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# X-chromosome-linked inhibitor of apoptosis as a key factor for chemoresistance in clear cell carcinoma of the ovary

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**Background:** X-chromosome-linked inhibitor of apoptosis (XIAP) is one of the anti-apoptotic proteins leading to chemoresistance in several cancers. The aim of this study is to evaluate the impact of XIAP expression upon ovarian clear cell carcinoma (CCC) that has a platinum-resistant phenotype.

**Methods:** Tissue microarrays made from 90 CCC patients were analysed for immunohistochemical expression levels of XIAP, c-Met, p-Akt and Bcl-XL. In addition, CCC cell lines were evaluated whether XIAP silencing could modulate sensitivity to platinum agent *in vitro*.

**Results:** High XIAP expression was observed in 30 (33%) of 90 CCC cases, and was associated with c-Met (<0.01) and Bcl-XL (<0.01) expression. Cases with high XIAP expression had lower response rate to primary platinum-based chemotherapy (10% vs 65%, P=0.02). In stages II–IV tumours, high XIAP expression was related with worse progression-free survival (PFS, P=0.02). Furthermore, high XIAP expression was identified as an independent worse prognostic factor for PFS and overall survival. Finally, downregulation of XIAP using XIAP-specific small interfering RNA increased sensitivity to cisplatin in human cancer cells derived from CCC.

**Conclusions:** X-chromosome-linked inhibitor of apoptosis expression was correlated with chemoresistance of primary chemotherapy, and identified as a prognostic marker for CCC. X-chromosome-linked inhibitor of apoptosis could be a candidate for new therapeutic target in CCC.

In 1973, clear cell carcinoma (CCC) of the ovary was pathologically defined by World Health Organisation as lesions characterised by clear cells growing in solid/tubular or glandular patterns as well as hobnail cells (Serov *et al*, 1973). Since then, many publications have identified the distinctive behaviour of CCC. First, CCC showed chemoresistance to primary chemotherapy using platinum-based therapy and combination with paclitaxel and platinum (Takano *et al*, 2012). Lower response was also reported in the second-line chemotherapy (Takano *et al*, 2008), and response duration was extremely short (Takano *et al*, 2013). Second, CCC

had significantly worse overall survival (OS) in advanced-staged tumours (Sugiyama *et al*, 2000; Winter *et al*, 2007). New strategy is needed for further treatment of CCC.

X-chromosome-linked inhibitor of apoptosis (XIAP) is overexpressed in several cancers including ovarian tumours (Lacasse *et al*, 1998; Tamm *et al*, 2000; Yang *et al*, 2003), and most reports showed that XIAP is a major contributor to chemoresistance to anticancer drugs. Caspase-3 had an important role for cisplatininduced apoptosis in ovarian cancer cells, and cisplatin induced apoptosis through the decrease of XIAP protein levels and the

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increase of caspases-3 activity in cisplatin-sensitive ovarian serous cancer cells in a dose-dependent manner, but the effect was not evident in cisplatin-resistant cells (Asselin et al, 2001). Overexpression of XIAP by adenoviral sense XIAP complementary DNA attenuated the ability of cisplatin to induce apoptosis (Li et al, 2001). On the other hand, downregulation of XIAP using adenoviral anti-sense XIAP infection or small interfering RNA (siRNA) increased sensitivity to cisplatin in resistant serous ovarian cancer cells (Sasaki et al, 2000; Ma et al, 2009). Thus, XIAP was strongly associated with platinum resistance of ovarian cancer cells. However, these investigations were performed by ovarian serous cancer cells, and the roles of XIAP in CCC have not been analysed. Also, downregulation of c-Met decrease XIAP protein levels in ovarian cancer cells, suggesting that XIAP was regulated by hepatocyte growth factor (HGF) /c-Met pathway through Akt (Bu et al, 2011). Although c-Met has been associated with tumourigenesis in CCC (Yamamoto et al, 2011), the relationship of XIAP expression and c-Met has not been studied.

Herein, we assessed the immunoreactivities of XIAP, c-Met, p-Akt and Bcl-XL, and analysed the correlation of XIAP expression and clinicopathologic characteristic of CCC. Moreover, we investigated whether XIAP downregulation could sensitise ovarian CCC cells to platinum agents.

## MATERIALS AND METHODS

**Patients and tissue microarray.** Tissue blocks from 92 patients with CCC who received primary debulking surgery at National Defense Medical College hospital between 1984 and 2008 were identified. In all, 1.5-mm cores were punched from donor blocks and inserted into a recipient block. All specimens were cut into 4- $\mu$ m-thick sections. Satisfactory immunohistochemical staining was obtained in 90 cases, and these cases were enrolled in this study. No patient had received neoadjuvant chemotherapy before primary surgery. The research was approved by the Ethic Committee of National Defense Medical College, Tokorozawa, Japan.

Immunochemistry and interpretation. We used mouse monoclonal antibody for MET4 (8G6 dilution 1:150; Knudsen et al, 2009), mouse monoclonal antibody for XIAP (48; dilution 1:300; BD Transduction, Franklin Lakes, NJ, USA), rabbit monoclonal antibody for phospho-Akt (p-Akt; Ser473; dilution 1:300; Cell Signaling Technology, Danvers, MA, USA) and rabbit monoclonal antibody for Bcl-XL (54H6; dilution 1:300; Cell Signaling Technology), respectively. Tissue microarray slides were deparaffinised in xylene and hydrated with alcohol and boiled in an autoclave at 121 °C for 15 min in 0.01 mol l<sup>-1</sup> citrate buffer (pH 6.0) and then allowed to cool at room temperature. Endogenous peroxidase was blocked by 0.3% H<sub>2</sub>O<sub>2</sub>/methanol. The slides were incubated at 4 °C overnight with primary antibodies and reacted with EnVision + system HRP-labelled polymer (DAKO Japan Inc., Bunkyo-ku, Tokyo) as secondary antibody for 30 min at room temperature. Specific antigen-antibody reactions were visualised with 0.2% diaminobenzine tetrahydrochloride and hydrogen peroxide, and counterstained with Mayer haematoxylin. For each antibody, negative control studies were performed without the primary antibody. No significant staining was observed in the negative control sections. As positive control, formalin-fixed paraffin-embedded tissue microarrays from 20 ovarian serous adenocarcinomas were stained. Ovarian serous adenocarcinomas were known to be stained previously by all antibodies (Bu et al, 2011; Yamamoto et al, 2011).

Immunoreactivity was scored according to intensity as negative (-), moderate (1 +) or strong (2 +). If cases had 2 + and >50%

of immunoreactive components, they were defined as high expression. Cases without high expression were defined as low expression. Two observers independently evaluated and interpreted the results of immunohistochemical staining without knowledge of the clinical data of each patient. In the interpretation of immunohistochemistry, any discrepancies between the two observers were resolved by discussion, and using a multiviewer microscope.

**Cell lines and culture conditions.** Ovarian cancer cell line derived from clear cell carcinoma, KK (Sasa *et al*, 1993), was used in this study. These cell lines were grown as monolayer cultures in RPMI-1640 (Life Technologies, Grand Island, NY, USA) + GlutmaxTM-I (Invitrogen Japan KK, Tokyo, Japan) medium supplemented with 10% foetal bovine serum (Invitrogen Japan KK), 100 U penicillin per ml, and 100 mg streptomycin per ml (Invitrogen Japan KK) in a humidified atmosphere of 5% CO<sub>2</sub> at 37 °C, and routinely tested for mycoplasma infection.

**Transient transfection.** Nonspecific control siRNA, XIAP-specific siRNA (Signal Science XIAP siRNA, #6446) and XIAP-specific siRNA II (Signal Science XIAP siRNA, #6550) were purchased from Cell Signaling Technology. KK cells cultured in 3.5 cm plates were transfected with 107 nM of XIAP siRNA or XIAP siRNA II and control siRNA using Lipofectamine 2000 (Invitrogen Japan KK) according to the manufacturer's specifications. X-chromosome-linked inhibitor of apoptosis knockdown was confirmed by western blot analysis in all the experiments. Names of the transfected cells using nonspecific control siRNA, XIAP-specific siRNA and XIAP siRNA II were determined as KK-C, KK-I and KK-II, respectively.

Cell proliferation and cytotoxicity assay. Cisplatin was obtained from Bristol Meier's Squib Oncology (Tokyo, Japan). KK-C, KK-I and KK-II were seeded onto 96-well plates at approximately  $1 \times 10^4$  or  $4 \times 10^4$  cells cm<sup>-2</sup> for cytotoxicity assays after 24 h from the treatment at different cisplatin concentrations (0, 10, 20 and  $30 \,\mu$ M). Cell viability was determined by MTT method using Tetra Color One (Seikagaku Corporation, Chiyoda-ku, Japan) according to the manufacturer's instructions.

Preparation of cell lysate for western blot analysis. Protein lysates were extracted in RIPA buffer (Wako Pure Chemical Industries Ltd, Osaka, Japan) according to the manufacturer's instructions. Protein concentrations were determined by Bradford assay (Bio-Rad Laboratories, Hercules, CA, USA). After electrophoresis of 10 µg cytosolic fractions loaded onto Mini-PROTEIN TGXTM gel (Bio-Rad Laboratories), proteins were transferred to PVDF membranes using Trans-Blot Turbo Transfer System Transfer Pack (Bio-Rad Laboratories). Subsequently, the membranes were blocked for 1 h in 4% BSA, and incubated overnight at 4 °C in primary antibodies. The following antibodies and concentrations were used: 1 out of 2000 rabbit anti-XIAP (Cell Signaling Technology), 1 out of 1000 rabbit monoclonal antibody for phospho-Akt (p-Akt; Ser473, Cell Signaling Technology), 1 out of 1000 rabbit monoclonal antibody for Bcl-XL (54H6, Cell Signaling Technology), 1 out of 1000 CONFIRM anti-total c-MET (SP44) rabbit monoclonal antibody for C-Met (Ventana Medical Systems, Tucson, AZ, USA), 1 out of 1000 rabbit monoclonal antibody for PTEN (138G6, Cell Signaling Technology) and 1 out of 5000 rabbit  $\beta$ -actin (Cell Signaling Technology). After washes with TBS-T, membranes were incubated for 1h at room temperature using horseradish peroxidase-conjugated anti-rabbit secondary antibody (Cell Signaling Technology), and were visualised using the ECL detection system (GE Healthcare UK Ltd, Buckinghamshire, England) by a LAS-3000 imaging system (Fujifilm, Minato-ku, Tokyo). Protein expression was determined

densitometrically and normalised against  $\beta$ -actin expression using the software Multi Gauge version 3.1 (Fujifilm).

Statistical analysis. The Stat View software ver.5.0 (SAS Institution Inc., Cary, NC, USA) was used for statistical analysis. Progression-free survival (PFS) was defined as the interval between the primary surgery and death or the date of progression disease. Overall survival was defined as the interval between the primary surgery and death. Staging was performed according to FIGO system. Performance status (PS) was evaluated by WHO criteria. Response rate was evaluated by using Response Evaluation Criteria in Solid Tumours (RECIST) criteria (Therasse et al, 2000). The  $\chi^2$ -test, Fisher's exact test and Mann-Whitney U-test were used to evaluate clinical significance of protein expression in clinicopathological parameters. Progression-free survival and OS curves were generated using the method of Kaplan-Meier. Comparisons of the survival distribution were made with log-rank test. Cox proportional hazards model was used for multivariate analysis of PFS and OS. All experiments were repeated independently at least four times. All values are presented as mean ± s.d. Statistical significance between two groups was determined by use of a two-tailed *t*-test. Statistical significance was defined as a P < 0.05.

## RESULTS

Patients' characteristics and XIAP expression. A total of 90 cases with CCC were enrolled in this study. Median age of the patients was 52 years (range 35-72). FIGO stage distribution was as follows: 46 cases (51%) in stage I, 9 cases (10%) in stage II, 31 cases (34%) in stage III and 4 cases (5%) in stage IV. Ten cases (11%) had <1 cm residual tumours and 17 cases (19%) had >1 cm residual tumours. Eighty-six cases (96%) underwent platinum-based chemotherapy as postoperative primary chemotherapy. Four cases with stage I disease refused to receive chemotherapy, despite physicians' recommendation. Distribution of XIAP immunoreactivity according to clinicopathological characteristics was shown in Table 1. X-chromosome-linked inhibitor of apoptosis expression was not influenced by age at diagnosis, WHO PS, FIGO stage and residual tumour diameter. Immunohistochemically, high expression levels of XIAP, c-Met, phospho-Akt and Bcl-XL were observed in 30 (33%), 31 (34%), 23 (26%) and 33 (37%), respectively. Representative staining of these proteins were shown in Supplementary Figure 1. The correlation of expression levels of c-Met, p-Akt and Bcl-XL in CCC tissues according to XIAP expression was also summarised in Table 1. High expression of XIAP was correlated with high expression of c-Met (P < 0.01) and Bcl-XL (P < 0.01). However, there was no significant association between expression levels of phospho-Akt and XIAP (P = 0.23).

XIAP expression in stage I clear cell adenocarcinoma of the ovary. Among 46 cases with stage I disease, there were no significant differences of age, physical status, FIGO stage and delivery of chemotherapy according to XIAP expression (Supplementary Table S1). In 46 cases with stage I disease, there were no statistical differences of PFS (P=0.22, Supplementary Figure 2A) and OS (P=0.99, Supplementary Figure 2B) according to XIAP expression. In multivariate analysis for PFS and OS using variables of age, peritoneal cytology, completion of surgical staging and XIAP expression, high expression of XIAP was not identified as a prognostic factor for PFS (P=0.29) and OS (P=0.99).

XIAP expression in stages II-IV clear cell adenocarcinoma of the ovary. Expression levels of XIAP, c-Met, phospho-Akt and Bcl-XL according to response of primary chemotherapy were assessable in 27 patients who had measurable disease. All cases received platinum-based chemotherapy after primary surgery. High XIAP expression was significantly correlated with response  
 Table 1. Distribution of XIAP immunoreactivity according to clinicopathological characteristics in 90 patients with clear cell carcinoma of the ovary

Clinicopathological variables	Number (%)	XIAP high	XIAP low	P-value			
Age at diagnosis (years)							
<50 ≥50	35 (39%) 55 (61%)	15 (50%) 15 (50%)	20 (33%) 40 (67%)	0.13			
WHO performance status							
0/1 ≥2	86 (96%) 4 (4%)	28 (93%) 2 (7%)	58 (97%) 2 (3%)	0.47			
FIGO stage							
         V	46 (51%) 9 (10%) 31 (35%) 4 (4%)	12 (40%) 2 (7%) 14 (46%) 2 (7%)	34 (57%) 7 (12%) 17 (28%) 2 (3%)	0.26			
Residual tumour diamete	er						
None ≤1 cm >1 cm	63 (70%) 10 (11%) 17 (19%)	19 (63%) 4 (14%) 7 (23%)	44 (73%) 6 (10%) 10 (17%)	0.62			
c-Met							
High Low	31 (34%) 59 (66%)	16 (53%) 14 (47%)	15 (25%) 45 (75%)	< 0.01			
p-Akt							
High Low	23 (26%) 67 (74%)	10 (33%) 20 (67%)	13 (22%) 47 (78%)	0.23			
Bcl-XL							
High Low	33 (37%) 57 (63%)	17 (57%) 13 (43%)	16 (27%) 44 (73%)	< 0.01			
Abbreviations: FIGO = International	Federation of O	bstetrics and G	iynecology; W	HO = World			

Abbreviations: HGO = International Federation of Obstetrics and Gynecology; WHO = World Health Organisation; XIAP = X-chromosome-linked inhibitor of apoptosis.

to primary chemotherapy in CCC patients (Table 2). Significantly more cases of responders had low expression of XIAP compared with non-responders (90% vs 35%, P < 0.01). Expression levels of c-Met, phospho-Akt, and Bcl-XL were not related with response to chemotherapy.

Among 44 cases with stages II–IV tumours, high XIAP expression was observed 18 cases (Supplementary Table S2). There were no statistical differences of XIAP expression according to age, WHO physical status, FIGO stage and residual tumours after primary surgery. There was a significant difference of PFS according to XIAP expression (P = 0.02, Figure 1A), however, significant difference was not observed in OS (P = 0.07, Figure 1B). In univariate analysis, there were no significant differences of PFS and OS according to expression of c-Met, phospho-Akt and Bcl-XL. Multivariate analysis for PFS and OS in 44 cases was shown in Table 3. In addition to residual tumour diameter, XIAP expression was identified as an independent prognostic factor for PFS (hazard ratio = 2.94, P = 0.02) and OS (hazard ratio = 2.70, P = 0.04).

**Downregulation of XIAP by siRNA and sensitivity to cisplatin.** Further, we investigated whether XIAP downregulation by siRNA could increase sensitivity to cisplatin in KK cells, which were derived from human ovarian clear cell carcinoma. First, XIAP expression ratio compared with no transfection was  $73.1 \pm 12.7\%$ 

Table 2. Expression levels of XIAP, c-Met, phospho-Akt and Bcl-XL according to response of primary chemotherapy in 27 patients with evaluable disease							
Proteins	Expression level	Number of the patients	Responders <sup>a</sup> (n = 10)	Non-responders <sup>b</sup> (n = 17)	P-value		
XIAP	High Low	12 15	1 (10%) 9 (90%)	11 (65%) 6 (35%)	0.02		
c-Met	High Low	11 16	2 (20%) 8 (80%)	9 (53%) 8 (47%)	0.20		
p-Akt	High Low	6 21	0 (0%) 10 (100%)	6 (35%) 11 (65%)	0.10		
Bcl-XL	High Low	10 17	2 (20%) 8 (80%)	8 (47%) 9 (53%)	0.32		

Abbreviations: CR = complete response; PD = progressive disease; PR = partial response; SD = stable disease; XIAP = X-chromosome-linked inhibitor of apoptosis. <sup>a</sup>The patients with CR and PR.

<sup>b</sup>The patients with SD and PD



Figure 1. Progression-free survival (PFS) and overall survival (OS) curves of the patients with stages II–IV clear cell carcinoma of the ovary according to XIAP expression. (A) Progression-free survival curves of the patients. The patients with high expression of XIAP had significantly worse PFS (P = 0.02). (B) Overall survival curves of the patients. Although high expression of XIAP was related with poor survival, significant difference was not observed in two groups (P = 0.07).

Variables		Progression-free survival			Overall survival		
		Hazard ratio (95% CI)		P-value	Hazard ratio (95% CI)		P-value
Age	≤50 vs >50	0.75	(0.32–1.76)	0.51	0.55	(0.20–1.48)	0.23
WHO performance status	0/1 vs ≥2	0.74	(0.14–3.79)	0.71	0.58	(0.11–2.95)	0.51
FIGO stage	II vs III/IV	0.58	(0.19–1.77)	0.37	0.60	(0.16–2.22)	0.44
Residual tumour diameter	>1 cm vs ≤1 cm	3.48	(1.40-8.62)	< 0.01	5.35	(1.98–15.0)	< 0.01
XIAP expression	High vs low	2.94	(1.22–7.09)	0.02	2.70	(1.02–7.16)	0.04

 $Abbreviations: CI = confidence \ interval; \ FIGO = International \ Federation \ of \ Obstetrics \ and \ Gynecology; \ WHO = World \ Health \ Organisation; \ XIAP = X-chromosome-linked \ inhibitor \ of \ apoptosis.$ 

in KK-C,  $20.6 \pm 3.9\%$  in KK-I and  $19.5 \pm 6.7\%$  in KK-II. X-chromosome-linked inhibitor of apoptosis expression was significantly downregulated in both KK-I and KK-II in comparison with KK-C (Figure 2A). Expression levels of c-Met, Bcl-XL and PTEN were similar in those cells, however, p-Akt expression was slightly decreased in KK-I and KK-II cells compared with KK-C:  $62.9 \pm 11.0\%$  in KK-I,  $64.5 \pm 4.6\%$  in KK-II. (Figure 2A). Next, these cells were treated with cisplatin for 24 h at a dose of 10  $\mu$ M. Expression levels of cleaved caspase-3 and cleaved PARP increased in both KK-I and KK-II cells (Figure 2B). Further, these cells were treated with cisplatin for 24 h in different dose (0, 10, 20 and  $30 \,\mu$ M). Apoptotic ratios of KK-I and KK-II were significantly higher compared with that of KK-C at each concentration in a dose-dependent manner (Figure 2C).

## DISCUSSION

Overall positive rate of XIAP in CCC was 33% in the present analysis. Previous report showed that serous adenocarcinoma of the ovary had approximately 85% positive rate of XIAP (Bu *et al*, 2011). Twenty samples used for positive control of XIAP were all judged as high expression of XIAP, implying that positive rate of XIAP in CCC was lower than that in serous ovarian cancers. Most of serous ovarian cancer had overexpression of p53; however, CCC had less involvement of p53 alteration and p53-independent mechanisms for chemoresistance in CCC was suggested (Eltabbakh *et al*, 2006). In this study, XIAP expression was not influenced by age at diagnosis, WHO PS, FIGO stage and residual tumour



Figure 2. Downregulation of XIAP by transfection with XIAP siRNA and sensitivity to cisplatin *in vitro*. (A) Clear cell carcinoma cell line, KK, was transfected with nonspecific siRNA (KK-C), XIAP-specific siRNA I (KK-I) and XIAP-specific siRNA II (KK-II). Expression ratio of XIAP protein compared with KK cells without transfection was as follows:  $73.14 \pm 12.7\%$  in KK-C,  $20.6 \pm 3.94\%$  in KK-I and  $19.54 \pm 6.67\%$  in KK-II, respectively. Phospho-Akt expression was decreased in KK-I and KK-II cells compared with KK-C:  $62.9 \pm 11.0\%$  in KK-I,  $64.5 \pm 4.6\%$  in KK-II. (B) After transfection, these cells were treated with for 24 h at a dose of  $10 \,\mu$ M. Expression levels of cleaved caspase-3 and cleaved PARP increased in both KK-I and KK-II cells compared with KK or KK-C. (C) Apoptosis fractions induced by 24-h treatment of cisplatin in KK-C, KK-I and KK-II were shown. Ratio of apoptotic fraction was higher in KK-I and KK-II compared with that of KK-C at each concentration. The data represented the mean  $\pm$  s.d. of at least four times. \**P*<0.01.

diameter. However, increased XIAP expression was significantly related with resistance to platinum-based chemotherapy and worse prognosis of CCC disease. To our knowledge, there have been no reports evaluating XIAP expression in a series of consecutive cases of ovarian clear cell adenocarcinomas. X-chromosome-linked inhibitor of apoptosis expression has been recognised as a predictor of platinum resistance in several cancers (Schimmer et al, 2006; Yang et al, 2012). In addition, high XIAP expression was related with worse prognosis in human neoplasms (Tamm et al, 2004a, b; Mizutani et al, 2007). Recently, inhibition of XIAP has been recognised as a potential target of cancer therapy. Downregulation of XIAP by transfection with XIAP siRNA resulted in decreased phospho-Akt expression, and increased chemosensitivity to anticancer drugs (Jiang et al, 2012). X-chromosome-linked inhibitor of apoptosis has been shown to act as an E3 ubiquitin ligase for PTEN and to promote Akt activity (Van Themsche et al, 2009). However, PTEN expression level was not affected in the present system, suggesting another signalling pathway to activate Akt signalling, which is not related with PTEN ubiquitination. In addition, XIAP downregulation by a small-molecule inhibitor abrogated XIAP/procaspase-9 interaction, and decreased cell viability that was resistant to anticancer agents (Aird et al, 2010). This study revealed that XIAP overexpression resulted in chemoresistance against platinum-based chemotherapy and worse PFS of CCC patients. In the present system, downregulation of XIAP by transfection with XIAP siRNA increased sensitivity to cisplatin in CC through downregulation of p-Akt. Although further study is needed, XIAP expression is identified as one of the fundamental mechanisms as to how CCC showed intrinsic chemoresistance.

Previous immunochemical analysis for ovarian cancers mainly consisting of serous adenocarcinomas revealed that c-Met expression was correlated with expression levels of phospho-Akt-Ser 473, XIAP and Bcl-XL (Bu *et al*, 2011). Significant correlation of XIAP with c-Met expression observed in this study suggested that XIAP was regulated by HGF/c-Met in CCC. In our study, XIAP expression was not related with phospho-Akt (Ser 473). This may be simply explained by a small sample number; however, there may be another HGF/c-Met/XIAP pathway that was not through activation of phospho-Akt at Ser 473.

In a report including a large case series of advanced-staged CCC, residual tumour was the only independent prognostic factor, and adjuvant chemotherapy was not identified as a prognostic factor (Takano *et al*, 2006). Multivariate analysis in stages II–IV CCC showed that XIAP expression was an independent prognostic factor for PFS and OS in CCC, in addition to residual tumour diameter. It was revealed that high expression of XIAP was significantly correlated with resistance to primary chemotherapy, and led to worse prognosis in CCC. So, inhibition of XIAP expression using a c-Met inhibitor (Appleman, 2011) or a XIAP inhibitor (Kamsteeg *et al*, 2003) could be a novel strategy for the treatment of CCC.

The limitation of this study included a retrospective investigation and a single-institutional analysis. Also, the results obtained by this study could potentially have a bias such as selection bias, and further prospective investigation is needed to confirm the impact of XIAP expression upon prognoses of CCC patients. Nevertheless, XIAP could have an important role in chemoresistant phenotype in CCC, and high expression of XIAP was identified as an independent prognostic factor in stages II–IV CCC tumours. X-chromosome-linked inhibitor of apoptosis should be further evaluated as one of promising targets in chemotherapy for CCC.

#### CONFLICT OF INTEREST

The authors declare no conflict of interest.

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