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An NQO1-selective Activated Prodrug of Triptolide: Synthesis and Anti-hepatocellular Carcinoma Activity Evaluation

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KEYWORDS: CX-23, triptolide (TP), anti-hepatocellular carcinoma, NAD(P)H:quinone oxidoreductase 1 (NQO1)

ABSTRACT: Hepatocellular carcinoma (HCC) is the leading cause of death in patients with cirrhosis. Due to its poor response to conventional chemotherapy drugs, the prognosis for its survival is the worst. NAD(P)H:quinone oxidoreductase 1 (NQO1) is an attractive anticancer target due to its overexpression in HCC. Although triptolide (TP) possesses potent anti-tumor activity, its clinical practice is greatly limited due to its general toxicities and narrow therapeutic window. Herein, we develop an NQO1-selective activated TP analog, named CX-23, which exhibited antiproliferation of HepG2 over normal hepatocytes in vitro. In vivo study shows that CX-23 can not only prevent the hepatocellular carcinoma progression but also migrate the liver and kidney toxicity. These findings indicate that NQO1 may serve as a targeted delivery system to release an antitumor reagent and CX-23 may be a promising lead for developing targeted anti-hepatocellular carcinoma drugs.

Primary liver cancer is one of the most common malignant tumors of the digestive tract in the clinic, 90% of which are hepatocellular carcinoma (HCC). Epidemiological surveys show that approximately 500,000 patients worldwide die of HCC every year, ranking fourth in the world for cancer mortality (1). For patients with early liver cancer, liver transplantation and resection are the preferred treatments for liver cancer, but most patients have reached late stage or metastasis at the time of diagnosis and cannot meet surgical criteria. For patients who cannot undergo surgery, chemotherapy is one of the important treatments for liver cancer(2, 3). Chemotherapy can not only control the growth and spread of tumors, but also transform inoperable tumors into operable tumors and increase the cancer cure rate. Conventional drugs used for anti-hepatocellular carcinoma are cytotoxic drugs, among which the more commonly used cytotoxic drugs are doxorubicin, cisplatin, docetaxel, and fluorouracil (4). These drugs are more toxic and less effective. Sorafenib is the first-line anti-hepatocellular carcinoma drug approved by Food and Drug Administration, however, some patients still have treatment failure or tolerability in clinical use. Compared with the control group, the sorafenib treatment group only prolonged the life of the patient for two months(5). Therefore, there is urgent need for anti-hepatocellular carcinoma reagents discovery and development.

Triptolide (TP) is a di-terpene tri-epoxide compound purified from *Tripterygium Wilfordii* Hook F (TWHF) (6) and possesses the ability to treat diseases such as tumor(7), immune system disorders(8), diseases of central nervous system (9) and cardiovascular (10). Among the pharmacological effects, its antitumor activity attracted the most attention of researchers. In vitro studies have shown the high potential antiproliferation of TP against all 60 NCI human tumor cells with the concentration

at the nanomolar (nM) levels. Moreover, TP can circumvent tumor drug-resistant tumor cells and sensitize cancer cells to other chemotherapeutic agents(11-13). The xeroderma pigmentosum type B (XPB/ERCC3) subunit (ATPase activity) of Transcription factor II Human (TFIIH) was generally accepted the principal TP target. TP can inhibit the ATPase activity of XPB by covalently binding with the thiol of Cys342 of XPB, which initiate transcription by RNA polymerase II (RNAPII) and nucleotide excision repair, offers a consistent mechanism that can be used to interpret most of the known pharmacologic activities of TP(14). Although triptolide has great potential as an antitumor agent, its application in the clinic is restricted due to its liver and kidney toxicities (15-17). The most common side effects of TP are observed in the liver and kidney, where the toxicities are mainly induced by the accumulation of parent drug(18).

As a great promising anticancer drug lead, several analogues of TP have been developed (Figure S1). Noteworthy all the analogues of TP in clinical development were adopted prodrug strategies and keep the intact core structure of TP. PG490-88 (omtriptolide), an esterification of 14-OH group of TP by a succinate acid, failed in phase I clinical trial for its slow and incomplete biotransformation in vivo(19). It indicated the modification of the 14-OH TP impede the enzymatic cleavage due to the hindrance of stereoselectivity. Minnelide is a phosphonoxyethyl derivative of TP and the latest TP analog which is under the phase II clinical trial. Minnelide can be metabolized by alkaline phosphatase and then spontaneously releases TP to carry out the antitumor activity (20). Therefore, simple esterification of 14-OH of TP may be not a good method, but spontaneously transformation is an effective way to achieve the successful goal.

Due to on-target toxicity of TP, it is necessary to develop a

cancer targeted pro-drug approach. NAD(P)H:quinone oxidoreductase 1 (NQO1) is a cytosolic reductase and a homodimeric flavoprotein that utilizes NAD(P)H to catalyze the two-electron reduction of a broad range of substrates most notably quinones(21). NQO1 is an attractive anticancer target for its higher levels expressed in many solid tumors (22). In hepatocellular carcinoma, the positive rate and highly positive rate of NQO1 protein were 86.42% and 58.97% in HCC tissues, and were significantly higher than in either adjacent nontumor tissues or normal liver tissues. Consistently, the five years survival rates of the cancer patients with over-expressed NQO1 are lower compared with those with low NQO1 levels(23, 24). Several classes of bio-reductive compounds that can undergo NQO1 reduction to active species have been developed (25-27). Mitomycin C is present in a clinically used antitumor antibiotic. Other quinones such as β -lapachone, EO9, AZQ and RH1 that show improved properties over mitomycin have not entered clinical research(22). Alternative strategy is that the antitumor reagents was selectively activated by NQO1 in cancer cells.

We have designed and screened TP analogs which contains quinone propionic acid structure (Figure S2). The resulting complexes were scored using the AutoDock Vina and the highest scoring compound (CX-23) was selected for further analysis. It can be clearly seen that quinone moiety of CX-23 binds with the residues of the NQO1 active site and flavin adenine dinucleotide (FAD) which is required for NQO1 catalytic activity (Fig. 1A). The leaving group triptolide is left outside the substrate binding pocket, suggesting that the interaction of quinone with NQO1 active site is not affected by the introduction of triptolide to quinone.

CX-23 contains a propionic acid, as a linkage between quinone and the C14 hydroxy group which is the most reactive and chemical modified functional group in TP. Reduction of the quinone propionic acid esters or amides bearing three methyl groups would lead to the corresponding hydroquinones in which severe steric repulsion between three methyl groups promotes rapid lactonization and release of an alcohol or amine. Several the quinone propionic acid compounds have been investigated for reductase-activated prodrugs such as those based on mustard and oxindoles for anticancer therapy (28). The quinone can be reduced by NQO1 and the propionic acid can facilitate the spontaneously transformation to release TP (Figure 1C), which can covalently bind to the XPB and exert its pharmacological activities.

To synthesis of CX-23, hydroquinone was first converted into CX-21 upon the reaction with methyl β , β -dimethylacrylate in the presence of methanesulfanic acid. CX-22 was synthesized using NBS and esterified with TP involving dicyclohexylcarbodiimide (DCC) and N, N-dimethylaminopyridine (DMAP) to yield CX-23 (Figure 1B). The structure of the CX-21, CX-22 and CX-23 were confirmed by mass spectrometry and NMR spectrometry (Figure S3).

First, we studied if CX-23 was targeted to XPB. The effect of CX-23 on XPB ATPase activity was evaluated (Figure S4A). CX-23 had no inhibitory effects on the XPB ATPase activity, whereas TP, which had been proved to target the XPB presented a higher inhibitory, indicating CX-23 cannot directly bind with XPB. Due to the same target for the toxicity and pharmacological activity of TP, the inability of CX-23 to inhibit

XPB makes it an ideal prodrug which can be converted to TP by cellular NQO1 but devoid of undesirable toxicity prior to entry into cells.

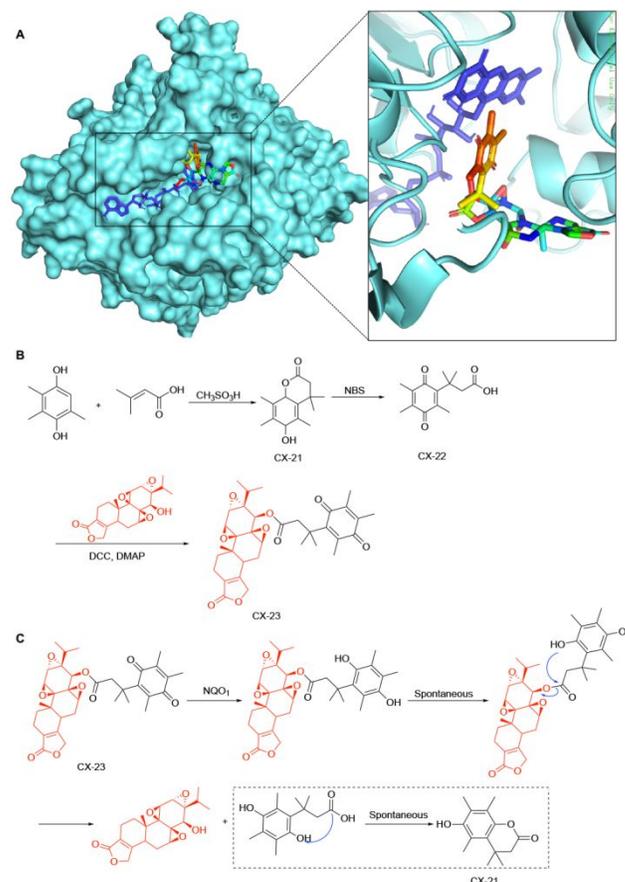


Figure 1. (A) the best pose of CX-23 bound NQO1 ($\Delta E = -12.2$ kcal/mol) using AutoDock Vina; The image was visualized using Pymol. Graphical representation of CX-23-NQO1 complex, TP Complex represented as capped sticks. FAD is expressed in blue. (B) synthesis of CX-23; (C) proposed metabolic and transformed pathways of CX-23 activated by NQO1.

We evaluated the effects of CX-23 and TP on the proliferation of HepG2 and normal hepatocytes. CX-23 presents a comparable antitumor cell activity with IC₅₀ value of 54.3 nM and TP retained the considerable antitumor cell activity with IC₅₀ value of 43.3 nM (Figure S4B). In contrast, in the case of normal hepatocytes with undetectable levels of NQO1, the IC₅₀ value was 540 nM, a value 9.89-fold above its anti-HepG2 activity and 14.3-fold higher than TP with the values of 37.7 nM (Figure S4C), which means a wide therapeutic index for CX-23. The levels of NQO1 activities both in normal hepatocytes and HepG2 cells were studied. The results show that the value of NQO1 activity is 361 ± 84 nmol/min/mg protein in HepG2 cells lysates, but undetected NQO1 activity in normal hepatocyte lysates.

To determine whether CX-23 depended on NQO1-activated for its antiproliferation, we test CX-23 in combination with a NQO1 inhibitor, dicoumarol(29), which alone shows no effect on the HepG2 with the concentration up to 20 μ M. From Figure S4B, we can see that CX-23 show the antiproliferation of

HepG2 when sole use. However, when in combination with dicoumarol, the antitumor efficacy of CX-23 has lost partly, suggesting that dicoumarol can impede the antiproliferation of HepG2 by CX-23 and CX-23 may depend on NQO1 to exert its anti-hepatocellular carcinoma activity.

A complementary assay was used to study the role of NQO1 in the transformation of CX-23 by investigating the metabolism of CX-23 in HepG2 lysates and recombinant NQO1. The metabolism half-life of CX-23 in HepG2 lysates is 7.27 min (Figure S5A) and two major metabolites (TP and CX-21) are identified (Figure 2A). Dicoumarol shows concentration-dependent inhibition of CX-23 metabolism in NQO1 and the transformation of CX-23 into TP was completely inhibited with the dicoumarol concentration of 20 μ M (Figure 2B). The recombinant NQO1 can also transform CX-23 into TP and CX-21 in the presence of NADPH with apparent K_m and V_{max} of 0.309 μ M and 0.223 pmol/mg/min, respectively (Figure 2C). Together, the results suggest that NQO1 can facilitate the transformation of CX-23 into TP. The reduction of CX-23 is NQO1-dependent and the products generated include CX-21 and TP, the latter being capable of inhibit ATPase activity.

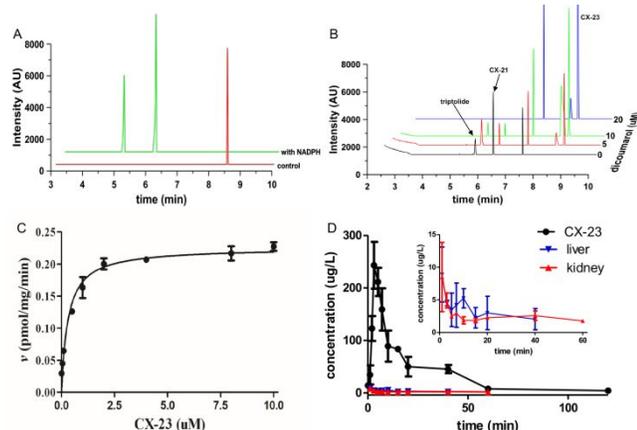


Figure 2. (A) the metabolism of CX-23 in HepG2 lysates with the presence of NADPH; (B) metabolism of CX-23 by NQO1 with the presence of dicoumarol; (C) formation rates of TP in recombinant NQO1 from CX-23; (D) pharmacokinetics of CX-23 in mice after 1.0 mg/kg ip dose.

To confirm that the antitumor activity of CX-23 was due to the activation by NQO1. We used a panel of primary and cancer cell lines which has high expression of NQO1 to obtain a profile of the general cytotoxicity of CX-23. For human keratinocytes and human skin fibroblast, the average IC_{50} value was above 1.0 μ M. In comparison, the IC_{50} value for Hela and A549 were 38.2 nM and 59.4 nM (Figure S4D), respectively, indicating that CX-23 was more selective for the highly-expressed NQO1 tumor cells.

Most drugs do not have specific tissue distribution and are easier to be eliminated in plasma and liver. Only a small number of drugs enter the treatment site, which can cause less effective. In this case, we carry out the metabolic stability of CX-23 both in human plasma and human liver microsomes. After 1 h incubation in vitro, the half-life of CX-23 in human plasma (Figure S5B) and liver microsomes (Figure S5C) are above 1 h. The plasma containing abundant carboxyl ester hydrolase and the microsomes containing cytochrome P450 enzymes

according for xenobiotic oxidation, indicating the carboxyl ester hydrolase-resistance of CX-23 and non-oxidized by CYP450 enzymes.

We have set up the analytical method which showed good linearity, selectivity, and reproducibility and this method (Table S1, S2) can be applied for the pharmacokinetics study of CX-23 (Figure 2D, Table S3). After a single intraperitoneal (ip) dose at 1.0 mg/kg, CX-23 was absorbed quickly with the absorption half-life value of 1.63 min and the time to reach the maximum plasma concentration (t_{max}) is 3 min with a maximum concentration of 243 μ g/L. CX-23 was also distributed quickly with the distribution half-life value of 15.1 min. The ratios of $AUC_{liver} / AUC_{plasma}$ and $AUC_{kidney} / AUC_{plasma}$ are less than 5%, which means less compound exposure in liver and kidney. The plasma concentration of CX-23 was sustained above IC_{50} for 40 min and declined slowly to 1.38 μ g/L at 2h. After 2h, the levels of CX-23 in plasma was under detectable. The terminal half-life was 154 min, indicating the parent drug was eliminated quickly without any cumulation in vivo. The in vivo anti-hepatocellular carcinoma activity and toxicity of CX-23 were evaluated by using HepG2 xenograft nude mice model. In preliminary experiment, we found there is no obvious toxic symptoms of CX-23 after a 1.0 mg/kg ip dose and TP after a 0.2 mg/kg ip dose. Accordingly, one week after HepG2 cell inoculated, mice were dosed once daily by ip over a period of two weeks. As shown in Figure 3, both TP and CX-23 caused an effective reduction in tumor growth. The equimolar dosage of CX-23 shows the similar antitumor effects (Figure.3A, 3B) as TP. During the whole treatment period, there are no significant body weight changes (Figure.3C) in the treated animals, suggesting there is no or minimal toxicity.

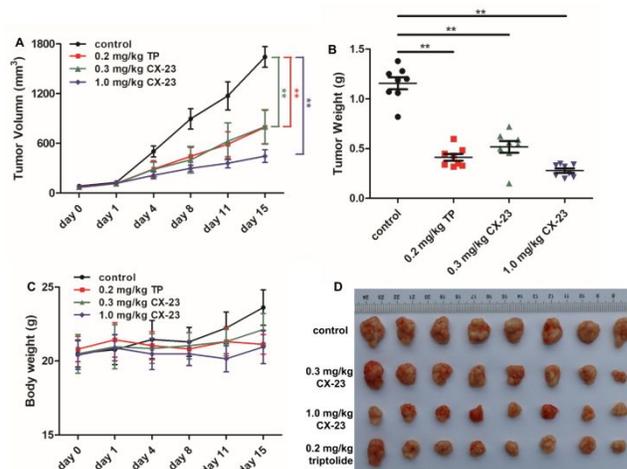


Figure 3. Anti-hepatocellular carcinoma activity of CX-23 compared with TP in nude mice. (A) tumor volume change during the experiment; (B) tumor weight at the end timepoint; (C) mice weight during the experiment; (D) images of tumor tissues at the end timepoint. (** $p < 0.01$ as compared with control; $n = 8$ mice/group). Student's t test was performed for the statistical analysis.

To further determine the toxicity of CX-23, Alanine aminotransferase (ALT), Aspartate aminotransferase (AST) and creatinine, Blood urea nitrogen levels were used to evaluate the liver and renal toxicity. As shown in Figure S6, there is no difference for TP and CX-23 group in the levels of ALT and AST, and no pathological abnormalities found in the liver compared with the control group. However, the levels of

creatinine and BUN in TP group are significantly increased compared with control group (Figure 4), indicating abnormal renal function. The histological examination (Figure 4B-E) showed that t glomerular capillary loop lesions were observed in TP treated group.

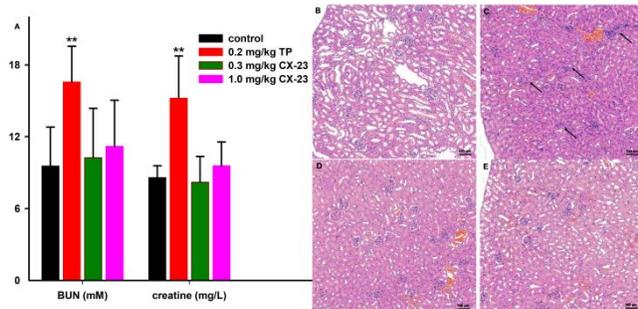


Figure 4. Effect of TP/CX-23 on kidney of nude mice. (A) the toxicity evaluation of TP and CX-23 on serum biochemical parameters. Mice was dosed every day for 15 days. * $p < 0.05$, ** $p < 0.01$ vs. control. (B) control group kidney section; (C) kidney section of TP (0.2 mg/kg)-treated group; (D) kidney section of CX-23 (0.3 mg/kg)-treated group; (E) kidney section of CX-23 (1.0 mg/kg)-treated group. (H&E staining, X200)

In conclusion, CX-23, a quinone propionic acid conjugated TP and prodrug has shown excellent anti-hepatocellular carcinoma activity, impeding the tumor cells growth meanwhile reducing the kidney toxicity. CX-23 represents a potent new targeted to NQO1-activated antitumor drugs.

ASSOCIATED CONTENT

Supporting Information

The supporting Information is available free of charge on the ACS Publications website.

Applied methods, Figures S1-S6 and Table S1-S3.

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‡ Meilin Liu and Wei Song have contributed equally.

Notes

The authors declare no competing financial interest.

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