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## Effects of cucurbitacins on cell morphology are associated with sensitization of renal carcinoma cells to TRAIL-induced apoptosis

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### Abstract

Cucurbitacins B and D were among the compounds identified as sensitizers of cancer cells to TRAIL-mediated apoptosis in a high-throughput screen. Therefore a series of cucurbitacins was further investigated for TRAIL sensitization and possible mechanisms of action. A total of six cucurbitacins promoted TRAIL-induced apoptosis (B, I, E, C, D, and K) and one (P) was inactive. Sensitization of renal adenocarcinoma cells to TRAIL was apparent after as little as 1–4 h

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pretreatment and did not require continued presence of cucurbitacin. Active cucurbitacins induced caspase-8 activation only after subsequent TRAIL addition and caspase activation was required for apoptosis suggesting amplified proximal signaling from TRAIL death receptors. Cucurbitacin-sensitized TRAIL-induced cytotoxicity was inhibited by *N*-acetyl cysteine. Structure–activity relationship analysis in comparison to published studies suggests that TRAIL-sensitizing and general cytotoxic activities of cucurbitacins may be decoupled. Cucurbitacins are reported to be inhibitors of STAT3 activation. However, their TRAIL-sensitizing activity is STAT3-independent. Treatment of renal carcinoma cells with active cucurbitacins produced rapid and dramatic changes in cell morphology and cytoskeletal organization (also prevented by NAC). Therefore, cucurbitacins may be useful as tools for investigating the molecular mechanism(s) of action of TRAIL sensitizers, particularly with regard to temporal aspects of sensitization and modulation of TRAIL signaling by cell morphology, and could form the basis for future therapeutic development in combination with TRAIL death receptor agonists.

## Keywords

Cucurbitacins; TRAIL; TRAIL sensitizers; Apoptosis; STAT3; Cell morphology

## Introduction

Induction of cancer cell-specific apoptosis via activation of TRAIL (tumor necrosis factor-related apoptosis-inducing ligand) signaling has become an important focus of cancer research [1–5]. Given the relatively common occurrence of TRAIL resistance in many types of cancer [6–9], the search for enhancers of TRAIL-induced cell killing is now widespread (see reviews cited above). This has resulted in the identification of many compounds with a very wide range of putative molecular mechanisms of action [1]. A recent high-throughput screening campaign utilizing the relatively TRAIL-resistant renal cell carcinoma line ACHN resulted in the identification of a number of synergistic TRAIL sensitizers, including cucurbitacins B and D [10]. Cucurbitacins, members of a group of tetracyclic triterpenoids produced by plants of the Cucurbitaceae family, have long been reported to have anticancer effects via several potential mechanisms of action [11]. Cucurbitacins are particularly well-characterized in terms of their ability to modulate JAK2/STAT3 signaling [11–13] and cucurbitacin B has recently been reported to activate generation of reactive oxygen species (ROS) in cells [14]. JAK2/STAT3 signal modulators, including cucurbitacin I [15], sorafenib [16], and tyrphostin AG490 [17, 18] have been reported to enhance TRAIL-induced cell killing in a variety of cancer cells. Similarly, certain other TRAIL sensitizers have been reported to require generation of ROS for activity [5, 19–21]. In order to further investigate the effects of cucurbitacins, a series of seven compounds were evaluated using renal carcinoma cells for their TRAIL sensitizing activity, effects on STAT3 activation, sensitivity to the ROS modulator *N*-acetyl cysteine (NAC) and effects on cell morphology.

## Materials and methods

### Chemicals and reagents

Cucurbitacins were obtained from the NCI Developmental Therapeutics Program (DTP) and/or from Chromadex (Irvine, CA—cucurbitacins B, D, E, and I only). 2,3-Bis (2-methoxy-4-nitro-5-sulfophenyl)-5-[(phenylamino)carbonyl]-2H-tetrazolium hydroxide (XTT; NSC 601519) was provided by the Drug Synthesis and Chemistry Branch, DTP/NCI (Frederick, MD). Bortezomib was purchased from the National Institutes of Health Pharmacy. Recombinant TRAIL ligand (168 amino acid TNF homologous extracellular domain) was purchased from Peprotech, Inc. (Rocky Hill, NJ). Cell culture media and additives were obtained from Cellgro (Manassas, VA), Hyclone (Logan, UT), Sigma (St.

Louis, MO) or Invitrogen (Carlsbad, CA). BCA protein assay kits were obtained from Pierce/Thermo (Rockford, IL). Other chemicals and reagents were obtained from Sigma (St. Louis, MO). Chemical structures were drawn using ChemDraw (CambridgeSoft Corp., Cambridge, MA) using structural information from the PubChem database (<http://pubchem.ncbi.nlm.nih.gov/>).

### **Antibodies, western blot reagents**

Anti-cleaved caspase-8 (18C8 rabbit monoclonal), and anti-phosphopaxillin (Tyr118) were obtained from Cell Signaling Technology (Danvers, MA). Anti-GAPDH (G8795 mouse monoclonal), conjugated with horse radish peroxidase, was from Sigma. Appropriate secondary antibodies conjugated with horse radish peroxidase were purchased from Thermo Scientific (Rockford, IL).

### **Cells and cell culture conditions**

ACHN and Caki-1 renal ACHN cells (National Cancer Institute, Frederick, MD) were maintained in culture and prepared for assays as described [10].

### **Cytotoxicity/TRAIL sensitization assays**

ACHN cells were trypsinized, transferred to clear 384-well tissue culture treated plates (BD Biosciences; San Jose, CA) at 3,500 cells per well in clear (phenol red-free) medium and allowed to attach overnight. Compounds were diluted into medium (at 10–20× final concentration) and added to plates containing ACHN cells. After 1–4 h incubation, TRAIL (40 ng/ml final concentration) or medium control was added. After 18–20 h, cell numbers were assessed using an XTT assay as described [10]. Cell numbers from the XTT assay were normalized to DMSO-treated controls. For the wash-out experiment, compound-containing medium was removed and the cells were washed (45 µl/well medium) before addition of fresh medium with or without TRAIL. In order to investigate the effect of ROS on sensitization, NAC (10 mM final concentration) was added to cells just before addition of compounds. NAC remained in the plates throughout the experiment. After 4 h with compounds followed by 18–20 h with (or without) TRAIL, cell numbers were estimated using the XTT assay. In this case, medium was removed and the cells washed with fresh medium before XTT addition in order to eliminate interference in the assay due to the NAC reagent. In experiments assessing the effects of the caspase inhibitor ZVAD-FMK and the inactive caspase inhibitor control ZFA-FMK, cell numbers were assessed using the Promega (Madison, WI) MTS assay and normalized to the ZVAD-FMK only or ZFAFMK only as appropriate. In some experiments, cell number was estimated using the CellTiter-Glo Luminescent Cell Viability Assay (Promega) following the manufacturer's instructions.

### **Caspase-8 activation assays**

The caspase-8 assay was performed, as previously described [10]. Briefly, cells were seeded at 7,000 cells/well in white luminescence tissue culture-treated 384-well plates. Alternatively in some experiments, ACHN cells were added to 96-well white luminescence tissue culture plates as previously described [22]. After treatment with cucurbitacins for various periods, TRAIL was added. After different time points, cells were lysed and caspase-8 activity was determined using the Caspase-Glo™ 8 Assay kit from Promega (Madison, WI) according to the manufacturer's instructions. This assay included MG132 in order to reduce background signal due to proteasome activity in the cells. The correlation of the luminescence assay with caspase-8 activation by western blotting analysis was previously demonstrated for these cells [10, 22].

## Western blot

ACHN cells were seeded in 6-well plates at  $2.5 \times 10^6$  cells/well. After overnight attachment, compounds were added followed by 4 h treatment at which point cells were lysed, proteins separated by SDS-PAGE (4–12% gels, 50  $\mu$ g protein/well), and transferred to PVDF membranes as described [10]. Lysis buffer was additionally supplemented with Halt Protease and Phosphatase Inhibitor (Pierce/Thermo) for phosphoSTAT3 blots. Western blots for detection of cleaved caspase-8 were prepared and analyzed as described previously [10, 22].

## Cell morphology

Changes in cell shape were observed under standard bright field microscopy. For changes in cytoskeleton, cells were fixed with 4% paraformaldehyde and permeabilized with 0.1% Triton X-100 followed by fluorescein-phalloidin (Invitrogen) to detect filamentous (F)-actin. Confocal images were collected on a Zeiss LSM 510 confocal microscope (Carl Zeiss MicroImaging GmbH, Jena, Germany).

## Data presentation

Unless otherwise noted, values reported represent average  $\pm$  standard deviation. Statistical significance was estimated by applying the Students *t* test.

## Results

### Cucurbitacins sensitize ACHN cells to TRAIL

A series of seven cucurbitacins were assessed using the previously described screening assay for the detection of compounds able to sensitize TRAIL-resistant ACHN renal carcinoma cells to TRAIL-induced apoptosis [10]. Results are shown in Fig. 1. Six cucurbitacins showed significant activity when combined with TRAIL, although all six also exhibited some toxicity in the absence of TRAIL. One, cucurbitacin P, had no significant effect on ACHN cells in the presence or absence of TRAIL. Half-maximal growth inhibition values for cucurbitacins in the presence of TRAIL were calculated from the data in Fig. 1 and are listed in Table 1. Cucurbitacin B, the most potent of the cucurbitacins as a TRAIL sensitizer and the cucurbitacin most commonly found in the literature, was also used in several additional experiments. Cucurbitacin also sensitized a number of other human cancer cells to TRAIL-induced apoptosis. A total of 20 cell lines (including ACHN) were tested, including six renal (in addition to ACHN), five breast, four colon, and four melanoma cell lines (see Supplemental Data).

Similar effects were observed when cucurbitacin pretreatment time was reduced from 4 to 1 h. Additionally, a much smaller, but detectable, TRAIL-specific reduction in cell numbers was observed even when cells were only treated with TRAIL for 4 h (rather than 20 h) after 4 h cucurbitacin pretreatment (data not shown). In all of these experiments, the cucurbitacins remained present throughout the TRAIL treatment. In order to assess whether the effects of the cucurbitacins on TRAIL sensitivity required the continued presence of cucurbitacin, a series of washout experiments were performed. After pretreatment of cells for 1 or 4 h with cucurbitacins (1–10  $\mu$ M—concentrations chosen based on their relative activities as TRAIL sensitizers), the compounds were removed, cells washed with medium, and treated in the presence or absence of TRAIL for 4 or 20 h. Figure 2 shows that after 1 h (A) or 4 h (B) cucurbitacin treatment, the TRAIL sensitization effect persisted even after removal of the compounds (followed by TRAIL for 20 h). In fact, even after only 1 h cucurbitacin followed by as little as 4 h of TRAIL treatment, sensitization could be observed (Fig. 2c).

### **Sensitization of ACHN cells by cucurbitacins leads to enhanced TRAIL-induced caspase-8 activation**

As previously determined for other TRAIL sensitizing compounds, including cucurbitacins B and D [10], the cucurbitacins were assessed for their ability to enhance TRAIL-dependent activation of caspase-8 (Fig. 3a). Cucurbitacins B, C, D, E, I, and K significantly increased caspase-8 activation in the presence of TRAIL. The inactive cucurbitacin P had no effect. None of the cucurbitacins tested increased caspase-8 activity in the absence of TRAIL. In each case (except P), cucurbitacins in the presence of TRAIL increased caspase-8 activity to 2.07–2.75 times the level induced by TRAIL alone (Fig. 3a). Consistent with the sensitization results (Fig. 2a, c), only a 1 h treatment with cucurbitacin B was necessary to observe a subsequent enhancement of TRAIL-induced caspase-8 activation (Fig. 3b). This enhanced TRAIL-induced caspase-8 activation by cucurbitacin B was confirmed by western blot analysis using an antibody specific for activated (i.e., cleaved) caspase-8 (Fig. 3c).

### **TRAIL-induced killing of cucurbitacin-sensitized cells is caspase-dependent**

Inclusion of the general caspase inhibitor ZVAD-FMK eliminated the reduction in cell numbers in response to the combination of cucurbitacin B and TRAIL compared to the inactive control inhibitor, ZFA-FMK (Fig. 3d). Interestingly, the reduction in cell number by cucurbitacin B alone was not affected by ZVAD-FMK, suggesting that cucurbitacin B alone may work via a non-apoptotic cell death mechanism or perhaps causes cytostasis rather than cytotoxicity.

### **TRAIL-induced killing of cucurbitacin-sensitized cells is eliminated by *N*-acetyl cysteine**

The redox state of a cell can affect STAT3 activity [23] and ROS generation has been reported to specifically mediate some of the effects of cucurbitacins [14]. Therefore, the effects of cucurbitacins on TRAIL-induced cell killing were assessed in the presence and absence of the ROS inhibitor NAC. Addition of NAC at 10 mM (final) eliminated the effects of all of the cucurbitacins (Fig. 4). By contrast, the TRAIL sensitizing effect of the proteasome inhibitor bortezomib was unaffected by NAC (Fig. 4). However, significant generation of ROS following cucurbitacin B treatment (with or without TRAIL) was not detected using MitoSOX or DCF for assessment of mitochondrial and cytosolic ROS respectively (data not shown).

### **Sensitization of renal carcinoma cells by cucurbitacin differs from bortezomib-induced sensitization**

A number of TRAIL-resistant cell lines can be sensitized to TRAIL-induced cell killing by cucurbitacin B, but not by the well-characterized TRAIL sensitizer bortezomib (Supplemental Data). ACHN cells were sensitized by both agents whereas only cucurbitacin B sensitized Caki-1 cells (Fig. 5). In addition, the TRAIL-sensitizing effect of cucurbitacin B was much more rapid than that of bortezomib (Fig. 5). In contrast with bortezomib, cucurbitacin B treatment did not affect the levels of the apoptosis proteins cFLIP, Mcl-1, or XIAP by western blot analysis nor did it affect cell surface levels of TRAIL receptors by FACS analysis (data not shown).

### **Cucurbitacins rapidly affect cell morphology**

Interestingly, sensitizing concentrations of active cucurbitacins always produced dramatic changes in cell shape. These effects were clearly visible by 1 h after drug exposure and were eliminated by inclusion of NAC (Fig. 6a). By contrast, inactive cucurbitacins, such as cucurbitacin P, did not produce this change in cell morphology. These changes in cell shape were accompanied by significant changes in the cellular distribution of actin. Clumping occurred in cells exposed to cucurbitacin B for 1 h (Fig. 6b). In addition, cucurbitacin B

treatment rapidly induced a significant decrease in the levels of phosphopaxillin in both ACHN and Caki-1 cells (Fig. 6c). Unlike cucurbitacins, bortezomib treatment did not result in rapid and dramatic changes in cell shape (data not shown).

## Discussion

Cucurbitacins B and D were previously identified in a highthroughput screen for TRAIL synergizing compounds [10]. Cucurbitacin I has been reported to sensitize mouse fibrosarcoma cells, human colorectal carcinoma cells, and human leukemia cells to TRAIL-induced apoptosis [15, 24]. Therefore, additional cucurbitacins were further investigated with regard to their effects on TRAIL-resistant ACHN cells. Based on IC<sub>50</sub> values (Fig. 1; Table 1), cucurbitacins B, C, D, E, and I have similar potencies while K is much less potent and P is inactive. The persistent TRAIL-sensitizing effects of cucurbitacins after removal of the compounds (Fig. 2) are consistent with rapid, irreversible effects on TRAIL-induced commitment step(s). It is possible that intracellular cucurbitacin concentrations remain high enough to retain activity even when they are removed from the extracellular medium. Cucurbitacins may participate in a rapid cell surface-reactive event, such as direct interaction with or modification of the TRAIL death receptors themselves, as part of their mechanism of action. The rapid time course also suggests that sensitization does not require changes in gene expression often implicated in sensitization of cells to TRAIL.

The pan-caspase inhibitor ZVAD-FMK blocked sensitization by cucurbitacin B, implying apoptotic cell death via caspase activation. Caspase-8 activation is among the earliest detectable responses of cells to TRAIL-mediated apoptosis [22, 25]. Enhancement of TRAIL-dependent caspase-8 activation by cucurbitacins correlated with cell death from the earliest time points tested. This is consistent with potential threshold effects for initiator caspases and enhanced signaling from the TRAIL death-inducing signaling complex (DISC) [5, 22, 25, 26]. A cell membrane-reactive activity of the cucurbitacins with TRAIL receptors would also be consistent with this observation. Caspase-8 activation and regulation are complicated processes that allow for multiple points of modulation [22, 27, 28]. Cucurbitacins may be useful for probing the molecular mechanisms underlying modulation of caspase activation by TRAIL sensitizers.

Generation of ROS is another mechanism proposed for many TRAIL sensitizers [5, 19–21] and cucurbitacin B as a single agent has been reported to induce ROS generation [14]. Although NAC, an inhibitor of ROS generation, blocked the TRAIL-sensitization effects of cucurbitacins, ROS generation in response to cucurbitacin B (in the presence or absence of TRAIL) was not detected (data not shown). Thus, the inhibitory effects of NAC on cucurbitacins and TRAIL could result from other mechanisms [29], including interference with a possible electrophilic mechanism of cucurbitacin action (via the C23–C24 double bond—Fig. 7). TRAIL sensitizers can be placed in at least two basic categories, those that are inhibited by NAC (including cucurbitacins) and those that are not (for example, bortezomib). The mechanistic bases for this dichotomy are beyond the scope of this report, but may provide important insights into TRAIL signaling pathways.

In addition to the differential effects of NAC, a number of other differences between cucurbitacins and bortezomib were observed. The pattern of sensitization of multiple cell lines by cucurbitacin B (see Supplemental Data) differs from those of other sensitizers including bortezomib (Fig. 5 and Supplemental Data). The TRAIL-sensitizing effect of cucurbitacins was much more rapid than that of bortezomib. Bortezomib also modulates the levels of a number of apoptotic proteins unaffected by cucurbitacin treatment. These differences, along with the effects on cell shape and modulation of activities by NAC,



suggest that the molecular mechanisms underlying bortezomib and cucurbitacin B sensitization of cells to TRAIL apoptosis are quite different and distinct.

The active cucurbitacins rapidly changed the morphology of both ACHN and Caki-1 cells, possibly via effects on F-actin [30, 31]. In fact, actin distribution was dramatically affected by cucurbitacin B as were levels of phosphopaxillin. Inactive cucurbitacins did not cause changes in cell morphology. In addition, these changes were completely blocked by the presence of NAC in the media. Paxillin phosphorylation is often associated with integrin signaling. Thus it seems likely that cucurbitacin B-induced changes the cell cytoskeleton may play an important role in the amplifying TRAIL apoptosis signaling. This potential role of cell adhesion on TRAIL signaling has recently been reported [32]. The involvement of such cytoskeletal effects on TRAIL signaling is worthy of further investigation. Cucurbitacins may be useful agents for more detailed investigations of the relationships between cell morphology and TRAIL apoptosis signaling.

The differences in structural features among the cucurbitacins assessed in this report (Fig. 7; Table 1—numbering follows Afifi et al. [33]) are subtle. Of particular note, however, is the C23–C24 double bond. Only cucurbitacin K (least potent of the active compounds) and the inactive cucurbitacin P lack this feature. Comparison with two published structure–activity relationship (SAR) analyses of the cytotoxic effects of cucurbitacins [34, 35] suggest both similarities and differences between structural features apparently important for toxicity and for sensitization to TRAIL-induced apoptosis. However neither of the cited studies included all of the cucurbitacins shown in Fig. 7. The cytotoxicity studies identified acetylation of the C25 hydroxyl as being important for increased cytotoxicity. This feature does not correlate with sensitization of ACHN cells to TRAIL. Similarly, the presence of the C1–C2 double bond does not appear to correlate with TRAIL-induced apoptosis, unlike the conclusions drawn by Bartalis and Halaweish [34] regarding cytotoxicity of the compounds alone. The effects of cucurbitacins alone on ACHN cell survival were minimal (Fig. 1), but complete killing curves were not obtained. The report that included cucurbitacin P [35] indicated that it was orders of magnitude less potent than the other cucurbitacins listed (including B, C, D, E, and I) suggesting that cucurbitacin P truly is an outlier (consistent with TRAIL sensitization results). Finally, neither of the cited reports identified the C23–C24 double bond as important in cytotoxicity, yet this correlates quite well with TRAIL sensitization. Future studies may further distinguish molecular features of the cucurbitacins required for TRAIL sensitization requirements in comparison to those that elicit nonspecific cytotoxicity.

Cucurbitacins are also reported to inhibit STAT3 activation and inhibition of STAT3 signaling has been identified as a means of sensitizing cells to TRAIL. However, three lines of evidence suggest that cucurbitacins are not sensitizing ACHN cells to TRAIL via inhibition of STAT3 activation. First, structural features of cucurbitacins identified as important for inhibition of STAT3 activation [36] are not implicated in TRAIL sensitization. This analysis [36] suggested that the presence of a hydroxyl group on C19 results in loss of the ability to block STAT3 activation (cucurbitacin A was used in that study). However, cucurbitacin C, which also has this feature, is a very good TRAIL sensitizer (although A and C differ in other ways). Unfortunately, cucurbitacin A was not available for these studies and Sun et al. [36] did not include cucurbitacin C in their report. There is minimal correlation between structural features required for inhibition of STAT3 activation and those required for sensitization of cells to TRAIL. Second, effects of STAT3 modulation are generally due to changes in STAT3-driven gene expression (e.g. Ref. [37]). However, as little as 1 h pretreatment with cucurbitacins was sufficient to sensitize ACHN cells to TRAIL and by 1–4 h the cells are committed to respond to TRAIL. This rapid sensitization of renal carcinoma cells to cucurbitacins suggests that this response does not rely on transcripto-driven changes in protein levels. Consistent with this conclusion is the

observation that the level of Bcl-xL, transcriptionally regulated by STAT3 [38], was unaffected by cucurbitacin B (see Supplemental Data). Therefore, based on the time courses reported here, the mechanisms of action of cucurbitacin sensitization of renal carcinoma cells to TRAIL are immediate and non-genomic, and may involve a direct interaction of the compound with a readily accessible target that is either on or in the cell. Third, the cucurbitacins tested had minimal effect on constitutively activated STAT3 as measured by western blots of phosphoSTAT3 (see Supplemental Data). Taken together, SAR comparisons (albeit limited), temporal aspects of sensitization, and western blot results for phosphoSTAT3 all suggest that cucurbitacins sensitize ACHN cells to TRAIL by a STAT3-independent mechanism.

Cucurbitacins are not currently clinically useful as single agents due to toxicity and low therapeutic indices [11]. Indeed, normal human renal epithelial cells are also sensitized to TRAIL-mediated apoptosis by cucurbitacin B (data not shown). This would obviously lead to significant off-target effects. Nonetheless, some recent reports have noted in vivo antitumor effects of cucurbitacins alone [39] or when combined with the chemotherapeutic drug gemcitabine in mouse cancer models [40]. Whether there is a therapeutic window for the clinical utilization of cucurbitacins for cancer therapy in humans is unknown. However, cucurbitacins may be useful precursors for clinically useful compounds following appropriate chemical modification (to reduce toxicity) or when used at low concentrations and brief exposures in conjunction with TRAIL. SAR analysis suggests that the TRAIL synergy effects of cucurbitacins are decoupled from their general cytotoxicity. As such, they could form the basis for development of less toxic TRAIL sensitizers.

As noted the rapidity of the response to cucurbitacins would tend to preclude mechanisms requiring changes in gene expression. This conclusion is consistent with the absence of effects of cucurbitacin B on cellular levels of c-FLIP, Mcl-1, and XIAP, or cell surface levels of TRAIL receptors (FACS analysis) (data not shown) as well as Bcl-xL (Supplementary Data). Instead, rapid effects on enzymatic activity and cytoskeletal organization are observed.

The mechanism(s) of cellular resistance to TRAIL and the mechanisms by which TRAIL sensitization can occur (both STAT3-related and STAT3-independent) are not well understood [1]. The temporal aspects of sensitization of cells to TRAIL also remain unclear. The TRAIL-sensitizing action of specific cucurbitacins in renal carcinoma cells requires the activation and cleavage of caspase-8 at very early time points, and suggest an important role for the cytoskeleton in controlling the extent of caspase-8 activation. Sensitization by cucurbitacins is rapid and persistent making them potentially useful reagents for developing increased understanding of the sequence(s) of molecular events that can lead to TRAIL sensitization and subsequent apoptosis. As a result cucurbitacins have potential as valuable tools for investigating early events in sensitization of cells to TRAIL.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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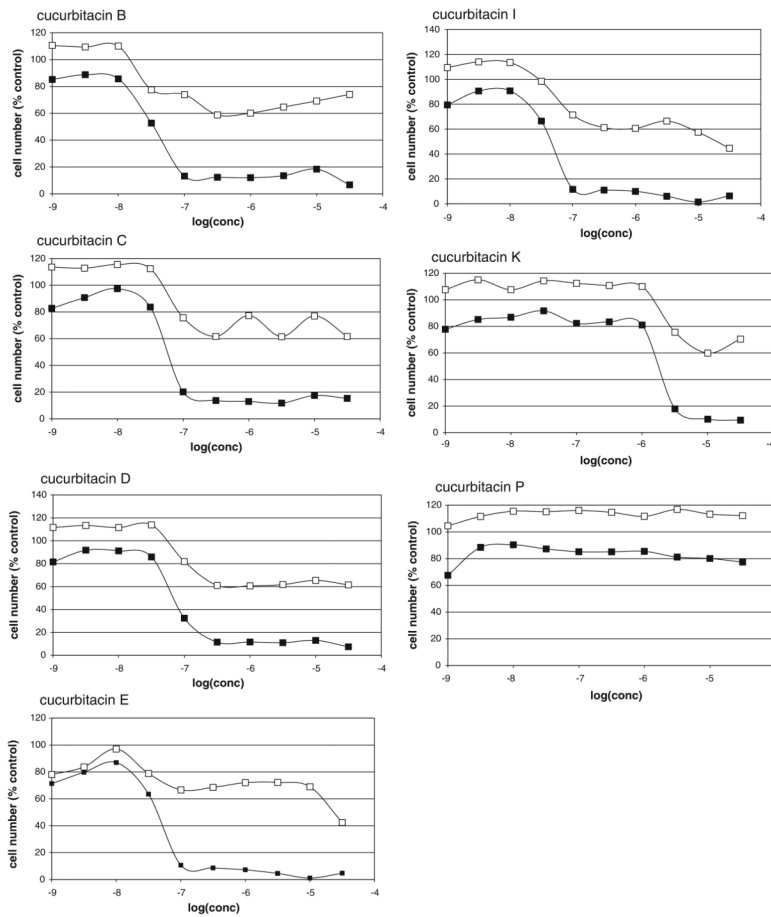
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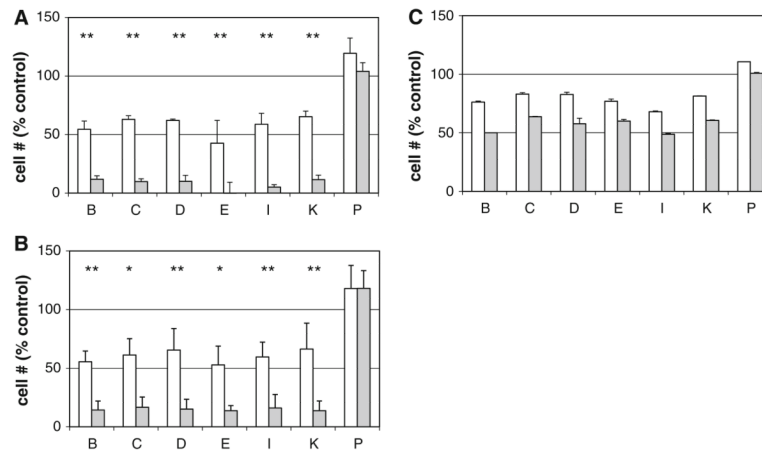
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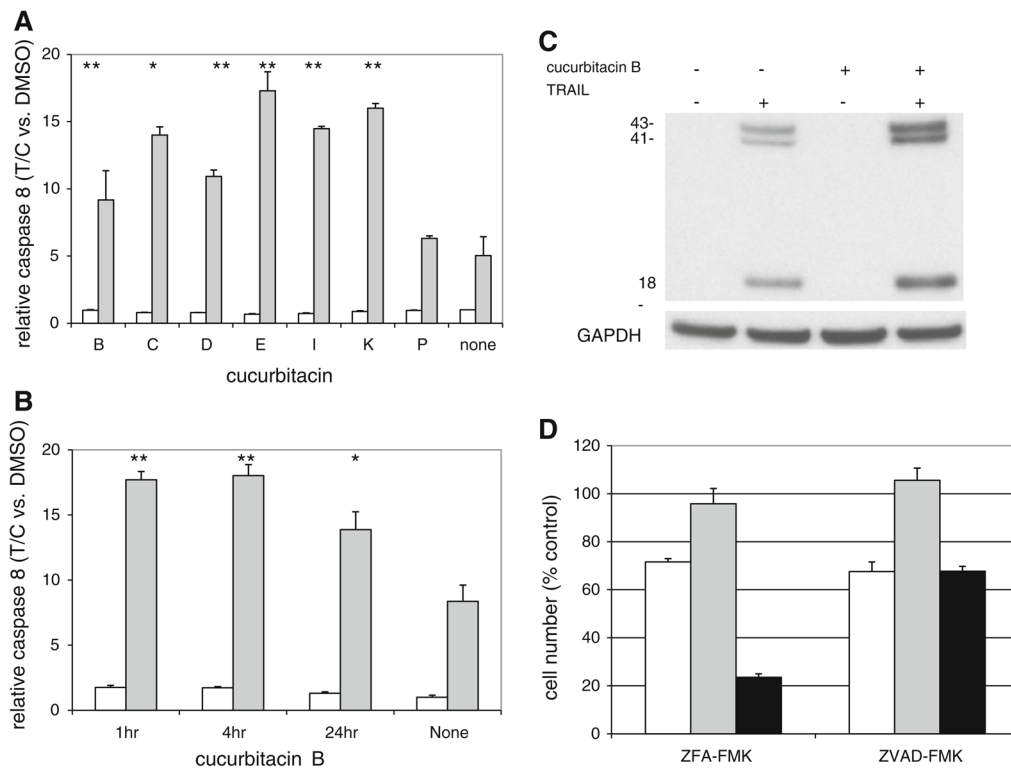


**Fig. 1.** Dose-dependent effects of cucurbitacins on ACHN cells. ACHN cells were allowed to attach overnight before treatment for 4 h with variable concentrations of the indicated cucurbitacin followed by addition of TRAIL (40 ng/ml) or culture medium. 20 h after TRAIL addition, viable cells were assessed by an XTT assay. Results were normalized to DMSO controls for each plate (control = 100%). Data points represent the average of 2 days of experiments, duplicate determinations each day. *Open symbols* represent compound without TRAIL, *closed symbols* represent compound followed by TRAIL



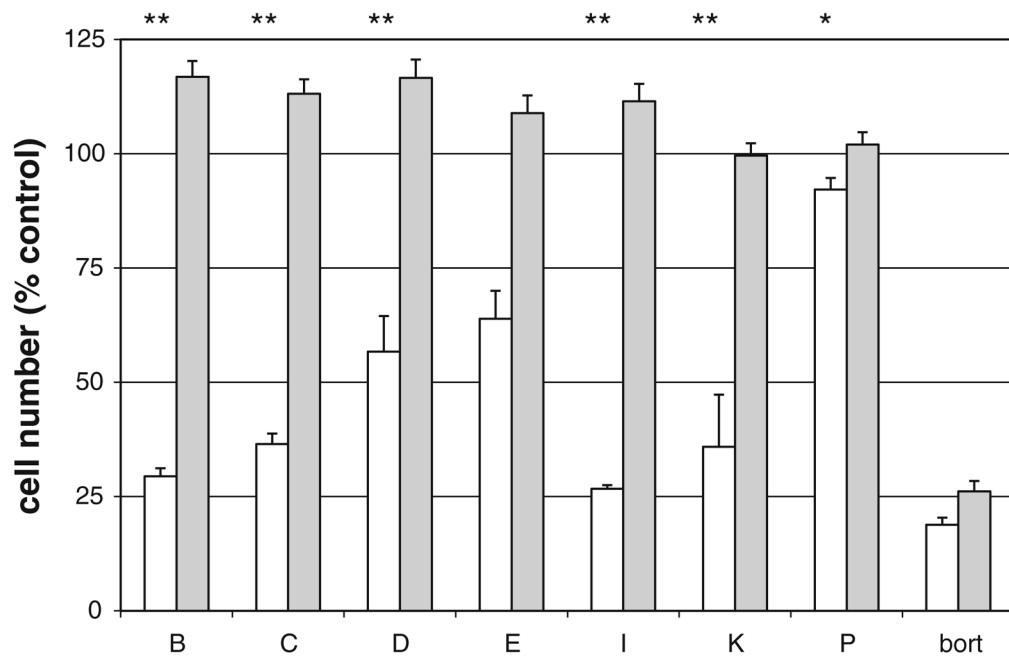
**Fig. 2.**

Effects of removal of cucurbitacins on enhancement of TRAIL-induced cell killing. ACHN cells were allowed to attach overnight before treatment for 1 or 4 h with 1  $\mu$ M (*cucurbitacins B, C, D, E, and I*) or 10  $\mu$ M (*cucurbitacins K and P*) followed by removal of drug-containing medium, washing with drug-free medium, and treatment in the presence (*gray bars*) or absence (*open bars*) of TRAIL (40 ng/ml). Viable cell numbers were assessed by XTT and normalized to untreated control (control = 100%) for each plate. **a** 1 h pretreatment followed by overnight incubation  $\pm$  TRAIL, **b** 4 h pretreatment followed by overnight incubation  $\pm$  TRAIL, **c** 1 h pretreatment followed by 4 h incubation  $\pm$  TRAIL:  $n = 4$  for (**a**),  $n = 3$  for (**b**) (*error bars* represent standard deviation), and  $n = 2$  for (**c**) (*error bars* represent range). \* $P < 0.05$ , \*\* $P < 0.01$ , Students *t* test comparing cucurbitacin alone to cucurbitacin + TRAIL



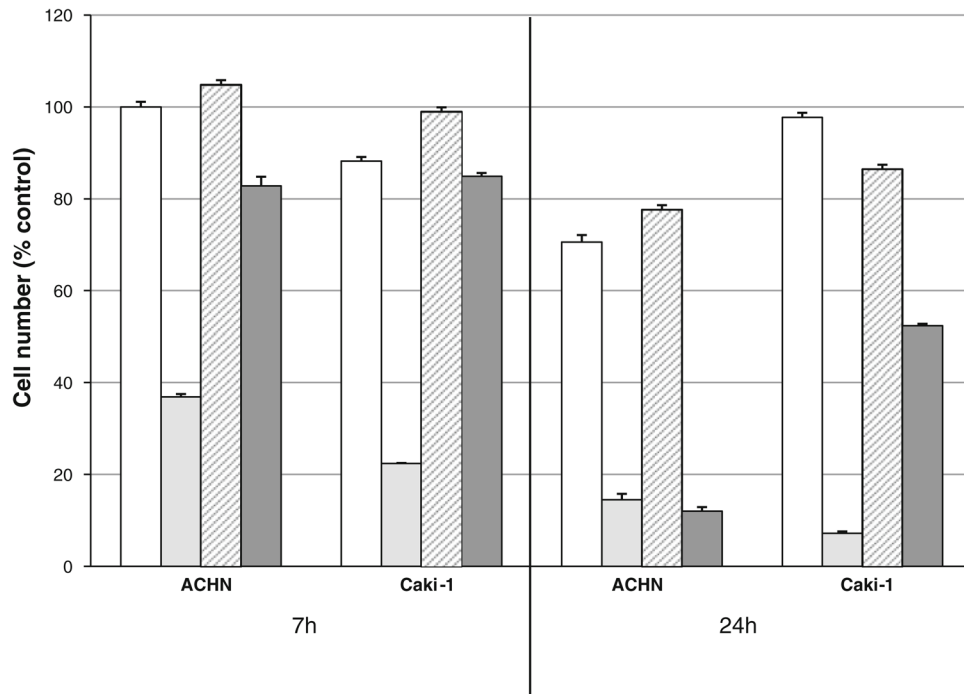
**Fig. 3.** Caspase activation in the presence of cucurbitacin B. **a** ACHN cells were treated for 4 h with DMSO (control) or the indicated cucurbitacin (10  $\mu$ M) followed by an additional 4 h in the presence (*gray bars*) or absence (*open bars*) of TRAIL (40 ng/ml) in the continued presence of cucurbitacin. Caspase-8 activity was determined using the Caspase-Glo™ 8 assay kit. *Error bars* represent standard deviation ( $n = 3-5$ ). *T/C* indicates treated/control ratio of caspase-8 activity, normalized to DMSO control. **b** ACHN cells were incubated in the presence of cucurbitacin B (1  $\mu$ M) for 1, 4, or 24 h followed by incubation in the presence (*gray bars*) or absence (*open bars*) of TRAIL (1000 ng/ml) for 1 h. Caspase-8 activity was assessed using the Caspase-Glo™ 8 assay kit. *T/C* indicates treated/control ratio of caspase-8 activity, normalized to DMSO control. *Error bars* represent standard deviation ( $n = 3$ ). **c** ACHN cells were incubated in the presence of cucurbitacin B for 1 h followed by incubation in the presence or absence of TRAIL. After 4 h, cell extracts were assessed by western blot analysis for activated (i.e. cleaved) caspase-8. \* $P < 0.05$ , \*\* $P < 0.01$ , Students *t* test comparing cucurbitacin alone to cucurbitacin + TRAIL. **d** ACHN cells were preincubated for 2 h with either ZFA-FMK (inactive control) or ZVAD-FMK (pan-caspase inhibitor). This was followed by incubations with cucurbitacin B (100 nM, *open bars*), TRAIL (40 ng/ml, *gray bars*) or cucurbitacin B (2–4 h) followed incubation with TRAIL (*black bars*). Cell numbers were assessed 18–20 h after addition of TRAIL. Data were normalized to ZFA-FMK alone or ZVAD-FMK alone. *Error bars* represent standard deviation ( $n = 3$ ).  $P < 0.01$  comparing cucurbitacin B + TRAIL + ZVAD-FMK to cucurbitacin B + TRAIL + ZFA-FMK



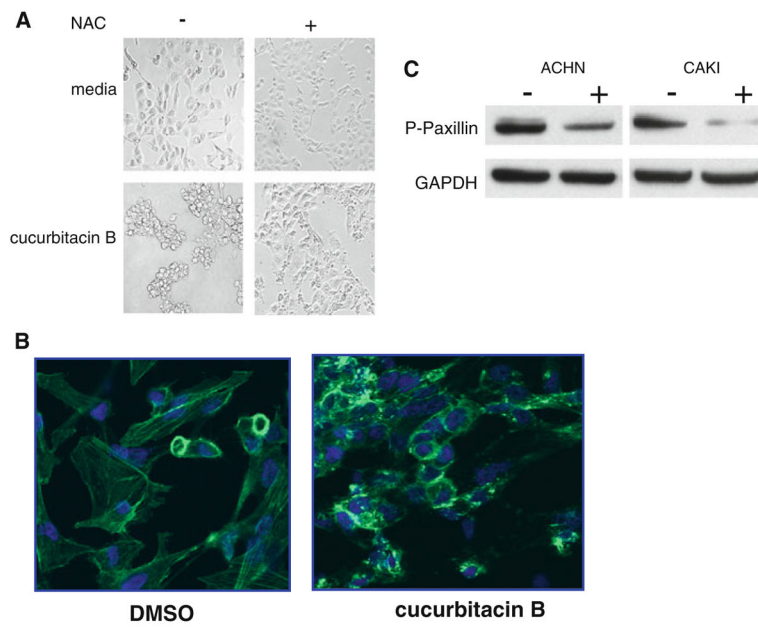


**Fig. 4.**

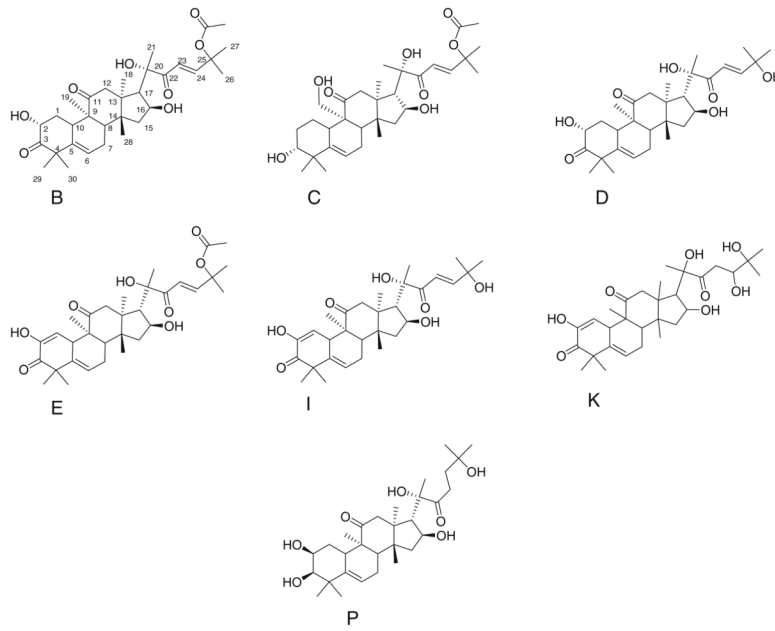
Effects of NAC on ACHN cell response to cucurbitacins. ACHN cells were plated at 3,500 cells/well in 384-well plates and allowed to attach overnight. NAC was added to half of the wells at a final concentration of 10 mM followed immediately by the indicated cucurbitacins (or DMSO control). After 4 h, TRAIL (or medium control) was added. After 20 h additional incubation, wells were washed ( $2 \times 100 \mu\text{l}$  per well), and cell numbers assessed by XTT. Results were normalized to values for cells in the absence of TRAIL or cucurbitacin. Bars represent compounds + TRAIL in the absence (*open bars*) or presence (*gray bars*) of NAC. Error bars indicate standard deviation ( $n = 3$ ). The following concentrations were used: cucurbitacins B, C, D, E, and F: 100 nM; cucurbitacins K and P: 10  $\mu\text{M}$ ; bortezomib (bort): 40 nM. \* $P < 0.05$ , \*\* $P < 0.01$ , Students *t* test comparing cucurbitacin + TRAIL with and without NAC



**Fig. 5.** Effects of cucurbitacin B and bortezomib on ACHN and Caki-1 cells. ACHN or Caki-1 cells were treated with or without cucurbitacin B (200 nM) or bortezomib (20 nM) in the presence of 100 ng/ml TRAIL for 7 or 24 h. Cell numbers were normalized to untreated controls. After treatment with TRAIL alone, cell numbers ranged from 84 to 96% of control. *Open bars* cucurbitacin B only; *light gray* cucurbitacin B + TRAIL; *cross-hatched* bortezomib only; *dark gray* bortezomib + TRAIL



**Fig. 6.** Effect of cucurbitacin B on cell morphology. **a** Caki-1 cells were treated with or without cucurbitacin B (200 nM) for 1 h in the presence or absence of 10 mM NAC. **b** Caki-1 cells were treated for 1 h with 200 nM cucurbitacin B followed by paraformaldehyde fixation and permeabilization with Triton X-100 before visualization of polymerized actin using fluorescein-phalloidin. **c** ACHN or Caki-1 cells were treated for 1 h with cucurbitacin B (200 nM), solubilized and subjected to western blot analysis using an antibody against phosphopaxillin



**Fig. 7.**  
Structures of cucurbitacins tested for sensitization of ACHN cells to TRAIL

**Table 1**

Structural features and TRAIL sensitizing activities of cucurbitacins

Feature	Cucurbitacin							
	B	C	D	E	I	K	P	
C1-C2	-	-	-	=	=	=	=	-
C2	-OH	-H	-OH	-OH	-OH	-OH	-OH	-OH
C3	=O	-OH	=O	=O	=O	=O	=O	-OH
C19	-H	-OH	-H	-H	-H	-H	-H	-H
C23-C24	=	=	=	=	=	=	=	-
C24	-H	-H	-H	-H	-H	-H	-OH	-H
C25	-OAc	-OAc	-OH	-OAc	-OH	-OH	-OH	-OH
IC <sub>50</sub> (nM) <sup>a</sup>	27.4	54.2	78.2	44.1	43.3	1970	1970	Inactive

<sup>a</sup>IC<sub>50</sub> values (average, *n* = 2) were calculated from the data in Fig. 1 (in the presence of TRAIL)