

# Analysis of prognostic factors of acute lymphoblastic leukemia in infants: report on CCG 1953 from the Children's Oncology Group

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**Infant acute lymphoblastic leukemia (ALL) has a poor therapeutic outcome despite attempts to treat it based on prognostic factor-guided therapy. This is the first cooperative group trial characterizing all infants at the molecular level for *MLL*/11q23 rearrangement. All infants enrolled on Children's Cancer Group (CCG) 1953 were tested for *MLL* rearrangement by Southern blot and the 11q23 translocation partner was identified (4;11, 9;11, 11;19, or "other") by reverse-transcriptase polymerase chain reaction (PCR).**

**One hundred fifteen infants were enrolled; overall event-free survival (EFS) was 41.7% (SD = 9.2%) and overall survival (OS) was 44.8% at 5 years. Five-year EFS for *MLL*-rearranged cases was 33.6% and for *MLL*-nonrearranged cases was 60.3%. The difference in EFS between the 3 major *MLL* rearrangements did not reach statistical significance. Multivariate Cox regression analyses showed a rank order of significance for negative impact on prognosis of CD10 negativity, age younger than 6 months, and *MLL* rearrangement,**

**in that order. Toxicity was the most frequent cause of death. Relapse as a first event in CCG 1953 was later (median, 295 days) compared with CCG 1883 historic control (median, 207 days). *MLL*/11q23 rearrangement, CD10 expression, and age are important prognostic factors in infant ALL, but molecular 11q23 translocation partners do not predict outcome. (Blood. 2006;108:441-451)**

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## Introduction

In the same treatment era that has seen the overall cure rate of childhood acute lymphoblastic leukemia (ALL) reach levels exceeding 70%, infants younger than 12 months of age with ALL continue to experience very low event-free survival (EFS). These infants comprise 2% to 5% of cases of childhood ALL.<sup>1</sup> Published results show 22% to 54% 3- to 6-year EFS.<sup>1-6</sup> Infants' high failure rate has been due to relapsed disease rather than toxicity.<sup>1,4,5</sup> Therapeutic efforts in infant ALL center on intensification and determining relevant prognostic factors to identify infants needing the most aggressive therapy.

ALL in infancy has been associated with unfavorable features such as hyperleukocytosis, hepatosplenomegaly, and central nervous system (CNS) disease.<sup>7,8</sup> The blast cells most commonly lack CD10 expression<sup>9</sup> and frequently coexpress lymphoid and myeloid antigens.<sup>10,11</sup> Cytogenetic 11q23 abnormalities, most commonly t(4;11)(q21;q23), and molecular genetic rearrangement of the *MLL* gene at chromosome band 11q23 have been associated with infant ALL.<sup>8,12</sup>

Correlation between various clinical and laboratory characteristics and outcomes has been studied in the context of a number of

clinical trials. These factors are interrelated, making it difficult to identify independent prognostic factors. Higher white blood cell (WBC) count has correlated with worse outcomes, but the WBC categories for comparison have varied in the different reports.<sup>1,5,6</sup> Even within the narrow one-year age range defined as infant ALL, the age in months at diagnosis shows a strong prognostic effect with increasingly poor prognosis for younger age. Infants younger than 6 months of age at diagnosis have had EFS from 8% to 40%, whereas those 6 to 12 months old at diagnosis have had EFS from 40% to 71%.<sup>1-3,5,6,13</sup>

Cytogenetic 11q23 rearrangement is found in 43% to 60% of infants with ALL<sup>12,14</sup>; *MLL* gene rearrangement is identified at the molecular level in 70% to 80% of all patients.<sup>13,15,16</sup> This difference in the level of detection of *MLL*/11q23 abnormalities makes it difficult to interpret outcomes when comparing various trials. The published 3- to 6-year EFS for infants with *MLL*/11q23-rearranged leukemia ranges from 5% to 28%.<sup>1,5,13,15-18</sup> Prior reports clearly suggest that infants with *MLL* translocation partner 4q21 do poorly.<sup>12,19</sup> The outcome for infants with 11q23 translocation partners other than 4q21 has been less clear.<sup>12,20</sup> The outcome for

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A complete list of the participating members of the Children's Oncology Group appears in "Appendix."

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P.A.D., F.O.S., W.G.W., Z.D., and G.H.R. designed the protocol; P.A.D., J.M.H., and H.S.J. ran the clinical trial; J.M.H., S.O.M., and P.A.D. wrote the manuscript; H.S. and D.V. performed the statistical analyses; N.A.H. reviewed the cytogenetics; R.M. performed the molecular analyses; and W.L.S. reviewed and summarized toxicity.

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infants without *MLL*/11q23 rearrangement has been better, with published 3- to 6-year EFS from 46% to 91%.<sup>1,5,13,15-18</sup>

CD10 negativity correlates strongly but not completely with *MLL*/11q23 rearrangement. Infants without CD10 expression have a reported EFS of 21% to 40% at 4 to 6 years, with an EFS of 45% to 73% at 4 to 6 years in infants with CD10<sup>+</sup> blasts.<sup>1-3,5,6,10</sup> CD10 positivity is therefore correlated with better outcome.

Given the overall poor outcome of infants with ALL and the problem of early relapse on the historic controls Children's Cancer Group (CCG) 107 and CCG 1883,<sup>1</sup> CCG 1953 was undertaken to evaluate intensive early therapy. The protocol was designed to assess the feasibility and outcome of induction/intensification followed by a reinduction, reintensification, and consolidation. The feasibility and outcome of bone marrow transplantation (BMT) using family or unrelated donors in infants with 11q23/*MLL* abnormality was also investigated. In order to identify prognostic factors, the trial was designed to include central analysis of Southern blot for detection of *MLL* rearrangement and reverse-transcriptase polymerase chain reaction (RT-PCR) to identify t(4;11)(q21;q23), t(11;19)(q23;p13), and t(9;11)(p22;q23).

## Patients, materials, and methods

### Patients

The protocol enrolled 115 infants (< 1 year of age) with newly diagnosed ALL from July 1996 to August 2000 onto CCG 1953. The diagnosis was based on morphology, cytochemistry, and immunophenotyping, all performed at individual CCG institutions. The protocol was approved by the National Cancer Institute and by Institutional Review Boards at CCG member institutions. Informed consent was obtained from parents according to the Code of Federal Regulations Guidelines and the Helsinki Protocol.

### Treatment

Treatment for patients on CCG 1953 consisted of 4 courses of an induction/intensification, reinduction, reintensification, and consolidation schema followed by 3 cycles of an intensified maintenance and 4 cycles of routine maintenance (Table 1). The protocol included high-dose methotrexate and G-CSF to decrease the length of neutropenia and facilitate the delivery of chemotherapy in a dose-intensive fashion.

Infants with 11q23/*MLL* abnormalities were allocated to BMT within 4 months if they had a protocol-defined 5-6/6 human leukocyte antigen matched related or unrelated donor. The protocol-specified preparative regimen included cytosine arabinoside 100 mg/kg/dose every 12 hours × 6 doses days -8, -7, -6; cyclophosphamide 45 mg/kg/dose intravenously daily × 2 doses days -7, -6; methylprednisolone 33 mg/kg/dose intravenous twice daily × 4 doses starting midnight day -2, -1; and total body irradiation (TBI) treatment dose was a total of 1200 cGy given in 150-cGy fractions (150 cGy twice daily × 8 doses days -3, -2, -1, 0) to the midline at the level of the umbilicus. Graft-versus-host disease (GVHD) prophylaxis consisted of single-agent cyclosporine 1.5 mg/kg intravenously every 12 hours starting day 1.

All infants received 7 doses of triple intrathecal therapy for CNS prophylaxis during the first 10 weeks of therapy, in order to maintain consistency with the concurrent Pediatric Oncology Group (POG) study 9407,<sup>21</sup> in the pre-BMT phase of the trial. Infants not undergoing BMT received an additional 14 doses of intrathecal therapy with either single-agent cytosine arabinoside (reintensification, consolidation, and intensified maintenance) or methotrexate (maintenance). Additionally, infants not undergoing BMT received very high-dose methotrexate as given in the CCG 107 and 1883 studies<sup>1</sup> in order to maintain the low (< 10%) CNS relapse rate attained in these studies. Patients with CNS disease at diagnosis received no additional therapy. CNS-directed radiation therapy was not administered, although patients receiving BMT received TBI.

The study opened for patient accrual in July of 1996. It was suspended in July of 1997 because of an induction mortality rate of 25% in the first 40 patients enrolled on both CCG 1953 and POG 9407, a POG study that used identical induction/intensification.<sup>21</sup> The deaths in CCG 1953 were 3 (60%) of 5 in those younger than 3 months and 4 (20%) of 20 in those older than 3 months. These deaths were due to infections associated with severe breakdown of skin and oral/gastric mucosa associated with daunomycin. The study was first amended with a change in anthracycline dosing based on weight rather than body surface area, by a graded dose schedule calculated to deliver an equivalent dose of daunomycin to older infants and de-escalate the dose for younger infants to a dose that was previously tolerated by infants on the CCG 192P protocol. The 48-hour continuous-infusion daunomycin dose was changed to 4 mg/kg for infants younger than 6 months; 5 mg/kg for infants 6 to 9 months; and 6 mg/kg for infants 9 months or older. The study reopened in November 1997. The induction mortality on CCG 1953 was reduced to 0 of 4 in those younger than 3 months and 1 (13%) of 8 in those older than 3 months, but unacceptable toxicity continued in those younger than 3 months on the POG 9407 study. As a result of the POG experience, a second modification was made in July of 1998, changing the daunomycin dosage to 2 mg/kg intravenous over 30 minutes daily for 2 days for infants younger than 90 days of age. Subsequent mortality was improved. A total of 115 patients were enrolled, and the study closed in August 2000.

The detailed results for infants receiving bone marrow transplantation will be described in a separate manuscript (P.A.D., J.M.H., H.S., D.V., N.A.H., R.M., F.O.S., W.G.W., W.L.S., H.S.J., Z.D., G.H.R., manuscript in preparation). This paper will include infants enrolled on CCG 1953 and the companion protocol POG 9407. A separate paper will report the long-term neuropsychologic outcomes of these infants (Thomas A. Kaleita, J.M.H., P.A.D., H.S., D.V., N.A.H., R.M., F.O.S., W.G.W., W.L.S., H.S.J., Z.D., G.H.R., manuscript in preparation).

### Historic controls

A group of 135 patients entered on CCG 1883 was the historic control for this protocol. The treatment and outcomes have been reported in detail (Table 2; Figure 1).<sup>1</sup> The historic controls and the CCG 1953 cohort were similar in presenting features with 2 main differences. First, there were slightly more infants younger than 90 days of age enrolled on CCG 1953 (20.9%) than on CCG 1883 (15.6%). Second, there were more CD10<sup>-</sup> infants on CCG 1953 (71.3% CD10<sup>-</sup> with all infants tested) than on CCG 1883 (57.7% CD10<sup>-</sup>, but no data are available for 38 of the 135 infants enrolled; Table 3). Thus, CCG 1953 had a population with poorer prognostic factors than the historic controls. Comparison of the frequency of *MLL*/11q23 gene rearrangement on these trials is problematic, as molecular and cytogenetic methods were employed on CCG 1953 but only cytogenetic methods were used on CCG 1883.

### Molecular analysis

All molecular analyses were performed in a Clinical Laboratory Improvement Amendments (CLIA)-certified laboratory (University of Minnesota Molecular Diagnostics Lab). Detection of *MLL* gene rearrangement by Southern analysis and detection of t(4;11), t(11;19), or t(9;11) fusion transcripts by RT-PCR were performed as previously published.<sup>22-24</sup>

### Cytogenetic analysis

Leukemic cell karyotype was determined by cytogenetic analysis at the individual institutions at the time of diagnosis. The protocol required both a direct preparation and a short-term unstimulated culture for bone marrow samples, with unstimulated peripheral blood as backup preparation. Complete analysis of 20 banded metaphases was required for each case. The definition of a clone was as follows: 2 or more metaphases with identical structural abnormalities or extra chromosomes, or 3 or more metaphases with identical missing chromosomes. Karyotypes were designated according to the ISCN 1995, An International System for Human Cytogenetic Nomenclature.<sup>25</sup>

Table 1. CCG 1953 therapy detail

Phase	Treatment regimen
<b>Induction, 4 wk</b>	
CPM	250 mg/m <sup>2</sup> every 12 h × 4 doses, days 2, 3
DXM	10 mg/m <sup>2</sup> orally 3 times daily, days 0-20, no taper
VCR	0.05 mg/kg, days 0, 14; 0.03 mg/kg day 7
DNM	2 mg/kg/dose intravenously over 30 min for patients age ≤ 90 d 2 mg/kg/d; 4 mg/kg over 48 h, age > 90 d to < 6 mo 2.5 mg/kg/d; 5 mg/kg over 48 h, age 6 to < 9 mo 3 mg/kg/d; 6 mg/kg over 48 h, age ≥ 9 mo, as a 48-h continuous infusion through central venous catheter, days 0, 1
L-ASP	6000 IU/m <sup>2</sup> × 8 doses, 3 times weekly
ITT	Intrathecal triple therapy (methotrexate 7.5 mg; hydrocortisone 7.5 mg; ARA-C 15 mg), days 0, 7, 14, 21, 28
MTX	4 g/m <sup>2</sup> , days 21, 28
CF rescue	10 mg/m <sup>2</sup> orally or intravenously every 6 h × ≥ 5 doses, start 18 h after MTX completion
VCR	0.05 mg/kg intravenous push
<b>Intensification, 4 wk</b>	
VP-16	100 mg/m <sup>2</sup> , days 36-40
DXM	10 mg/m <sup>2</sup> orally 3 times daily, days 0-20, no taper
CPM	300 mg/m <sup>2</sup> /day intravenously, days 36-40
<b>Reinduction, 4 wk</b>	
CPM	250 mg/m <sup>2</sup> every 12 h × 4 doses, days 2, 3
VCR	0.05 mg/kg days 0, 14; 0.03 mg/kg day 7
DNM	45 mg/m <sup>2</sup> /d × 2 days
DXM	10 mg/m <sup>2</sup> divided 3 times daily, days 0-20, no taper
L-ASP	6000 IU/m <sup>2</sup> intramuscularly × 8 doses, 3 times daily, days 0-20
ITT	Days 0, 14
BMT for 11q23/MLL	
<b>Reintensification, 6 wk</b>	
VCR	0.75 mg/m <sup>2</sup> intravenously, days 0, 7
VHMTX	6000 mg/m <sup>2</sup> in 1 h, followed by 1200 mg/m <sup>2</sup> /h × 23 hours × 2 doses, days 0, 14
CF rescue	200 mg/m <sup>2</sup> intravenously over 1 h, then 12 mg/m <sup>2</sup> intravenous push every 3 h × 6 doses, then 12 mg/m <sup>2</sup> intravenously or orally every 6 h until plasma MTX level < 0.1 μM; start 12 h following MTX completion (36 h after start)
VP-16	100 mg/m <sup>2</sup> /d intravenously, days 28-32
CPM	300 mg/m <sup>2</sup> /d intravenously over 30 min, days 28-32
IT ARA-C	15 mg intrathecally, days 7, 21
<b>Consolidation, 9 wk</b>	
VHMTX	6000 mg/m <sup>2</sup> intravenously in 1 h, followed by 1200 mg/m <sup>2</sup> /h intravenously for 23 h, days 28, 42
CF rescue	10 mg/m <sup>2</sup> orally or intravenously every 6 h × ≥ 5 doses, start 18 h after MTX completion
IT ARA-C	15 mg intrathecally, days 35, 49
VCR	0.75 mg/m <sup>2</sup> intravenously, days 28, 42
IV ARA-C	3000 mg/m <sup>2</sup> intravenous 3 h infusion every 12 h, days 0, 2, 7, 8 for total of 8 doses
L-ASP	6000 IU/m <sup>2</sup> intramuscularly 3 h after intravenous ARA-C on days 1, 8
<b>Intensified maintenance, 72 d × 3</b>	
VCR	0.75 mg/m <sup>2</sup> intravenously, days 0, 28
DXM	10 mg/m <sup>2</sup> divided orally 3 times daily, days 1-4 and 28-32
6-MP	75 mg/m <sup>2</sup> orally, days 0-48
MTX	20 mg/m <sup>2</sup> intramuscularly every week, days 0, 7, 14, 21, 28, 35, 42
VP-16	100 mg/m <sup>2</sup> /d, days 49-53
CPM	300 mg/m <sup>2</sup> /d intravenously over 30 min, days 49-53
IT ARA-C	15 mg intrathecally, days 0, 28
<b>Routine maintenance, 4 cycles, 1 y</b>	
VCR	1.5 mg/m <sup>2</sup> intravenously every 4 wk × 12 doses, days 0, 28, 56
PRED	40 mg/m <sup>2</sup> orally divided in 3 doses for 5 d with each VCR dose
MTX	20 mg/m <sup>2</sup> orally weekly, days 7, 14, 21, 28, 35, 42, 49, 56, 63, 70, 77
6-MP	75 mg/m <sup>2</sup> orally daily for 12 weeks per cycle
IT MTX	8 mg age 12-23 mo, 10 mg age 24-35 mo, day 0

CPM indicates cyclophosphamide; DXM, dexamethasone; VCR, vincristine; DNM, daunomycin; L-ASP, L-asparaginase; ITT, intrathecal triple therapy (methotrexate; hydrocortisone; ARA-C); BMT, bone marrow transplantation; ARA-C, cytosine arabinoside; MTX, methotrexate; CF, citrovorum factor (leucovorin); VP-16, etoposide; VHMTX, very high dose methotrexate; IT, intrathecal; IV, intravenous; 6-MP, mercaptopurine; and PRED, prednisone.

## Statistical methods

Life-table estimates were calculated by the Kaplan-Meier (KM) procedure<sup>26</sup> and the standard deviation (SD) of the life-table estimate was obtained using Peto et al's estimate for the variance.<sup>27</sup> The relative hazard rate (RHR) was estimated by the ratio of observed to expected events in the comparison groups.<sup>27,28</sup> The primary end points examined were EFS and

overall survival (OS) from study entry. EFS events included induction failure (nonresponse to therapy or death during induction), leukemic relapse at any site, death during remission, and second malignant neoplasm (SMN), whichever event occurred first. Disease-free survival (DFS) is also provided for some comparisons; this uses only the subset of patients who achieved an initial remission and includes the postinduction events (relapse, death, SMN). For all the life-table analyses, patients not experiencing the event of

**Table 2. Cumulative dose comparison of key drugs**

Drug	CCG 1883	CCG 1953
Cyclophosphamide, g/m <sup>2</sup>	11	9.5
<b>Daunomycin*, mg/m<sup>2</sup></b>		
3 mo or less	87.5	160
4 to less than 6 mo	125	166
6 to less than 9 mo	125	193
9 to less than 12 mo	125	218
L-asparaginase, units/m <sup>2</sup>	132 000	108 000
Etoposide, mg/m <sup>2</sup>	0	2 500
<b>Cytosine arabinoside, g/m<sup>2</sup></b>		
Less than 6 mo	12	24
6 to less than 12 mo	24	24
Dexamethasone, mg/m <sup>2</sup>	0	700
Prednisone, mg/m <sup>2</sup>	8 820	2 400
Intravenous methotrexate, g/m <sup>2</sup>	134.6	142.4

\*CCG 1953 values are estimated conversions based on final protocol dosages.

interest were censored in the analysis at the time of their last contact. Life-table comparisons of patient treatment groups or patient prognostic factors generally used the log-rank statistic<sup>29,30</sup> to examine differences among the groups. Cox regression analysis was used to assess the relative importance of certain patient prognostic characteristics in a multivariate analysis for EFS outcome.<sup>31</sup> Hypothesis testing in the regression analyses used likelihood ratio tests. Simple chi-square tests were used to assess similarity of patient characteristics when compared with the historic control study.

## Results

### Patient characteristics

The study population characteristics are detailed in Table 3. Twenty-four patients (20.9%) were younger than 3 months of age. Hepatomegaly and splenomegaly were present in 77.0% and 80.9% of patients, respectively, both defined as either enlarged but not below umbilicus (moderately enlarged) or below umbilicus (markedly enlarged). CNS disease (blasts present on cytospin examination of cerebrospinal fluid in which the WBC count was at least  $0.005 \times 10^9/L$  [5 cells/ $\mu L$ ] or clinical signs of CNS leukemia) was present in 16.4%. There were 32.2% with a presenting WBC count under  $50 \times 10^9/L$ , 30.4% between  $50 \times 10^9/L$  and  $199 \times 10^9/L$ , and 37.4% had a WBC count of  $200 \times 10^9/L$  or higher. *MLL*/11q23 abnormality was present in 68.7%, and 71.3% of the infants had CD10<sup>-</sup> blasts.

### Molecular and cytogenetic testing for *MLL*/11q23 rearrangement

Seventy-nine (68.7%) of 115 cases showed *MLL* gene rearrangement by Southern blot, RT-PCR, and/or cytogenetics (Table 4). There were 35 identified by cytogenetics and/or RT-PCR as having t(4;11)(q21;q23), 20 found to have t(11;19)(q23;p13), and 9 found to have t(9;11)(p22;q23). Fifteen cases were *MLL* rearranged but with 11q23 rearrangements other than these 3. This "other" 11q23 translocation partner was occult (*MLL* detected by Southern blot but not detected cytogenetically) in 9 patients. These 9 were *MLL* rearranged by Southern blot, with cytogenetics reported as either normal (4), inadequate (3), or with another karyotypic abnormality [one del(11)(p11.2p13) and one t(10;11)(p11;q13)]. The remaining 6 "other" *MLL* abnormalities/translocation partners were as follows: t(1,11)(p32;q23); t(4;1)(1;q35;p13p32;q23); der(11)t(11;12)(q23;q15); cytogenetic del(11)(q23q25) and der(19)ins(19;

**Table 3. Presenting features of infants with ALL treated on CCG 1953 and 1883**

Variable and category*	CCG 1953, no. of patients (%)	CCG 1883, no. of patients (%)	P
<b>Age</b>			.224
Less than 3 mo	24 (20.9)	21 (15.6)	
3 to less than 6 mo	37 (32.2)	36 (26.7)	
At least 6 mo	54 (47.0)	78 (57.8)	
<b>Race</b>			.002
White	70 (62.5)	96 (71.1)	
Hispanic	34 (30.4)	22 (16.3)	
Black	0 (0.0)	10 (7.4)	
Other/unknown	8 (7.1)	7 (5.2)	
<b>Sex</b>			.94
Male	54 (47.0)	64 (47.4)	
Female	61 (53.0)	71 (52.6)	
<b>WBC count</b>			.58
Less than $50 \times 10^9/L$	37 (32.2)	52 (38.5)	
$50 \times 10^9/L$			
to $199 \times 10^9/L$	35 (30.4)	37 (27.4)	
At least $200 \times 10^9/L$	43 (37.4)	46 (34.1)	
<b>Liver</b>			.91
Normal	25 (22.9)	33 (24.4)	
Moderately enlarged	70 (64.2)	83 (61.5)	
Markedly enlarged	14 (12.8)	19 (14.1)	
<b>Spleen</b>			.66
Normal	22 (19.1)	31 (23.0)	
Moderately enlarged	65 (56.5)	69 (51.1)	
Markedly enlarged	28 (24.4)	35 (25.9)	
<b>CNS disease</b>			.18
Yes, CNS-3	18 (16.4)	34 (25.4)	
Maybe, CNS-2	18 (16.4)	24 (17.9)	
No, CNS-1	74 (63.7)	76 (56.7)	
<b>CD10</b>			.04
CD10 <sup>-</sup>	82 (71.3)	56 (57.7)	
CD10 <sup>+</sup>	33 (28.7)	41 (42.3)	
<b><i>MLL</i>/11q23 abnormality</b>			.04
Present	79 (68.7)	35 (53.0)†	
Absent	36 (31.3)	31 (47.0)	

For the CCG 1953 study, N = 115; for the CCG 1883 study, N = 135.

\*For some variables, some data were not submitted for individual patients (liver size, race, CNS, CD10), so totals do not add up to 115.

†11q23 info for 1883 is based solely on karyotypes accepted by central review. Information was not complete on all 135 patients. Percentages were calculated excluding patients for whom data were unknown.

11)(p13.1;q13q23) with *MLL* rearrangement by Southern blot; add(11q23); and t(10;11)(p13;q23).

Cases categorized as "BMT ineligible" were those with no *MLL* rearrangement or 11q23 abnormality identified by molecular and cytogenetic testing. In 12 cases, testing did not include the entire triad of cytogenetics, Southern blot, and RT-PCR. In these 12, the sample was inadequate for Southern blot testing, and cytogenetics were either inadequate (2) or normal (10). In 10 of these 12 cases, RT-PCR was completed and was negative for the 3 11q23 translocations for which testing was done.

**Table 4. *MLL* gene rearrangement and 5-year EFS**

<i>MLL</i> /11q23 abnormality	CCG 1953	
	No. of patients	5-year EFS, %
t(4;11)(q21;q23)	35	29.0
t(11;19)(q23;p13)	20	30.0
t(9;11)(p22;q23)	9	22.2
Other 11q23	15	53.3
Nonrearranged	36	60.3

For the CCG 1953 study, N = 115.

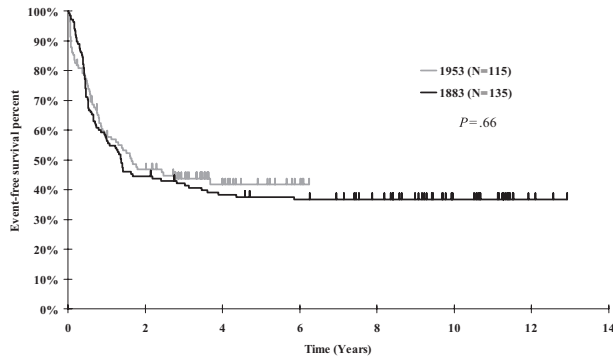


Figure 1. Kaplan-Meier estimates of event-free survival for infant ALL patients treated on CCG 1953 and CCG 1883.

### Treatment outcomes

**Remission induction.** Induction was defined as the length of time it took to get through that phase, just before day 35. Data to assess remission induction were available for 103 of 115 patients entered. There were 10 patients who died prior to day 35. There were 85 patients who were M1 (< 5% blasts) at end of induction, although 3 of these died of complications incurred during induction. There were 3 patients with M2 (5%-25% blasts) marrows at the end of induction, and one with M3 (> 25% blasts). The complete remission (CR) rate was 82.5% (85/103). The remission induction rate for the historic control was 94%.

There were 12 patients who were not evaluable for remission induction because a marrow aspiration was not done on day 35. Ten of these 12 patients had M1 marrows on either day 7 or day 14 or both. Of these 10, 8 patients recovered counts and received further study therapy. One patient received 16 days of systemic therapy and was taken off study and subsequently died of disseminated fungal disease with evidence of peripheral blasts 71 days from study entry. One patient died 54 days from study entry of pulmonary actinomycosis and mucormycosis with an empty marrow at autopsy. One patient with an M2 marrow aspirate on day 7 was removed from study on day 19 of induction; a marrow aspirate was done on 25 days after study enrollment and this marrow aspirate documented a remission. One patient had no evaluable marrow aspirates done during induction; this patient recovered counts and received further study therapy.

**First remission transplantation.** Patients with *MLL* rearrangement identified by central review were assigned to BMT if a related or unrelated matched or 1-antigen-mismatched donor was identified, and transplantation could be scheduled by 4 months of study entry.

There were 79 patients with *MLL* rearrangements and 37 of these received a transplant in first remission. One patient without central identification of *MLL* rearrangement underwent transplantation in first remission. Compliance with the protocol-specified BMT regimen was poor. Sixteen patients were treated according to the protocol, and there is a group of 22 *MLL*-rearranged chemotherapy patients that made it out to the median time to transplantation (125 days) and serve as the "control" group for this comparison. In comparing the EFS outcome subsequent to transplantation for the group that received *MLL* transplants per protocol to the post-125-day outcome in the *MLL* control, the 4-year EFS is 58.7% (SE = 12.6%) for the chemotherapy control and 25.0% (SE = 10.8%) for the BMT group, with a log rank *P* value of .006. Nineteen patients received transplants with non-protocol-specified preparative regimens (to be reported in detail in a separate BMT

manuscript [P.A.D., J.M.H., H.S., D.V., N.A.H., R.M., F.O.S., W.G.W., W.L.S., H.S.J., Z.D., G.H.R., manuscript in preparation]).

**Study outcome and comparisons to the historic control.** EFS for the 115 infants was 41.7% (SD = 9.2%) and OS was 44.8% (SD = 5.6%) at 5 years. EFS for CCG 1883 at 5 years was 37.6% (Figure 1; *P* = .66, RHR = 1.07 for CCG 1883 vs CCG 1953). Comparing EFS for the infants younger than 90 days of age, the current study had a much better EFS at 5 years of 41.7% versus 9.5% for CCG 1883 (*P* = .06, RHR = 1.89 for CCG 1883). Outcomes within CCG 1953 were examined for cohort 1, who received the original daunomycin dosing, versus cohort 2, who received a modified daunomycin dosing (including those treated after both the first and second modification). EFS outcome was not significantly improved for cohort 2 versus cohort 1 at 5 years, 42.3% versus 36.0%, respectively (*P* = .37, RHR = 1.30 for cohort 1). Reasons for failure were as follows: cohort 1, relapse in 7 (28%) of 25 and toxic death in 9 (36%) of 25; cohort 2, relapse in 17 (19%) of 90 and toxic death in 31 (34%) of 90. The patterns of failure were not statistically significantly different (*P* = .52, chi square).

For those alive at the end of induction, 5-year DFS was 49.2% (38.5% for those infants in CCG 1883, *P* = .09). The difference in DFS was especially dramatic for infants younger than 90 days of age at diagnosis. The 5-year DFS for infants younger than 90 days of age at diagnosis on CCG 1953 was 56.3% and on CCG 1883 was 11.1% (*P* = .009). There was no difference for the infants older than 90 days of age at diagnosis, with 5-year DFS for infants in remission at end induction of 43.0% on CCG 1953 and 47.6% on CCG 1883 (*P* = .40).

**Event-free survival by individual prognostic factors.** The 5-year EFS for *MLL*-rearranged cases was 33.6% and for *MLL*-nonrearranged cases was 60.3% (Figure 2; *P* = .006, RHR = 2.29 for *MLL* rearranged). The difference in EFS between the different *MLL* rearrangements using a global log rank test (used to compare all 4 groups simultaneously) did not reach a conventional statistical significance level (Figure 3; *P* = .12). The 5-year EFS for t(4;11) was 29.0%, t(11;19) 30.0%, t(9;11) 22.2%, and other 11q23 53.3%. Outcomes for the t(4;11), t(11;19), and t(9;11) groups were similar (*P* = .77). When the remaining *MLL*/11q23 partners ("other 11q23") were compared with the combination of the t(4;11), t(11;19), and t(9;11) *MLL* groups, there was a better outcome for the "other 11q23" group (*P* = .03).

Outcomes by age and WBC count are presented with CD10 and *MLL* data in Table 5. Outcome by CD10 expression was very similar to the *MLL* data. Five-year EFS for CD10<sup>-</sup> infants was 31.4% and for CD10<sup>+</sup> infants was 66.7% (*P* < .001). Examining age, the 5-year EFS for infants 3 months of age or younger was

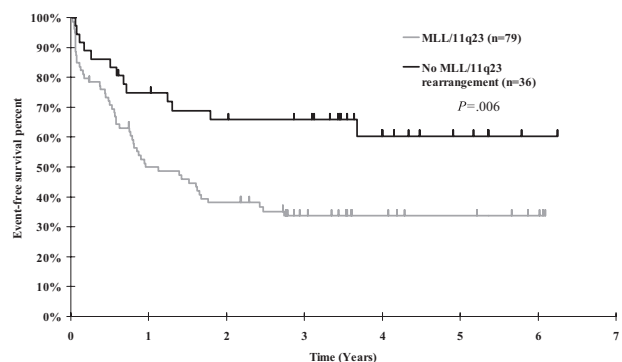
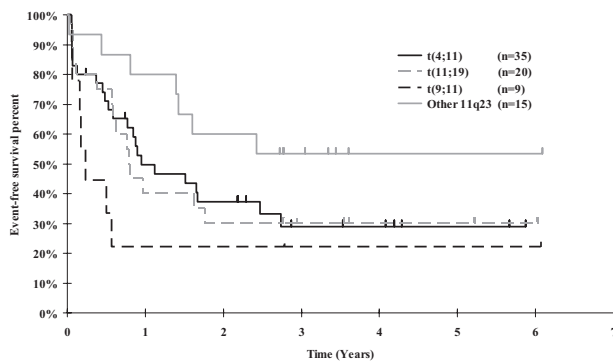


Figure 2. Kaplan-Meier estimates of event-free survival for CCG 1953 patients by *MLL* status.



**Figure 3.** Kaplan-Meier estimates of event-free survival for CCG 1953 patients in *MLL*-rearranged subgroups. Comparison of t(4;11), t(11;19), and t(9;11) groups,  $P = .77$ . Comparison of "other 11q23" group versus combination of t(4;11), t(11;19), and t(9;11) groups,  $P = .03$ .

41.7% but for those 3 to less than 6 months was 23.4% and for those 6 months or older was 53.9% ( $P = .01$ ). Five-year EFS by WBC count was as follows: 54.0% for infants with WBC count less than  $50 \times 10^9/L$ , 37.2% for WBC count  $50 \times 10^9/L$  to  $199 \times 10^9/L$ , and 31.1% for WBC count of at least  $200 \times 10^9/L$  (global  $P = .19$ ;  $P = .09$  for ordered trend).

**Early marrow response.** Most patients had a rapid early response (RER) status by day 7 (82 [82%] of 101 with results available compared with 60% in the historic control). The effect of early marrow response on outcome was examined separately for *MLL*/11q23-rearranged versus -nonrearranged patients who were in remission at the end of induction. For the infants with *MLL*/11q23 rearrangement, rapid early responders ( $n = 32$ ) had a 5-year DFS of 42.9%; the DFS of intermediate responders (M2/3 day 7, M1 day 14) was 44%. There were only 2 slow responders, both of whom had very early disease events at 7 and 9 months. Among the *MLL*/11q23-nonrearranged group, rapid early responders had a 5-year DFS of 65.8%, the intermediate responders had 50% DFS ( $P = .4$ ), and there were no slow responders.

Multivariate Cox regression analyses were used to determine the relative prognostic influence of *MLL*/11q23 status, CD10 expression, age at diagnosis, and WBC count. In a univariate Cox regression for each of these factors individually, the rank order of significance would be (1) CD10 expression,  $P = .001$  (RHR of 3.10 for CD10<sup>-</sup> compared with CD10<sup>+</sup>); (2) age,  $P = .012$  (RHR of 1.99 for age 6 months and older versus age 0-5 months); (3) presence or absence of *MLL*/11q23 abnormality,  $P = .013$  (RHR of

2.25 for patients with *MLL*/11q23 present versus those with *MLL*/11q23 absent); and (4) WBC count,  $P = .18$  (RHR of 1.51 for WBC count  $\geq 50,000 \times 10^9/L$  versus WBC count  $< 50,000 \times 10^9/L$ ). Most patients that were CD10<sup>-</sup> also had *MLL*/11q23 present, and patients that were CD10<sup>+</sup> usually had *MLL*/11q23 absent. Thus, once one of these factors is included in the regression model, the second factor would have little statistical significance on outcome after adjustment for the first. Hence, with CD10 present in the regression model, *MLL*/11q23 did not retain sufficient additional prognostic effect to meet a typical significance criterion ( $P = .38$ ). Age had a marginally significant effect ( $P = .07$ ) after adjusting for CD10 expression, but WBC count failed to demonstrate a significant effect ( $P = .83$ ) after adjusting for CD10 expression. Thus, in this particular infant study, only CD10 expression maintained a strong prognostic effect in multivariate analysis ( $P = .004$ ), although age did also show some effect on outcome.

In a stratified life-table analysis, data were analyzed for outcomes within the 2 *MLL* groups (rearranged and nonrearranged) varying by WBC count, age, and CD10 expression. Outcomes were looked at only for those groups with reasonable numbers of patients. The group that was *MLL* rearranged had only CD10<sup>-</sup> infants in significant numbers and among those, WBC count and age did not significantly influence outcome. Among the group without a detected *MLL*/11q23 abnormality, CD10<sup>-</sup> infants had poor outcomes ( $n = 14$ ), with 5-year EFS of 37% compared with CD10<sup>+</sup> infants ( $n = 22$ ) having a 5-year EFS of 73.6%.

**Patterns of relapse and events.** First events on CCG 1953 were death in 40, isolated marrow relapse in 19, CNS relapse in 1, isolated testicular relapse in 2, concurrent marrow plus CNS relapse in 1, and concurrent marrow plus testicular relapse in 1. Among the 40 patients who experienced death as a first event, the median time to death was 137 days (approximately 20 weeks); range, 9 to 901 days. For the historic control CCG 1883, 8 (of 135) patients experienced death as a first event; the median time to death was 170 days (approximately 24 weeks); range, 3 to 1026 days. Among the 24 patients who experienced relapse as a first event in CCG 1953, the median time to relapse was 295 days (approximately 42 weeks); range, 63 to 1343 days. For CCG 1883, 74 patients experienced relapse as a first event; the median time to relapse was 207 days (approximately 30 weeks); range, 49 to 2138 days.

In *MLL*/11q23-rearranged patients with events, first event was relapse in 33% and death in 67%. For *MLL*/11q23-nonrearranged infants with events, first event was death in 46%, relapse in 46%,

**Table 5.** CCG 1953 and CCG 1883: five-year EFS according to prognostic factors

Variable	CCG 1953			CCG 1883		
	No. of patients	5-year EFS, %	<i>P</i>	No. of patients	5-year EFS, %	<i>P</i>
<b>Age</b>			.012			< .001
Less than 3 mo	24	41.7		21	9.5	
3 to less than 6 mo	37	23.4		36	27.8	
At least 6 mo	54	53.9		78	49.7	
<b>CD10</b>			< .001			.002
CD10 <sup>+</sup>	33	66.7		41	55.3	
CD10 <sup>-</sup>	82	31.4		56	28.6	
<b>WBC count</b>			.191			< .001
Less than $50 \times 10^9/L$	37	54.0		52	57.6	
$50 \times 10^9/L$ to $199 \times 10^9/L$	35	37.2		37	34.8	
At least $200 \times 10^9/L$	43	31.1		46	17.4	

For the CCG 1953 study,  $N = 115$ ; for the CCG 1883 study,  $N = 135$ .

and induction failure in 8% (1 patient). The median time to death was 110 days for *MLL*-rearranged babies and was 320 days for *MLL*-nonrearranged patients. The median time to relapse was 291 days. Forty-seven percent of relapses in *MLL*-rearranged patients had occurred by day 300 (median 319 days); all relapses in *MLL*-nonrearranged infants occurred by this day (median 230 days). (These figures are not available for the historic control, as *MLL* testing was not broadly performed.)

### Toxicities

The median number of hospital days through all phases was 87, with the majority during induction/intensification. With the modification of therapy, decreasing induction anthracycline from body surface area to weight dosing, a decrease in induction hospital days was not seen (median 48 days for age < 90 days and 49.5 days for age ≥ 90 days before the amendments compared with 59.5 days for age < 90 days and 49 days for age ≥ 90 days after the amendments). Grades 3 and 4 nonhematologic toxicities are shown in Table 6.

Overall, 495 infectious complications occurred among 109 patients. Bacterial infections occurred most frequently (74%), followed by viral (13%), fungal (11%), and protozoan (1%). The most common sites of infection were blood (57%), cutaneous (12%), stool (9%), and lower respiratory (8%). Infectious complica-

tions continued through all phases: induction/intensification (84%), reinduction (35%), reintensification (41%), consolidation (72%), intensified maintenance (32%), and maintenance (28%). Despite the high rate of infectious complications, infection-related mortality was greatly reduced after induction/intensification: induction/intensification (16.5%), reinduction (0.9%), reintensification (0%), consolidation (0.9%), intensified maintenance (1.7%), and maintenance (3.5%). Significantly, reinduction, which included both continuous infusion daunomycin and dexamethasone, was not associated with the infection-related mortality of induction/intensification. In addition, although infectious complications were high during consolidation (very high-dose methotrexate and high-dose cytosine arabinoside), infection-related mortality was acceptable.

There were 19 infection-related deaths (8 bacterial, 5 fungal, and 6 viral) during induction/intensification among the 115 patients (16.5%). There were 7 deaths in induction/intensification among 25 patients (28%) in cohort 1 and 12 deaths among 90 patients (13%) treated in cohort 2. Despite the decrease in mortality, infectious-related complications did not change significantly in numeric incidence after induction treatment modifications.

Twenty-nine percent (n = 6) of infants younger than 3 months of age at diagnosis died of infection-related causes during induction/intensification compared with 14% (n = 13) of infants 3 months of

**Table 6. Grades 3 and 4 nonhematologic toxicities by phase of therapy**

	Induction/ intensification, no. of patients	Reinduction, no. of patients	Reintensification, no. of patients	Consolidation, no. of patients	Intensified maintenance, no. of patient exposures	Maintenance, no. of patient exposures
<b>Total</b>	115	91	66	53	120	126
<b>Cardiac</b>						
Cardiac function	2	0	0	0	0	1
Hypertension	12	2	1	0	0	1
Hypotension	9	0	0	0	1	1
<b>Coagulation</b>						
Fibrinogen	8	2	0	0	0	1
PT	7	0	0	2	0	1
PTT	10	1	0	1	0	0
Hemorrhage	6	1	0	0	0	0
<b>Gastrointestinal</b>						
Stomatitis	54	3	10	0	1	4
Abdominal pain	5	1	0	0	0	3
Diarrhea	24	9	5	7	9	9
Nausea/vomiting	11	1	6	1	2	5
<b>Liver</b>						
AST	18	1	13	10	3	2
ALT	25	4	12	11	8	8
Bilirubin	18	2	8	6	1	2
<b>Nervous system</b>						
Peripheral	8	0	0	0	1	4
Central	5	1	3	0	3	5
<b>Pancreas</b>						
Glucose	17	2	2	0	1	2
<b>Pulmonary</b>						
PaO <sub>2</sub>	8	0	0	0	0	0
Functional	28	2	2	2	3	2
Clinical	16	0	0	0	1	1
<b>Renal</b>						
BUN	6	1	0	0	0	0
Creatinine clearance	2	1	1	0	0	0

PT indicates prothrombin time; PTT, partial thromboplastin time; AST, aspartate aminotransferase; ALT, alanine aminotransferase; PaO<sub>2</sub>, oxygen saturation; and BUN, blood urea nitrogen.

**Table 7. Infection-associated mortality during phase 1 by age at diagnosis**

	Age less than 3 mo		Age 3 to less than 6 mo		Age at least 6 mo	
	%	No. out of total	%	No. out of total	%	No. out of total
Before 11/97 amendment	50	2/4	30	3/10	18	2/11
After 11/97 amendment	0	0/4	11	3/28	11	5/45
After 7/98 amendment	31	4/13	NA	NA	NA	NA
Total	29	6/21	16	6/38	13	7/56

Phase 1 indicates induction/intensification; NA, not applicable.

age or older at diagnosis (Table 7). Corresponding with the daunomycin dosage amendments, a decrease in mortality was seen across all age groups. However, no clear correlation with rates of grades 3 and 4 mucositis or skin breakdown was seen (Table 8) nor was there a relationship to any other recorded grades 3 and 4 toxicities.

## Discussion

Poor outcomes for infants with ALL led to the development of protocols that simultaneously intensified therapy and attempted to define prognostic factors. In the current report, we present data on overall disease outcome and prognostic factor analysis for the largest series of infants in a single clinical trial studied at the molecular level for *MLL* gene rearrangement.

The CCG clinical trials preceding this one (CCG 107 and CCG 1883) reported 5-year EFS for infants of 33% and 38%, respectively.<sup>1</sup> The EFS for CCG 1953 was 41.6% at 5 years. Thus, while there are fewer relapses and relapse occurs later (207 days on CCH 1883 vs 295 days on CCG 1953), ultimate outcome has not improved to a level of statistical significance despite intensified therapy. With the remission induction rate inferior to the historic control due to early toxicity and the 5-year EFS higher, the benefit of intensified therapy is lost in early toxicity. The prognostic factor analysis must be taken in the context of the treatment failure patterns observed. In CCG 107 and 1883, the major cause of treatment failure was early marrow relapse.<sup>1</sup> CCG 107 and 1883 reported 2 (2.0%) and 5 (3.7%) deaths in induction, and 65% and 72% of relapses were within the first year after induction.<sup>1</sup>

The CCG 1953 protocol was thus developed to address early relapse and to test the hypothesis that intensification of early therapy would improve outcome for these patients. The induction chemotherapy of the protocol was modeled on the CCG 192P protocol, which used intensive multi-agent chemotherapy. Infants treated on this protocol had an EFS of 36% at 4 years, with minimal treatment-related morbidity.<sup>32</sup> Dosing of chemotherapy for infants in previous protocols was determined by weight not body surface area. Using weight rather than body surface area results in

significantly lower doses of drugs for these patients. Therefore, to intensify the protocol therapy, body surface area was used to calculate doses of all agents except vincristine. Dexamethasone at 10 mg/m<sup>2</sup> was chosen as the steroid during the first year of therapy, based on reports of improved outcome using this steroid and particularly better CNS disease control. Additional consideration in the development of the protocol was that *MLL*-rearranged ALL in infants may have a biology similar to *MLL*-rearranged acute myeloid leukemia (AML). High-dose cytosine arabinoside is an active agent in the consolidation of AML, including AML in infants with *MLL* rearrangements. This biology also contributed to the rationale for BMT for infants with *MLL* rearrangement. The CCG 107 and 1883 studies used very high-dose methotrexate as CNS prophylaxis and these studies demonstrated that this approach led to acceptable CNS prophylaxis with the toxicity of central nervous prophylaxis.

The early relapse pattern is reversed with the intensified therapy of CCG 1953, with 19 (16.5%) deaths related to induction, and less early relapse. The early toxicity seen in CCG 1953 resulted in amendments decreasing anthracycline intensity. The subsequent cohorts had decreased induction deaths but EFS only marginally increased. The concurrent POG 9407 trial was similarly amended, and remains open (COG P9407), and was further amended with the removal of dexamethasone in induction replaced by prednisone.

In CCG 107 and 1883, age, WBC count, French-American-British morphology, hepatosplenomegaly, CD10 expression, and t(4;11) status (as opposed to any 11q23 rearrangement) were all significant prognostic factors.<sup>1</sup> However, it must be emphasized that 11q23 status was determined strictly by cytogenetics and results were available to report on only 38 of 99 patients on CCG 107 and on 66 of 135 patients on CCG 1883<sup>1</sup> (additional cases reviewed since publication). Thus the impact of particular 11q23 partners on outcome was not determined for CCG 1883 (N.A.H., unpublished data, December 1988–August 1993).

For the CCG 1953 cohort, paired molecular and cytogenetic determination of *MLL*/11q23 status yielded the expected outcomes: the rate of *MLL* rearrangement (69%), that most of these are t(4;11), and that a significant number of t(11;19) are detected with this methodology. *MLL*/11q23 abnormality was an important

**Table 8. Grades 3 and 4 mucositis and skin breakdown during phase 1**

	Age less than 3 mo		Age 3 to less than 6 mo		Age at least 6 mo	
	%	No. out of total	%	No. out of total	%	No. out of total
<b>Mucositis</b>						
Before 11/97 amendment	50	2/4	30	3/10	27	3/11
After 11/97 amendment	50	2/4	NA	NA	NA	NA
After 7/98 amendment	31	4/13	57	16/28	5	24/45
<b>Skin breakdown</b>						
Before 11/97 amendment	50	2/4	10	1/10	9	1/11
After 11/97 amendment	25	1/4	43	12/28	31	14/45
After 7/98 amendment	69	9/13	NA	NA	NA	NA

Phase 1 indicates induction/intensification; NA, not applicable.



predictor of outcome; the 5-year EFS for *MLL*-rearranged infants was significantly less (33.6%) than that of *MLL*-nonrearranged cases (60.3%). With molecular testing completed on the great majority of cases, we showed that the individual *MLL* rearrangements t(4;11), t(11;19), and t(9;11) do not have different effects on prognosis. Specifically, t(4;11) did not have outcomes worse than t(11;19) and t(9;11). The group of infants with t(11;19) (n = 20) having outcomes equivalent to those with t(4;11) stands in contrast to data from the previous CCG trials<sup>12,19</sup> but is in agreement with data from Rubnitz et al.<sup>20</sup>

The outcomes of infants for whom *MLL* rearrangement was indeterminate but CD10 was negative were as poor as those of *MLL*-rearranged infants. This suggests that CD10<sup>-</sup> infants without sufficient material for molecular testing, who have inadequate or normal cytogenetics, either had undetected *MLL*/11q23 rearrangement or, as suggested by multivariate analysis, CD10 negativity stands alone as a negative prognostic factor. Such infants should be treated as *MLL*-rearranged infants in protocols using *MLL*/11q23 status to dictate therapy.

Age and CD10 expression remain individual adverse prognostic factors, but WBC count at diagnosis did not emerge as prognostically significant (Table 5). The 5-year EFS for infants younger than 90 days of age (n = 24; 41.7% in CCG 1953) is a significant improvement from the 9.5% 5-year EFS seen in that age group in CCG 1883 (n = 21). It may be that for those infants younger than 90 days of age surviving induction, the increase in intensity of CCG 1953 compared with CCG 1883 was sufficient to overcome late relapse.

CD10 expression and *MLL*/11q23 status correspond so closely in infants with ALL that it is difficult to assess their independent prognostic significance. In multivariate analysis, age had reduced prognostic significance after adjustment for either CD10 status or *MLL*/11q23 status with the close association between age under 6 months, CD10 negativity, and *MLL*/11q23 rearrangement making it difficult to discern the relative impact of these factors on disease outcome. Future clinical trials should continue to assess these factors.

Evaluation of the impact of early marrow response shows that while early response status is important, it does not override the impact of *MLL*/11q23 rearrangement. As was seen in CCG 1883,<sup>1</sup> the majority of infants were rapid early responders. *MLL*/11q23-rearranged early responders had an EFS of 46% compared with the overall 37% EFS of this group. Slow early response in *MLL*/11q23-rearranged infants was particularly ominous. Interestingly, rapid early marrow response was a better prognostic sign in CCG 1953 than CCG 1883, where there were still poor outcomes seen in *MLL*/11q23-rearranged infants even if the marrow was in early response. Specifically, infants in CCG 1883 with cytogenetically detected 11q23 rearrangements and rapid early response had a 5-year EFS of 29% (n = 24). Thus, for those infants surviving induction, the intensified CCG 1953 therapy improved outcomes for *MLL*/11q23-rearranged infants. While this reflects the fact that poor-risk patients were able to tolerate the less intense CCG 1883 induction and survive ultimately to relapse, it also points out that intensified therapy can change the natural history of *MLL* disease.

There was only one isolated CNS relapse as a first event in this group of infants (0.9%) and only 4 isolated CNS relapses (3%) in CCG 1883.<sup>1</sup> The CNS-directed therapy in these protocols was nearly identical and consisted of intrathecal chemotherapy (21 total doses in CCG 1953, 18-22 in CCG 1883), high-dose methotrexate (delivered 4 times in each protocol), high-dose Ara-C (8 doses in CCG 1953, 4 in CCG 1883), and L-asparaginase therapy (18 doses

each protocol) without CNS radiotherapy. It is intriguing to note the similar low CNS relapse rate in 2 protocols that have had such dissimilar toxicity/relapse patterns. The current open protocol for infant ALL in the COG P9407 study<sup>21</sup> includes these treatments but does not include “very high-dose” methotrexate (33.6 g/m<sup>2</sup>).

The intensity of therapy has resulted in regimen-related toxicity in a pattern that is predominantly infectious. In many cases the infections seen were the type of infection expected in immune incompetent patients, such as pneumocystis and viral infections. These were felt to be due to dexamethasone in induction. Consequently, the currently open successor protocol substitutes the high-dose induction dexamethasone with standard doses of prednisone to determine whether disease control will be maintained with less-prolonged immune suppression and less infection-related mortality. This remains an outcome to be examined in the ongoing study COG P9407.

With the design of future protocols for infant ALL will come the opportunity to evaluate innovative therapies. Traditionally such therapies are ethically reserved for those with EFS less than 50%.<sup>33</sup> *MLL*-rearranged patients on this study had a 5-year EFS of 34%, clearly challenging us to come up with new therapy plans. With CCG 1953 therapy, *MLL*-nonrearranged infants have an EFS of 60.3%. While not “excellent,” these patients too present us with the challenge of improving outcomes while decreasing toxicity. Stratification of infants using multiple prognostic factors may help to better select subsets of patients requiring more intensive therapy. At the same time, even if *MLL*/11q23-rearranged infants were to achieve outcomes better than an EFS of 50%, the prospect of less toxic therapy with innovative or targeted therapies challenges us to look beyond the traditional paradigms of therapy design. In light of this, there is a great need to increase the availability of preclinical data to guide the selection of new agents for infants with ALL.

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## Appendix

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*Note:* Institutions listed without a principal investigator may have served as treatment facilities collaborating with a principal investigator at a nearby institution or are no longer members of the COG.

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