] leading to
enhanced tumor cell growth and poor prognosis [25].
The hypothesis of this study is that β-catenin is a target
of MSeA and its inhibition would translate into
enhanced drug effect. Our results (Figure 1) established
that MSeA is a potent inhibitor of β-catenin in various
cancer types. This broad inhibitory effect of MSeA on
the expression of β-catenin is pivotal for explaining the
established synergy between selenium and various che-
motherapeutic agents against multiple cancers.

The data in Figure 2 demonstrate that the inhibition
of β-catenin level after MSeA is due to the inhibition of
the active form of β-catenin in the nucleus. Thus, these
data indicate that pharmacologic doses of MSeA offer
effective inhibition of β-catenin activation.

Recent findings by Zhang et al demonstrate that sele-
nium effect against esophageal squamous cell carcinoma
is correlated with its inhibition on β-catenin/TCF path-
way [39]. Our results confirm this finding in various
human cancers and prove that the decreased level of
β-catenin is associated with enhanced efficacy of various
classes of chemotherapy. Thus, indicating the impor-
tance of β-catenin inhibition in drug response.

To determine whether the inhibition of β-catenin is
due to a decrease in synthesis or an increase in degrada-
tion, de novo protein synthesis was inhibited using CHX
in the presence and absences of MSeA. The data in
Figure 3 indicate that MSeA inhibition of β-catenin is
due to increase in degradation but not decrease in synth-
esis in both CRC cells (HT-29 and HCT-8). Many studies
have showed that cytoplasm β-catenin is degraded by an
axin/GSK-3β/APC complex [12-14] and the degradation
is a GSK-3β phosphorylation dependent process [14,15].
The degradation of the cytoplasm β-catenin prevents its
accumulation and translocation into the nucleus. Data in
Figure 4 demonstrated that the inhibition of β-catenin by
MSeA is GSK-3β dependent phosphorylation in HT-29
but independent in HCT-8. The GSK-3β independent
degradation of β-catenin is a novel finding in HCT-8
cells and indicates that the MSeA effect involves other
signaling pathways than Wnt/β-catenin, which will be
investigated in future studies.

In preclinical models, sequential combination treat-
ment of selenium compounds (MSC, SLM or MSeA)
and various chemotherapeutic agents (SN-38 or doc-
etaxel) were proven synergistic against various cancers
including HNSCC, CRC and CaP [32,33]. Studies were
carried out to determine whether the combination treat-
mant of MSeA and chemotherapeutic agents affect
β-catenin level in those cell lines. Our results in Figure 5
showed that treatment with MSeA in combination with
SN-38 or docetaxel down regulated the total and the
nuclear β-catenin. These results confirm that the
chemotherapeutic agent did not interfere with selenium
inhibition of the level and activity of β-catenin. However,
neither SN-38 nor docetaxel alone affect the expression
level of β-catenin (Figure 5).

To determine that inhibition of β-catenin by MSeA
will translate into enhanced drug-cytotoxicity, cells
knocked down β-catenin by ShRNA were more sensitive
to growth inhibition by SN-38 than wild type. Collect-
ively, this study indicates that the decreased level of
β-catenin is associated with enhancement of drug
induced inhibition of cell growth. (table 1).

Further, the data in Figure 6 indicate that silencing
β-catenin increases the cytotoxicity of various chemother-
apeutic agents. The efficacy of SN-38, docetaxel, paclitaxel,
oxaliplatin, 5-FU and topotecan was significantly increased in HCT-8R when compared with control groups (Figure 6B).

Conclusions
These results support the hypothesis that β-catenin is a target of MSeA and its inhibition results in enhanced drug-cytotoxicity in multiple cancers. Degradation of β-catenin by GSK-3β is not a general mechanism but is cell type dependent.

Although Selenium is a multi-target agent [32,33,40], inhibition of β-catenin is a critical determinant of drug response. These preclinical results provided the rationale for validation of this new and innovative approach in a clinical setting.

Materials and methods
Cell lines and drugs
Human cancer cell lines of colorectal (HCT-8 and HT-29), head and neck (FaDu and A253) and prostate (PC3 and C42) were purchased from American type cell culture (ATCC, Manassas, VA) and maintained in RPMI 1640 with 10% fetal bovine serum (FBS). The cell lines were tested regularly using Stratogene mycoplasma plus PCR Primer set (La Jolla, CA) and they were free from Mycoplasma. SN-38, docetaxel, 5-FU, paclitaxel, oxaliplatin, lithium chloride (LiCl) and cycloheximide (CHX) were purchased from Sigma Aldrich (St. Louis, MO). MSeA (CH₃SeO₂H) was purchased from PharmaSe Inc. (Lubbock, TX). Topotecan was obtained from GlaxoSmithKline (Durham, NC). Puromycin dihydrochloride was used at concentration of 5 μM alone or in combination with MSeA (5 μM). Puromycin dihydrochloride was used at concentration of 20 μM for clones’ selection.

Preparation of cytoplasm and nuclear extract
Cytoplasm and nuclear extracts were prepared as previously described [41]. Briefly, to obtain cytoplasm extract, untreated and treated HCT-8 cells were harvested and suspended in lysis buffer (0.08 M KCl, 35 mM HEPES, pH 7.4, 5 mM potassium phosphate, pH 7.4, 5 mM MgCl₂, 25 mM CaCl₂, 0.15 M sucrose, 2 mM PMSF, 8 mM dithiothreitol). After overnight storage at -80°C, cells were passed through a 28-gauge needle, centrifuged and the supernatant collected to represent the cytoplasm extract. The remaining pellet was re-suspended in lysis buffer, sonicated, centrifuged and the supernatant collected to represent the nuclear extract.

Silencing the expression of β-catenin
HCT-8 cells were utilized to generate a stable transfection using small hairpin β-catenin RNA (β-catenin shRNA) purchased from Santa Cruz Biotechnology Inc. (Santa Cruz, CA). Transfection was carried out following the manufacture’s instructions. Briefly, HCT-8 cells were plated 5 × 10⁵ cells/well (6-well plate) one day before the transfection. A well with 70-80% cells confluence was transfected with control shRNA Plasmid-A (a negative control) that encodes a scrambled shRNA sequence that does not inhibit β-catenin to generate HCT-8 scrambled control (HCT-8SC). Another well with the same cells confluence, was transfected with β-catenin shRNA plasmid DNA, a β-catenin-specific lentiviral vector plasmid to knock down expression and generate HCT-8 recombinant clones (HCT-8R). Clones of stable transfectants were selected using 20 μM of puromycin dihydrochloride. After selection, 10 individual clones were evaluated using western blots and the 2 clones (HCT-8RH7 and HCT-8RF4) that demonstrated the most effect of survivin suppression were selected for further studies.

Western blots analyses
Western blots were performed as described previously [33] to determine the effects on the intracellular protein levels. Briefly, untreated and treated cells were collected and digested using RIPA buffer (1 M Tris, 1 M NaCl, Triton X-100 and distilled water) with fresh protease inhibitor cocktail. Protein level was measured using Bio-Rad DC protein assay and a synergy HT spectrophotometer (BioTek Instruments, Winooski, VT). Equal amount of protein (50 μg) was loaded on 4-20% SDS-PAGE. After transfer, nitrocellulose membrane was

Cycloheximide (CHX) was applied for 5, 10, 20, 30 minutes and 24 hours at a nontoxic concentration of 100 μM alone or in combination with MSeA (5 μM). Puromycin dihydrochloride was used at concentration of 20 μM for clones’ selection.
rinsed with PBS-T, blocked with 5% milk and hybridized with the selected antibody. The following primary antibodies were used: anti-β-catenin, anti-GSK-3β (BD Biosciences, San Jose, CA) and anti-p-GSK-3β (cell signaling technology, Danvers, MA). The following secondary antibodies were used: goat anti-mouse IgG and goat anti-rabbit IgG (Santa Cruz Biotechnology, Santa Cruz, CA). After incubation with the primary and secondary antibodies, membrane was rinsed and incubated with chemiluminescence or enhanced chemiluminescence technology, Danvers, MA). The following secondary bodies were used: anti-

Cell growth assay

Cell growth was evaluated using sulforhodamine B (SRB) assay as previously described and performed [33]. Briefly, after drug treatment, HCT-8WT, HCT-8SC and HCT-8RH7 and HCT-8RF4 cells were incubated in a drug-free medium for 5 days, fixed, washed and stained with SRB dye. The optical density of bound dye was measured at 570 nm using synergy HT multi-mode microplate reader (BioTek Instruments, Winooski, VT).

Statistical analyses

Each experiment has been repeated at least 3 times. Values were presented as the mean plus or minus standard deviation. Statistical analyses were performed comparing all treatments groups using unpaired t-student test. Significant difference between groups was noted when the p value was less than 0.05.

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Authors’ contributions

MSS performed and designed experiments, prepared and wrote the manuscript. DRR performed cytotoxicity experiments. YMR participated in study design, data interpretation and preparation of the manuscript. RGA designed the research strategy, supervised the project, assisted in data generation, results interpretation and correction of the manuscript. All authors read and approved the final manuscript.

Competing interests

The authors declare that they have no competing interests.

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