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A Phase 1 trial of temsirolimus and intensive re-induction chemotherapy for 2nd or greater relapse of acute lymphoblastic leukaemia: a Children's Oncology Group study (ADVL1114)

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Summary

The phosphatidylinositol 3-kinase (PI3K)/mammalian (or mechanistic) target of rapamycin (mTOR) signalling pathway is commonly dysregulated in acute lymphoblastic leukaemia (ALL). A phase 1 trial of the mTOR inhibitor temsirolimus in combination with UKALL R3 re-induction chemotherapy was conducted in children and adolescents with second or greater relapse of ALL. The initial temsirolimus dose level (DL1) was 10 mg/m² weekly × three doses. Subsequent patient cohorts received temsirolimus 7.5 mg/m² weekly × three doses (DL0) or, secondary to toxicity, 7.5 mg/m² weekly × two doses (DL-1). Sixteen patients were enrolled, 15 were evaluable for toxicity. Dose-limiting toxicity (DLT) occurred at all three dose levels and included hypertriglyceridaemia, mucositis, ulceration, hypertension with reversible posterior leucoencephalopathy, elevated gamma-glutamyltransferase or alkaline phosphatase and sepsis. The addition of temsirolimus to UKALL R3 re-induction therapy resulted in excessive toxicity and was not tolerable in children

Authorship

Conflict of interest

All other authors declare no competing financial interest.

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with relapsed ALL. However, this regimen induced remission in seven of fifteen patients. Three patients had minimal residual disease levels <0.01%. Inhibition of PI3K signalling was detected in patients treated at all dose levels of temsirolimus, but inhibition at an early time point did not appear to correlate with clinical responses at the end of re-induction therapy.

Keywords

acute lymphoblastic leukaemia; mTOR inhibitor; clinical trials; pharmacodynamics; relapse

Introduction

Mammalian (or mechanistic) target of rapamycin (mTOR)/phosphatidylinositol 3-kinase (PI3K) signalling, a critical pathway in cell proliferation, metabolism and apoptosis, is commonly dysregulated in acute lymphoblastic leukaemia (ALL), which may confer chemotherapy resistance (Tasian, *et al* 2014). While mutations in the *MTOR* gene, also known as *FRAP* (*FKBP-12-rapamycin associated protein*), itself are less common, constitutive activation of PI3K/mTOR signal transduction proteins or loss of negative regulation of mTOR occurs in many haematological malignancies (Bjornsti and Houghton 2004, Grabiner, *et al* 2014, Smolewski 2006, Tasian, *et al* 2014). Constitutive signalling activation may result from increased surface expression of growth factor receptors on leukaemia cells or from mutation of intracellular downstream effectors, including PI3K/mTOR pathway proteins (Chapuis, *et al* 2010). Additionally, loss of tumor suppressors that normally regulate PI3K signalling, such as *PTEN* (*phosphatase and tensin homolog*), can induce dysregulation of normal cellular equilibrium and facilitate aberrant signalling activation (Gutierrez, *et al* 2009).

Targeting constitutive PI3K/mTOR activation with small molecular inhibitors has been studied in ALL (Tasian, *et al* 2016). mTOR inhibitors (MTIs), such as sirolimus (rapamycin), inhibit growth of B-ALL and T-ALL cell lines *in vitro* and induce synergistic cell death in combination with anthracyclines, L-asparaginase and dexamethasone (Houghton, *et al* 2008, Teachey, *et al* 2006, Teachey, *et al* 2008). Further, mTOR inhibition can reverse glucocorticoid resistance in ALL, partly due to modulation of anti-apoptotic proteins, such as MCL1 (Wei, *et al* 2006). *In* murine models of human ALL, MTIs result in inhibition of leukaemia proliferation and prolong animal survival compared to vehicle-treated controls (Crazzolara, *et al* 2009, Houghton, *et al* 2008, Teachey, *et al* 2006, Teachey, *et al* 2008).

Despite aggressive retrieval strategies, the prognosis for children with refractory ALL is poor (Chessells 1998, Gaynon, *et al* 1998, Nguyen, *et al* 2008, Parker, *et al* 2010). Molecularly-targeted agents, including kinase inhibitors such as MTIs, have shown promise in treating some patients with haematological malignancies (Tasian, *et al* 2014). We conducted a trial to define the maximum tolerated dose (MTD) of temsirolimus in combination with intensive re-induction chemotherapy in children or adolescents with second or greater relapse of ALL. Additional exploratory aims included preliminary

assessment of treatment response within the context of a phase 1 trial and pharmacodynamic measurement of PI3K/mTOR pathway inhibition.

Methods

Patient eligibility

Patients 1 and <22 years of age in second or greater relapse of ALL were eligible. Relapsed leukaemia was defined as >25% blasts in bone marrow (M3) or 5–25% blasts in bone marrow (M2) with evidence of extramedullary disease. Patients with an isolated central nervous system relapse or extramedullary disease requiring immediate radiation therapy were not eligible.

Eligibility criteria included a Lansky/Karnofsky performance score 50, recovery from acute toxic effects of prior therapy and lack of active infections. Patients had to be 2 weeks from prior cytotoxic therapy with the exception of maintenance-type ALL therapy for which there was no washout period or a single intrathecal methotrexate within 72 h of systemic therapy initiation. Patients had to be 7 days from short-acting growth factor therapy or 14 days if long-acting, 7 days from biological therapy, >42 days from immunotherapy and > 3 half-lives from prior therapy with a monoclonal antibody. Patients had to be 3 months from prior HSCT and without evidence of graft-versus-host disease. At least two weeks from local radiation had to have elapsed. Patients receiving corticosteroids must have been taking a stable or decreasing dose for 7 days prior to enrolment. Hydroxycarbamide (hydroxyurea) use was permitted until 24 h prior to the first dose of study chemotherapy.

Other eligibility requirements included a normal age-adjusted serum creatinine or glomerular filtration rate (GFR) 70 ml/min/1.73 m², normal cardiac function defined by shortening fraction 27% or ejection fraction 50%, adequate pulmonary function with a baseline oxygen saturation >94% on room air, and adequate liver function defined as total bilirubin $1.5 \times$ the institutional upper limit of normal for age, normal gamma-glutamyltransferase (GGT) for age, alanine aminotransferase 225 u/l and albumin 20 g/l. Serum triglyceride and cholesterol were required to be 3.89 mmol/l and 7.77 mmol/l, respectively, and a fasting glucose had to be within normal limits for age.

This clinical trial was registered at www.clinicaltrials.gov (NCT01403415). Local institutional review board approval of the protocol was required. Written informed consent from patients 18 years or from parents/legal guardians of children age <18 years was obtained (and assent as appropriate) according to institutional policies. All authors had access to primary clinical trial data.

Drug administration and study design

The primary objectives of the study were 1) to estimate the MTD and/or recommended phase 2 dose of temsirolimus administered in combination with UKALL R3 re-induction chemotherapy (Parker, *et al* 2010) in children with relapsed ALL and 2) to define and describe the toxicities of temsirolimus in combination with intensive re-induction chemotherapy. The secondary objectives were 1) to determine the complete response rate

(CR), 2) to determine minimal residual disease (MRD) levels at end induction and 3) to evaluate the responsiveness of ALL cells to mTOR inhibition.

A rolling six dose escalation trial design was used (Skolnik, *et al* 2008) in which two to six patients can be concurrently enrolled onto a dose level, dependent upon (1) the number of patients enrolled at the current dose level, (2) the number of patients who have experienced DLT at the current dose level, and (3) the number of patients entered but with tolerability data pending at the current dose level. The MTD is the maximum dose at which fewer than one-third of patients experience DLT, therefore, if two or more of a cohort of up to six patients experience DLT at a given dose level, then the MTD has been exceeded and dose escalation is stopped. Temsirolimus was supplied by the Cancer Therapy Evaluation Program (National Cancer Institute; Bethesda, MD).

Each temsirolimus dose was administered intravenously 1 h post-mitoxantrone on day 1 and in proximity to vincristine on days 8 and 15. In DL-1 the day 15 temsirolimus dose was excluded. The starting dose of temsirolimus at dose level 1 (DL1) was 10 mg/m²/dose, two-thirds of the Federal Drug Administration-approved monotherapy dose in adults (Hudes, *et al* 2007) and well below the single agent dose of 150 mg/m²/dose identified in children with relapsed/refractory solid tumours (Spunt, *et al* 2011). Dose escalation to 15 mg/m²/dose (DL2) and 20 mg/m²/dose (DL3) was planned with one dose de-escalation to 7.5 mg/m²/ dose (DL0) if DL1 was deemed too toxic.

No intra-patient dose escalation was allowed. Each patient was enrolled for a single 36-day cycle of reinduction therapy and was monitored for toxicity to at least day 42. Due to DLTs at both DL1 and DL0, the protocol was amended to add a DL-1 cohort that received weekly temsirolimus 7.5 mg/m²/dose for two doses (in lieu of the three doses administered at DL0 and DL1).

Toxicities were graded according to the Common Terminology Criteria for Adverse Events version 4.0 (http://ctep.cancer.gov). International Consensus Conference on Toxicity Assessment guidelines were utilized to identify expected toxicities of multi-agent chemotherapy backbones and to define DLTs of combination therapy for the current study (Horton, et al 2010). Per these guidelines, specific non-haematological toxicities occurring in >6% of patients with similar chemotherapy were not considered DLTs if they returned to grade 2 by Day 42 of protocol therapy. These Grade 3 and 4 laboratory abnormalities included electrolyte abnormalities, elevated hepatic function tests and hypofibrinogenaemia. In addition, grade 3 fasting hyperglyceamia, hypercholesterolaemia, and hypertriglyceridaemia were not considered DLTs and use of medications to decrease hyperglycaemia or hyperlipidaemia was permitted. Other toxicities not considered DLTs included grades 3 or 4 nausea, vomiting, fatigue, anorexia, malaise or weight loss, hypertension, fever, infection and febrile neutropenia that were anticipated sequelae of the chemotherapy regimen. Any toxicity resulting in temsirolimus dose omission was considered a DLT. Haematological DLT for patients with ALL was defined as bone marrow aplasia at day 42 not attributable to leukaemia progression.

Complete response (CR) for patients with ALL was defined as bone marrow morphology with <5% blasts (M1), no evidence of extramedullary disease, and recovery of peripheral blood counts (absolute neutrophil count (ANC) > 0.5×10^9 /l and a platelet count > 50×10^9 /l). A CR with incomplete blood count recovery (CRi) was a CR without normalization of ANC and/or platelet count by Day 43 of re-induction. Partial response (PR) was defined as clearance of peripheral blasts with 5–25% blasts in bone marrow (M2) or as an M1 bone marrow without complete eradication of extramedullary disease. Patients who failed to qualify as a CR, CRi or PR were defined as stable or progressive disease (SD/PD). Disease evaluations were obtained at baseline and at the end of the 36-day cycle of therapy.

Pharmacodynamic analyses

In consenting patients, peripheral blood specimens were obtained to assess *in vivo* inhibition of PI3K pathway targets using phosphoflow cytometry as previously described (Loh, *et al* 2015). Samples (3–5 ml) were obtained at three time points: immediately prior to temsirolimus therapy (baseline, day 0), at day 3–5 of therapy (after the first dose of temsirolimus) and at day 36 (at end of re-induction).

Results

Patient Characteristics

Sixteen patients, aged 1–21 years, were enrolled between February 2012 and August 2014 (Table I). One patient did not initiate protocol therapy due to patient choice and was thus not evaluable. One evaluable patient had T-ALL, 14 evaluable patients had B-lineage ALL including infant *KMT2A* (*MLL*)-rearranged ALL (n=3), or *BCR-ABL1*-rearranged (Ph⁺) ALL (n=2) (Table II). Twelve patients were in second relapse, and three were in third relapse. Patients had previously undergone a mean of three prior salvage chemotherapy regimens prior to study entry (range 2–5), and 7 of the 15 patients had relapsed after a bone marrow transplant. All evaluable patients completed the full cycle of re-induction therapy.

Toxicity

Four patients were enrolled at DL1, two of whom experienced DLTs. One patient had persistence of Grade 3 hypertension, elevated GGT and mucositis at Day 42. The other patient had grade 4 fasting hypertriglyceridaemia (Table II). Per protocol design, the dose for the subsequent cohort was reduced to DL0. Of the six patients enrolled at DL0, three experienced DLTs. One patient had reversible posterior leucoencephalopathy syndrome, acidosis and mucositis, and died of sepsis at Day 33 prior to end of reinduction bone marrow evaluation. Another patient had persistently elevated GGT, and a third patient had grade 2 perirectal ulceration that did not resolve by day 42. The study was temporarily closed while clinical and correlative biology data were evaluated. Pharmacodynamic assays demonstrated inhibition of PI3K signalling at a temsirolimus dose of 7.5 mg/m², confirming that this dose had clinical potential for benefit, and the trial was amended to include DL-1 as outlined in the Methods.

Three of five patients treated at DL-1 experienced DLTs. Two patients had asymptomatic grade 4 hypertriglyceridemia, one of whom required omission of the second dose of

Table III delineates the non-haematological non-dose limiting toxicities at least possibly attributed to temsirolimus that occurred in >10% of patients. The most common non-haematological toxicities were electrolyte abnormalities, transaminitis and/or elevated GGT levels, and fever. Hypertriglyceridaemia and hyperglycaemia were more commonly reported than hypercholesterolaemia. Renal and pulmonary toxicities were rare.

A grade 5 sepsis event was the only infection-related toxicity that met the protocol definition of DLT. Other infectious toxicity was observed in this highly immunocompromised patient population including 11 of 15 patients (73%) with febrile neutropenia. Of the ten evaluable patients treated at DL1 or DL0, six had microbiologically-documented bacteraemia at day 10–12 of therapy or grade 3 catheter-related infections. Infectious organisms included gramnegative rods (*Klebsiella, E. coli, Citrobacter, Pseudomonas*), gram-positive cocci (*S. epidermis* or *S. aureus, Enterococcus*) and fungus (*Candida*). Four patients had >1 documented infection during their time on study. Three patients had grade 3 bacterial colitis/ enterocolitis. The protocol amendment to decrease temsirolimus administration to two doses per cycle (DL-1) also included recommendations for hospitalization and broad-spectrum anti-bacterial and anti-fungal prophylaxis in the cohort of patients treated at DL-1.

Response

Clinical responses were observed in patients treated at each dose level (Table II). Seven of the 15 evaluable patients had a CR or CRi. Three of these had flow cytometric MRD <0.01% and two others had MRD 0.01–0.03%, thereby allowing potential eligibility for subsequent bone marrow transplantation beyond the scope of this trial (Pulsipher, *et al* 2014). Six of the seven responders had undergone a prior bone marrow transplant and 1 was in 3^{rd} relapse at time of study entry.

Pharmacodynamic assessments

Peripheral blood samples from patients with B-ALL (n=8) or T-ALL (n=1) were obtained from patients treated at DL0 (n=5) or DL-1 (n=4) and analysed by phosphoflow cytometry. Two patients without constitutive activation of PI3K/mTOR signalling did not demonstrate signalling inhibition post-treatment. Seven of nine patients demonstrated elevated pretreatment (Day 0) levels of phosphorylated (p) PI3K pathway phosphoproteins (pPI3K, pmTOR, pS6, p4EBP1, pAkt, and/or pERK) in gated ALL blasts in comparison to controls (Figure 1A) (Tung, *et al* 2007). Compared to baseline, the mean inhibition of one or more PI3K phosphoproteins decreased in all Day 3–5 post-treatment blood specimens tested (Figure 1B). Insufficient numbers of residual ALL cells were present in most Day 36 peripheral blood samples for reliable measurement of leukaemia-specific signal transduction (data not shown). No differences in signalling inhibition were observed between patients treated at DL0 vs. DL-1. *In vivo* inhibition of PI3K/mTOR proteins at the Day 3–5 time point after temsirolimus and cytotoxic chemotherapy initiation was not predictive of clinical responses at Day 36.

Discussion

Therapy for children with relapsed ALL is hampered by low remission rates and high toxicity, especially in second and subsequent relapses. Many promising molecularly targeted agents are in development, but probably need to be combined with cytotoxic chemotherapy to improve cure rates and overall survival. This trial was conducted to evaluate the safety and tolerability of the MTI temsirolimus in combination with re-induction chemotherapy.

In this trial, the addition of temsirolimus to UKALL R3 re-induction therapy resulted in excessive toxicity and was deemed not tolerable in children with relapsed ALL. Dose limiting toxicity occurred at all dose levels. Most DLTs, such as hypertriglyceridaemia, mucositis and poor wound healing, were predictable based upon prior reports of mTOR inhibition in other cancers (Bagatell, et al 2014, Hudes, et al 2007, Spunt, et al 2011), as well as expected toxicities of similar intensive ALL re-induction chemotherapy backbones (Horton, et al 2010, Raetz, et al 2008, Sun, et al 2014). The use of high-dose dexamethasone pulses and asparaginase probably exacerbated the hypertriglyceridaemia often reported with MTI monotherapy. Similarly, a recent phase 1 trial of temsirolimus combined with irinotecan and temozolomide in children with relapsed solid tumours required modification to exclude patients on chronic steroids due to hyperlipidaemia (Bagatell, et al 2014). Temporary hypertriglyceridaemia may be considered an acceptable therapy-associated sequela in patients with multiply relapsed leukaemia. However, combination of MTIs with alternative salvage chemotherapy backbones that evoke fewer overlapping metabolic toxicities may increase successful delivery of this therapeutic strategy and reinduction of leukaemia remission. For example, in patients with first ALL relapse, preliminary data from a phase 1 trial combining the oral MTI everolimus with a 4-drug ALL induction has reported CR2 rates approaching 90% with less severe mucosal and gastrointestinal toxicity than was observed in our trial (Place, et al 2015).

Children and adolescents with multiply relapsed ALL have a significant risk of morbidity and mortality with intensive salvage therapy. Several recent relapsed ALL trials have reported a 4–5% induction mortality rate and 45–50% rate of grade 3 or 4 infections during re-induction therapy (Messinger, *et al* 2012, Raetz, *et al* 2008). Data from this study are highly concordant with reported infectious toxicity. We observed a 73% febrile neutropenia rate (11/15) and a 53% (8/15 patients) grade 3 or 4 documented infection rate and one grade 5 bacterial sepsis. The Therapeutic Advances in Childhood Leukaemia (TACL) consortium reported Grade 3 infections in 92% of patients with relapsed/refractory ALL treated with the R3 re-induction platform and 31% of patients had polymicrobial infection (Sun, *et al* 2014). More stringent hospitalization recommendations and infectious prophylaxis guidelines may abrogate some infectious complications (Messinger, *et al* 2012). Our study indicates that the addition of temsirolimus may increase infection risk of the UKALL R3 backbone, however, the presence or magnitude of increased risk cannot be fully assessed due to the infectious complication rate of the UKALL R3 backbone and risk of infection in this patient population.

Despite the excessive toxicity of combination therapy, 7 of the 15 evaluable patients (47%) were in CR/CRi at the end of re-induction therapy with five of these patients having MRD < 0.03%. Six of the 12 patients with a second relapse achieved a CR3, and 1 of the 3 patients with a 3^{rd} relapse achieved a CR4. Reismuller et al (2013) reported 30–40% CR3 and CR4 rates in a similar patient population. While remission rates of patients in second or greater ALL relapse treated with UKALL R3 therapy alone are currently unknown, our study with combination therapy demonstrates that a third or greater CR is achievable in nearly half of patients.

Pharmacodynamic assays performed during our study demonstrated baseline PI3K pathway signalling activation in most patients treated on this trial and *in vivo* target inhibition is measurable after first doses of temsirolimus and chemotherapy.

Temsirolimus with UKALL R3 re-induction chemotherapy led to unacceptable toxicity in this trial, however, alternative chemotherapy backbones in combination with other MTIs have demonstrated safety and tolerability (Daver, *et al* 2015, Tasian, *et al* 2014). A phase 1 trial (NCT01614197) evaluating the safety of temsirolimus in combination with 5 days of etoposide and cyclophosphamide is currently accruing. The study is based on the hypothesis that the combination of temsirolimus, etoposide and cyclophosphamide will result in less additive toxicity, such as hyperglycaemia, hypertriglyceridaemia and/or poor wound healing. In addition, oral formulations of MTIs, such as everolimus (NCT01523977) are being evaluated (Place, *et al* 2015). Furthermore, targeting of more proximal or multiple proteins in the PI3K/mTOR pathway with alternative tyrosine kinase inhibitors may result in greater sustained signalling inhibition and, possibly, to less dependence on conventional cytotoxic chemotherapy to achieve cure (Reismuller, *et al* 2013, Tasian, *et al* 2016).

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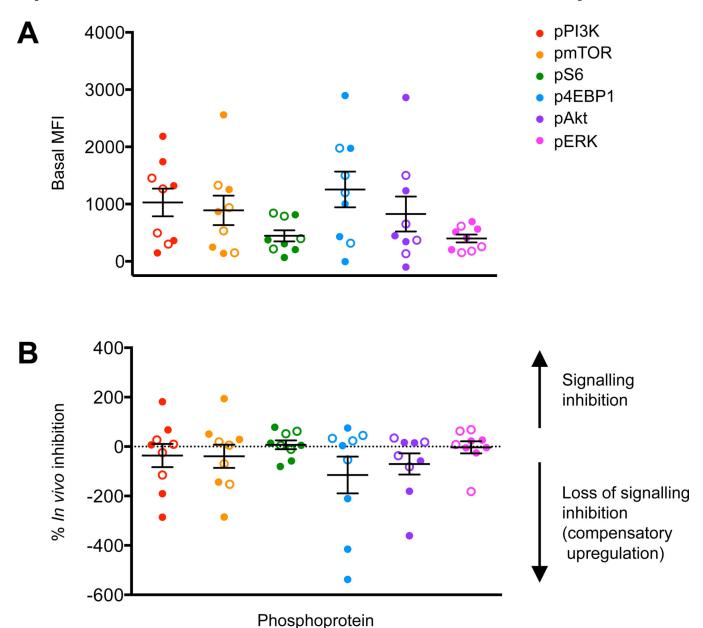


Figure 1. Phosphoflow cytometric analyses of *in vivo* signalling effects of temsirolimus and UKALLR3 chemotherapy

(A) Basal (pre-treatment) levels of PI3K pathway phosphoproteins were measured in gated ALL cells in peripheral blood specimens from ADVL1114 patients treated at DL0 or DL-1.
(B) Phosphoprotein inhibition at Day 3–5 of therapy after first dose of temsirolimus in comparison to basal phosphoprotein levels for each patient. Solid symbols = patients with complete response / complete response with incomplete blood count recovery, open symbols = stable disease/progressive disease. MFI = median fluorescence intensity. Data are displayed with means (central horizontal lines) and standard errors of the mean (whiskers).

Table I

Characteristics of patients enrolled on ADVL1114.

Characteristic	Number (%)		
	Number (%)		
Age (years)			
Median	9		
Range	1 – 21		
Sex			
Male	8 (50)		
Female	8 (50)		
Race			
White	13 (81)		
Asian	0 (0)		
American Indian or Alaska Native	0 (0)		
Black or African American	2 (13)		
Unknown	1 (6)		
Ethnicity			
Non-Hispanic	12 (75)		
Hispanic	4 (25)		
Prior Chemotherapy Regimens			
Median	3		
Range	2 – 5		

Table II

Dose-limiting toxicities and clinical responses of patients treated on ADVL1114.

Dose	Patient	Dx	Response	Dose-Limiting Toxicity
DL1 10 mg/m ² × 3	1	KMT2A	PD	
	2	B-ALL	Inevaluable	
	3	KMT2A	PD	Hypertension, mucositis, and GGT not resolved by day 42
	4*	B-ALL	CRi3	
	5	B-ALL	PD	Hypertriglyceridaemia
DL0 7.5 mg/m ² × 3	6*	B-ALL	CR4	
	7	B-ALL	PD	
	8	B-ALL	died	Sepsis, mucositis, acidosis, RPLE
	9	B-ALL	PD	Perianal ulcer not resolved by day 42
	10*	B-ALL	CRi3	
	11*	T-ALL	PD	GGT elevation not resolved by day 42
DL-1 7.5 mg/m ² × 2	12*	Ph^+	CR3	
	13	KMT2A	PD	
	14*	Ph^+	CRi3	Anorexia, GGT elevation, alkaline phosphatase elevation not resolved by day 42
	15*	B-ALL	CR3	Hypertriglyceridaemia
	16	B-ALL	CR3	Hypertriglyceridaemia

* = undergone prior haematopoetic stem cell transplantation; B-ALL = B cell acute lymphoblastic leukaemia; CR = complete response; CR3 = 3^{rd} complete response; CR4= 4^{th} complete response; CRi3 = 3^{rd} complete response with incomplete blood count recovery; DL = dose level; Dx = diagnosis; GGT= gamma glutamyltransferase; KMT2A = *KMT2A* (*MLL*)-rearranged acute lymphoblastic leukaemia (infant)PD = progressive diseasePh⁺ = *BCR-ABL1*-rearranged acute lymphoblastic leukaemia; RPLE = reversible posterior leukoencephalopathy; T-ALL = T cell acute lymphoblastic leukaemia.

Table III

Non-dose-limiting non-haematological toxicities related to protocol therapy and observed in >10% of evaluable patients (n=15).

-	Maximum Grade of Toxicity					
Toxicity	Grade 1	Grade 2	Grade 3	Grade 4		
Abdominal pain	1	1				
Alanine aminotransferase increased	2	7	4			
Alkaline phosphatase increased	3		1			
Anorexia	3		2			
Aspartate aminotransferase increased	3	3	2			
Blood bilirubin increased	3	2	1			
Catheter-related infection			4			
Cholesterol high	2					
Diarrhoea	4	3	1			
Fatigue	2	3				
Febrile neutropenia			7	1		
Fever		2	1			
GGT increased	1		2	1		
Hyperglycaemia	1	4	3			
Hypertension		3	2			
Hypertriglyceridaemia	1	3	4	4		
Hypoalbuminaemia	1	4	1			
Hypocalcaemia	2	1	1	1		
Hypokalaemia	2		2	1		
Hyponatraemia	3		2			
Hypophosphataemia	2	4	2			
Infections and infestations - other		1	1			
Mucositis oral		3	2	1		
Nausea	2	2	1			
Oral pain	1	1				
Pain	1	2				
Peripheral sensory neuropathy	1	1	1			
Sepsis				1		
Vomiting	4					