

# A phase I study of tasisulam sodium (LY573636 sodium), a novel anticancer compound in patients with refractory solid tumors

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## Abstract

**Purpose** This phase I study was carried out to determine the phase II recommended dose of tasisulam sodium (hereafter, tasisulam), a novel anticancer agent with a unique mechanism of action.

**Methods** Tasisulam was administered intravenously, every 21 days, in patients with refractory solid tumors using a three-plus-three dose-escalation schema.

**Results** Fifty-three patients were enrolled; the first 34 were treated with a flat dose of tasisulam of up to 2,400 mg, the dose level at which all three patients had dose-limiting toxicity (DLT). Controlling for  $C_{\max}$  proved important to reduce the risk of toxicity; therefore, we initially focused on identifying which parameters explained  $C_{\max}$

(end-of-infusion concentration) variability. Pharmacokinetic analysis indicated that  $C_{\max}$  negatively correlates with lean body weight (LBW). Thus, the dosing regimen was revised using a LBW-based algorithm targeting a specific  $C_{\max}$ . A loading/chronic dose paradigm was then implemented as pharmacokinetic results revealed a long terminal half-life of tasisulam, likely because of its high-affinity albumin binding.  $C_{\max}$ -based dose escalation was stopped at the 420- $\mu\text{g}/\text{mL}$  cohort, in which one of the 16 patients had DLT (transient hepatic transaminase elevation); grade 3/4 hematologic toxicity was noted in later cycles in three patients. Although response was not a primary objective, 33% of heavily pretreated patients with post-dose radiological assessments had stable disease.

**Conclusion** Implementation of a novel targeted  $C_{\max}$ -based dosing regimen allowed for the recommendation of a phase II tasisulam dose (loading dose of 420  $\mu\text{g}/\text{mL}$  targeted  $C_{\max}$  with all subsequent doses administered at 65% of chronic dose given every 21 days) despite pharmacological challenges posed by high albumin binding.

**Keywords** Tasisulam · Solid tumors · Phase I · Chemotherapy · LY573636

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## Introduction

Tasisulam sodium (benzamide, *N*-[(5-bromo-2-thienyl)sulfonyl]-2,4-dichloro-, sodium salt; hereafter referred to as tasisulam), an acylsulfonamide, is a member of a new class of antineoplastic agents [1]. An *in vitro*, antitumor-screening analysis conducted by the National Cancer Institute (NCI COMPARE analysis) [2] demonstrated a broad range of activity for tasisulam in 60 tumor cell lines including leukemia, melanoma, non-small cell lung (NSCLC), colon, ovarian, renal, and breast cancers. In addition, the activity profile of tasisulam in the COMPARE analysis did not match that of any other known anticancer compound [2], which suggests a mechanism of action for tasisulam that is distinct from currently available agents.

Preclinical *in vitro* studies indicate that tasisulam induces apoptosis in cancer cells by the mitochondrial (intrinsic) cell death pathway [3], and also induces a block at the G2/M phase of the cell cycle. However, the precise cellular or mitochondrial target that leads to the induction of apoptosis remains unknown. Tasisulam has also demonstrated anti-angiogenic activity *in vitro* (Lilly data on file).

*In vivo* pharmacokinetic studies in rats and dogs indicate that tasisulam is metabolized primarily by the liver, and has low total plasma clearance with a relatively long half-life (approximately 10 h in rats and 20 h in dogs). *In vitro* measurements also showed that tasisulam is highly protein bound (~99%) in humans, mice, rats, and dogs (Lilly data on file). In addition, there was preclinical evidence of a correlation between the maximum plasma concentration ( $C_{max}$ ) of tasisulam and toxicity (Lilly data on file).

The broad spectrum of preclinical antitumor activity of tasisulam and its potentially unique mechanism of action prompted this phase I dose-escalation study in patients with advanced or refractory solid tumors. The primary objective of the study was to determine the recommended phase II dose of tasisulam; secondary objectives were to characterize the toxicity and pharmacokinetic profiles of tasisulam, and to discern whether there was any antitumor activity.

## Patients and methods

### Eligibility criteria

Patients  $\geq 18$  years of age with histologically or cytologically documented advanced and/or metastatic malignancy for which no proven effective therapy exists were eligible to participate in this study. Patients were required to have a performance status of 0 or 1 on the Eastern Cooperative Oncology Group (ECOG) scale, adequate bone marrow reserve, kidney and liver function. Discontinuation of all

the previous therapies (including chemotherapy, radiotherapy, or other investigational therapy) at least 4 weeks before study enrollment and recovery from all toxicities were also required. Patients who had hematologic malignancies or a history of central nervous system neoplasm or who required palliative radiotherapy at study entry were excluded from study participation.

Written informed consent was obtained from each patient before study participation. The study was approved by the institutional review board(s) of the participating institutions, and was conducted according to applicable laws and regulations, good clinical practices, and the ethical principles of the Declaration of Helsinki.

### Treatment plan and dose-escalation schema

For the purpose of this open-label, single-arm, dose-escalating, phase I study, tasisulam was supplied by Eli Lilly and Company (Indianapolis, IN) in single-use, 20-mL glass vials containing tasisulam sodium, sodium chloride, and water or as a freeze-dried formulation in a 50-mL-vial equivalent to 500 mg of base compound. Tasisulam sodium was diluted in 500 mL of 0.9% sodium chloride, and was stored at room temperature for up to 8 h. This compound was administered as an intravenous infusion over 2 h ( $\pm 10$  min) on day 1 of a 21-day cycle.

Dose escalation and the maximum tolerated dose (MTD) were based on toxicity and pharmacokinetic results observed in cycle 1. Dose-limiting toxicity (DLT) was defined as possibly drug-related grade  $\geq 3$  non-hematologic toxicity (except for nausea/vomiting without maximal symptomatic/prophylactic treatment and alopecia) according to the NCI Common Toxicity Criteria (CTC) version 2.0 or grade 4 hematologic toxicity of  $>5$  days duration.

Initially, dose escalation was based on a flat-dose escalation schema ranging from 100 to 2,400 mg (100, 200, 400, 660, 1,000, 1,400, 1,800, and 2,400 mg). Early in the study, inpatient dose escalation to the next higher cohort was permitted if a patient completed two cycles of treatment without a DLT; however, the toxicities of these patients were not included in the DLT evaluation. The flat-dose escalation schema was later revised to one based on lean body weight (LBW; discussed further in “Results”).

The dose-escalation schema followed a three-plus-three design [4]. Per the protocol, if  $\geq 2$  patients in a cohort had DLT, then the MTD was surpassed and dose escalation was stopped. The MTD was then identified as the next lower cohort in which less than two of six patients had DLT, but if the dose exposure was nonlinear, patient recruitment was allowed into additional cohorts between the highest cohort reached and the previous cohort evaluated to identify the recommended phase II dose.

## Baseline and treatment assessments

Medical history, physical examination, performance status assessment, and tumor measurement of palpable or visual lesions were all carried out at baseline and at every cycle. Complete blood counts, serum chemistry, and urinalysis were performed at baseline, before each cycle, and post-therapy. Twelve-lead electrocardiograms were conducted at baseline, during cycle 1, and post-therapy. Clinically indicated radiologic imaging (computerized tomography scan, magnetic resonance imaging, or chest radiograph) was performed at baseline and at every other cycle to assess response to treatment. Tumor responses were followed and recorded using Response Evaluation Criteria in Solid Tumors (RECIST) [5]; however, the study was not designed to assess treatment efficacy.

## Bioanalytical methods

Plasma samples containing ethylenediaminetetraacetic acid (EDTA) were analyzed for total tasisulam concentrations using two validated liquid chromatography-tandem mass spectrometry (LC-MS/MS) methods. A dual-range method (that is, high-range and low-range) was needed due to the large peak to trough plasma concentrations observed. The calibration range for the high-range method was 2.50–500.0 µg/mL, and for the low-range method, it was 0.025–5.00 µg/mL. Samples above the upper limit of quantification were diluted and reanalyzed to yield results within the calibrated range. Samples below the lower limit of quantification using the high-range method were reanalyzed using the low-range method. Tasisulam and internal standard were extracted from EDTA plasma by protein precipitation using a 96-well plate format.

For the high-range method, accuracy (% relative error) for the original validation ranged from –3.73 to 12.52% for the intra-assay accuracy and between –1.93 and 8.36% for the inter-assay accuracy. Precision (% relative standard deviation) ranged from 1.18 to 6.72% for intra-assay precision and between 2.095 and 7.27% for inter-assay precision.

For the low-range method, accuracy (% relative error) for the original validation ranged from –2.88 to 13.84% for the intra-assay accuracy and between 5.77 and 7.69% for the inter-assay accuracy. Precision (% relative standard deviation) was between 2.38 and 11.54% for the intra-assay precision and between 0.760 and 8.080% for the inter-assay precision.

## Pharmacokinetic evaluations

Blood samples were collected predose, 0.5, 1, 2 (end of infusion), 2.25, 2.5, 3, 4, 5, 6, 8, 12, 24, 48, 72, and 168 h

(7 days) after the start of the infusion to evaluate the pharmacokinetic parameters of tasisulam. In addition, EDTA plasma samples (approximately 7 mL) were obtained predose from the first 12 patients in the study to assess protein binding. Pharmacokinetic parameters were computed by standard noncompartmental methods using WinNonlin Enterprise 5.0.1.

## Statistical methods

Dose proportionality was assessed by fitting the power model to pharmacokinetic parameters against dose and summarized using the methods described by Smith et al. [6]. Body-size parameters were added to the power model to determine which of these best explained the variability in pharmacokinetic parameters. The choice was based on selecting the model with the smallest Akaike Information Criterion (AIC) [7] for those covariates that significantly reduced the underlying interpatient variability.

## Results

### Patient characteristics

This study was conducted at the H. Lee Moffitt Cancer Center, Tampa, FL and the Memorial Sloan-Kettering Cancer Center, New York, NY, in collaboration with Eli Lilly and Company. From August 2003 to February 2007, 62 patients were entered into the study and 53 received at least one dose of tasisulam (seven patients did not meet enrollment criteria and two withdrew consent). Patient demographics are summarized in Table 1.

### Dose-escalation

On the basis of preclinical toxicity and efficacy testing (Lilly data on file), tasisulam was administered as a flat dose, ranging from 100 to 2,400 mg, to 34 patients enrolled in cohorts 1 through 9 (Table 2). DLTs were not observed in patients enrolled in cohorts 1 through 6 (100–1,400 mg). One of the three patients treated in cohort 7 (1,800 mg) was discontinued from the study in cycle 1 because of tumor-related spinal cord compression and was replaced; this patient had a grade 3 DLT (diarrhea). Three additional patients were then enrolled in this cohort (for a total of seven patients), and no additional DLTs were observed. In cohort 8 (2,400 mg), all three enrolled patients experienced DLT. One patient with metastatic NSCLC and a history of inflammatory colitis developed grade 3 stomatitis, grade 3 thrombocytopenia, and gastrointestinal bleeding, and later developed grade 4 febrile neutropenia and fatal acute respiratory distress syndrome (ARDS) in the setting of bilateral

**Table 1** Baseline patient demographics and disease characteristics (N = 53)

Sex, n (%)	
Female	37 (69.8)
Male	16 (30.2)
Ethnicity, n (%)	
Caucasian	47 (88.7)
Hispanic	3 (5.7)
African descent	2 (3.8)
East/Southeast Asian	1 (1.9)
Age, years	
Median	57
Range	18–83
ECOG performance status, n (%)	
0	28 (52.8)
1	22 (41.5)
2	2 (3.8)
Unknown <sup>a</sup>	1 (1.9)
Tumor sites/histology, n (%)	
Lung <sup>b</sup>	14 (26.4)
Sarcoma <sup>c</sup>	10 (18.9)
Carcinoma <sup>d</sup>	8 (15.1)
Ovarian	6 (11.3)
Melanoma	3 (5.7)
Colon	2 (3.8)
Breast	2 (3.8)
Other <sup>e</sup>	8 (15.1)
Median no. prior medications (range) <sup>f</sup>	6 (1–16)

ECOG Eastern Cooperative Oncology Group

<sup>a</sup> Patient was subsequently identified as having a performance status of 0 at baseline

<sup>b</sup> Non-small cell lung cancer (4 patients), adenocarcinoma (4 patients), mesothelioma (4 patients), bronchioloalveolar carcinoma (1 patient), and squamous (1 patient)

<sup>c</sup> Sarcoma (4 patients), uterine leiomyosarcoma (3 patients), osteogenic sarcoma (2 patients), and angiosarcoma (1 patient)

<sup>d</sup> Carcinoma (2 patients), thyroid carcinoma medullary (2 patients), adenocarcinoma unknown primary (1 patient), adenoid cyst carcinoma (1 patient), adrenal gland carcinoma (1 patient), and cholangiocarcinoma (1 patient)

<sup>e</sup> One patient each: endometrial, endocervical, bladder, pancreas, papillary adenocarcinoma, parotid gland, thymoma, and urothelium

<sup>f</sup> N = 52; prior medication(s) in one patient was not available

pulmonary nodules and pleural effusions. Another patient developed a prolonged prothrombin time (16-day duration without significant bleeding) that was later attributed to an interaction between tasisulam and the concomitantly administered medication, warfarin. The third patient developed hemoptysis from a tracheostomy site and grade 3 bilateral pneumonia 3 days after administration of tasisulam; the patient's white blood cell count was 12.4 k/ $\mu$ L and

platelet count was 159 k/ $\mu$ L. Bronchoscopy was negative for active bleeding, and no further hemoptysis was noted, but secretions were considered consistent with alveolar hemorrhage (grade 3 pulmonary hemorrhage). Because the MTD had been surpassed at the 2,400-mg flat-dose cohort, three patients were enrolled at an intermediate flat dose of 2,000 mg (cohort 9), and three additional patients were enrolled at the 1,800-mg flat dose (cohort 7). No additional DLTs were observed.

One patient each in cohorts 7 and 9 had grade 4 hematologic toxicity (thrombocytopenia/pancytopenia, respectively) in cycle 2 that lasted at least 8 days in one patient, suggesting the possibility of cumulative toxicity from the accumulation of bound drug with repeated flat dosing.

### Pharmacokinetics

Of the 34 patients administered flat doses of 100–2,400 mg (cohorts 1–9), 33 had full pharmacokinetic profiles from cycle 1. The analysis of pharmacokinetic data showed high interpatient variability (Table 3), a low total plasma clearance (approximately 0.03 L/h), a volume of distribution that was likely limited to the distribution volume of albumin (approximately 9 L) [8], and a terminal elimination half-life that corresponded to the turnover rate of albumin (approximately 11 days) [9]. Values of  $C_{\max}$  increased less than proportionately with dose, with coefficients of variation of 22% after adjusting for dose (Fig. 1a). On the basis of preclinical animal studies, in which the degree of  $C_{\max}$  variability appeared to be associated with the risk of toxicity (Lilly data on file), it was thought that controlling  $C_{\max}$  variability in patients should reduce their risk of toxicity. Therefore, the relationship between body size-related parameters and  $C_{\max}$  was explored. LBW, which was calculated using an algorithm incorporating the patient's height, weight, and gender [10, 11], was identified as an important cofactor that demonstrated a clear inverse relationship between LBW and  $C_{\max}$  (Fig. 1b) and reduced the interpatient variability of  $C_{\max}$  (from 22 to 13%) (Fig. 1c). Thus, pharmacokinetic analysis revealed that the  $C_{\max}$  for tasisulam (defined as its concentration at the end of the 2-h infusion) negatively correlated with the LBW of the patient. This led to a change from the flat-dose paradigm to an LBW-based dosing paradigm targeting an individualized  $C_{\max}$ .

### Phase II dose determination

Although the dose-escalation findings suggested an MTD for tasisulam of approximately 2,000 mg, the significant interpatient and inpatient variability of  $C_{\max}$  in the flat-dose cohorts and the high degree of albumin binding of tasisulam (99.7–99.9%) suggested that the flat-dosing schema was unlikely to provide consistent drug exposure.

**Table 2** Dose-limiting toxicity (DLT) and treatment cycles administered by cohort

Cohort	Dose	No. DLTs/No. pts in cycle 1	Number cycles administered	
			Median (range)	Total
1	100 mg	0/3	2 (2–4)	8
2	200 mg	0/3	2 (2–4)	8
3	400 mg	0/3	3 (2–4)	9
4	660 mg	0/3	2 (1–2)	5
5	1,000 mg	0/3	3 (2–4)	9
6	1,400 mg	0/3	2 (1–10)	13
7	1,800 mg	1/10 (G3 diarrhea)	2 (1–8)	28
8	2,400 mg	3/3 (G3 stomatitis + gastrointestinal bleeding + G4 febrile neutropenia; increased PT/INR for 16 days <sup>a</sup> ; G3 pneumonia + G3 pulmonary hemorrhage)	1 (1–2)	4
9	2,000 mg <sup>b</sup>	0/3	2 (2–4)	8
10	400 µg/mL	0/3	2 (1–6)	9
11	420 µg/mL	1/16 (transient grade 3 hepatic transaminase elevation) <sup>c</sup>	3 (1–14)	72

DLT was defined as any drug-related grade  $\geq 3$  non-hematologic toxicity according to the NCI Common Toxicity Criteria (CTC) Version 2.0, 1998 (except for nausea/vomiting without maximal symptomatic/prophylactic treatment and alopecia) or grade 4 hematologic toxicity of  $>5$  days duration in cycle 1

G grade, PT/INR prothrombin time/international normalized ratio

<sup>a</sup> The elevated PT/INR was later attributed to the interaction of tasisulam and the concomitant medication, warfarin; the protocol was subsequently amended to exclude patients who were actively receiving warfarin therapy as tasisulam and warfarin compete for the same albumin binding site

<sup>b</sup> An intermediate cohort of 2,000 mg (cohort 9) was explored. This cohort represented a dose between 1,800 mg (cohort 7) and 2,400 mg (cohort 8)

<sup>c</sup> Three additional patients developed hematologic toxicity (grade 3/4 thrombocytopenia/neutropenia) in later cycles ( $\geq$  cycle 2)

These findings led to a change from flat-dosing to a LBW-based dosing paradigm targeting specific  $C_{\max}$  values in cohorts 10 and 11.

Because of the hematologic toxicity observed in later cycles in cohorts 7 and 9, a loading dose was implemented in cycle 1 followed by a lower chronic dose in subsequent cycles in cohorts 10 and 11 to decrease the risk of drug accumulation due to the relatively long terminal half-life of tasisulam. The loading dose in cycle 1 for the LBW-based cohorts targeted a  $C_{\max}$  of 400 µg/mL (cohort 10) or 420 µg/mL (cohort 11). The chronic doses were 70 and 65% of the loading doses for cohorts 10 and 11, respectively. The initial LBW-based  $C_{\max}$  dose of 400 µg/mL was considered equivalent to the flat doses that were considered tolerable and showed antitumor activity (disease stabilization).

None of the three patients enrolled in cohort 10 had DLT (Table 2). In cohort 11, one of the six enrolled patients had DLT (transient grade 3 hepatic transaminase elevation); however, two of the six patients developed grade 3 or 4 hematologic toxicity in cycle 2, suggesting that escalation to a higher targeted  $C_{\max}$  was unlikely to identify a dose suitable for administration in phase II. Therefore, cohort 11 was expanded for further study. Of the 16 patients treated

in cohort 11, one patient had DLT, and three patients developed grade 3/4 thrombocytopenia/neutropenia in later cycles (cycle 2 or more); thus, dose escalation was stopped.

The median number of treatment cycles in cohort 11 was 3 (range 1–14 cycles) (Table 2). Of the 16 patients in cohort 11, 14 completed at least two cycles, and eight completed at least four cycles. Therefore, considering the heavily pre-treated patient population and the tolerability of the 420 µg/mL  $C_{\max}$  targeted dose, this dose was recommended for phase II study.

All of the 19 patients enrolled in cohorts 10 and 11 had full pharmacokinetic profiles in cycle 1. The pharmacokinetic parameters (CL and  $V_{ss}$ ) were similar in these cohorts compared with those of the flat-dose cohorts 1–9; however, comparison of the 1,800-mg flat-dose group and the 420-µg/mL dose group revealed that the LBW-dosing schema decreased the variability of  $AUC_{(0-\infty)}$  and CL by approximately 50%, and the variability of  $C_{\max}$  from 28 to 19% (Table 3). Comparisons of concentration versus time profiles for six pairs of patients, who were similar in their LBW difference, indicated smaller differences between  $C_{\max}$  values in patients dosed to a targeted  $C_{\max}$  of 420 µg/mL (Fig. 2a) compared with those given a flat dose (Fig. 2b).

**Table 3** Summary of noncompartmental pharmacokinetic parameters following a 2-h intravenous infusion of tasisulam

PK parameter	Geometric mean (CV%) <sup>a</sup>	
	1,800 mg ( <i>n</i> = 8)	420 µg/mL targeted <i>C</i> <sub>max</sub> ( <i>n</i> = 13)
<i>t</i> <sub>1/2</sub> (h) <sup>a</sup>	213 (74–942)	269 (154–617)
<i>C</i> <sub>max</sub> (µg/mL) <sup>b,c</sup>	322 (28)	359.83 (19)
AUC <sub>(0–∞)</sub> (h µg/mL)	70,900 (106)	98,100 (51)
CL (L/h)	0.026 (107)	0.024 (51)
<i>V</i> <sub>ss</sub> (L)	7.98 (29)	9.28 (24)
	Arithmetic mean (CV%) (Range)	
LBW (kg)	53.8 (39) (29.2–87.4)	48.6 (22) (35.9–71.7)
Dose per kg LBW (mg/kg)	38.4 (39) (20.6–61.7)	49.7 (8) <sup>d</sup> (42.0–55.6)

PK pharmacokinetic, CV coefficient of variation, *t*<sub>1/2</sub> terminal elimination half-life, *C*<sub>max</sub> maximum concentration (end-of-infusion concentration), AUC<sub>(0–∞)</sub> area under the concentration curve, CL clearance; *V*<sub>ss</sub> steady-state volume of distribution, LBW lean body weight

<sup>a</sup> The minimum and maximum values are reported if *N* ≤ 3; geometric mean and CV% values are reported if *N* > 3, except for *t*<sub>1/2</sub> for which geometric mean and range are reported

<sup>b</sup> Ten patients (for 1,800 mg) and 15 patients (for 420 µg/mL) were used to calculate the geometric mean for *C*<sub>max</sub>

<sup>c</sup> *C*<sub>max</sub> is reported as received

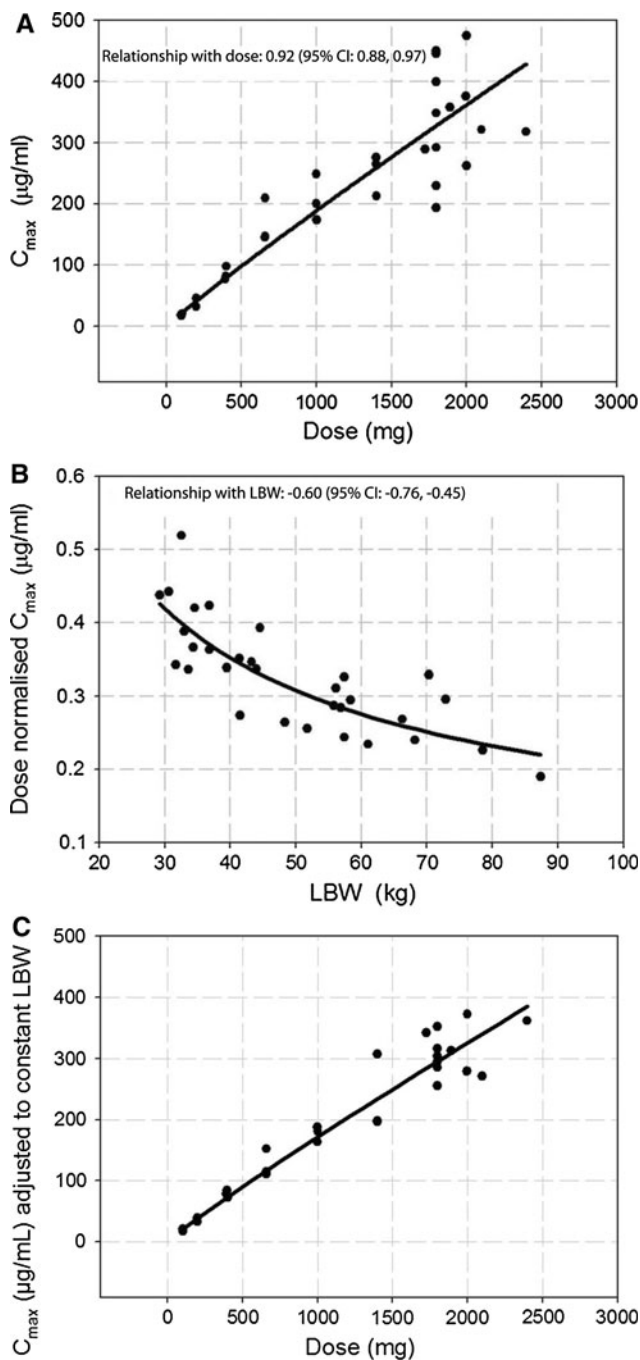
<sup>d</sup> Variation in mg/kg as the relationship between LBW and *C*<sub>max</sub> is not directly proportional

### Overall safety

The median number of cycles administered was 2 (range 1–18 cycles). The most common grade 3/4 toxicities considered possibly related to study drug were hematologic (thrombocytopenia, anemia, and neutropenia; Table 4). Fifteen patients received one or more transfusions (six red blood cells; seven red blood cells and platelets; one red blood cells, platelets, and plasma; and one plasma).

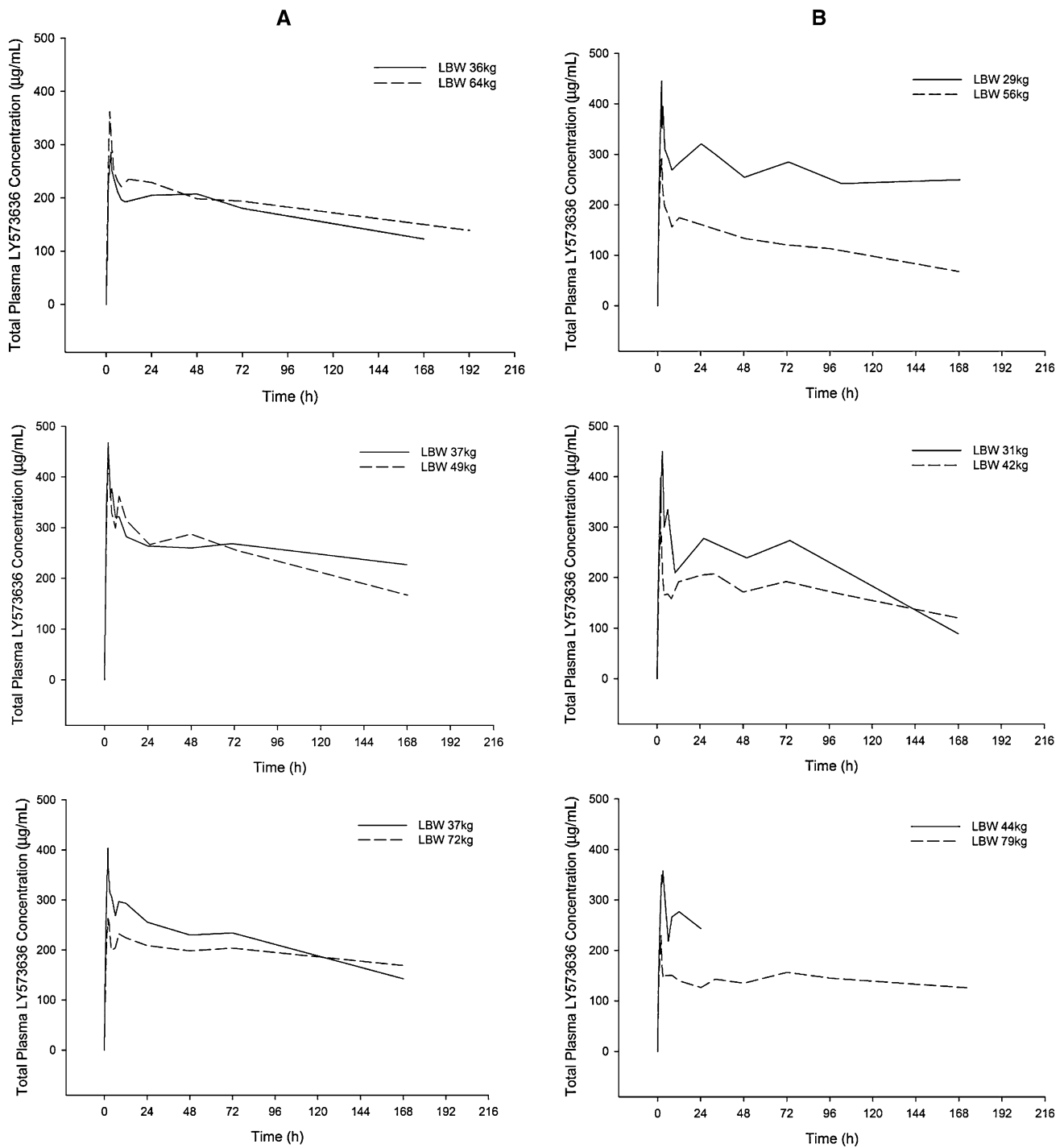
Of the six patients who died during the study, one patient in cohort 8 died because of complications related to bone marrow suppression (febrile neutropenia and ARDS during cycle 1), and one patient in cohort 11 died because of an outpatient cardiopulmonary arrest after receiving one cycle of study drug (for which drug relatedness could not be ruled out). An additional patient in cohort 11 died in cycle 2 from NSCLC in the setting of post-obstructive pneumonia; the patient had concurrent grade 2 neutropenia that was considered possibly related to study drug. All other patients died because of causes related to their malignancy.

Six patients (cohort 7 or above) were discontinued from the study because of adverse events considered possibly related to study drug and/or complications: coagulopathy



**Fig. 1** Results from assessment of dose proportionality showing *C*<sub>max</sub> values increasing less than proportionately with dose (a); the relationship between dose-normalized *C*<sub>max</sub> and LBW, demonstrating an inverse relationship between LBW and *C*<sub>max</sub> (b); and dose proportionality adjusted for LBW, showing reduced interpatient variability of *C*<sub>max</sub> (c). Interpatient CV% in *C*<sub>max</sub> after accounting for each covariate (cohorts 1–9): *C*<sub>max</sub> vs. dose: CV% = 22%; *C*<sub>max</sub> vs. dose + LBW: CV% = 13%

with prolonged prothrombin time (patient had been receiving warfarin) associated with rectal bleeding; vaginal bleeding; hypokalemia; thrombocytopenia; febrile neutropenia; and pneumonia and diffuse pulmonary hemorrhage.



**Fig. 2** Total plasma tasisulam (LY573636) concentrations using lean body weight (LBW) and targeted  $C_{max}$  dosing at 420 µg/mL (a) and flat dosing at 1,800 mg (b). The similar differences in LBW indicate that

the variability in pharmacokinetic parameters (both AUC and  $C_{max}$ ) was reduced with LBW dosing

**Efficacy**

Eight patients were discontinued from the study or died before cycle-2 response assessment because of ARDS and cardiopulmonary arrest (one patient, each); pneumonia and pulmonary hemorrhage (one patient); lost to follow-up (one

patient); withdrawal of consent (one patient); and disease progression (three patients, two of whom were objectively confirmed to have disease progression in follow-up visits). Of the 45 patients assessed for response, there were no objective responses; 25 had progressive disease, 15 had stable disease, and five had an unknown response because of

**Table 4** CTC grade 3 and 4 possibly drug-related toxicities ( $N = 53$ )

Toxicity	Grade 3 <sup>a</sup> <i>n</i> (%)	Grade 4 <i>n</i> (%)
Thrombocytopenia	7 (13.2)	3 (5.7) <sup>b</sup>
Anemia	5 (9.4)	1 (1.9)
Neutropenia	2 (3.8)	4 (7.5) <sup>c</sup>
Leukopenia	2 (3.8)	1 (1.9)
Hemorrhage/bleeding	4 (7.5)	0
Hypokalemia	4 (7.5)	0
Prothrombin time increased <sup>d</sup>	2 (3.8)	0
Diarrhea	2 (3.8)	0
Epistaxis	2 (3.8)	0
Cardiopulmonary arrest	0	1 (1.9)
Adult respiratory distress syndrome	0	1 (1.9)
Febrile neutropenia	1 (1.9)	1 (1.9)
Pneumonia	1 (1.9)	0

CTC Common Toxicity Criteria (Version 2.0, 1998)

<sup>a</sup> Additional grade 3 toxicities observed in one patient include lymphopenia, alkaline phosphatase, alanine transferase, aspartate transferase, abdominal pain/cramping, infection, nausea, vomiting, pleural effusion, and stomatitis/pharyngitis

<sup>b</sup> Platelet recovery times were available for two patients with platelet count nadirs  $<15 \times 10^3/\mu\text{L}$ , and were approximately 11 days for each

<sup>c</sup> Absolute neutrophil count (ANC) recovery times were available for two patients with ANC nadirs of approximately  $0.2 \times 10^3/\mu\text{L}$ , and were 12 and 14 days

<sup>d</sup> The prolonged prothrombin time of one patient was associated with grade 3 rectal bleeding/hematochezia and that of the other was a DLT without significant bleeding; both patients were co-administered warfarin

incomplete evaluation. However, five patients who received extensive prior therapy (median 6, range 5–12 prior cancer treatments) for NSCLC (two patients), ovarian cancer (two patients), or thymoma (one patient) received at least 10 cycles of tasisulam (minimum of 7 months) before disease progression.

## Discussion

This is a first-in-human, phase I, dose-escalation study of tasisulam, a novel anticancer agent with a potentially unique dual-faceted mechanism of action (apoptosis via intrinsic mitochondrial death pathway and anti-angiogenic activity). The recommended phase II dosing regimen for tasisulam, administered as an intravenous 2-h infusion on day 1 of 21-day cycles, was identified as a targeted  $C_{\text{max}}$  of 420  $\mu\text{g}/\text{mL}$  (based on LBW). The most common toxicities considered possibly related to study drug were those associated with bone marrow suppression, specifically, thrombocytopenia and neutropenia. There was no grade 3/4 renal toxicity or neuropathy. Grade 3 nausea, vomiting, and stomatitis were observed in one patient each.

Extensive pharmacokinetic analysis of the flat doses (100–2,400 mg) administered to the first 34 patients revealed wide variability in tasisulam plasma concentrations ( $C_{\text{max}}$ ). Preclinical animal studies suggested a correlation between plasma  $C_{\text{max}}$  and toxicity, which provided the rationale for identifying covariates that affected  $C_{\text{max}}$  in humans. LBW was identified in this study as an important covariate that significantly reduced interpatient variability in  $C_{\text{max}}$ . These findings led to a novel dosing strategy using LBW to achieve specific  $C_{\text{max}}$  targets. Despite the novelty of this dosing schema, dose calculation proved straightforward in this clinical study. Investigators entered patient height, weight, sex, and cycle number into a CD-ROM-based dosing calculator to determine the appropriate patient dose.

Laboratory studies performed on human plasma and serum confirmed that approximately 99.7% of tasisulam is bound to albumin and may be saturable across the presumed therapeutic dose range. Tasisulam showed bi-exponential disposition with a median terminal elimination half-life of approximately 11 days. Therefore, given the extensive albumin binding of tasisulam and its long terminal half-life, a loading dose in cycle 1 followed by a lower chronic dose in subsequent cycles was implemented to decrease the risk of a progressive rise in drug concentrations in later cycles, which could increase the risk of saturating albumin binding.

Although response was not a primary study objective, 15 of the 45 patients (33%) assessed for response at the end of cycle 2 (28% of all patients dosed) had stable disease as the best response despite having been heavily pretreated for various refractory solid tumors. In addition, five patients received at least 10 cycles of tasisulam before disease progression. The results of this trial and the findings from another phase I trial evaluating tasisulam [12] led to the initiation of phase II studies in relapsed/refractory solid tumors, including NSCLC, platinum-resistant ovarian cancer [13], soft tissue sarcoma, and malignant melanoma [14]. Pharmacokinetic and safety data from these phase II studies will provide additional opportunities to identify other potentially important laboratory or pharmacokinetic factors that could optimize dosing. In addition, preclinical and clinical investigations continue in an effort to provide additional insights on the mechanism of action of tasisulam.

In summary, analysis of pharmacokinetic and safety in this phase I study allowed for the development of a novel dosing algorithm for tasisulam, a first-in-class anticancer compound with a unique mechanism of action. Implementation of a LBW-based, targeted  $C_{\text{max}}$  dosing regimen and a loading/chronic dosing paradigm enabled the determination of the phase II recommended dose for tasisulam, despite pharmacological challenges posed by a high degree of high-affinity albumin binding and a long terminal half-life.



Although this study was not designed to assess efficacy, stable disease was observed in a reasonable proportion of extensively pretreated patients with solid tumors, suggesting the possibility of antineoplastic activity for tasisulam; testing of this hypothesis is being pursued in currently ongoing phase II clinical trials.

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