Phase II and UGT1A1 genotype study of irinotecan dose escalation as salvage therapy for advanced gastric cancer

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BACKGROUND: To assess the efficacy and safety of individualised dose optimisation of irinotecan monotherapy as salvage treatment for advanced gastric cancer (AGC).

METHODS: A total of 43 patients were enrolled. Intravenous irinotecan (350 mg m^{-2}) was administered every 3 weeks. The dose was increased (425 mg m^{-2} and 500 mg m^{-2}) or decreased (250 mg m^{-2}) depending on patient tolerance. UGT1A1 genotypes were determined by direct sequencing of genomic DNA extracted from peripheral blood.

RESULTS: A total of 183 cycles of irinotecan were administered, with a median of four cycles per patient. The overall response rate was 9.3%, and the disease control rate was 62.8%. Median time to disease progression was 2.8 months, and median overall survival was 8.0 months. Grade 3–4 neutropenia was the most common toxicity (53.5%), and febrile neutropenia was the least common toxicity (4.6%). Compared with defective allele groups, UGT1A1 *1/*1 was associated with a lower incidence of grade 3–4 neutropenia during the first cycle ($P = 0.018$).

CONCLUSION: Individualised irinotecan dose escalation based on patient tolerance was not associated with increased toxicity and shows modest activity as salvage chemotherapy for AGC. The role of UGT1A1 genotype in clinical toxicity requires further evaluation.

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Gastric cancer is the fourth most common type of cancer and the second most common cause of cancer deaths world-wide (Parkin *et al*, 2005). Although improvements in early diagnosis have increased the number of curative resections, many patients present with locally advanced unresectable lesions or distant metastases at the initial diagnosis. Even after complete resection, local and distant relapse are common. In these patients, palliative chemotherapy can improve the length and quality of life compared with the best supportive care alone (Murad *et al*, 1993; Pyrhonen *et al*, 1995; Glimelius *et al*, 1997).

Fluoropyrimidine and platinum combination chemotherapy (with or without anthracycline) is a commonly used first-line treatment for advanced gastric cancer (AGC) (Kim *et al*, 1993, 2002; Webb *et al*, 1997; Vanhoefer *et al*, 2000; Koizumi *et al*, 2003; Kang *et al*, 2010). Our group examined the efficacy and safety of capecitabine plus cisplatin (XP) (Kim *et al*, 2002) and XP with docetaxel regimens (Kang *et al*, 2011). Results from the REAL-2 and ML17032 trials indicate that capecitabine could replace infusional 5-fluoracil (5-FU), demonstrating equal efficacy with the advantage of convenience (Cunningham *et al*, 2008; Kang *et al*, 2009).

No standard chemotherapy regimen for AGC currently exists, but more patients are given fluoropyrimidine, a platinum compound,

and a taxane (concomitantly or sequentially). However, about 50% of gastric cancers do not respond to chemotherapy. Most patients who achieve response eventually experience disease progression, with unsatisfactory treatment results. Thus, there is an urgent need for an effective and less toxic regimen for patients with AGC who do not respond to a chemotherapy regimen consisting of fluoropyrimidine, a platinum compound, and a taxane.

Irinotecan is an S-phase-specific, semisynthetic derivative of camptothecin that interferes with DNA replication and cell division by inhibiting topoisomerase I. Its mechanism of action differs from those of fluoropyrimidines, platinum compounds, and taxanes. Pharmacokinetic studies of irinotecan have reported large variations among individuals, as assessed by area under the concentration-time curves of the active metabolite 7-ethyl-10-hydroxycamptothecin (SN-38) and the inactive glucuronide metabolite (SN-38G), which is conjugated to UDP-glucuronosyltransferase (UGT; Gupta *et al*, 1994). UGT1A1 is believed to be the isoform primarily responsible for SN-38G formation, and inter-individual variability in SN-38G formation is due to various UGT1A1 genotypes (Iyer *et al*, 1998). A phase I trial by Abigerges *et al* (1995) reported all objective responses at dose levels above 350 mg m^{-2} irinotecan. One patient responded to a dose of 260 mg m^{-2} , and one patient with progressive gastric cancer treated with 350 mg m^{-2} irinotecan achieved partial response with 600 mg m^{-2} .

We conducted a prospective phase II study of irinotecan monotherapy for the treatment of patients with AGC who

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previously failed chemotherapy consisting of a fluoropyrimidine, a platinum compound, and a taxane. We used a patient-specific dose escalation scheme based on tolerance during the preceding cycle and evaluated the relationship between UGT1A1 genotype with irinotecan toxicity.

PATIENTS AND METHODS

Patients

All enrolled patients received a diagnosis of advanced, unresectable, histologically confirmed adenocarcinoma of the stomach. The inclusion criteria were as follows: (i) 18 to 65 years old; (ii) Eastern Cooperative Oncology Group performance status of 0 or 1; (iii) prior treatment with chemotherapy consisting of a fluoropyrimidine (5-FU, capecitabine, doxifluridine, S1, or tegafur-uracil), a platinum compound (cisplatin or oxaliplatin), and a taxane (paclitaxel or docetaxel); (iv) presence of measurable lesion(s) based on criteria of the Response Evaluation Criteria in Solid Tumors (RECIST, version 1.0); (v) no previous radiotherapy; (vi) estimated life expectancy >3 months; (vii) adequate bone marrow function (leukocyte count $\geq 4000 \mu\text{l}^{-1}$, absolute neutrophil count $\geq 1500 \mu\text{l}^{-1}$, haemoglobin $\geq 9.0 \text{ g dl}^{-1}$, and platelet count $\geq 100\,000 \mu\text{l}^{-1}$); and (viii) adequate renal and hepatic function (serum creatinine $< 1.5 \text{ mg dl}^{-1}$, bilirubin $< 1.2 \text{ mg dl}^{-1}$, and serum transaminases (aspartate transaminase and alanine transaminase) less than three times the upper normal limit and less than five times the upper normal limit for patients with liver metastases). Patients were excluded if they had brain metastases, significant gastrointestinal bleeding or obstruction, or a serious comorbid condition; concomitantly used any drug that was a possible substrate, inducer, or inhibitor of UGT1A1; or lacked the ability to comply with the requirements of the protocol. All patients provided written informed consent, and the Asan Medical Centre institutional review board approved this study (approval No. 2005-0215).

Treatment protocol and dose adjustment

Patients received irinotecan at a starting dose of 350 mg m^{-2} in 250 ml 5% dextrose-water over 90 min every 3 weeks. At the second cycle, the dosage was decreased or increased depending on individual patient tolerance in the preceding cycle. In the absence of moderate toxicity (grade 2 or worse non-haematological toxicity (except alopecia) or grade 3 or worse haematological toxicity (except anaemia)), the dose was increased to 425 mg m^{-2} at cycle 2, and then to 500 mg m^{-2} at cycle 3 and subsequent cycles. If a patient experienced significant toxicities (grade 3 or worse non-haematological toxicities, grade 4 neutropenia lasting ≥ 7 days, grade 3 thrombocytopenia with bleeding, or grade 4 thrombocytopenia), the dose was reduced to that of the previous cycle and re-escalation of irinotecan was not allowed.

For patients who experienced significant toxicity in the first cycle, irinotecan was reduced to 250 mg m^{-2} for cycle 2 and every cycle thereafter. Otherwise, irinotecan was maintained at the same dose as the previous cycle. Patients who experienced significant toxicity despite dose reduction to 250 mg m^{-2} were withdrawn from the study. A chemotherapy cycle was delayed if the granulocyte count was $< 1500 \mu\text{l}^{-1}$ or if the platelet count did not return to a minimum of $100\,000 \mu\text{l}^{-1}$ on the day of infusion. Other non-haematological toxicities (except alopecia) were required to be grade 1 or better prior to cycle initiation. A treatment delay of up to 1 week was permitted without dose modification. If treatment was delayed more than 1 week, the dose for the following cycle was reduced by 1 dose level; if treatment was delayed by more than 3 weeks, the patient was withdrawn from the study. Colony-stimulating factors (e.g., G-CSF, GM-CSF) and antibiotics were not given prophylactically or to treat

uncomplicated neutropenia. Irinotecan treatment was administered for a maximum of nine cycles unless there was disease progression, withdrawal of consent, or unacceptable toxicity.

Efficacy and safety assessment

Tumour response was evaluated every two treatment cycles according to the RECIST criteria (version 1.0). The imaging technique used at baseline was used at subsequent examinations, and the investigators for this trial assessed response to chemotherapy. A patient was considered eligible for tumour assessment if he or she received at least two cycles of chemotherapy or experienced rapid tumour progression (per-protocol population). For intention-to-treat (ITT) analysis, those who failed to return to the clinic for any reason after receiving at least one cycle of chemotherapy were included in the denominator to calculate the response rate (RR). Toxicity was evaluated according to the National Cancer Institute Common Toxicity Criteria for Adverse Events scale (version 3.0).

UGT1A1 genetic assay

Genomic DNA was extracted from blood leukocytes using a DNA Blood Mini Kit (Qiagen, Hilden, Germany). The first exon and flanking region of the human *UGT1A1* gene were amplified by polymerase chain reaction (PCR) using the following primers: forward 5'-CATGATACAAGTGAGCAGGC-3' and reverse 5'-GGGGCTAGTTAATCGATCCA-3'. The 30- μl reaction mixture contained $1 \times$ buffer, 0.2 mmol l^{-1} dNTPs, 50–100 ng genomic DNA, $0.5 \mu\text{mol l}^{-1}$ primers, and 1 unit Taq DNA polymerase (TaKaRa, Shiga, Japan). After a 3-min hot start at 94°C , 30 cycles were performed on a GeneAmp PCR 9700 system (PerkinElmer, Norwalk, CT, USA). The amplified DNA product was purified using QIAquick columns (Qiagen) according to the manufacturer's recommendations. Nucleotide sequences of the PCR products were determined by an automated sequencer (ABI3130XL, Applied Biosystems, Foster City, CA, USA).

Genotyping of -3279T>G (*UGT1A1**60 allelic variant) was determined by pyrosequencing. The 227-bp fragment containing -3279 T>G was amplified using the following primers: forward 5'-TTAACCAAAGAACATTCTAACGG-3' and reverse 5'-biotin-TGCTGTTCCCCAACTTC-3'. The PCR mixture included 0.2 mmol l^{-1} dNTPs, $0.2 \mu\text{mol l}^{-1}$ each primer, 150-ng human genomic DNA, and 2 U Taq polymerase (TaKaRa) in a 20- μl volume. After an initial denaturation at 95°C for 5 min, amplification was carried out with 30 cycles of denaturation at 95°C for 30 s, annealing at 55°C , and extension at 72°C for 30 s. Sepharose beads (40 μl per well; Streptavidin Sepharose High Performance, Amersham Biosciences, Uppsala, Sweden) were diluted with binding buffer, mixed with the PCR product, transferred to a filter plate, and incubated for 10 min at room temperature on a shaking device with continuous mixing. Then, the liquid was removed by vacuum filtration (Multiscreen Resist Vacuum Manifold, Billerica, MA, USA), and the double-stranded DNA was denatured in 0.5 mol l^{-1} NaOH. The immobilised template was treated for 5 s each with 70% ethanol, denaturation buffer, and then washing buffer, and transferred to a PSQ 96 plate (Biotage, Uppsala, Sweden). The washed template was resuspended in annealing buffer (20 mmol l^{-1} Tris-acetate, pH 7.6) that contained the sequencing primers (5'-CCAAGGGTAGAGTTAG-3'). Primer and template mixtures were heated at 90°C for 3 min, and the reactions were allowed to cool to room temperature. The resulting mixture was analysed on a PSQ96MA pyrosequencer (Biotage).

Statistical analysis

Overall RR was the primary end point of this study. Simon's optimal two-stage design was used to test the null hypothesis

($P_0 \leq 0.05$) vs the alternative hypothesis ($P_1 \geq 0.2$). The first stage required that at least 1 patient in 10 had a confirmed response (with $\alpha = 0.05$ and $\beta = 0.2$) before the second stage began. The second stage required the treatment of an additional 19 assessable patients. If four or more patients achieved a confirmed response, then the primary end point would be met.

The TTP was measured from the first day of irinotecan treatment until disease progression was noted, and OS was measured from the first day of irinotecan treatment until death from any cause. Kaplan–Meier estimates were used to analyse all time-event variables, and curves were compared by the log-rank test. Patient characteristics and toxicities were evaluated by descriptive methods.

If the phase II part of the study indicated that our regimen was effective, then we planned to carry out a prospective investigation of the relationship between UGT1A1 diplotype and the severity of neutropenia during the first cycle. We compared the percentage of grade 3–4 neutropenia in patients carrying the *1/*1 diplotype with those of patients carrying a defective allele using the Fisher's exact test. If we assume that 90% of patients with the *1/*1 diplotype will not experience grade 3–4 neutropenia, and 50% of patients with defective alleles (*28, *6, or *60) will not experience a grade 3–4 neutropenia, then a sample size of 40 gives 80% power (at $\alpha = 0.1$) to detect a difference based on the known frequency of the UGT1A1 *1/*1 diplotype in the Korean population (30%) (Ki et al, 2003). SPSS for Windows (SPSS Inc., Chicago, IL, USA) was used for all statistical analyses.

RESULTS

Patient characteristics

From September 2005 to June 2007, 43 eligible patients who were previously exposed to a fluoropyrimidine, a platinum compound, and a taxane were enrolled (Table 1). The median age was 51 years (range: 31–64 years), 15 patients underwent curative gastrectomy, and 11 patients received adjuvant chemotherapy including mitomycin C and oral 5-FU. Most patients (77%) had two or more sites of metastasis. In all, 37 patients (86%) received second-line chemotherapy, and docetaxel monotherapy was the most common second-line regimen (26 out of 37, 70%). A total of 32 patients (74%) had disease progression during previous chemotherapy, and 41 patients (95%) had disease progression within 6 months after withdrawal of the previous chemotherapy regimen. The median TTP (from day 1 of the first cycle of the previous chemotherapy) was 2.8 months (range: 1.2–12.8 months). The median treatment-free interval (TFI) was 31 days (range: 17 days–7.2 months).

Drug delivery

The median number of chemotherapy cycles was four (range: 1–9; total: 183 cycles), with 26.3% of cycles delivered at 250 mg m^{-2} , 55.6% at 350 mg m^{-2} , 12.8% at 425 mg m^{-2} , and 5.0% at 500 mg m^{-2} . Five patients did not proceed to the second cycle because of patient refusal ($n = 2$) or disease progression ($n = 3$). Overall, 10 of the 38 patients (26.3%) who received more than one cycle of chemotherapy received a dose of 425 mg m^{-2} at least once, and 3 of the 38 patients (7.9%) received a dose of 500 mg m^{-2} . The irinotecan dose was reduced to 250 mg m^{-2} for 9 of these 38 patients (23.7%). Treatment was delayed for 11 patients (25.6%) and in 23 cycles (12.6%). Dose reduction occurred for 12 patients (27.9%) and in 48 cycles (26.2%). Reasons for dose reduction were fatigue (83.3%), granulocytopenia (14.6%), and diarrhoea (6.3%). The most frequent reason for dose delay was neutropenia (63.6%). The median dose intensity per patient was 116.7 mg m^{-2} per week (range: 71.2–152.6), and the mean dose of irinotecan at each cycle

Table 1 Baseline clinicopathological characteristics of patients with advanced gastric cancer

	No. of patients	%
Total no.	43	100
Age, years (median, range)	51 (31–64)	
Gender		
Male	35	81.4
Female	8	18.6
ECOG performance status		
0	17	39.5
1	26	60.5
Histology		
Well/moderately differentiated	15	34.9
Poorly differentiated or signet-ring cell type	21	48.9
Unknown	7	16.3
Metastasis sites		
Liver	22	51.2
Peritoneum	20	46.5
Distant abdominal lymph nodes	29	67.4
Lung	4	9.3
Bone	4	9.3
No. of metastatic sites		
1	10	23.3
2	22	51.2
≥ 3	11	25.6
Prior gastrectomy	15	34.9
No. of previous treatment lines		
1	6	13.9
2	37	86.1
First-line chemotherapy regimen		
T	2	4.6
FP	26	60.4
FT	3	7.0
TP	6	14.0
FTP	6	14.0

Abbreviations: ECOG = Eastern Cooperative Oncology Group; T = taxane (docetaxel); F = fluoropyrimidine (5-FU, capecitabine, or S-1); P = platinum (cisplatin or oxaliplatin).

was $351.9 \pm 8.7 \text{ mg}$ (mean $\pm$ s.e.) during the second cycle ($n = 38$), $347.1 \pm 14.6 \text{ mg}$ during the third cycle ($n = 26$), $350.0 \pm 17.0 \text{ mg}$ during the fourth cycle ($n = 22$), $360.0 \pm 20.5 \text{ mg}$ during the fifth cycle ($n = 15$), and $366.0 \pm 21.0 \text{ mg}$ during the sixth cycle ($n = 14$).

Efficacy

Of the first 29 patients enrolled, 3 patients achieved a partial response. Fourteen more patients were subsequently included in the study. Overall, four patients achieved confirmed partial response, resulting in objective RRs of 9.3% (95% CI: 0.6–18.0) for the ITT population and 9.8% (95% CI: 0.4–18.2) for the per-protocol population. The duration of objective response in these four patients was 2.5, 2.8, 5.8, and 11.0 months. A total of 23 patients had stable disease, and the overall tumour control rate was 62.8% for the ITT population. The tumour control rate was 80% for the 10 patients who received an increased dose at cycle 2, and 67.9% for the remaining 28 patients. Two patients were not assessable for response because they refused further treatment after the first cycle of chemotherapy. Response to previous chemotherapy was not correlated with response to irinotecan monotherapy ($P = 0.537$).

At a median follow-up time of 18.4 months (95% CI: 8.7–28.1), the median TTP was 2.8 months (95% CI: 2.3–3.4), and the median OS was 8.0 months (95% CI: 6.2–9.8; Figure 1). The TTP and OS were not significantly associated with patient age (age ≤ 51 years

vs age >51 years), sex, histological differentiation, number of metastatic sites, prior response to previous chemotherapy, TTP in the previous chemotherapy (TTP ≤ 2.8 months vs TTP > 2.8 months), or TFI (TFI ≤ 31 days vs TFI > 31 days). The median TTP for the 10 patients who received an increased dose at cycle 2 was 4.1 months; the median TTP for the remaining 28 patients was 2.8 months (hazard ratio: 0.91, 95% CI: 0.41–2.04, *P* = 0.833).

Safety

All 43 patients were evaluable for adverse events. Table 2 shows the patient distribution according to the highest observed toxicity grade and incidence of adverse events per chemotherapy cycle. The most common haematological toxicity was granulocytopenia; grade 3–4 granulocytopenia occurred in 23 patients (53.5%) and in 27.5% of the cycles. Febrile neutropenia was the least common toxicity, developing in two patients (4.6%) who were treated successfully with antibiotics and G-CSF, with no recurrence after adequate dose modification. The most common grade 3–4 non-haematological toxicities were fatigue (32.6% of patients; 10.1% of cycles), nausea (16.3% of patients; 4.5% of cycles), vomiting

(16.3% of patients; 3.9% of cycles), and anorexia (14.0% of patients; 4.5% of cycles). Severe diarrhoea occurred in only four patients (9.3%) and in 2.3% of cycles. All other toxicities were observed in less than 5% of patients. No treatment-related deaths were directly attributable to irinotecan chemotherapy. Among the 10 patients who underwent dose escalation from cycle 2, grade 3 or worse granulocytopenia occurred in five patients, vomiting in two patients, and diarrhoea in one patient (Table 3). The prevalence of severe toxicities for patients who underwent irinotecan dose escalation (*n* = 10) did not differ from that of the other patients (*n* = 28).

UGT1A1 genotype and allele frequencies

Table 4 shows the UGT1A1 variants and allele frequencies of the 43 enrolled patients. UGT1A1 allele frequencies were 0.278 for –3279T>G (*60) and 0.222 for 211G>A (*6). This finding

Table 3 Grade 3 or worse toxicity related to irinotecan dose escalation in patients proceeding to the second cycle (*n* = 38)

Grade 3 or worse	Dose escalation	
	Yes (<i>n</i> = 10)	No (<i>n</i> = 28)
Anaemia	0	5 (17.8%)
Leukopenia	3 (30%)	9 (32.1%)
Neutropenia	5 (50%)	15 (53.5%)
Thrombocytopenia	0	1 (3.5%)
Bleeding	0	0
Febrile neutropenia	0	1 (3.5%)
Fatigue	0	10
Anorexia	0	4 (14.2%)
Nausea	0	5 (17.8%)
Vomiting	2 (20%)	2 (7.1%)
Stomatitis	0	0
Constipation	0	0
Diarrhoea	1 (10%)	2 (7.1%)
Abdominal pain	0	0
Transaminitis (ALT)	0	2 (7.1%)
Transaminitis (AST)	0	1 (3.5%)
Hyperbilirubinemia	0	1 (3.5%)
Cholinergic symptoms	0	0

Abbreviations: ALT = alanine transaminase; AST = aspartate transaminase.

Figure 1 Time to progression (dotted line) and overall survival (solid line) of patients with advanced gastric cancer (*n* = 43).

Table 2 Haematological and non-haematological toxicities associated with irinotecan monotherapy (*n* = 43)

Category	Grade (no. of patients, %)				Grade (no. of cycles, %) ^a			
	1	2	3	4	1	2	3	4
Anaemia	10 (23.3)	25 (58.1)	3 (7.0)	3 (7.0)	66 (37.1)	72 (40.4)	14 (7.9)	3 (1.7)
Leukopenia	11 (25.6)	11 (25.6)	13 (30.2)	2 (4.7)	44 (24.7)	32 (18.0)	24 (13.5)	2 (1.1)
Neutropenia	7 (16.3)	8 (18.6)	7 (16.3)	16 (37.2)	30 (16.9)	35 (19.7)	24 (13.5)	25 (14.0)
Thrombocytopenia	6 (14.0)	1 (2.3)	2 (4.7)	2 (4.7)	25 (14.0)	4 (2.2)	3 (1.7)	3 (1.7)
Bleeding	1 (2.3)	2 (4.7)	0	0	2 (1.1)	2 (1.1)	0	0
Febrile neutropenia	NA	NA	1 (2.3)	1 (2.3)	NA	NA	1 (0.6)	1 (0.6)
Fatigue	5 (11.6)	24 (55.8)	14 (32.6)	0	76 (42.7)	73 (41.0)	18 (10.1)	0
Anorexia	16 (37.2)	18 (41.9)	6 (14.0)	0	79 (44.4)	41 (23.0)	8 (4.5)	0
Nausea	14 (32.6)	18 (41.9)	7 (16.3)	0	67 (37.6)	55 (30.9)	8 (4.5)	0
Vomiting	5 (11.6)	17 (39.5)	7 (16.3)	0	26 (14.6)	49 (27.5)	7 (3.9)	0
Stomatitis	14 (32.6)	9 (20.9)	0	0	36 (20.2)	14 (7.9)	0	0
Constipation	14 (32.6)	2 (4.7)	0	0	29 (16.3)	2 (1.1)	0	0
Diarrhoea	24 (55.8)	10 (23.3)	3 (7.0)	1 (2.3)	82 (46.1)	14 (7.9)	3 (1.7)	1 (0.6)
Abdominal pain	9 (20.9)	6 (14.0)	0	0	12 (6.7)	6 (3.4)	0	0
Transaminitis (ALT)	9 (20.9)	1 (2.3)	2 (4.7)	0	14 (7.9)	3 (1.7)	0	0
Transaminitis (AST)	12 (27.9)	2 (4.7)	1 (2.3)	0	22 (12.4)	3 (1.7)	1 (0.6)	0
Hyperbilirubinemia	1 (2.3)	0	2 (4.7)	0	1 (0.6)	1 (0.6)	1 (0.6)	0
Cholinergic symptoms	24 (55.8)	3 (7.0)	0	0	57 (32.0)	4 (2.2)	0	0

Abbreviations: ALT = alanine transaminase; AST = aspartate transaminase; NA = not applicable. ^aToxicity evaluation was available for 178 cycles of chemotherapy.

is consistent with that reported by a previous study of Asian populations. The frequency of the -40insTA (*28) allele was 0.095, and none of the enrolled patients were homozygous for the *28 allele.

Association of UGT1A1 genotype with toxicity and irinotecan dose escalation

The proteins encoded by UGT1A1 *60, *28, and *6 are known to possess different enzymatic activities. Thus, we categorised patients as having no defective alleles ($n=11$), one defective allele ($n=15$), or more than one defective allele ($n=14$; Table 5). The results indicate that patients with no defective allele had a lower incidence of grade 3–4 neutropenia than other patients ($P=0.018$, Figure 2). Severe (grade 3 or worse) neutropenia developed more frequently in patients who were homozygous for the $-3279\text{T}>\text{G}$ allele 4 out of 4 (100%) than in patients homozygous for the wild-type allele 7 out of 20 (35%) or those who were heterozygous 6 out of 16 (38%). Severe leukopenia and neutropenia was also more common in patients carrying UGT1A1*6 or *28.

DISCUSSION

Therapeutic options such as salvage chemotherapy are limited for patients with AGC. In this study, we evaluated individualised dosing of irinotecan monotherapy given every 3 weeks after failure with fluoropyrimidine, a platinum agent, and a taxane. We found that individualised dose optimisation produced an overall RR of 9.3%, disease control rate of 62.8%, and a tolerable toxicity profile.

Several phase II studies have evaluated irinotecan monotherapy as first-line chemotherapy for gastric cancer. In an European trial, Kohne *et al* (2003) administered irinotecan (350 mg m^{-2}) every 3 weeks and reported an RR of 20%. In a Japanese study, Futatsuki *et al* (1994) reported an RR of 23% in 60 evaluable patients who

received a dose of 100 mg m^{-2} every week or 150 mg m^{-2} every 2 weeks. As a second-line treatment for AGC, irinotecan produced a positive response in 16–20% of patients who did not respond to previous fluoropyrimidine or cisplatin-based chemotherapy (Futatsuki *et al*, 1994; Chun *et al*, 2004; Wesolowski *et al*, 2009). In this study, we observed an RR of 9.3% and TTP of 2.8 months, despite intensive previous treatment in our study population; 37 of the 43 enrolled patients (86%) underwent second-line chemotherapy before enrolment. Although the RR and TTP in this study were modest, the OS associated with individualised dosing of irinotecan monotherapy appeared promising compared with results of previous trials. Asian studies assessing the efficacy of salvage regimens for metastatic gastric cancer have reported an RR of 6.7–23.1%, TTP of 2.2–4.4 months, and OS of 6.2–10 months (Yoshida *et al*, 2006; Jeung *et al*, 2007; Jo *et al*, 2007; Lee *et al*, 2008; Shin *et al*, 2008; Sym *et al*, 2008; Zhong *et al*, 2008). We should be cautious when comparing these results because of differences in patient characteristics, inclusion/exclusion criteria, and further treatment after the clinical trial. For example, in this study all patients had a good performance status, and 28 of 43 patients (65.1%) underwent further treatment with chemotherapy after stopping irinotecan chemotherapy. In this context, there could be a discrepancy between the short progression-free interval and longer survival after disease progression.

In this study, the most common haematological toxicity was grade 3–4 granulocytopenia ($n=23$, 53.5%), and the least common

Figure 2 Number of defective alleles (*6, *28, and *60) and grade 3 or 4 neutropenia (%) during the first cycle of irinotecan monotherapy ($n=40$).

Table 4 UGT1A1 genotypes and allele frequencies

UGT1A1	–3279 T>G	–364 C>T	–64 G>C	–40·39 insTA	211 G>A	686 C>A	1456 T>G
Amino-acid change					G71R	P229Q	Y486D
Functional change	Reduced			Reduced	Reduced	Reduced	Reduced
Marker allele	*60			*28	*6	*27	*7
Frequency	0.278	0.095	0.056	0.095	0.222	0.008	0.000

Table 5 Relationship between the number of defective alleles and toxicity ($n=40$)

UGT1A1	No. of defective alleles	No. of patients	Irinotecan dose escalation	Grade 3–4 neutropenia	Febrile neutropenia	Treatment delay
*1/*1	0	($n=11$)				
		11	4/11 (36%)	1/11 (9%)	0/11	2/9 (22%)
*1/*60	1	($n=15$)	3/9 (33%)	3/9 (33%)	0/9	4/9 (44%)
*1/*6	1	6	1/6 (17%)	4/6 (67%)	1/6 (17%)	2/5 (40%)
*60/*60	2	($n=14$)	0/3	3/3 (100%)	0/3	0/3
*6/*60	2	1	0/1	0/1	0/1	0/1
*6/*6	2	4	1/4 (25%)	2/4 (50%)	0/4	1/3 (33%)
*28/*60	2	5	1/5 (20%)	3/5 (60%)	0/5	1/5 (20%)
*6/*28/*60	3	1	0/1	1/1 (100%)	0/1	1/1 (100%)

toxicity was febrile neutropenia ($n = 2$, 4.6%). The most common grade 3–4 non-haematological toxicities were fatigue (32.6%), nausea (16.3%), and vomiting (16.3%). The frequency of grade 3–4 granulocytopenia in our population (53.5%) was higher than that reported by previous studies (22–47%) that used standard irinotecan dosing (Futatsuki *et al*, 1994; Ychou *et al*, 2002; Van Cutsem *et al*, 2005). Few toxicities affecting the digestive system, such as severe diarrhoea (9.3%), were observed in our study. This result was likely due to appropriate management of delayed diarrhoea (Duffour *et al*, 2003). The safety profile was acceptable despite dose escalation. In fact, the 10 patients who underwent dose escalation had rates of grade 3 or worse neutropenia, vomiting, and diarrhoea that were similar to those of the other patients.

A European phase I study that evaluated irinotecan administered once every 3 weeks to treat AGC reported higher RRs at higher doses (450–750 mg m⁻²; Abigerges *et al*, 1995). Ychou *et al* (2002) administered an initial irinotecan dose of 350 mg m⁻², increasing the dose to 500 mg m⁻² in subsequent cycles for patients without grade 2 or worse non-haematological toxicities. In that study, the increased dose given to 63% of patients in the first-line setting was associated with increased activity. The RR was 35.5% for patients who received the high dose and 24.5% for patients who received the low dose. Van Cutsem *et al* (2005) reported that individualised dose escalation of irinotecan (up to 500 mg m⁻² based on patient tolerance) allowed more patients to receive higher doses without increased toxicity. This study demonstrated that individualised dose escalation based on toxicity in the preceding cycle could be a feasible salvage strategy for treating AGC. In addition, the RR and disease control rate in this study were similar to those obtained with standard dosing of irinotecan in a second-line setting (Futatsuki *et al*, 1994; Chun *et al*, 2004; Wesolowski *et al*, 2009) and irinotecan-based combination chemotherapy as salvage treatment (Sym *et al*, 2008).

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Pharmacogenetic features, such as UGT1A1 profile, are believed to have an important role in the highly variable toxicity to irinotecan (Ando *et al*, 2000). Among UGT1A1 genotypes, *60 is common in Asian populations, as seen in this study. In Caucasian populations, allele frequencies have been reported as 0.550 for *60, 0.388 for *28, and 0.007 for *6 (Kaniwa *et al*, 2005). We found that the number of UGT1A1 defective alleles was associated with the severity of neutropenia. The clinical implications of this finding are limited, because severe neutropenia did not result in febrile neutropenia or treatment delay. However, our study may provide information about the influence of UGT1A1 genotype on the severity of neutropenia.

In conclusion, our study demonstrated the safety of individualised irinotecan dose escalation based on patient tolerance. In addition, the modest anticancer activity of irinotecan indicates its feasibility as salvage chemotherapy for AGC after failure of previous treatment with fluoropyrimidine, a platinum drug, and a taxane. UGT1A1 genotyping may be a useful tool to identify the most appropriate starting dose.

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Conflict of interest

The authors declare no conflict of interest.

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