

# Allylated Curcumin Analog CA6 Inhibits TrxR1 and Leads to ROS-Dependent Apoptotic Cell Death in Gastric Cancer Through Akt-FoxO3a

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Vinothkumar Rajamanickam\*  
Tao Yan\*  
Liangrong Wu  
Yanni Zhao  
Xiaohong Xu  
Heping Zhu  
Xi Chen  
Meihong Wang  
Zhouli Liu  
Zhiguo Liu  
Guang Liang  
Yi Wang 

Chemical Biology Research Center,  
School of Pharmaceutical Sciences,  
Wenzhou Medical University, Wenzhou,  
Zhejiang 325035, People's Republic of  
China

\*These authors contributed equally to  
this work

**Background:** Gastric cancer is one of the leading causes of cancer-related deaths. Allylated monocarbonyl analogs of curcumin (MACs) have been reported to selectively inhibit a broad range of human cancers including gastric cancer. However, the precise molecular mechanisms underlying the inhibitory activities of MACs are not fully known.

**Methods:** In this study, we examined the anti-tumor activities of an allylated MAC, CA6, on gastric cancer cells and gastric cancer xenograft mouse model. The potential molecular anti-tumor mechanisms of CA6 were also elucidated.

**Results:** Our data show that CA6 exhibited significant cytotoxicity in gastric cancer cells, which was seen as an induction of G2/M cell cycle arrest and apoptosis. These activities were mediated through an elaboration of ROS levels in gastric cancer cells and induction of endoplasmic reticulum stress. CA6 increased ROS levels through directly binding to and inhibiting thioredoxin reductase R1 (TrxR1). Also, CA6-generated ROS inhibited Akt and activated forkhead O3A (FoxO3a), causing cytotoxicity in gastric cancer cells. Finally, CA6 treatment dose-dependently reduced the growth of gastric cancer xenografts in tumor-bearing mice, which was associated with reduced TrxR1 activity and increased ROS in the tumor.

**Conclusion:** In summary, our studies demonstrate that CA6 inhibited gastric cancer growth by inhibiting TrxR1 and increasing ROS, which in turn activated FoxO3a through suppressing Akt. CA6 is a potential candidate for the treatment of gastric cancer.

**Keywords:** gastric cancer, curcumin analog, reactive oxygen species, TrxR1, FOXO3a

## Introduction

Gastric cancer is one of the leading causes of cancer-related death worldwide.<sup>1,2</sup> Many patients have inoperable disease at diagnosis.<sup>3</sup> Typically, locally advanced disease is treated with adjuvant or neoadjuvant therapy, in possible combination with surgery. However, the metastatic disease outcome is extremely poor.<sup>4</sup> In addition, development of drug resistance, disease relapse, and severe side effects of therapies contribute to morbidity and mortality in gastric cancer patients.<sup>5</sup> Therefore, these major clinical challenges urgently require new and safe treatment options.

Cancer cells may be distinguished from healthy cells in part by their metabolic phenotype. Metabolic perturbations of cancer cells result in higher level of endogenous reactive oxygen species (ROS). Studies have reported high levels of hydrogen peroxide in a panel of human tumor cell lines.<sup>6</sup> Increased levels of ROS may be derived through NADPH oxidases<sup>7,8</sup> and loss of tumor suppressor pathways altering antioxidant networks.<sup>9</sup> Cancer cells may utilize elevated ROS for progression, increasing the rate

Correspondence: Yi Wang  
Chemical Biology Research Center,  
School of Pharmaceutical Sciences,  
Wenzhou Medical University, Wenzhou,  
Zhejiang 325035, People's Republic of  
China  
Tel/Fax +86- 577-85773060  
Email yi.wang1122@wmu.edu.cn

of DNA mutations. One possible mechanism by which cancer cells escape the cytotoxic effects of elevated ROS is to maintain cell viability through the upregulation of reductive pathways. The thioredoxin (Trx) system using NADPH channeled through thioredoxin reductase 1 (TrxR1) is one of the major redox systems. Increasing evidence suggests that many redox regulators are involved in resistance to anticancer drugs.<sup>10</sup> Targeting TrxR1 has been shown to occur with many different electrophiles with anticancer potential.<sup>11</sup>

Recent studies have shown that compounds that further enhance ROS levels in cancer may bypass the elevated reductive pathways in cancer cells and unmask ROS cytotoxicity.<sup>12</sup> One of such ROS altering compounds is curcumin.<sup>13</sup> However, curcumin failed to demonstrate an efficacy profile in clinical trials because of its highly reactive  $\beta$ -diketone moiety.<sup>14</sup> In our previous studies, we synthesized several allylated monocarbonyl analogs of curcumin (MACs) with enhanced selective toxicity and improved pharmacokinetic properties.<sup>15</sup> Among these analogs, CA6 (Figure 1A) showed significant growth inhibition in gastric and colon cancer cells, while exhibiting markedly lower cytotoxicity in normal gastric epithelial cells.<sup>16</sup> In the present study, we aimed to identify the mechanisms underlying the inhibitory effects of CA6 in gastric cancer cells. We show that CA6 inhibited the growth of gastric cancer cells through ROS-dependent activation of the endoplasmic reticulum (ER) stress and induction of apoptosis. Importantly, our results show that CA6 increased ROS by directly binding to and inhibiting the activity of TrxR1. Elevated ROS inhibited Akt and activated forkhead O3A (FoxO3a) in gastric cancer cells to induce cytotoxicity.

## Materials and Methods

### Chemicals and Reagents

Curcumin, methyl thiazolyl tetrazolium (MTT), and antioxidant N-acetyl cysteine (NAC) were purchased from Sigma-Aldrich (St. Louis, MO). Cell cycle and apoptosis detection kits, Annexin V/Propidium iodide (PI) were purchased from BD Biosciences (Franklin Lakes, NJ). Reactive oxygen species probes 2,7-Dichlorodihydrofluorescein diacetate (DCFH-DA), dihydroethidium (DHE), and lipid peroxidation byproduct malondialdehyde (MDA) assay kits were obtained from Beyotime Institute of Biotechnology (Nantong, China). Caspase-9 activity assay, cytosolic and nuclear protein extraction kits, crystal violet dye, Hoechst 33258, and 4',6-diamidino-2-phenylindole (DAPI) staining solutions were purchased from Beyotime Institute of

Biotechnology. Antibodies against Akt, phosphorylated Akt (p-Akt, Thr208 and Ser473), activating transcription factor-4 (ATF4), eukaryotic initiation factor 2 $\alpha$  (eIF2 $\alpha$ ), phosphorylated eIF2 $\alpha$ , CCAAT-enhancer-binding protein homologous protein (CHOP), FOXO3a, and p-FOXO3a were purchased from Cell Signaling Technology (Danvers, MA). Antibodies against p-FOXO3a, lamin B1 and cell proliferation marker Ki67 were purchased from Abcam (Cambridge, UK). Antibodies against murine double minute 2 (MDM2), cell division cycle-2 (CDC2), Cyclin B1, cleaved-poly (ADP ribose) polymerase (PARP), phosphatase and tensin homolog deleted on chromosome 10 (PTEN), TrxR1, GAPDH, and horseradish/fluorophore-conjugated secondary antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). CA6 was synthesized as outlined previously.<sup>15</sup>

### Gastric Cancer Cell Lines

Human gastric cancer cell lines BGC-823 and SGC-7901 were obtained from the Institute of Biochemistry and Cell Biology, Chinese Academy of Sciences (Shanghai, China). Cells were cultured in RPMI-1640 medium (Gibco, Eggenstein, Germany) supplemented with 10% heat-inactivated fetal bovine serum (FBS; Hyclone, Logan, UT), 100 units/mL penicillin, and 100  $\mu$ g/mL streptomycin (Gibco, Eggenstein, Germany).

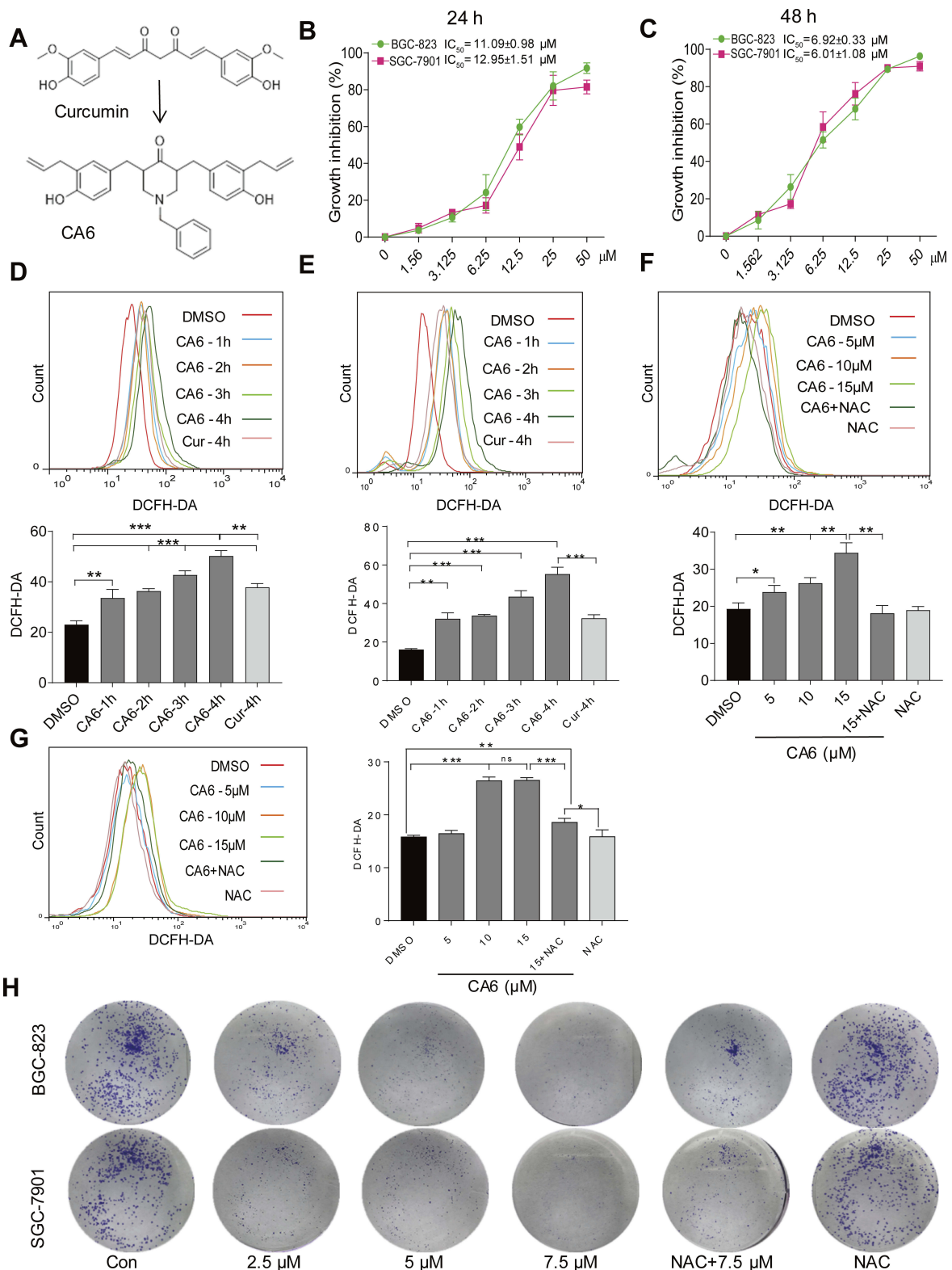
### Assessment of Cell Viability

Cell viability was measured using MTT assay. Cells were seeded in 96-well plates at a density of  $8 \times 10^3$  cells per well in 100  $\mu$ L complete growth medium. Next day, cells were treated with increasing concentration of CA6 for 24 h or 48 h. CA6 was dissolved in DMSO and DMSO alone was used as vehicle control. MTT solution was then added and cells were incubated for 4 hrs at 37°C. DMSO was added to each well. The absorbance was measured at 490 nm using Spectra MAX190 (Molecular Device, USA). Cell viability was calculated using the following formula:

$$\text{Growth inhibition (\%)} = \frac{(\text{absorbance of control} - \text{absorbance of CA6})}{\text{absorbance of control}} \times 100\%$$

### Measurement of Intracellular Reactive Oxygen Species (ROS) in Cultured Cells

Gastric cancer cells were seeded in 6-well plates at a density of  $3 \times 10^5$  cells per well and allowed to attach overnight. Cells were then exposed to CA6 (15  $\mu$ M) or



**Figure 1** CA6 inhibits gastric cancer cell growth and increases ROS levels. **(A)** The chemical structures of parental compound curcumin and its analog CA6. **(B and C)** BGC-823 and SGC-7901 cells were challenged with increasing concentrations of CA6 for 24 h **(B)** or 48 h **(C)**. Cell viability was measured by the MTT assay. IC<sub>50</sub> values for CA6 in the two different cell lines are presented [Mean ± SEM; n = 3]. **(D and E)** Cells were treated with CA6 (15 μM) for different time periods and intracellular ROS levels were measured by flow cytometry. Data showing DCFH-DA staining in BGC-823 **(D)** and SGC-7901 **(E)** cells. Curcumin (20 μM) was used as the positive control [Mean ± SEM; n = 3; \*\*P<0.01, \*\*\*P<0.001]. **(F and G)** Cells were treated with CA6 at indicated concentrations for 4 h. NAC was administered at 5 mM alone or as a 2 h pretreatment before exposure of cells to CA6 (15 μM). Data showing DCFH-DA staining of BGC-823 **(F)** and SGC-7901 **(G)** cells [Mean ± SEM; n = 3; \*P<0.5, \*\*P<0.01, \*\*\*P<0.001, ns = not significant]. **(H)** Effect of CA6 on the colony-forming ability of gastric cancer cells. BGC-823 and SGC-7901 cells were exposed to various concentrations of CA6 for 24 h, with or without pretreatment with NAC (5 mM) for 2 h. Colony detected by crystal violet staining (blue). Representative images shown.

curcumin (20  $\mu$ M) for various time points. In some experiments, cells were pretreated with NAC (5 mM) for 2 h prior to the CA6 treatment. Cells were then stained with DCFH-DA and fluorescence was measured using FACS Calibur flow cytometer (BD Biosciences, CA).

## Colony Formation Assay

BGC-823 and SGC-7901 cells were seeded in 6-well plates and incubated for 24 h. Cells were then exposed to various concentrations of CA6 for 24 h, with or without pretreatment with NAC (5 mM) for 2 h. Medium was replaced with fresh culture medium and cells were further cultured for another 6 days. Cells were washed with PBS, fixed in 4% paraformaldehyde and stained with crystal violet.

## Cell Cycle and Apoptosis Analysis

BGC-823 and SCG-7901 cells were seeded in 6-well plates at a density of  $3 \times 10^5$  cells per well and allowed to attach overnight. The following day, cells were challenged with CA6 at 5, 10 or 15  $\mu$ M for 16 h (cell cycle analysis) or 24 h (apoptosis analysis), with or without NAC pretreatment (5 mM) for 2 h. Cells were fixed in ice-cold 75% ethanol. Cells were labeled with PI at a final concentration of 0.05 mg/mL for cell cycle phase detection. For apoptosis detection, cells were harvested in 500  $\mu$ L binding buffer and labeled with FITC-conjugated Annexin V for 10 min and PI for 5 min. Analysis was performed using FACS Calibur flow cytometer. Data for cell cycle distribution and apoptosis were analyzed using FlowJo7.6 software (TreeStar, San Carlos, CA).

## Western Blot Analysis

Protein lysates from cells and tumor tissues were prepared using RIPA lysis buffer supplemented with protease inhibitor cocktail. Cytosolic and nuclear proteins were extracted using fractionation kits from Beyotime. Protein concentrations were determined via Bradford protein assay (Bio-Rad, Hercules, CA). Proteins were separated in 10–15% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to polyvinylidene difluoride (PVDF) membranes. The membranes were then blocked with 5% skim milk in tris-buffered saline containing 0.1% Tween 20 (TBST) for 90 min at room temperature. Subsequently, primary antibodies (1:1000) in TBST were added and membranes incubated overnight at 4°C. After washing with TBST, horseradish peroxidase (HRP)-conjugated secondary antibodies (1:3000) were added for 1 h at room temperature. After additional washing with

TBST, immunoreactivity was detected by enhanced chemiluminescence detection system (ECL, GE Healthcare). Densitometric quantification was performed using Image J software (National Institute of Health, MD). Protein levels were normalized to loading control GAPDH. In the case of phosphorylated proteins, levels were normalized to respective total proteins determined by antibodies recognizing both phosphorylated and unphosphorylated forms.

## Cell Transfection for Gene Silencing

Sequences of siRNA targeting ATF4 and TrxR1 are shown in [Supplementary Table S1](#). siRNA oligonucleotides were synthesized by GenePharma (Shanghai, China). BGC-823 cells were seeded at a density of  $1 \times 10^5$  in 6-well plates for 24 hrs. siRNA against human ATF4, TrxR1, or non-targeting control siRNA (GenePharma) were transfected at a final concentration of 50 pmol (ATF4) or 100 nmol (TrxR1) using lipofectamine 3000 reagent (Invitrogen, CA) in serum-free medium for 6 hrs. Complete growth medium was then added and the cells were cultured for an additional 24 hrs. Levels of silenced genes were determined by Western blotting.

## Determination of Caspase-9 Activity

Caspase-9 activity in cells and tissue lysates was determined using Caspase-9 Assay Kit (Beyotime). Caspase-9 activity was normalized to total protein concentration in the lysates and expressed as percentage of control.

## CA6 and TrxR1 Interaction Studies

To identify the interaction between CA6 and TrxR1, three approaches were used. Firstly, molecular docking study was carried out using Tripos molecular modeling package Sybyl-x.v1.1.083 (Tripos, St. Louis, MO, USA). The crystal structure of human TrxR1 used for docking was obtained from the Protein Data Bank (PDB ID 2ZZ0, chain A). AutoDock Tools version 1.5.6 and PyMol was used to analyze the docking results.

Secondly, the binding of CA6 with recombinant human (rh) TrxR1 (Sigma-Aldrich) protein was determined using ProteOn XPR36 Protein Interaction Array system (Bio-Rad Laboratories, Hercules, CA) with a GLH sensor chip (ProteOn). Briefly, rhTrxR1 was loaded to the sensor, which was activated with 10 mM NiSO<sub>4</sub>, and the CA6 samples were prepared with running buffer (PBS, 0.1% SDS, 5% DMSO). Sensor and sample plates were placed on the instrument. CA6 samples were then captured in the

first flow cells, and the second flow cell was left as a blank. Six concentrations of CA6 were simultaneously injected at a flow rate of 30 mM/min for 120 s of association phase, followed by 120 s of dissociation phase at 25°C. The final graphs represent the difference between the duplex or quadruplex sensorgrams and the blank sensorgrams. Data analysis was done using the ProteOn Manager Software, and the KD values were calculated using aligning the kinetic data from various concentrations of CA6 to the 1:1 langmuir binding model.

Finally, CA6 binding to TrxR1 was determined using 1.1'-bis(anilino)-4,4'-bis(naphthalene)-8,8'-disulfonate (bis-ANS) displacement assay. Briefly, bis-ANS (5 µM) and rhTrxR1 proteins (5 nM) were mixed in PBS (pH 7.4) and incubated for 15 min to reach stable relative fluorescence units (RFUs) emitted at 430–590 nm under the excitation at 385 nm. Then, non-fluorescent CA6 at 1, 2, 4, 6, 8 and 18 µM was added. RFUs at 430–590 nm were measured.

### TrxR1 Activity Assays

TrxR1 activity was determined at room temperature using SpectraMax M5 microplate reader (Molecular Devices, USA). NADPH-reduced TrxR1 at 160 nM was incubated with various concentrations of CA6 for 2 h at room temperature in 96-well plates. A mixture of TE buffer (50 mM Tris-HCl, pH 7.5; 1mM EDTA, 50 mL) containing 5,5'-dithiobis(2-nitrobenzoate) (DTNB) and NADPH was added to achieve final concentrations of 2 mM and 200 mM, respectively. The linear increase in absorbance at 412 nm during the initial 3 min was recorded. The same amounts of DMSO (1%, v/v) were added to the control experiments and the activity was expressed as the percentage of the control.

TrxR1 activity in lysates was measured by end-point insulin reduction assay as described previously.<sup>17,18</sup> Briefly, 100 µg total proteins were incubated in a final reaction volume of 50 µL, containing 100 mM Tris-HCl (pH 7.6), 0.3 mM insulin, 660 µM NADPH, 3 mM EDTA, and 15 µM E. coli-derived Trx (Sigma-Aldrich) for 1 hr at 37°C. The reaction was terminated by adding 200 µL of 1 mM DTNB. A blank sample, containing everything except Trx, was treated in the same manner. The absorbance at 412 nm was measured, and the blank value was subtracted from the corresponding absorbance value of the sample. The activity was expressed as the percentage of the control.

### In vivo Gastric Cancer Xenograft Model

Seven-week-old athymic BALB/c nu/nu female mice weighing 20–22 g were purchased from Vital River Laboratories (Beijing, China). Animal studies conducted are in compliance with the ARRIVE guidelines.<sup>19,20</sup> All experimental protocols involving animal were approved by the Wenzhou Medical University Policy on the Care and Use of Laboratory Animals (Approved documents: wyd2014-0059). Animals were housed (6 animals/cage) at a constant room temperature with a 12/12 h light/dark cycle and fed a standard rodent diet and water ad libitum.

BGC-823 cells were injected subcutaneously into the right flank of each mouse at  $1 \times 10^6$  cells in 0.1 mL PBS. Once the tumors reached a volume of 100–160 mm<sup>3</sup>, mice were randomly distributed into 4 groups (n=6) as follows: 1) control mice treated with vehicle only; 2) mice treated with 10 mg/kg CA6; 3) mice treated with 20 mg/kg CA6; 4) mice treated with 50 mg/kg curcumin. All treatments were carried out with intraperitoneal injections every day. Tumor volumes were determined every day by measuring length (l) and width (w) and calculated as  $0.5 \times l \times w^2$ . Body weights were also determined every day. On day 12, mice were euthanized and the tumor specimens were harvested and weighed. Tumor specimens were then processed for histological analyses and protein measurement studies. MDA content in tumor tissue lysates was analyzed using lipid peroxidation MDA assay (Beyotime).

### Determination of Tumor Superoxide Production

Portion of tumor tissues extracted from mice was immediately embedded in OCT compound and cut into 5 µm-thick sections. Tissue sections were then incubated with dihydroethidium (DHE, 10 mM) in PBS at 37°C for 30 min in dark. DHE is oxidized upon reaction with superoxide to ethidium bromide, which then binds to DNA. Red fluorescence from ethidium bromide was viewed and captured using a fluorescence microscope ( $\lambda_{ex}$  490 nm,  $\lambda_{em}$  610 nm, Nikon, Japan).

### Immunohistochemical Staining of Tissues

Harvested tumor tissues were fixed in 10% formalin and embedded in paraffin. Specimens were sectioned at 5 µm thickness. Tumor sections were stained using routine immunohistochemical techniques with primary antibodies against Ki67 (1:200), PTEN (1:100), p-Akt (1:200), or FoxO3a (1:200), respectively. HRP-conjugated secondary antibodies

and diaminobenzidine (DAB) were used for detection. Tumor specimens as well as heart, liver and kidney tissues were also stained with hematoxylin and eosin (H&E).

## Statistical Analysis

Statistical analyses were performed only when a minimum of  $n = 3$  independent samples were acquired. All data are reported as mean  $\pm$  SEM. Statistical analysis was performed using GraphPad Prism 6.0 software (GraphPad, San Diego, CA). One-way ANOVA followed by Dunnett's post hoc test was used when comparing more than two groups of data, and one-way ANOVA, non-parametric Kruskal–Wallis test followed by Dunn's post hoc test was used when comparing multiple independent groups. When comparing two groups, the unpaired Student's *t*-test was used. A *P* value  $< 0.05$  was considered statistically significant.

## Results

### CA6 Reduces Cell Viability of Gastric Cancer Cells via Inducing Intracellular ROS

We firstly measured the viability of gastric cancer cells upon exposure to CA6. BGC-823 and SGC-7901 cells were challenged with increasing concentrations of CA6 and cell viability was measured using MTT assay. As shown in [Figure 1B](#) and [C](#), CA6 dramatically decreased cell viability of both gastric cancer cell lines after 24- and 48-h treatment. At 24-h post-exposure, we obtained the half-maximal inhibitory concentration ( $IC_{50}$ ) values of  $11.09 \pm 0.98$  and  $12.95 \pm 1.51 \mu\text{M}$  for BGC-823 and SGC-7901 cells, respectively. Longer exposure at 48 h appeared to be more effective, as seen by  $IC_{50}$  values of  $6.92 \pm 0.33$  and  $6.01 \pm 1.08 \mu\text{M}$  for BGC-823 and SGC-7901 cells, respectively.

Previously, we have reported that elevated ROS is the primary mediator of cytotoxicity induced by many curcumin analogs.<sup>16</sup> Therefore, we evaluated whether the inhibitory effect of CA6 on gastric cancer cells involved intracellular ROS accumulation. As expected, CA6 increased ROS levels in both BGC-823 ([Figure 1D](#)) and SGC-7901 cells ([Figure 1E](#)). Curcumin, used as a positive control, also increased ROS levels ([Figure 1D](#) and [E](#)). These results suggest that CA6 is an inducer of ROS in gastric cancer cells. Next, we pretreated BGC-823 and SGC-7901 cells with NAC (N-acetyl cysteine, 5 mM), a specific ROS inhibitor, for 2 h prior to CA6 exposure.

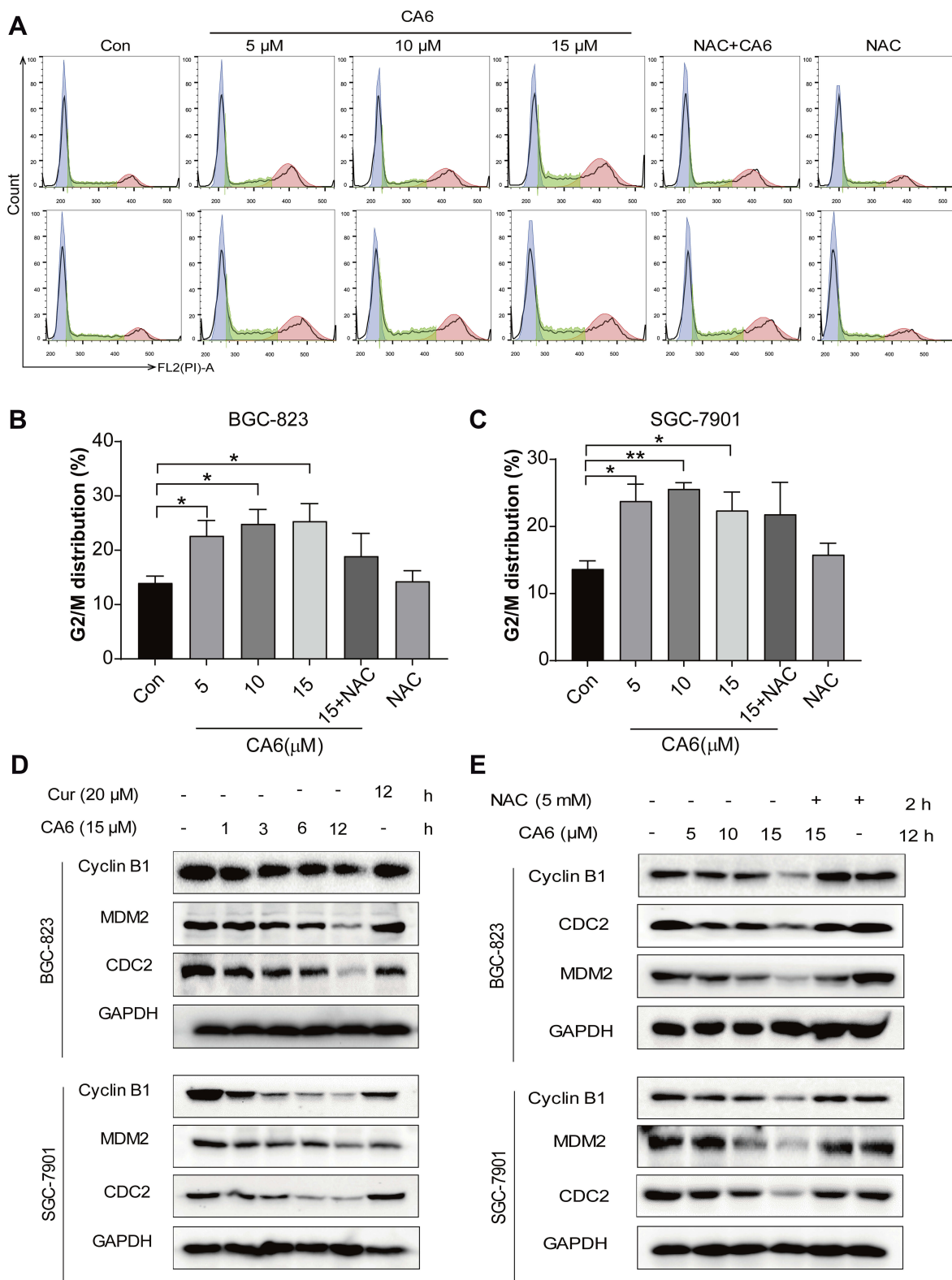
Our results show that NAC pretreatment decreased the levels of ROS in both tested gastric cancer cells ([Figure 1F](#) and [G](#)). In addition, colony-forming ability of gastric cancer cells was also suppressed by CA6 ([Figure 1H](#)). Whereas, pretreatment with NAC significantly reversed the inhibitory effect of CA6 ([Figure 1H](#)). These findings suggest that CA6-induced intracellular ROS accumulation may be a major cellular mechanism of its inhibitory activity against gastric cancer cells.

### CA6-Induced ROS Causes G2/M Cell Cycle Arrest

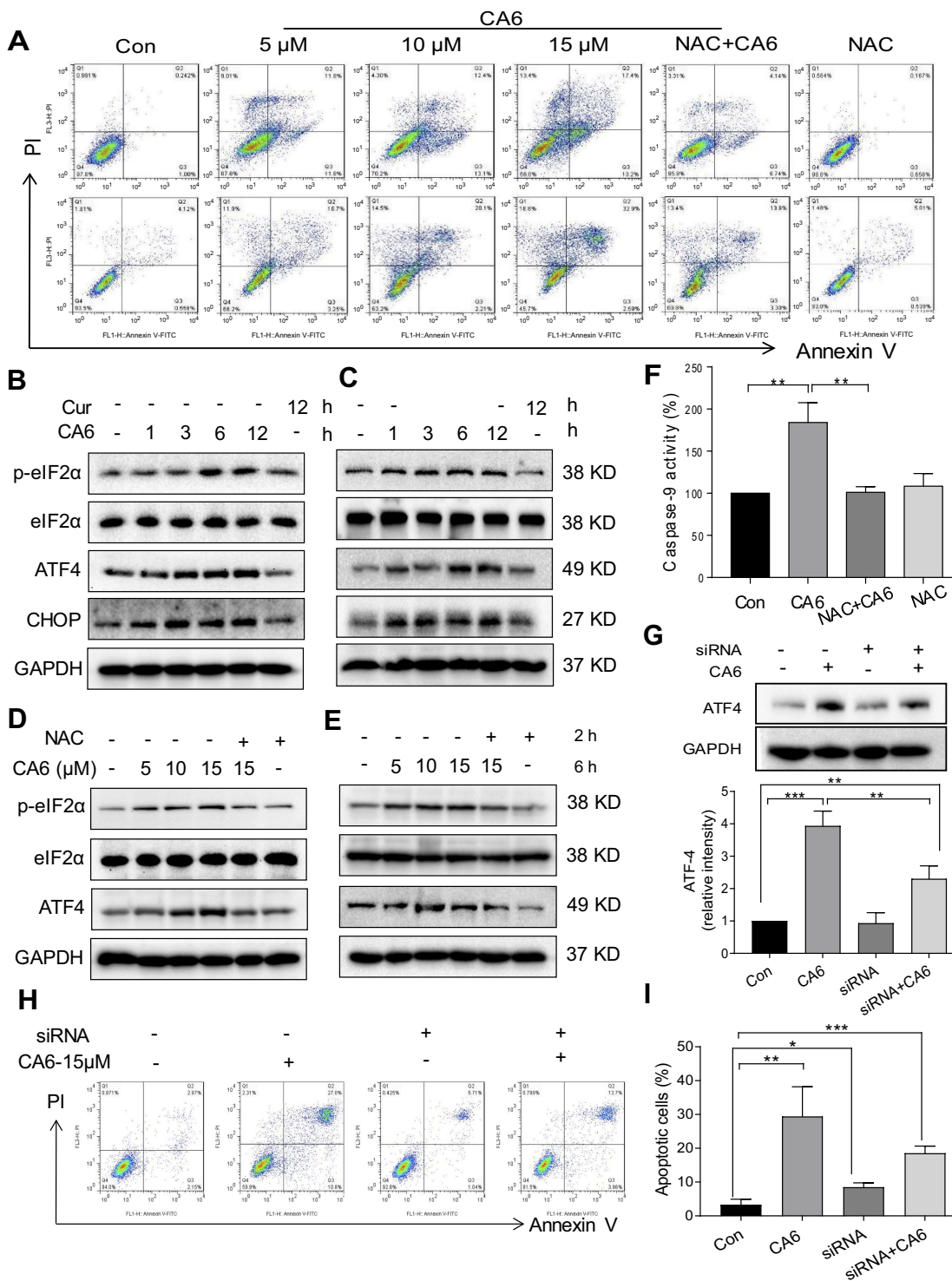
We next examined the possible effect of CA6 on cell cycle regulation. Flow cytometric analysis revealed an accumulation of cells in the G2/M phase after CA6 exposure ([Figure 2A–C](#)). However, NAC pretreatment significantly reduced CA6-induced cell arrest in the G2/M phase ([Figure 2A–C](#)). These results show that CA6 reduced cell viability in part through halting cycle progression. We confirmed these results by measuring G2/M cell cycle-associated proteins cyclin B1, murine double minute (MDM2) and cell division cycle protein 2 (CDC2). Consistent to the data of cell cycle analysis, CA6 treatment reduced the protein levels of cyclin B1, MDM2 and CDC2 ([Figure 2D](#)). The inhibitory effects of CA6 on the expression of these proteins were much stronger than those of curcumin ([Figure 2D](#)). Furthermore, NAC pretreatment prevented CA6-mediated decrease of cell cycle regulating proteins ([Figure 2E](#)). These results suggest that the cell cycle arrest effect of CA6 is partially through the induction of ROS.

### CA6 Induces ROS-Dependent ER Stress Response

To investigate the underlying mechanism associated with CA6-induced gastric cancer cell growth inhibition, we examined whether CA6-induced apoptosis. Our data showed that CA6-induced apoptosis in both BGC-823 and SGC-7901 cells ([Figure 3A](#), [Supplementary Figure S1](#)). NAC pretreatment decreased the number of apoptotic cells, confirming a ROS-mediated activity of CA6 ([Figure 3A](#), [Supplementary Figure S1](#)). Morphological examination also shows apoptotic changes, including cell shrinkage, enlarged nuclei, nuclear fragmentation and chromatin condensation after CA6 treatment ([Supplementary Figure S1C–D](#)). Furthermore, CA6 treatment significantly increased the activity of caspase-9 in BGC-823 cells, while



**Figure 2** CA6 induces ROS-dependent G2/M cell cycle arrest. **(A)** BGC-823 (first row) and SGC-7901 (second row) were challenged with CA6 for 16 h, with or without pretreatment with NAC (5 mM) for 2 h. Cell cycle distribution was analyzed by PI staining. Representative histograms are shown [n = 3]. **(B and C)** Quantification of cells in the G2/M phase cells following exposure to CA6. Cells were treated as indicated in panel A [Mean ± SEM; n = 3; \*P<0.05, \*\*P<0.01, \*\*\*P < 0.001]. **(D)** BGC-823 and SGC-7901 cells were exposed to 15 μM CA6 for indicated time points. Levels of cell cycle regulating proteins Cyclin B1, MDM-2 and CDC-2 were determined by Western blot. Curcumin (20 μM) was used as the positive control and GAPDH as the loading control. Representative blots shown [n = 3]. **(E)** BGC-823 and SGC-7901 cells were exposed to CA6, with or without NAC pretreatment (5 mM). Levels of Cyclin B1, MDM-2 and CDC-2 were determined. GAPDH was used as loading control. Representative blots shown [n = 3].



**Figure 3** CA6 induces ER stress and gastric cancer cell apoptosis. **(A)** BGC-823 and SGC-7901 cells were pretreated with NAC (5 mM) for 2 h before exposure to CA6 for 24 h. Percentage of cell apoptosis was determined by Annexin-V/PI staining. Representative dot plots shown [n = 3]. **(B, C)** BGC-823 (B) and SGC-7901 (C) cells were challenged with CA6 (15  $\mu$ M) for indicated time points. Levels of ER stress-related proteins p-eIF2 $\alpha$ , ATF4, and CHOP were determined. Curcumin was used as a positive control at the dose of 20  $\mu$ M. GAPDH was used as the loading control. Representative blots shown. **(D, E)** BGC-823 (D) and SGC-7901 (E) cells were exposed to CA6 at the dose of 5, 10 and 15  $\mu$ M for 6 h. Where indicated, NAC pretreatment (5 mM) was carried out for 2 h. Cells treated with curcumin (20  $\mu$ M) for 12 h were used as the positive control. Protein levels of p-eIF2 $\alpha$ , ATF4 and CHOP were determined. **(F)** BGC-823 cells were transfected with siRNA against ATF4. Cells were then exposed to CA6 (15  $\mu$ M) for 6 h. Levels of ATF4 were determined. GAPDH was used as loading control. Representative immunoblot (upper panel) and densitometric quantification (lower panel) is shown [Mean  $\pm$  SEM; n = 3; \*\*P<0.01]. **(G)** BGC-823 cells were exposed to CA6 (15  $\mu$ M) for 12 h, with or without pretreatment with NAC (5 mM) for 2 h. The activity of caspase-9 was determined [Mean  $\pm$  SEM; n = 3; \*\*P<0.01, \*\*\*P<0.001]. **(H, I)** Flow cytometric analysis of apoptotic cells following knockdown of ATF4 and exposure to CA6. BGC-823 cells transfected with ATF4 siRNA were exposed to CA6 (15  $\mu$ M) for 24 h. Dot plots (H) and quantification of apoptotic cells (I) are shown [Mean  $\pm$  SEM; n=3; \*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001].



pretreatment with NAC markedly reversed this effect (Figure 3F). These results show that CA6-induced apoptosis in gastric cancer cells through ROS elevation.

Previous studies have shown that oxidative stress can cause cellular apoptosis by multiple mechanisms including both extrinsic and intrinsic pathways. We have reported that ROS-mediated endoplasmic reticulum (ER) stress induces apoptosis in gastric cancer cells.<sup>21</sup> Therefore, we evaluated the induction of ER stress in response to CA6 challenge. Our results show that CA6 increased ER stress-specific proteins phosphorylated-eukaryotic initiation factor-2 $\alpha$  (p-eIF2 $\alpha$ ), activating transcription factor-4 (ATF4), and CCAAT-enhancer-binding protein homologous protein (CHOP) in both BGC-823 (Figure 3B, D) and SGC-7901 cells (Figure 3C, E). These inductions were seen in cells at 6 h post CA6 exposure. Since CA6 increased ROS production in gastric cancer cells rapidly in 1 h (Figure 1D and E), we evaluated whether ROS was necessary for ER stress induction. We observed decreased levels of ER stress-related proteins in cells pretreated with NAC before exposure to CA6 for 6 h (Figure 3D and E). These results suggest that CA6-induced ER stress activation was dependent on increased intracellular ROS generation. Finally, to confirm if ER stress plays an essential role in response to CA6-induced cell death, ATF4 was silenced in BGC-823 cells (Figure 3G). ATF4-silencing led to a significantly reduced number of apoptotic cells upon CA6 exposure (Figure 3H and I). These results suggest that CA6-mediated cytotoxicity in gastric cancer cells through ROS-mediated ER stress activation.

## CA6-Induced ROS Elevation Promotes Apoptosis Through ROS-Dependent Modulation of Akt/FoxO3a/PTEN Signaling Axis

Multiple signaling pathways that regulate cell survival can be altered by increased ROS levels in cells. These signaling pathways include mitogen-activated protein kinases<sup>22</sup> and phosphoinositide 3-kinase (PI3K)/Akt pathway.<sup>23</sup> Akt regulates cell survival by inactivating downstream proteins such as FoxO3a transcription factors.<sup>24</sup> Akt phosphorylation, a proxy for activation, induces nuclear export of FoxO proteins through the nuclear pore complex.<sup>25</sup> As FoxO proteins exhibit tumor suppression functions and regulate the expression of genes involved in apoptosis and cell cycle arrest,<sup>24</sup> we reasoned that the inhibitory activity of CA6 on gastric cancer growth may involve alteration of the Akt-FoxO3a pathway. Our results show that CA6 decreased

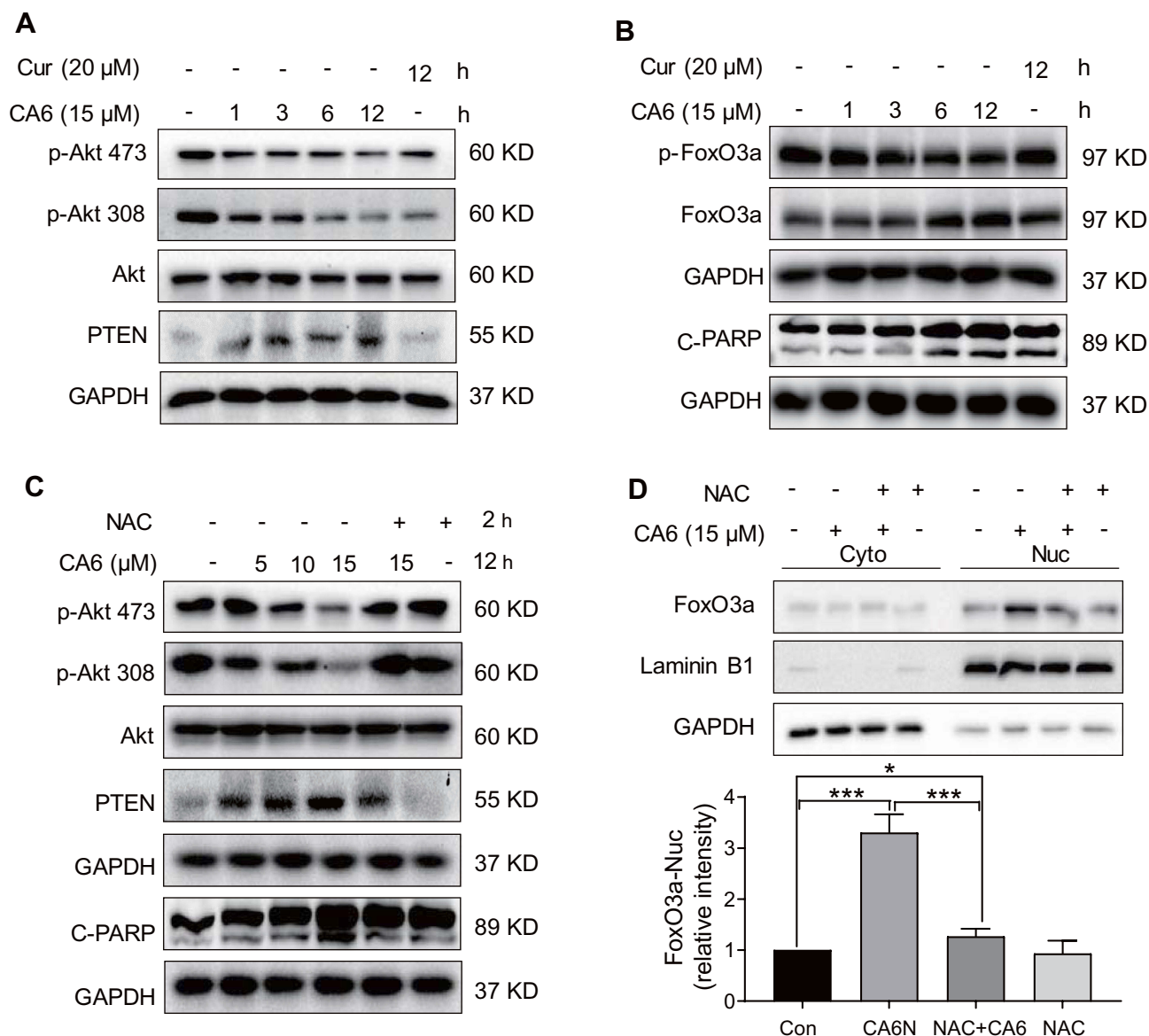
the levels of p-Akt (Ser473 and Thr308) and p-FoxO3a (Figure 4A and B). We also found that the levels of FoxO3a proteins were increased by CA6 (Figure 4B). Increased levels of phosphatase and tensin homolog deleted on chromosome 10 (PTEN), the main negative regulator of the PI3K-Akt pathway, were also noted in cells exposed to CA6 (Figure 4A). Also, these alterations in Akt-FoxO3a were associated with increased levels of apoptosis marker, cleaved-poly (ADP ribose) polymerase (Figure 4B).

Next, to explore the possible link between CA6-induced ROS and Akt-FoxO3a, we eliminated ROS in CA6-challenged cells using NAC. As shown in Figure 4C, depletion of ROS reduced CA6-induced apoptosis, as evident by reduced cleaved-PARP levels. NAC also increased p-Akt levels in gastric cancer cells and reduced PTEN (Figure 4C). Furthermore, neutralizing ROS prevented CA6-induced nuclear accumulation and dephosphorylation of FoxO3a (Figure 4D, [Supplementary Figure S2](#)). Taken together, these results confirmed that CA6 amplified ROS generation, leading to growth inhibition and induction of apoptosis, possibly through activating FoxO3a.

## CA6 Targets TrxR1 and Suppresses Its Enzymatic Activity

Studies have shown that both curcumin<sup>26,27</sup> and some curcuminoids<sup>28,29</sup> inhibit TrxR1. To determine whether CA6 targets TrxR1, we first performed a molecular simulation using a docking software. Our docking results showed that CA6 interacted with TrxR1 protein by covalent bonds on residues Cys-494 and Tyr-405 of the C-terminal redox center (Figure 5A). We validated the potential interaction of CA6 with TrxR1 by surface plasmon resonance (SPR). Our SPR data show that CA6 directly bound to recombinant human (rh) TrxR1 protein with a high affinity (KD value = 12.2  $\mu$ M) (Figure 5B). We then used bis-ANS probe to assess CA6 TrxR1 binding, which interacts with proteins through its hydrophobic phenyl and naphthyl rings. We mixed rhTrxR1 with bis-ANS to obtain the stable fluorescence levels. CA6 was then added to observe the displacement of TrxR1. Fluorescence units were increased when bis-ANS was mixed with rhTrxR1 protein, whereas CA6 decreased the signal indicating competitive binding of CA6 to the hydrophobic pocket of rhTrxR1 protein (Figure 5C).

Above findings suggest that TrxR1 may be the direct target of CA6. Therefore, we evaluated whether CA6 could effectively inhibit TrxR1 enzymatic activity. We evaluated the direct inhibitory effects of CA6 on rhTrxR1 protein activity



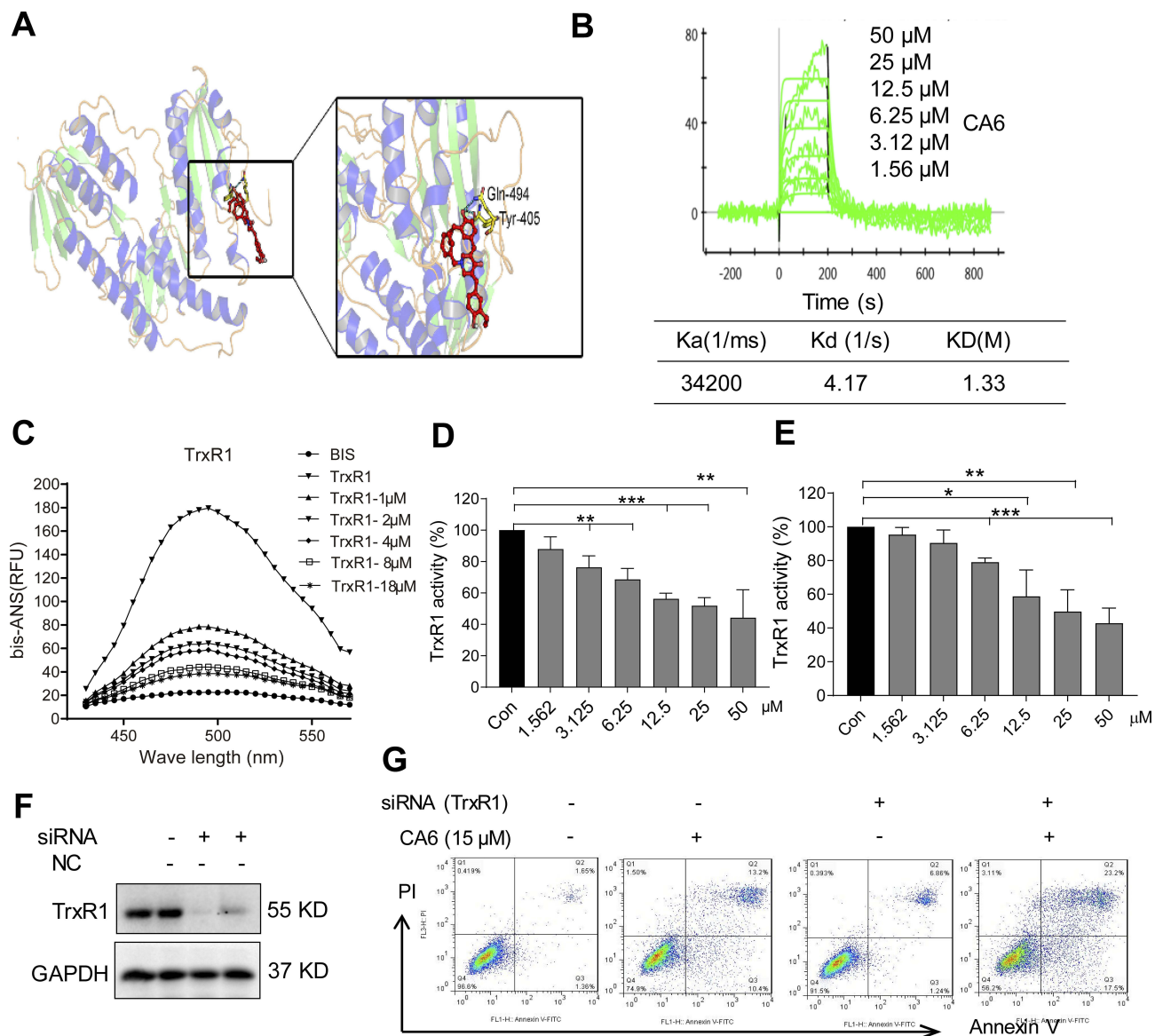
**Figure 4** CA6 induces apoptosis in gastric cancer cells through ROS-dependent modulation of AKT and FOXO3a. **(A)** BGC-823 cells were treated with CA6 (15  $\mu$ M) for indicated time points. Curcumin was used at the dose of 20  $\mu$ M as positive control. Levels of phosphorylated Akt (Thr308 and Ser473) and PTEN were determined. GAPDH was used as loading control. Representative immunoblots shown [n = 3]. **(B)** Cells treated as indicated in panel A were probed for levels of phosphorylated-FoxO3a and cleaved-PARP. **(C)** BGC-823 cells were exposed to CA6 (15  $\mu$ M) for 12 h, with or without pretreatment with NAC (5 mM) for 2 h. Levels of p-Akt473, p-Akt308, PTEN, and cleaved-PARP were determined. Representative immunoblots shown [n = 3]. **(D)** Cytosolic and nuclear fractions from BGC-823 cells treated as indicated in panel C. Protein levels of FoxO3a were examined. GAPDH and Laminin B1 were used as loading control. Representative blots shown in the upper panel and densitometric quantification are shown in the lower panel [Mean  $\pm$  SEM; n = 4; \*P < 0.05; \*\*\*P < 0.001].

using the 5,5'-dithiobis(2-nitrobenzoate) (DTNB) assay. As expected, CA6 dose-dependently inhibited the activity of rhTrxR1 (Figure 5D). Furthermore, addition of CA6 to lysates prepared from BGC-823 cells also showed reduced TrxR1 activity (Figure 5E). To examine whether inhibition of TrxR1 was involved in CA6-induced gastric cancer apoptosis, we silenced TrxR1 in cells (Figure 5F) and exposed the cells to CA6 (Figure 5G). As shown in Figure 5G, TrxR1 silencing significantly enhanced CA6-induced BGC-823 cell apoptosis

when compared to CA6 treatment alone. Collectively, our findings show that CA6 targeted TrxR1 and induced apoptotic cell death by reducing TrxR1 activity.

### CA6 Inhibits BGC-823 Xenograft Tumor Growth in vivo by Reducing TrxR1 Activity and Enhancing ROS

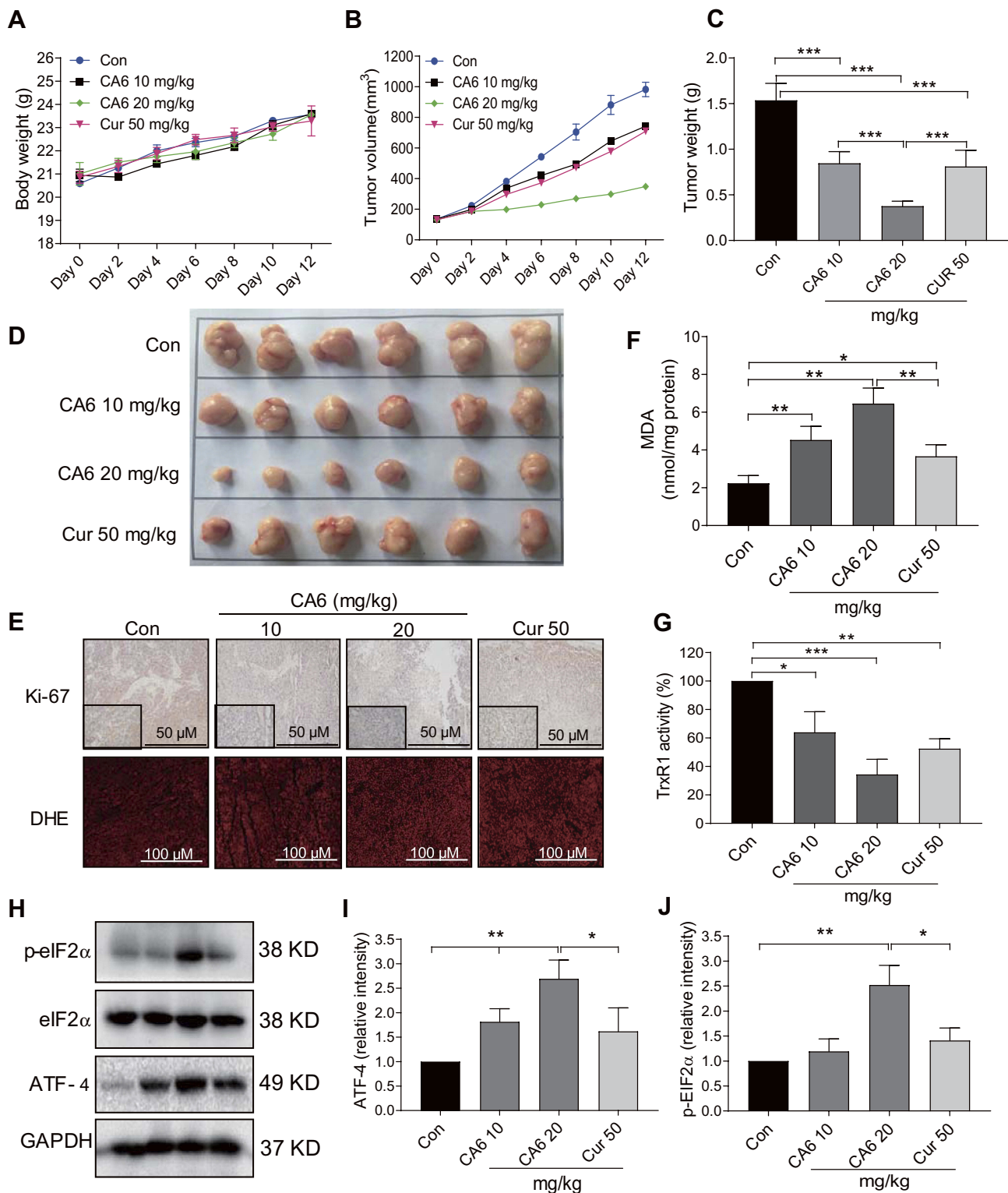
Our in vitro studies suggest that CA6 inhibited TrxR1 activity and induced apoptosis through the elaboration of



**Figure 5** CA6 binds to TrxR1 and suppresses its enzymatic activity. **(A)** Molecular docking between CA6 and TrxR1 was simulated with Tripos molecular modeling package Sybyl-x.v1.1.083. **(B)** Recombinant human (rh) TrxR1 was added to different concentration of CA6 and binding affinities were measured using SPR. **(C)** Binding between CA6 and rhTrxR1 was analyzed through bis-ANS displacement assay. **(D)** The inhibitory effect of CA6 on rhTrxR1 enzymatic activity was determined using DTNB assay [Mean  $\pm$  SEM; n=3; \*\*P<0.01, \*\*\*P<0.001]. **(E)** TrxR1 enzymatic activity in BGC-823 cell lysates was determined using end-point insulin reduction assay [Mean  $\pm$  SEM; n=3; \*P<0.05, \*\*P<0.01, \*\*\*P<0.001]. **(F)** BGC-823 cells were transfected with siRNA against TrxR1 and levels of TrxR1 were determined via Western blotting. GAPDH was used as loading control. Representative immunoblot shown [n = 3]. **(G)** BGC-823 cells transfected with TrxR1 siRNA were exposed to CA6 (15  $\mu$ M) for 24 h. Apoptotic cells were determined using Annexin V/PI staining. Representative dot plots from 3 independent experiments were shown.

ROS. To confirm whether these mechanisms in vivo, we evaluated the anti-tumor effect of CA6 in BGC-823 xenograft tumor mouse model. We used CA6 at two different doses (10 and 20 mg/kg) and curcumin at the dose of 50 mg/kg as control. As shown in Figure 6A, CA6 or curcumin treatment did not affect the mice body weight. No histological alteration was noted in heart, liver, and kidney tissues of mice in any experimental group (Supplementary Figure S3). However, CA6- and curcumin

treatments resulted in a significant reduction in both tumor volumes and tumor weights (Figure 6B–D). CA6 at the dose of 20 mg/kg showed a greater inhibitory effect on tumor growth compared to that of CA6 at 10 mg/kg or curcumin at 50 mg/kg (Figure 6B–D). Reduced tumor growth upon CA6 treatment was associated with reduced ki-67 immunoreactivity in harvested tumor specimens (Figure 6E, upper row). We further show that CA6 increased ROS levels in tumor specimens as detected by



**Figure 6** CA6 inhibits BGC-823 xenograft tumor growth by elevating ROS and reducing TrxR1 activity. BGC-823 cells were implanted in nude mice. Mice were then treated with CA6 (10 mg/kg or 20 mg/kg) for 12 days. Curcumin (50 mg/kg) was used as positive control. **(A)** Body weights of mice implanted with BGC-823 cells at indicated intervals [n=6]. **(B)** Tumor volumes determined at indicated time points following tumor cell injection [n=6]. **(C)** Measurement of tumor, harvested tumor weights on day 12 [Mean  $\pm$  SEM; n=6; \*\*\*P<0.001]. **(D)** Images of harvested tumors from mice on day 12. **(E)** Staining of gastric tumor specimens for cell proliferation marker Ki-67 (top row). Ki-67 was detected using DAB chromogen staining (brown) [n = 6; scale bar = 50  $\mu$ m]. Lower panel showing DHE staining (red) of tumor specimens [n = 6; scale bar = 100  $\mu$ m]. **(F)** Measurement of oxidative stress marker MDA in tumor tissue lysates [Mean  $\pm$  SEM; n=6; \*P<0.05, \*\*P<0.01]. **(G)** TrxR1 enzymatic activity of tumor tissue lysates as measured via endpoint insulin reduction assay [Mean  $\pm$  SEM; n=6; \*P<0.05, \*\*P<0.01, \*\*\*P<0.001]. **(H)** Levels of ER stress-related proteins p-eIF2 $\alpha$  and ATF4 in tumor lysates. GAPDH was used as loading control. Representative immunoblots were shown. **(I, J)** Densitometric quantification of ATF4 and p-eIF2 $\alpha$ . ATF4 was normalized to GAPDH and p-eIF2 $\alpha$  to eIF2 $\alpha$  [Mean  $\pm$  SEM; n=6; \*P<0.05, \*\*P<0.01].

dihydroethidium (DHE) staining (Figure 6E, bottom row). Increased oxidative stress induced by CA6 is also evident through the measurement of lipid peroxidation product malondialdehyde (MDA) (Figure 6F). Also, our results show that CA6 significantly decreased TrxR1 enzyme activity in gastric cancer tissues (Figure 6G).

Lastly, we confirmed the involvement of ER stress pathway activation and altered Akt-FoxO3a signaling in tumor specimens. As shown in Figure 6H and I, ATF4 levels were induced by CA6 at both 10 and 20 mg/kg doses. However, p-eIF2 $\alpha$  only showed an increase with CA6 at 20 mg/kg dose (Figure 6H and J). These data show that CA6-induced ER stress in gastric cancer tissues. Immunohistochemical staining of tumor tissues from CA6-treated mice also shows decreased levels of p-Akt (Ser473) and p-FoxO3a, and increased levels of FOXO3a and PTEN (Figure 7A). Finally, we found that CA6 treatment increased the activity of caspase-9 in tumors (Figure 7B). Taken together, our *in vivo* studies mirror our *in vitro* findings and show that CA6 exhibited inhibitory activities in gastric cancer cells through inhibiting TrxR1 and inducing ROS-mediated apoptotic cell death (Figure 7C).

## Discussion

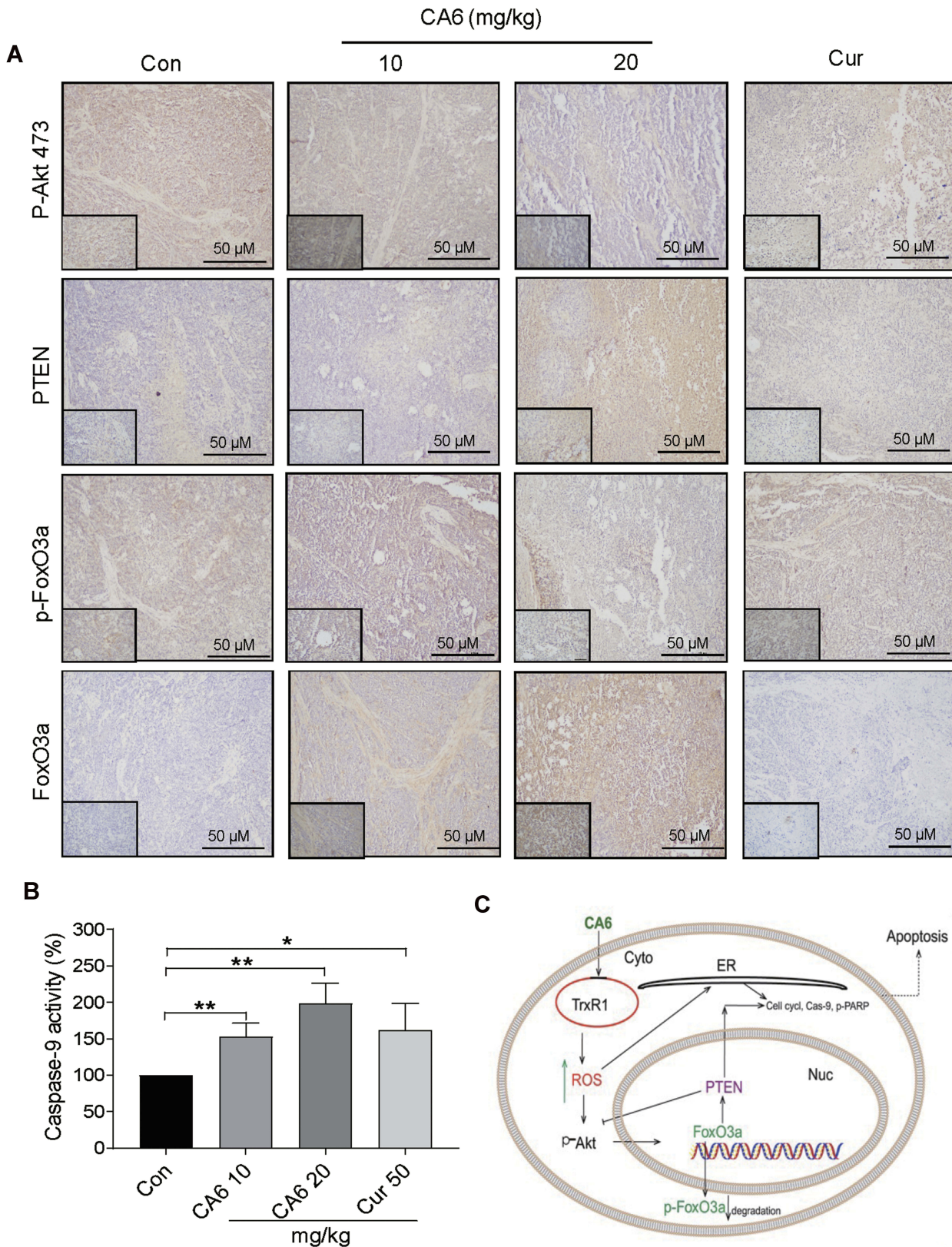
In the present study, we found that the allylated MAC, CA6, inhibited the growth of gastric cancer cells *in vitro* and *in vivo*. We show that CA6 directly bound to TrxR1 and inhibited its activity, leading to increased accumulation of ROS in gastric cancer cells. Furthermore, CA6-induced ROS-dependent apoptosis by inhibiting Akt survival pathway and activating FoxO3a. Collectively, our studies show that TrxR1 could be the potential therapeutic target for the treatment of gastric cancer, and we have provided preclinical evidence for the potential use of CA6 in clinic.

Enhancing ROS generation has been identified as a potential reason to induce apoptosis and curb cancer growth.<sup>30,31</sup> Even though cancer cells basally generate high levels of ROS,<sup>6</sup> which may facilitate proliferation and genomic instability. There appears to be a balance in the chaotic redox system of cancer cells. Agents that further enhance ROS or disrupt the redox state of cancer cells show selective cancer cell cytotoxicity.<sup>32</sup> Curcumin and its derivatives have been reported to increase ROS levels in various cancer cell types.<sup>29,33,34</sup> In the present study, we found that curcumin analog CA6 also increased ROS generation rapidly in both cultured gastric cancer cells and gastric tumors in mice. These findings suggest that CA6-induced cytotoxicity seen in gastric cancer cells may partly be

mediated through elevated ROS levels. ROS-mediated oxidative stress has been shown to be critical in inducing cell death and cell cycle phase arrest.<sup>29</sup> Consistent with this notion, CA6 treatment caused G2/M phase arrest in gastric cancer cells and significantly hampered the colony-forming ability of cells. These effects were intricately linked to ROS levels as NAC was able to neutralize CA6-induced G2/M phase arrest, restore colony growth potential, and prevented gastric cancer apoptosis. Our studies support the use of ROS enhancers as potential treatment options for gastric cancer cells.

There is a bidirectional relationship between ROS and ER stress.<sup>35</sup> Elevated ROS inducing ER-stress-mediated apoptosis has also been demonstrated in different cancers.<sup>35,36</sup> ER stress manifests as increased phosphorylation of eIF2 $\alpha$ , which is required for transient downregulation of protein synthesis and selective induction of ATF4.<sup>37</sup> ATF4 also induces the expression of CHOP, which is involved in the initiation of cell death.<sup>38,39</sup> This link is abundantly displayed in gastric cancer cells exposed to CA6. CA6 increased the levels of p-eIF2 $\alpha$  and induced ATF-4 and CHOP. These changes were found to be quite rapid. It was not surprising that eliminating ROS by NAC or knocking down the expression of ATF4 reduced ER-stress-mediated apoptosis in gastric cancer cells. Findings presented here describe a stress-mediated mechanism in which ER-stress is secondary to CA6-induced ROS.

ROS detoxifying enzymes that maintain intracellular ROS homeostasis are regulated by transcription factors, including the FoxO proteins.<sup>40</sup> One member of the FoxO family, FoxO3a, plays a vital role in oxidative stress resistance in cells.<sup>41</sup> FoxO3a is a downstream substrate of Akt and other growth factors. Activated Akt phosphorylates FoxO3a at multiple sites, leading to nucleus exclusion of FoxO3a and retention in the cytoplasm.<sup>42</sup> Aberrantly activated Akt and suppressed FoxO3a have been reported in a number of cancers.<sup>43</sup> There is also increasing evidence that many cytotoxic drugs induce apoptosis in cancer cells by altering the Akt-FoxO3a pathway. For example, docetaxel induces cancer cell apoptosis by enhancing FoxO3a activity and stimulating its nuclear translocation.<sup>44</sup> A recent study also shows that increased expression of ER-stress protein CHOP may trigger FoxO3a-DNA binding, suggesting a potential link between FoxO3a accumulation and ER stress.<sup>45</sup> In our present study, we found that CA6 inhibited Akt in gastric cancer cells. This Akt inhibition likely relieved the negative regulation of Akt on FoxO3a. We observed decreased FoxO3a phosphorylation



**Figure 7** CA6 inhibits tumor growth by modulating AKT and FoxO3a in vivo. **(A)** Immunohistochemical staining of harvested tumor specimens for p-Akt (Ser473), PTEN, p-FoxO3a and FoxO3a. Antigens were detected by DAB chromagen (brown). Tissues were counterstained with hematoxylin (blue) [scale bar = 50  $\mu$ m]. **(B)** Caspase-9 activity in tumor tissue lysates [Mean  $\pm$  SEM; n=6; \*P<0.05, \*\*P<0.01]. **(C)** Schematic illustration of the inhibitory activity of CA6 on gastric cancer growth.

and increased nuclear FoxO3a levels in cells challenged with CA6. One unanswered question and an important direction for future studies are identifying FoxO3a target genes in gastric cancer cells. Although we anticipate that these targets include growth factor signaling, apoptosis, p53 signaling, and oxidative stress and metabolism,<sup>46</sup> identification of gastric cancer-specific targets is likely to generate a list of potential treatment targets.

One of the key findings of our study is that CA6 targeted TrxR1. The Trx system plays a crucial role in coordinating innate antioxidant defense.<sup>47</sup> Trx/TrxR system is upregulated in many human cancers<sup>10</sup> and believed to facilitate tumor development and metastasis owing to their growth-promoting, apoptosis-resisting, and angiogenesis-supporting functions.<sup>48</sup> Moreover, Trx/TrxR system confers an aggressive tumor phenotype, poorer prognosis, decreased patient survival and resistance to programmed cell death.<sup>10</sup> Many clinically used drugs have been reported to inhibit TrxR including mitomycinC<sup>49</sup> and alkylating and platinum compounds.<sup>50</sup> Not surprisingly, many compounds are being developed to target TrxR.<sup>51</sup> Herein, using various approaches, we found that CA6 directly bound and targeted TrxR1. The specific interaction of CA6 with TrxR1 was demonstrated by SPR and Bis-ANS displacement assays. More importantly, CA6 inhibited TrxR1 enzymatic activity in BGC-823 cells and BGC-823-derived tumor tissues. Depleting TrxR1 in BGC-823 cells also potentiated the effect of CA6 and increased apoptotic cell death. Although the data strongly support CA6 as an inhibitor of TrxR1, much work still needs to be done to clarify the interaction between CA6, TrxR1 and ROS-dependent apoptosis in gastric cancer cells. For example, a kinetic enzymatic study of the type of inhibition by CA6 on TrxR1 is lacking in the present study. Also, further experiments are needed to evaluate the contribution of other possible TrxR1-independent ROS sources in the metabolism of gastric cancer cells after CA6 treatment.

In summary, the current study demonstrates that CA6 exhibited significant cytotoxicity in gastric cancer cells, including inducing G2/M cell cycle arrest and apoptosis. We found that these activities were mediated through increased ROS levels in gastric cancer cells, both in culture and in the mouse xenograft tumor model. Elevated ROS led to ER stress-dependent cancer cell apoptosis. More importantly, we discovered that CA6 increased ROS levels in gastric cancer cells through directly binding to and inhibiting the activity of TrxR1. Overall, these studies implicate TrxR1 in the pathogenesis of gastric

cancer and provide support for further development and testing of CA6 in combating gastric cancer.

## Abbreviations

Akt, Protein kinase B; ATF-4, Activating transcription factor-4; bis-ANS, 1,1'-bis(anilino)-4,4'-bis(naphthalene)-8,8'-disulfonate; CDC2, Cell division control protein kinase-2; CHOP, CCAAT/enhancer-binding protein homologous protein; DAPI, 4',6-diamidino-2-phenylindole; DCFH-DA, 2,7-Dichlorodihydrofluorescein diacetate; DHE, Dihydroethidium; DMSO, Dimethylsulfoxide; DTNB, 2,4-dinitrochlorobenzene; eIF2, Eukaryotic initiation factor-2; ER, Endoplasmic reticulum; FITC, Fluorescein isothiocyanate; FoxO3a, Forkhead box O3; GAPDH, Glyceraldehyde 3-phosphate dehydrogenase; H&E, Hematoxylin & eosin; HRP, Horseradish peroxidase; IC50, Half maximal inhibitory concentration; MAC, Allylated mono-carbonyl analog of curcumin; MDA, Malondialdehyde; MDM2, Mouse double minute 2 homolog; MTT, Methyl thiazolyl tetrazolium; NAC, N-acetyl cysteine; NADPH, Nicotinamide adenine dinucleotide phosphate.; OCT, Optimal cutting temperature; OD, Optical density; PARP, Poly (ADP ribose) polymerase; PI, Propidium iodide; PI3K, Phosphoinositide 3-kinase; PTEN, Phosphatase and tensin homolog; RFU, Relative fluorescence units; ROS, Reactive oxygen species; RPMI, Roswell Park Memorial Institute media; siRNA, Small interfering RNA; Trx, Thioredoxin; TrxR1, Thioredoxin reductase-1.

## Supporting Information

[Supplementary methods](#) information, including 1 table and 3 figures, are hyperlinked to this article at each citation of the materials in the main text.

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## Author Contributions

All authors contributed to data analysis, drafting or revising the article, gave final approval of the version to be published, and agree to be accountable for all aspects of the work.

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## Disclosure

The authors report no conflicts of interest in this work.

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