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Sulforaphane metabolites cause apoptosis via microtubule disruption in cancer

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Abstract

Sulforaphane (SFN) inhibited growth in many cancers, but its half-life is 2 h in circulation. However, its metabolites, sulforaphane-cysteine (SFN-Cys) and sulforaphane-N-acetyl-cysteine (SFN-NAC) had longer half-lives and decreased the cell viability in both dose- and time-dependent manners in human prostate cancer. Flow cytometry assay revealed that these two SFN metabolites induced apoptosis with the features such as vacuolization, disappeared nuclear envelope, nuclear agglutination and fragmentation via transmission electron microscopy observation. Western blot showed that the sustained phosphorylation of ERK1/2 mediated by SFN metabolites caused activation and upregulation of cleaved Caspase 3 and downregulation of α -tubulin. High expression of α -tubulin was demonstrated to be positively correlated with cancer pathological grading. Both co-immunoprecipitation and immunofluorescence staining implicated the interaction between SFN metabolite-induced phosphorylated ERK1/2 and α -tubulin, and Caspase 3 cleavage assay showed that α -tubulin might be the substrate for cleaved Caspase 3. More, the SFN metabolite-mediated reduction of α -tubulin increased the depolymerization and instability of microtubules by microtubule polymerization assay. Reversely, microtubule-associated protein Stathmin-1 phosphorylation was increased via phosphorylated ERK1/2 and total Stathmin-1 was reduced, which might promote over-stability of microtubules. Immunofluorescence staining also showed that SFN metabolites induced the 'nest-like' structures of microtubule distribution resulting from the disrupted and aggregated microtubules, and abnormal nuclear division, suggesting that the disturbance of spindle formation and mitosis turned up. Thus, SFN-Cys and SFN-NAC triggered the dynamic imbalance of microtubules, microtubule disruption leading to cell apoptosis. These findings provided a novel insight into the chemotherapy of human prostate cancer.

Key Words

- ▶ sulforaphane metabolites
- ▶ Caspase 3
- ▶ α -tubulin
- ▶ Stathmin-1
- ▶ apoptosis

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Introduction

Sulforaphane (SFN) effectively inhibited cell growth in a variety of cancers (Clarke *et al.* 2008, Ahn *et al.* 2010). Due to a short half-life in metabolism, SFN has not been used for clinical treatment yet. However, SFN metabolites,

sulforaphane-cysteine (SFN-Cys) and sulforaphane-N-acetyl-cysteine (SFN-NAC) had longer half-lives in circulation; they were cleared from the body within 72 h (Clarke *et al.* 2011). Nevertheless, we do not know whether

SFN-Cys and SFN-NAC inhibit cancer growth although these two metabolites are proved to be the inhibitors of histone deacetylase (Rajendran *et al.* 2013).

SFN metabolites induced the phosphorylation of ERK1/2 leading to the inhibition of tumorigenesis (Dickinson *et al.* 2015). Recently, we found that SFN-Cys suppressed invasion via ERK1/2-mediated downregulation of Galectin-1 in human prostate cancer cells (Tian *et al.* 2016). More, MEK1/2 colocalized and interacted with α -tubulin in cultured human fibrosarcoma cells (Cao *et al.* 2010). Studies showed that sustained ERK1/2 phosphorylation by SFN induced apoptosis (Li *et al.* 2014, Peng *et al.* 2015), and transient ERK1/2 activation by SFN contributed to cancer progression *in vitro* (Hsu *et al.* 2013). Studies showed that SFN bound to α -tubulin, and downregulation of α -tubulin resulted in cell apoptosis (Mi *et al.* 2008). N-acetyl-S-(N-allylthiocarbamoyl)cysteine (NAC-AITC) downregulated α -tubulin and activated Caspase 3 in bladder cancer (Bhattacharya *et al.* 2012). Besides, we demonstrated that SFN-Cys downregulated α -tubulin via phosphorylated ERK1/2, and α -tubulin was a substrate for cleaved Caspase 3 in human non-small-cell lung cancer cells (Lin *et al.* 2017). Therefore, α -tubulin might be a target for SFN metabolites, which might induce α -tubulin cleavage, aggregation leading to disequilibrium of microtubule dynamics.

The dynamics of microtubules can be regulated by microtubule-destabilizing proteins (Bhat & Setaluri 2007), and their high expression in cancer tissues contributed to the grading of malignancy and poor prognosis (Nemunaitis 2012). Reversely, low expression and phosphorylation of microtubule-destabilizing protein Stathmin-1 might cause microtubule over-stability and apoptosis (Ghosh *et al.* 2007). Activated ERK1/2 might be increased in microtubules, and activated p38 promoted the microtubule depolymerization through phosphorylation of Stathmin-1 in hypoxic cells (Hu *et al.* 2010). Additionally, activated ERK1/2 might phosphorylate Ser 25 of Stathmin-1 in response to growth factor (Marklund *et al.* 1993a,b). Tumor necrosis factor-induced microtubule stabilization mediated by hyperphosphorylation of Stathmin-1 promoted cell death (Vancompernelle *et al.* 2000). Therefore, Stathmin-1 might also be an attractive target for cancer therapy to disturb microtubule dynamics. It was reported that overexpression of Stathmin-1 decreased polymerization of microtubules and diminished sensitivity to Taxol; inhibition of Stathmin-1 expression sensitized K562 leukemic cells to killing by Taxol (Bhat & Setaluri 2007). Hence, if SFN metabolites lower Stathmin-1 expression,

we can combine them with Taxol to reduce the doses of drugs and to enhance the efficiency of treatment.

Collectively, we will investigate if SFN metabolites induce apoptosis in human prostate cancer cells and characterize the underlying mechanisms to develop novel anti-cancer therapeutics with high efficiency and safe doses.

Materials and methods

Reagents

SFN-Cys, SFN-NAC, anti- β -actin, anti- α -tubulin, anti-Caspase 3, anti- β -tubulin and Protein A/G PLUS Agarose were purchased from Santa Cruz Biotechnology. Anti-ERK1/2 and anti-pERK1/2 (Thr202/Tyr204) were obtained from Cell Signaling Technology. Anti-Stathmin-1 and anti-pStathmin-1 (Ser 25) were ordered from Sangon Biotech Co., Ltd. (Shanghai, China). Caspase 3 colorimetric assay kit was purchased from Clontech Laboratories (TaKaRa). Annexin V-FITC/PI apoptosis assay kit was purchased from NeoBioscience (Shenzhen, China).

Cell culture

Human prostate cancer DU145 and PC3 cells were purchased from the Cell Resource Center, Peking Union Medical College. The species origin was confirmed with PCR. The identity of the cells was authenticated with STR profiling (FBI, CODIS). RPMI-1640 medium (Hyclone) containing 10% (v/v) fetal bovine serum and 1% penicillin–streptomycin was used for routine cell culture. Cancer cells were incubated in routine condition and cells at 70–80% confluence were treated with the indicated concentrations of SFN metabolites at the indicated time points.

Morphological observation

Cells (2×10^5 per well) were seeded into 6-well plates, and exposed to SFN-Cys or SFN-NAC (0, 15, 30, 45 μ M) for 24 h. Images were photographed via a phase-contrast microscope (Leica) linked to a digital camera (Olympus) at $\times 100$ magnification.

Apoptosis detection by transmission electron microscopy

Cells were treated by 20 μ M SFN-Cys for 24 h first, and then cells were sequentially fixed with 3% glutaraldehyde for 3 h

at 4°C, washed with 0.1 M PBS, post-fixed with 1% osmium tetroxide, dehydrated with alcohol, embedded in epoxy resin and stained with uranium acetate and lead nitrate. Finally, the samples were examined with the transmission electron microscope (JEM-2100, JEOL, Japan).

TMA and immunohistochemistry

Human Prostate Cancer Tissue Microarrays (TMA) with 64 patient samples and different Gleason patterns were established by Shanghai Biochip (Shanghai, China). Briefly, slides (4 µm) were deparaffinized, hydrated and subjected to antigen retrieval and peroxidase blocking according to the UltraSensitive S-P detection kit (Maixin, Fuzhou, China). Then, slides were incubated with anti- α -tubulin (1:500) and HRP-labeled secondary antibody. Further, freshly prepared System-HRP (DAB) detection kit (Maixin) was applied to view the pathological results. The slides were counterstained with hematoxylin. Then, the stained slides were observed under microscope LEICA DM6000 B and images were acquired. Images were manually scored based on no signal, weak, moderate and strong staining criteria and ultimately calculated as a histopathologic score (H-score). H-score with a range of 0–300 was calculated according to the following formula: (3 × percentage of cells with strong staining) + (2 × percentage of cells with moderate staining) + (1 × percentage of cells with weak staining). The difference between H-score was determined by Kruskal–Wallis rank-sum test via SAS software.

Western blot

Cells were treated with desired concentrations of SFN metabolites at the certain time, then the whole cell lysate was collected, and the procedures were followed as that described previously (Tian *et al.* 2016).

Flow cytometry analysis

Apoptosis was determined by dual staining with FITC-annexin V and propidium iodide (PI) via the kit (NeoBioscience) according to the manufacturer's instruction. Briefly, after treatment with various concentrations of SFN-Cys or SFN-NAC, both floating and adherent cells were collected and washed twice with pre-cold PBS and were resuspended in 250 µL 4 × binding buffer. Then, Annexin V-FITC and PI were added. After 15-min incubation in the dark, samples were analyzed via the flow cytometer (FACSAria, BD). The absolute

numbers and the apoptosis index were calculated by BD FACSDiva 7.0 software.

Caspase 3 colorimetric assay

Caspase 3 activity was determined by Caspase 3 colorimetric assay kit (Clontech) according to the manufacturer's instruction. Briefly, cells were treated with different concentrations of SFN-Cys for 24 h. Then, cytosolic extract was incubated with the reaction mixture and colorimetric substrate of Caspase 3 at 37°C for 1 h. The absorbance was measured by BioTek microplate reader (Synergy HT, USA) at 405 nm and represented the activity of Caspase 3. While the change of Caspase 3 activity was defined as the absorbance ratio of the treated group vs the control (set as 1).

Immunofluorescence staining of α -tubulin

Cells were incubated on chamber slides for 2 days, fixed with 4% paraformaldehyde/PBS and treated with 0.1% Triton X-100/PBS. Mouse anti- α -tubulin and rabbit anti-pERK1/2 were used at 1:100 dilution overnight at 4°C. Then, washes were done for three times; either Alexa Fluor488-conjugated goat anti-mouse IgG secondary antibody or Alexa Fluor594-conjugated goat anti-Rabbit IgG (H+L) secondary antibody was used at 1:500 dilution and incubated for 1 h at room temperature. Nuclei were stained with 4,6-diamidino-2-phenylindole (DAPI) (Sigma). Slides were mounted in ProLong antifade reagent (Invitrogen-Molecular Probes) and viewed in a Zeiss LSM 510 confocal microscope equipped with a 63 × 1.4 numerical aperture.

Microtubule polymerization assay

Cells (4×10^5) were seeded into 6 cm plate and exposed to SFN-NAC (30 µM) for 24 h. Then, cells were collected and washed twice with PBS, then lysed at 37°C for 30 min with 300–500 µL lysis buffer (20 mM Tris–HCl, pH 6.8, 1 mM MgCl₂, 2 mM EGTA, 1% NP-40) with Protease Inhibitor Cocktail (Roche). The cell lysate was centrifuged at 14,000g for 15 min at 25°C. The supernatant containing soluble tubulin (depolymerization/free) was collected, while the pellet containing assembled tubulin (polymerization/microtubule) was suspended in 40 µL of pellet lysis buffer (20 mM Tris–HCl, pH 6.8, 1 mM MgCl₂, 2 mM EGTA, 2% SDS) after washed with PBS. Then, the precipitate was heated at 100°C for 30 min until the pellet was solved.

The two fractions (soluble and insoluble) were separated by Western blot.

Co-immunoprecipitation

Cells were treated with 30 μ M SFN-NAC for 24 h and then lysed with Non-denaturing Lysis Buffer (Applygen) for 30 min on ice and the protein was extracted by centrifugation at 12,000 *g* for 30 min. The anti- α -tubulin (Ruiying Biological) was added to the protein lysate, incubated overnight at 4°C. The α -tubulin complexes were pulled down with protein A/G agarose for 3 h, and the proteins were purified by centrifugation and boiled for 5 min to reverse the crosslink. Western blot was used to recognize the conjugated proteins.

Caspase 3 cleavage assay

Cells were treated with 30 μ M SFN-NAC for 24 h and the harvested cells were lysed in 25 mM Tris-HCl, 150 mM NaCl, 1 mM EDTA, 1% NP-40, 5% glycerol, pH 7.4 and 12 μ g cell extract was incubated with 4 μ L recombinant Caspase 3 (Sino Biological Inc.) in 50 μ L reaction buffer containing 25 mM Hepes, pH 7.5, 0.1% (w/v) Chaps, 10 mM DTT, at 37°C for 6 h. After incubation, Western blot analysis was used to detect the expression of α -tubulin and β -actin.

Statistical analysis

Data were expressed as mean \pm s.e.m. Multiple groups were evaluated via Kruskal-Wallis rank-sum test and two groups were compared by Student *t* test. Statistical significance was determined at the 0.05 or 0.01 level.

Results

SFN metabolites induced apoptosis in DU145 and PC3 cells

We observed cell morphological changes after the cells were treated with SFN-Cys and SFN-NAC. Results showed that the normal spindle-shaped cells developed into round-shaped ones with higher transparency, short processes and unclear cell border, suggesting that the cell viability decreased with SFN metabolites treatment (Fig. 1A). Further, cell proliferation assay results showed that the cell viability gradually decreased as the increasing concentrations of SFN metabolites. After treatment with either 20 μ M SFN-Cys or 30 μ M SFN-NAC, both DU145 and

PC3 cells showed a remarkable reduction in cell viability, respectively (Fig. 1B). Meanwhile, the ultrastructural features of apoptosis were viewed by transmission electron microscopy. After treatment with 20 μ M SFN-Cys for 24 h, the cells exhibited cytoplasmic vacuolization (black arrow), disappeared nuclear envelope and nuclear agglutination and fragmentation (white arrow) (Fig. 1C). Flow cytometry assay showed that apoptosis occurred in a concentration-dependent manner (Fig. 1D).

SFN metabolites induced apoptosis via sustained ERK1/2 phosphorylation in both time- and concentration-dependent manners

Here, we discovered that either SFN-Cys or SFN-NAC persistently induced the phosphorylation of ERK1/2 following a concentration-dependent manner (Fig. 2A and B). After the treatment with either 20 μ M SFN-Cys or 30 μ M SFN-NAC, the results showed a sharp increase of phosphorylated ERK1/2 (pERK1/2) in the curves. In order to investigate the mechanisms of apoptosis in living cells, we chose either 20 μ M SFN-Cys or 30 μ M SFN-NAC for further assays. Likewise, the levels of pERK1/2 were up to peak at 24 h (Fig. 2C and D). Therefore, we defined 24 h as the optimal treatment time. Besides, the addition of PD98059 (25 μ M), the inhibitor of pERK1/2, reversed either SFN-Cys-induced or SFN-NAC-induced phosphorylation of ERK1/2 (Fig. 2E and F), indicating that PD98059 successfully inhibited ERK1/2 phosphorylation. Meanwhile, flow cytometry assay showed that SFN-Cys-induced ERK1/2 phosphorylation contributed to apoptosis (Fig. 2G and H). However, pERK1/2 might further activate multiple signaling molecules, such as cytoskeletal proteins.

SFN metabolites upregulated cleaved Caspase 3 and downregulated α -tubulin via sustained ERK1/2 phosphorylation

Here, we first examined the expression of cleaved Caspase 3 protein, and this antibody recognized both full-length and cleaved forms of Caspase 3 proteins. Western blot showed that SFN-Cys upregulated the expression of cleaved Caspase 3 in a concentration-dependent manner, while the full-length Caspase 3 levels were not affected in these two cell types (Fig. 3A). Especially after the treatment with PD98059, the upregulation of cleaved Caspase 3 was significantly reversed (Fig. 3B), indicating that SFN-Cys upregulated cleaved Caspase 3 via sustained ERK1/2 activation.

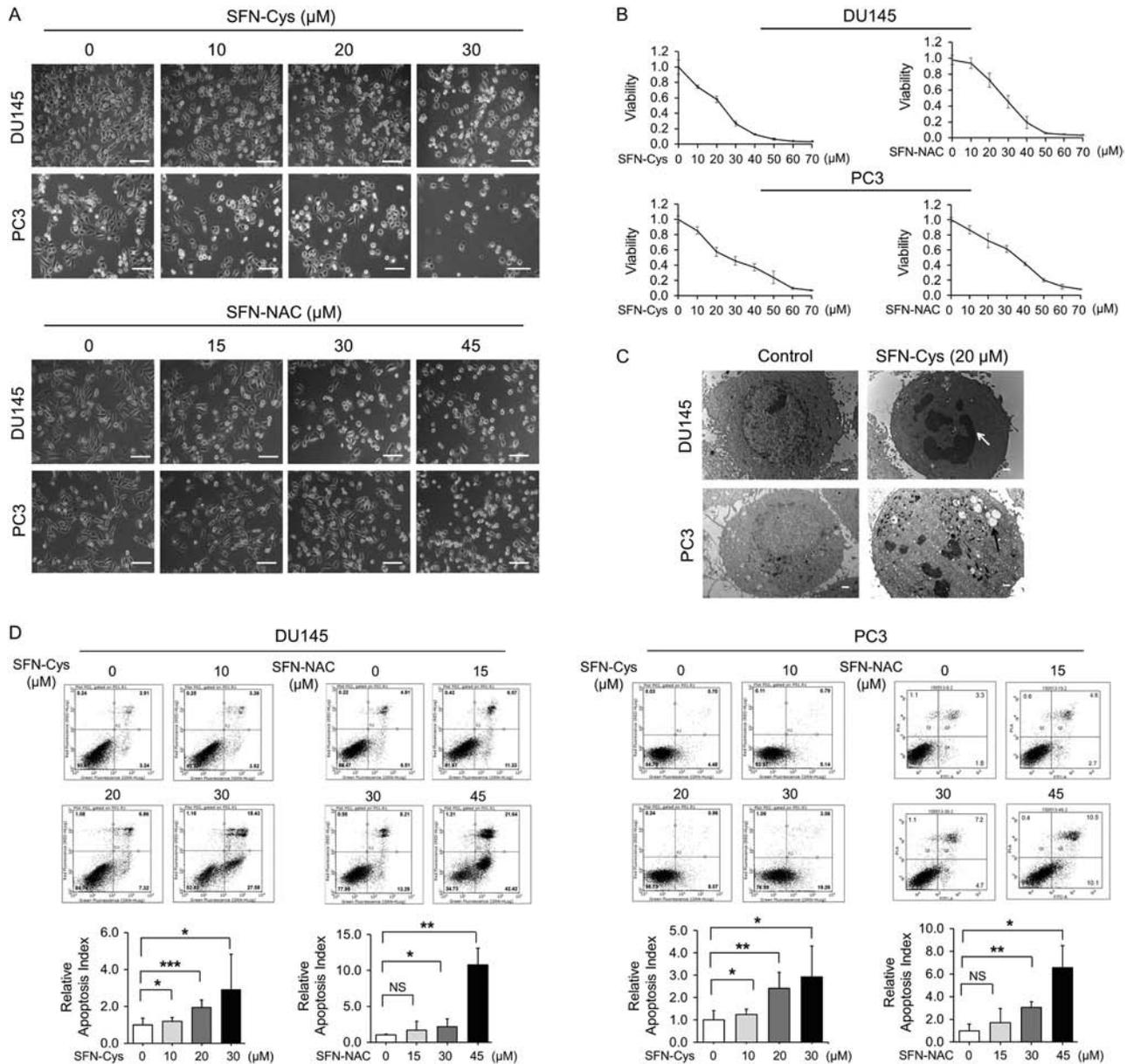


Figure 1

SFN-Cys and SFN-NAC induced apoptosis in prostate cancer cells. (A) DU145 and PC3 cells were treated with SFN-Cys (0, 10, 20, 30 μM) and SFN-NAC (0, 15, 30, 45 μM) for 24 h. Representative phase-contrast images were taken by a Leica DMIRB Microscope at $\times 100$ magnification. Scale bars, 100 μm . (B) The viability of cells treated with SFN-Cys or SFN-NAC (10–70 μM) was detected by MTS assays. Y-Axis means the percentage of viability ($\text{OD}_{490\text{nm}}$) vs that in the control group. (C) DU145 and PC3 cells were treated with 20 μM SFN-Cys for 24 h. Representative electron microscope images showed the features of apoptosis. White arrow: chromosome agglutination and nuclear fragmentation; black arrow: cytoplasmic vacuolation, scale bars, 1 μm . (D) DU145 and PC3 cells were treated with different concentrations of SFN-Cys and SFN-NAC for 24 h. Apoptosis was measured by FITC-annexin V and PI staining via the flow cytometry. Relative apoptosis index was represented by the ratio of the apoptosis in the treated cells vs the untreated cells (set as 1). * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, NS, no significance. Data were shown as means \pm s.e.m. ($n = 3$).

In addition, we further found that both SFN-Cys and SFN-NAC significantly reduced the expression of α -tubulin (Fig. 3C and D), but no significant changes were determined on β -tubulin (Fig. 3C and D). Nevertheless, after PD98059 was applied to block ERK1/2 phosphorylation, the downregulation of α -tubulin was remarkably reversed

(Fig. 3E and F), indicating that these two SFN metabolites significantly downregulated α -tubulin via sustained ERK1/2 phosphorylation. Taken together, we concluded that SFN metabolites induced the phosphorylation of ERK1/2 to upregulate cleaved Caspase 3 and downregulate α -tubulin leading to apoptosis.

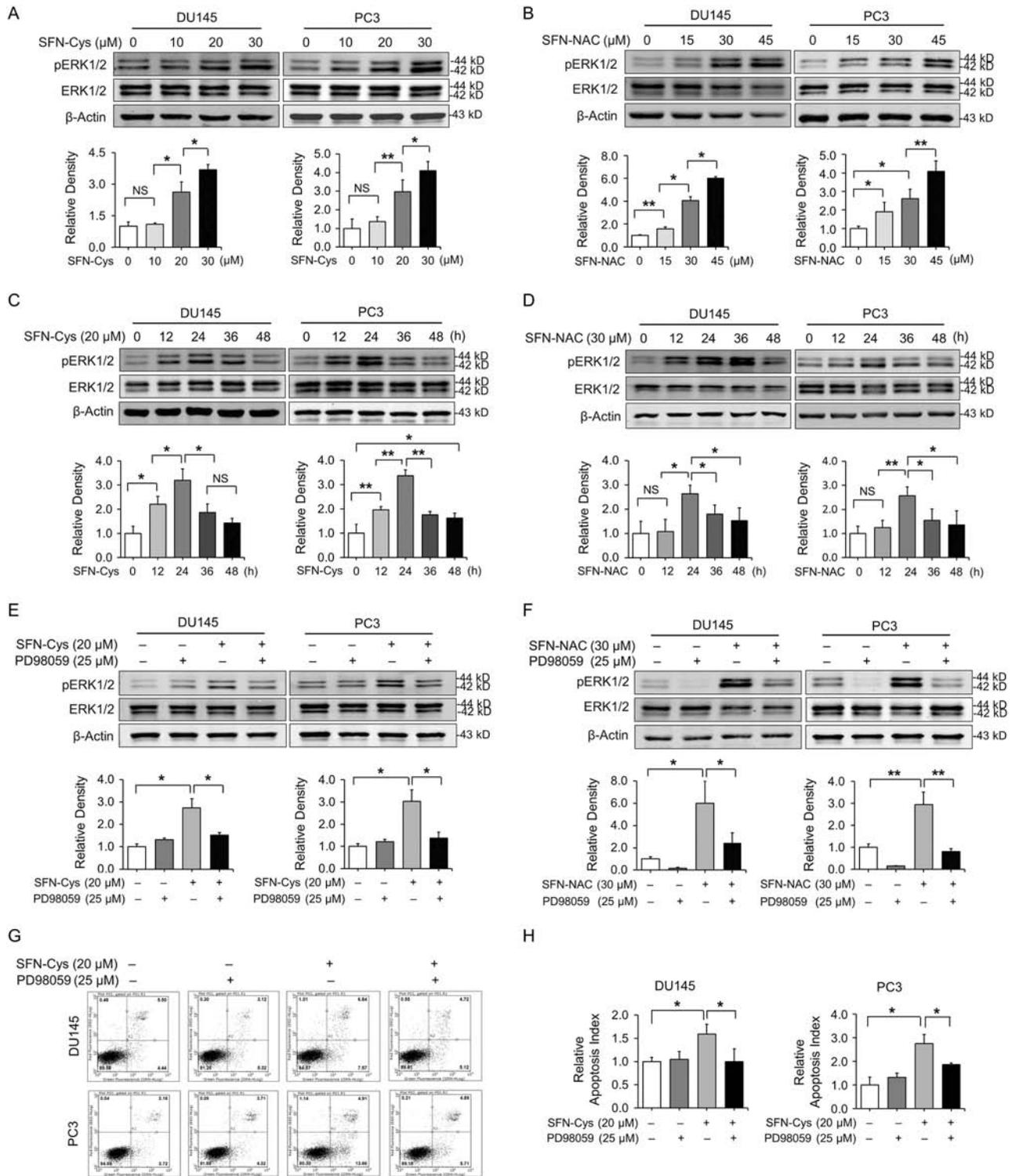


Figure 2

SFN metabolites induced apoptosis via sustained activation of ERK1/2. Phosphorylated ERK1/2 was detected at indicated concentrations of (A) SFN-Cys and (B) SFN-NAC. Phosphorylated ERK1/2 was detected at indicated incubation times by (C) SFN-Cys and (D) SFN-NAC. Phosphorylated ERK1/2 was detected after treatment with (E) 20 μM SFN-Cys or (F) 30 μM SFN-NAC for 24 h in the presence or absence of 25 μM PD98059. The density ratio of pERK/total ERK represented the level of phosphorylated ERK1/2. (G) DU145 and PC3 cells were treated by SFN-Cys plus PD98059, and the apoptosis assays were performed via FITC-annexin V and PI staining with flow cytometry detection. (H) The statistical analysis of the apoptosis index. * $P < 0.05$, ** $P < 0.01$, NS, no significance. Data were shown as means \pm s.e.m. ($n = 3$). β -Actin was probed for normalization and equal loading.

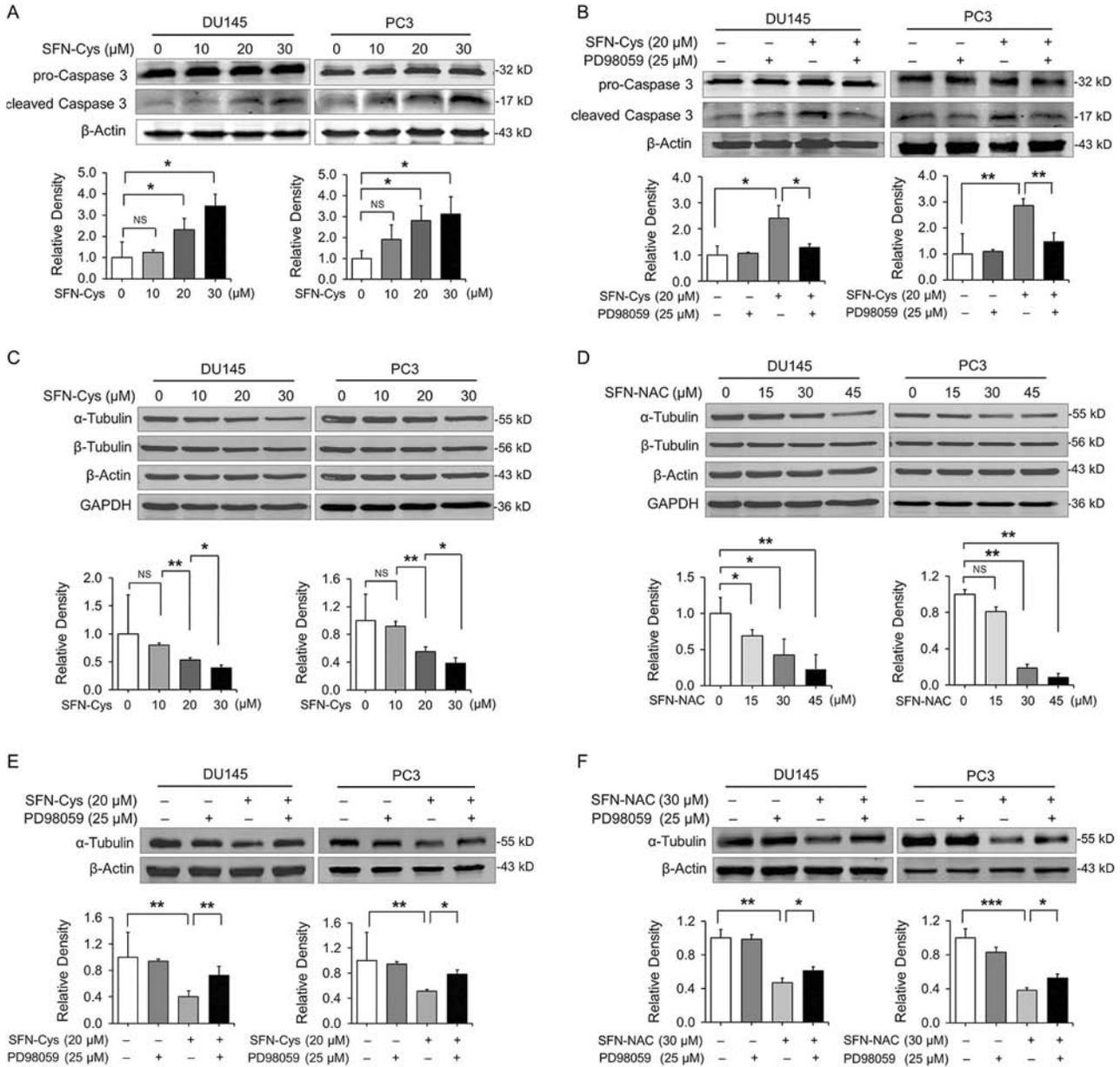


Figure 3

SFN metabolites upregulated cleaved Caspase 3, downregulated α-tubulin via sustained ERK1/2 phosphorylation. (A) The expression of cleaved and full-length Caspase 3 was detected by immunoblotting with SFN-Cys treatment for 24 h in both DU145 and PC3 cells. The Y-axis represented the relative density of cleaved Caspase 3 vs control (set as 1). (B) The expression of cleaved Caspase 3 was detected by immunoblotting at 20 μM SFN-Cys treatment for 24 h with/without 25 μM PD98059 in DU145 and PC3 cells. The Y-axis represented the relative density of cleaved Caspase 3 vs the control (set as 1). The expression of α-tubulin was detected by immunoblotting with different concentrations of (C) SFN-Cys and (D) SFN-NAC treatment for 24 h in DU145 and PC3 cells. The Y-axis represented the relative density of α-tubulin vs the control (set as 1). The expression of α-tubulin were detected by immunoblotting with the treatment of (E) SFN-Cys and (F) SFN-NAC for 24 h combined with PD98059 in DU145 and PC3 cells. The histogram exhibited the relative levels of α-tubulin vs the control. *P < 0.05, **P < 0.01, ***P < 0.001, NS, no significance. Data were shown as means ± s.e.m. (n = 3). β-Actin was used as an internal control.

SFN metabolites raised Caspase 3 activity leading to the downregulation of α-tubulin

The tumor samples with different pathological grading and adjacent tissues were laid in the arrays. As shown in

Fig. 4A, α-tubulin was weakly stained in adjacent tissues, while in tumor tissues, it was remarkably stained. As the malignancy of tumor increased, the expression of α-tubulin was significantly increased (dark staining) and gradually pervaded the whole prostate tissues (Fig. 4A). Similarly,

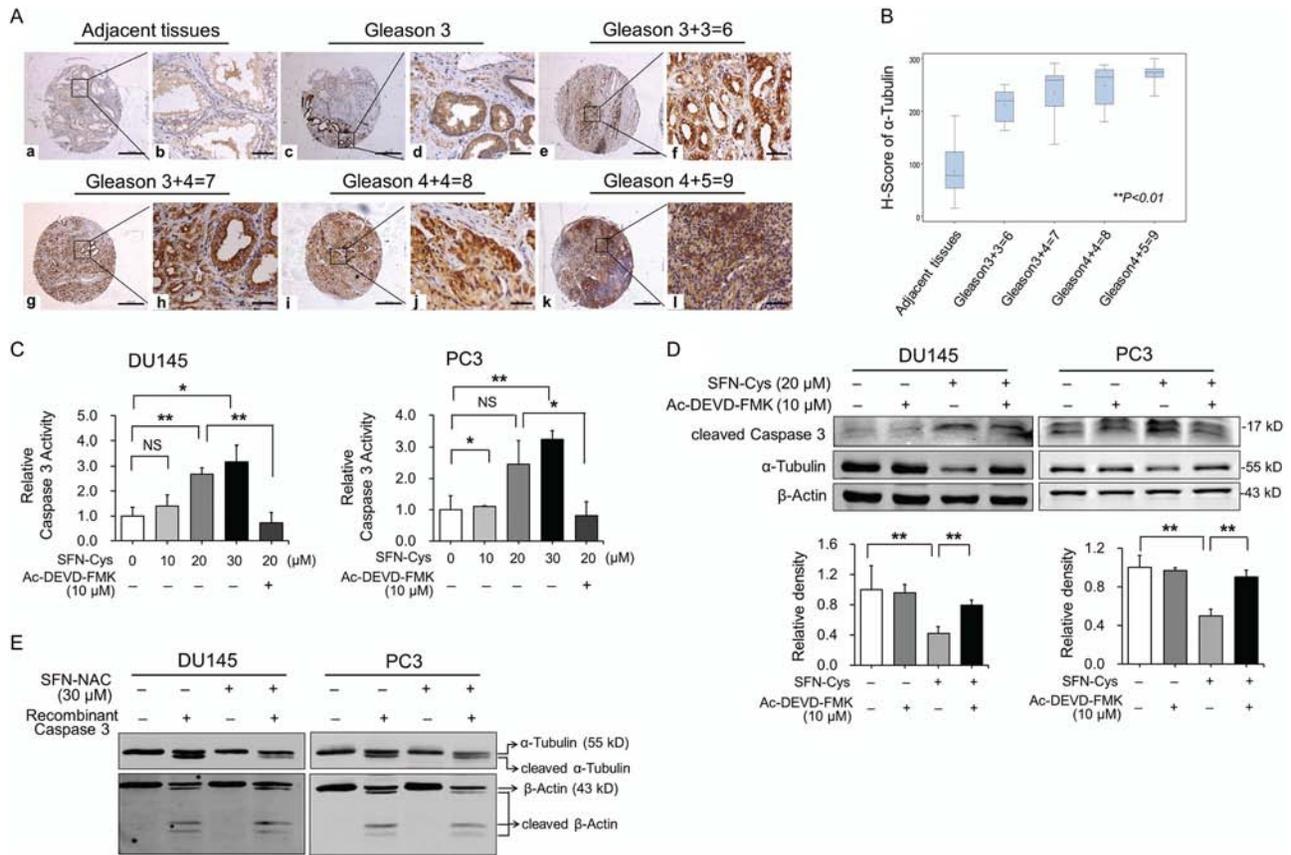


Figure 4

SFN metabolites induced Caspase 3 activation to cleave α -tubulin. (A) α -Tubulin was expressed in human prostate cancer tissues associated with various histopathological grading by IHC staining. The expression of α -tubulin in the adjacent tissues was observed as the control (a, b). IHC, magnification $\times 50$, scale bars, $500\mu\text{m}$, in a, c, e, g, i, k; magnification $\times 400$, scale bars, $50\mu\text{m}$, in b, d, f, h, j, l. (B) Boxed plot of α -tubulin expression levels among the adjacent tissues ($n=36$) and prostate cancer tissues with different Gleason grading (X -axis) ($n=48$). Box plots represented the 25th and 75th interquartile ranges and the median α -tubulin score by Kruskal–Wallis rank sum test. The horizontal line in the graph indicated the grand mean. (C) Caspase 3 activity was measured via Caspase 3 colorimetric assay kit *in vitro*. Cytosolic extract was incubated with the reaction mixture and colorimetric substrate of Caspase 3 at 37°C for 1 h. The absorbance was measured by BioTek microplate reader at 405nm . The cell lysate treated by $20\mu\text{M}$ SFN-Cys incubated with Caspase 3 inhibitor Ac-DEVD-FMK to confirm the specificity of the assay. (D) The cells were treated by $20\mu\text{M}$ SFN-Cys and/or $10\mu\text{M}$ Ac-DEVD-FMK to grow for 24 h. Western blot showed the expression of cleaved Caspase 3 and α -tubulin, and α -tubulin was quantified by gray scale analysis. Y-Axis represented the relative levels of α -tubulin (control was set as 1). Data were shown as means \pm s.e.m. ($n=3$). β -Actin was used as an internal control. (E) Caspase 3 cleaved α -tubulin *in vitro*. The cleaved α -tubulin represented the fragments of α -tubulin that was cleaved by Caspase 3. The cleaved β -actin represented the three fragments of cleaved β -actin that were cleaved by Caspase 3. * $P<0.05$, ** $P<0.01$, NS, no significance. A full colour version of this figure is available at <http://dx.doi.org/10.1530/ERC-17-0483>.

Table 1 Analysis of the expression level of α -tubulin by immunohistochemical staining of a TMA.

Variable	Case					P	
	Adjacent	Gleason 3+3=6	Gleason 3+4=7	Gleason 4+4=8	Gleason 4+5=9		
Kruskal–Wallis rank sum test for H-score comparison							
H-Score	N	36	12	21	10	5	
	Mean	83.89	212.07	236.64	249.14	269.62	
	s.d.	45.44	31.28	45.25	39.61	26.19	
	Median	75.55	220.13	258.99	264.92	274.10	<0.01
	Min	15.85	163.50	137.25	179.98	228.90	
	Max	191.23	250.94	291.30	287.85	300.00	
	q1	51.68	180.47	209.88	213.34	264.67	
	q3	104.84	236.77	267.37	279.14	280.45	

higher α -tubulin score was obtained statistically in prostate cancer tissues (Table 1), as the higher of Gleason grade, the median score was significantly increased ($P < 0.01$) (Fig. 4B). Together, α -tubulin expression was enhanced and the levels were positively related to the pathological grading, suggesting that α -tubulin was a target for prostate cancer treatment.

After the cells were treated with 20 μ M SFN-Cys for 24h, the Caspase 3 activity was significantly increased in a concentration-dependent manner (Fig. 4C). However, Caspase 3 inhibitor Ac-DEVD-FMK (10 μ M) decreased the SFN-Cys-induced activity of Caspase 3 after it was added to the lysate of SFN-Cys-treated cells, suggesting that the Caspase 3 activity was specifically inhibited by SFN-Cys

(Fig. 4C). Reversely, we added 10 μ M Ac-DEVD-FMK to SFN-Cys-treated cells for 24h, Western blot showed that the downregulation of α -tubulin was reversed (Fig. 4D), indicating that SFN-Cys downregulated α -tubulin by activated Caspase 3.

Besides, we proved that Caspase 3 degraded α -tubulin. Total protein lysates from SFN-NAC-treated and -untreated cells were incubated with or without recombinant Caspase 3 at 37°C, respectively. Only one band of α -tubulin was viewed by Western blot in the non-recombinant Caspase 3-treated cells; however, an α -tubulin doublet was clearly seen in the recombinant Caspase 3-treated cells, and the density of the doublet in SFN-NAC-treated cells was also reduced (Fig. 4E). These results implied that SFN-NAC

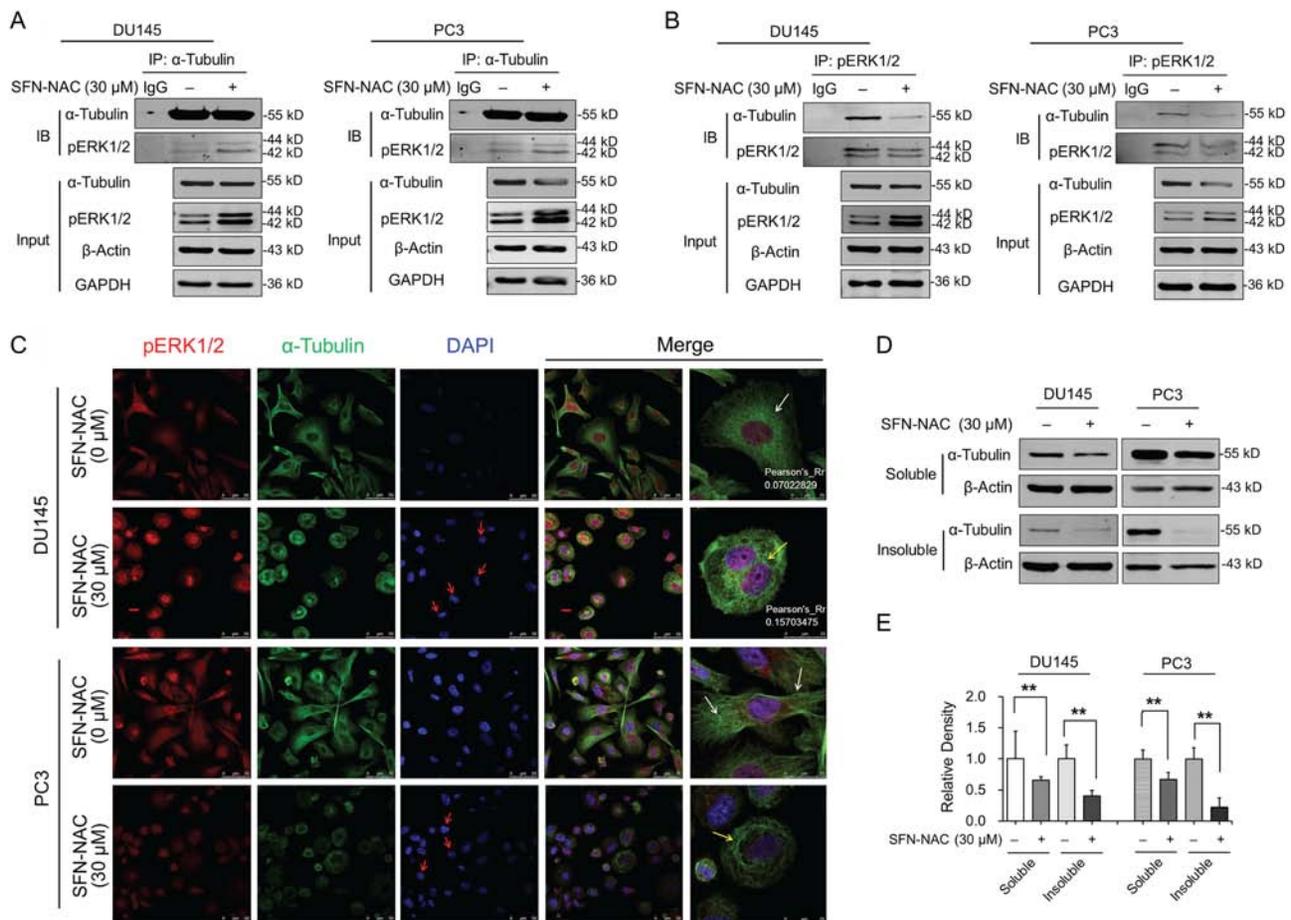


Figure 5

SFN-NAC promoted phosphorylated ERK1/2 binding to α -tubulin leading to the alteration of the microtubule morphology and disturbed microtubule assembly. Cells were treated with 30 μ M SFN-NAC for 24h. The binding of phosphorylated ERK1/2 to α -tubulin was detected in PC3 and DU145 cells by (A) forward and (B) reverse Co-immunoprecipitation (Co-IP). β -Actin and GAPDH were used to be the loading controls for input proteins. The immunoprecipitation via normal mouse IgG was used as the negative control. (C) Immunofluorescence staining of pERK1/2 (Red) and α -tubulin (Green) showed the raising co-localization in cells and the changes of microtubule morphology treated with SFN-NAC. Blue: DAPI-stained DNA; White arrows: normal microtubules; yellow arrows: the abnormal microtubules; red arrows: double nuclei in cells. Scale bars, 25 μ m. The images in last row exhibited the zoom-in merged results. (D) The dynamics of microtubules was measured by microtubule polymerization assay *in vivo*, and β -actin as the loading control. (E) The histogram showed the quantification of soluble and insoluble α -tubulin (the control groups were set as 1). These results were from three independent experiments. * $P < 0.05$, ** $P < 0.01$, NS, no significance.

downregulated α -tubulin, and recombinant Caspase 3 was responsible for the cleavage of α -tubulin in the cell lysates. Endogenously activated Caspase 3 might also degrade α -tubulin (Lin *et al.* 2017). Interestingly, Western blot analysis also showed that the β -actin was clearly cleaved (Fig. 4E). However, β -actin levels were not changed when it was used as an internal control. Thus, we concluded that β -actin might not be a target for activated Caspase 3 *in vivo*.

SFN metabolites mediated the binding of phosphorylated ERK1/2 to α -tubulin alternating the microtubule morphology and disturbing the microtubule assembly

We detected the interaction between α -tubulin and pERK1/2 by endogenous co-immunoprecipitation (Co-IP) after the cells were treated with 30 μ M SFN-NAC for 24 h. Anti- α -tubulin immunoprecipitated with the conjugate of α -tubulin and pERK1/2 in either PC3 or DU145 cells. The amount of pERK1/2 (Co-IP) in SFN-NAC-treated cells was more than that in the untreated cells (Fig. 5A), suggesting that pERK1/2 bound to α -tubulin and SFN-NAC increased the binding significantly. Simultaneously, the reversed Co-IP assay was also performed (Fig. 5B). Results showed that anti-pERK1/2 immunoprecipitated with both pERK1/2 and its interacted α -tubulin protein; the amount of α -tubulin (Co-IP) in SFN-NAC-treated cells was much more than that in the control cells (Fig. 5B). These results indicated that SFN-NAC significantly increased the interaction between pERK1/2 and α -tubulin.

We used immunofluorescence and confocal analysis to verify the colocalization of ERK1/2/ α -tubulin after the cells were treated with 30 μ M SFN-NAC for 24 h (Fig. 5C). The overlapped parts (yellow color) are stronger in the treated DU145 cells than those of the control cells (Fig. 5C). Nevertheless, since large amount of α -tubulin was absent in PC3 cells, the yellow parts are weaker in color in the treated PC3 cells (Fig. 5C). Results showed that SFN-NAC promoted the colocalization of pERK1/2 and α -tubulin, which matched the binding of pERK1/2 to α -tubulin in Co-IP assay. Also, the confocal analysis exhibited more significant morphological changes on microtubules following the treatment with SFN-NAC (Fig. 5C). The control cells exhibited the spindle shape with integrated microtubules, which were orderly distributed in the cytoplasm (white arrows). After treatment with SFN-NAC, abnormal nuclear division showed up (red arrows) and microtubule proteins were disrupted and aggregated around the cell nuclei

(yellow arrows), which constituted 'nest-like' structures of microtubule distribution. Therefore, we concluded that SFN-NAC disrupted microtubules, which might interfere with spindle formation and mitosis leading to apoptosis.

The cytoplasmic α -tubulins are divided into soluble (depolymerization/free) and insoluble (polymerization/microtubule) contents. Here, we discovered a reduction of both soluble α -tubulin and insoluble α -tubulin after the cells were exposed to 30 μ M SFN-NAC (Fig. 5D), and the insoluble α -tubulin in the treated cells were decreased more than that in control cells (Fig. 5E). As mentioned earlier, 30 μ M SFN-NAC downregulated total α -tubulin by analyzing the whole-cell lysates (Fig. 3D). Consequently, SFN-NAC reduced insoluble α -tubulin and downregulated free α -tubulin resulting in microtubule disassembly and 'nest-like' structure of microtubule distribution, which were consistent with the results in immunofluorescence assay.

SFN metabolites regulated the expression and phosphorylation of Stathmin-1, promoting microtubule polymerization

Here, we investigated whether SFN metabolites regulated Stathmin-1 causing the disturbance of microtubule dynamics. Western blot analysis showed that SFN-NAC significantly downregulated Stathmin-1, while phosphorylated Stathmin-1 was elevated (Fig. 6A). Nonetheless, after treatment with PD98059, we found that the decrease of Stathmin-1 was not blocked ($P > 0.05$), while the phosphorylation of Stathmin-1 that SFN-NAC mediated was reduced ($P < 0.05$). These data indicated that SFN-NAC negatively regulated Stathmin-1, while positively regulated the phosphorylation of Stathmin-1 by activating ERK1/2, and both of which promoted the polymerization of microtubules.

However, the natural assembly of microtubules is in a balance of dynamics between polymerization and depolymerization. Although SFN metabolites had the double effects on both polymerization (by Stathmin-1) and depolymerization (by α -tubulin) of microtubules, the overlapping results showed a dynamic imbalance. Finally, SFN metabolites-triggered microtubule disruption contributed to cell apoptosis (Fig. 6B).

Discussion

Previous studies showed that SFN induced apoptosis and inhibited invasiveness in a variety of cancer cells (Zhang *et al.* 1992, 1994, Chiao *et al.* 2002,

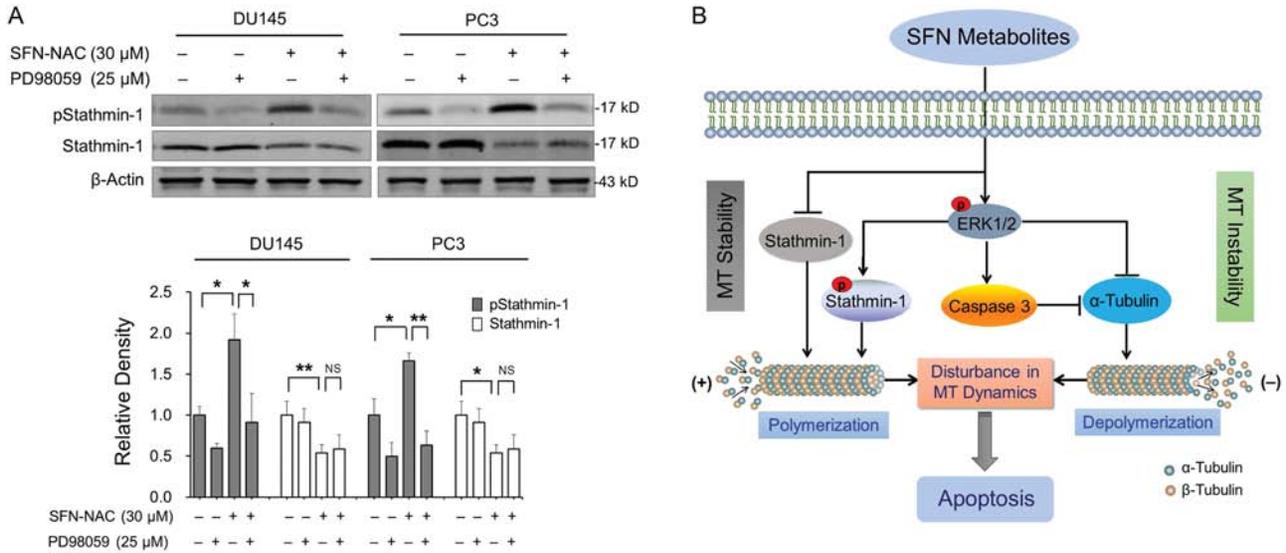


Figure 6 SFN-NAC downregulated Stathmin-1 and promoted Stathmin-1 phosphorylation might link to microtubule polymerization. (A) Total Stathmin-1 and phosphorylated Stathmin-1 (pStathmin-1) were tested by immunoblot after cells were exposed to 30 μM SFN-NAC and 25 μM PD98059 in PC3 and DU145 cells. β-Actin was used as a loading control. **P*<0.05, ***P*<0.01, NS, no significance. Data were shown as means ± s.e.m. (*n*=3). (B) A schematic of the involved signal pathways that SFN metabolites disturbed microtubule dynamics leading to apoptosis in prostate cancer cells. MT, microtubule. A full colour version of this figure is available at <http://dx.doi.org/10.1530/ERC-17-0483>.

Kim *et al.* 2010). However, the half-life of SFN in plasma is too short to execute a persistent action in targeting cancer cells. Inversely, SFN-Cys and SFN-NAC have much longer retention time and might play more effective roles in cancer suppression. Herein we uncovered that two SFN metabolites induced the phosphorylation of ERK1/2 and regulated the downstream effectors leading to microtubule disruption and apoptosis. The microtubule disruption might be triggered by Caspase 3-mediated α-tubulin downregulation as well as Stathmin-1 downregulation and phosphorylation.

Studies showed that SFN interacted with α-tubulin by covalently binding leading to microtubule depolymerization and apoptosis in human lung cancer A549 cells (Xiao *et al.* 2012). Here, we also demonstrated that the increasing expression of α-tubulin contributed to cancer progression and grading of malignancy. Therefore, α-tubulin might be a therapeutic target of SFN metabolites. We discovered that phosphorylated ERK1/2 colocalized and interacted with α-tubulin after the cells were treated with SFN-NAC by both immunofluorescence and co-immunoprecipitation assay. Meanwhile, both SFN-Cys and SFN-NAC downregulated α-tubulin, which was mediated by phosphorylated ERK1/2. These data might also result from ERK1/2-mediated Caspase 3 activation and upregulation. Apparently, ERK1/2 played a pivotal role in α-tubulin-linked microtubule disturbance.

It was reported that extracellular signal-regulated kinase 1 (MEK1) interacted with α-tubulin to determine the mitotic stability in cultured human fibrosarcoma cells (Cao *et al.* 2010). Further, MEK1/2 regulated microtubule assembly, spindle pole tethering and asymmetrical division by altering the stability of microtubule dynamics during meiotic maturation of mouse oocytes (Yu *et al.* 2007). These results were also consistent with the results in the SFN-treated neuroblastoma cells; ERK activation by sulforaphane contributed to growth inhibition and apoptosis (Yao *et al.* 2015, Mondal *et al.* 2016). Thus, the activated ERKs are the key triggers to induce downstream cascades. These regulations of α-tubulin-associated proteins constitutively might contribute to microtubule disruption. It was well known that activation of Caspase 3 might involve either the mitochondrial or the death receptor pathway leading to cell death (Singh *et al.* 2004). Caspases were shown to be capable of cleaving cytoskeletal proteins, such as Tau, α-tubulin, Drebrin, Spinophilin, etc. (Klaiman *et al.* 2008). Here, we found that SFN metabolites activated Caspase 3 and the cleaved Caspase 3 was elevated. Cellular α-tubulin was cleaved in the cancer cells with intracellular treatment by SFN metabolites combined with extracellular Caspase 3 digestion. These data indicated that Caspase 3 activation induced by SFN metabolites had the potential to cleave α-tubulin because no α-tubulin was digested by recombinant Caspase 3 in

the control cells. These procedures provided necessary support for degradation of cell skeleton. As a SFN analogue, NAC-allyl isothiocyanate was shown to induce dose-dependent activation of Caspase 3 and cleavage of α -tubulin in human bladder cancer (Bhattacharya *et al.* 2012). Coincidentally, we demonstrated that SFN-Cys activated Caspase 3 and cleaved α -tubulin leading to cell apoptosis in human non-small lung cells (Lin *et al.* 2017). Here, we further demonstrated that both α -tubulin and β -actin were cleaved by recombinant Caspase 3 in the SFN metabolite-treated cells although β -actin was not changed as an internal control. The treated cells might contain more particular mechanisms to protect β -actin from protease degradation.

Stathmin-1 downregulation might reverse the malignant phenotypes (Jackson *et al.* 2007, Azarenko *et al.* 2008, 2014). Stathmin-1 either sequestered tubulin dimers or stimulated microtubule catastrophes leading to promotion of microtubule disassembly (Jackson & Singletary 2004). Also, Stathmin-1 bound to α -tubulin stimulating α -tubulin's intrinsic GTPase activity, promoting α -tubulin hydrolysis (Belmont & Mitchison 1996). The microtubule-destabilizing activity of Stathmin-1 can be turned off by phosphorylation in response to the stimulating signals, including those for cell growth and cell cycle progression (Andersen *et al.* 1997). Herein, we found that SFN metabolites induced the phosphorylation of Stathmin-1 and downregulated the total Stathmin-1. These data provided the evidence that SFN metabolites might play key roles in microtubule inhibition by modulating the interaction between Stathmin-1 and α -tubulin. Studies showed that mutation of Stathmin-1 phosphorylation site decreased sensitivity to Bortezomib and influenced α -tubulin polymerization (Ge *et al.* 2010). Besides, the phosphorylation of Stathmin-1 reduced α -tubulin binding and inhibited Stathmin-1-induced destabilization of microtubules *in vitro* (Bhat & Setaluri 2007, Wegiel *et al.* 2016). Studies also showed that the anti-microtubule drug Taxol stabilized microtubule assembly and blocked cell cycle progression (Van Vaerenbergh *et al.* 1998), therefore, SFN metabolites might inhibit cell growth via microtubule disruption.

SFN-bound α -tubulin might form aggresome-like protein aggregates, and these aggregates can be degraded via the ubiquitin-proteasome system (UPS) (Mi *et al.* 2009a,b). Report also showed that Hsp27 upregulation induced by SFN might increase the proteasome activity in HeLa cells (Gan *et al.* 2010). The alteration of either soluble or insoluble α -tubulin in microtubule might influence microtubule treadmilling and shortening

(Mi *et al.* 2009a,b). The drugs that either stabilize or destabilize microtubules have potential to bind to soluble or insoluble α -tubulin to induce apoptosis (Mukhtar *et al.* 2014). Taxol bound to β -tubulin, while SFN metabolites bound to α -tubulin, and these two tubulins are the targets of anti-cancer drugs (Jordan 2002). Nonetheless, taxol might generate drug resistance (Jordan & Wilson 2004), while SFN metabolites, as plant-derived components might have less toxicity and side effects. It has been reported that Estramustine binds to a novel site in α -tubulin; the combination of Estramustine and either Vinblastine or Taxol might cause synergetic effects in cancer inhibition (Panda *et al.* 1997, Chen *et al.* 2013). Therefore, combination of these drugs with SFN metabolites might produce more efficient results of treatment with a lower dose, toxicity and less resistance.

Taken together, SFN metabolites promoted activation of ERK1/2 and Caspase 3 and regulated the expression of α -tubulin, total Stathmin-1 and phosphorylated Stathmin-1 leading to microtubule disruption and apoptosis. These results might help us understand the mechanisms of carcinogenesis to establish a novel therapeutic approach to target prostate cancer cells.

Declaration of interest

The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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Author contribution statement

Conception and design: W W, Y Z, G X Y and H T. Acquisition of data: Y Z, G X Y and H T. Analysis and interpretation of data: Y Z, G X Y, H T, Y B H, S W, Y G and K L. Writing, review and/or revision of the manuscript: W W and Y Z. Administrative, technical or material support: Y Z, G X Y, H T, Y B H, S W, Y G and K L. Study supervision: W W.

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