

ARE THERE REGIONAL VARIATIONS IN THE PRESENTATION OF CHILDHOOD LEUKEMIA?

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ABSTRACT

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Introduction: Treatment of childhood acute lymphoblastic leukemia (ALL) is based on risk stratification. This study aimed to assess the agreement between risk group classifications in the different childhood ALL treatment protocols used in a referral hospital in southern Brazil.

Methods: We retrospectively reviewed the medical records of patients aged 1 to 18 years with B-cell ALL treated at a hospital from January 2013 to April 2017. Agreement between risk classifications was assessed by the kappa coefficient.

Results: Seventy-five patients were analyzed. There was poor agreement between risk stratification by GBTLI 2009 and BFM 95 protocols ($\kappa=0.22$; $p=0.003$) and by GBTLI 2009 and IC-BFM 2002 protocols ($\kappa=0.24$; $p=0.002$). Risk group distribution was 13.3% for low risk, 32.0% for intermediate risk, and 54.7% for high risk based on stratification by the GBTLI 2009 protocol, and 28.0% for low risk, 42.7% for intermediate risk, and 29.3% for high risk based on stratification by the IC-BFM 2002 protocol. Overall survival was 68.6%.

Conclusion: This study provides numerous points to ponder about the treatment of leukemia in Brazil. The percentage of patients classified as high risk in our sample was higher than that reported in the international literature. This difference, however, had no impact on overall survival, which was shorter than that reported in the international literature.

Keywords: Childhood ALL; Risk factors; Immunophenotyping; Minimal residual disease

INTRODUCTION

Leukemia accounts for 30% of all cancers diagnosed in children younger than 15 years, and 75% of these cases are diagnosed as acute lymphoblastic leukemia (ALL)¹. As in other childhood cancers, the survival of children and adolescents with ALL has improved significantly over time². The 5-year survival rate for ALL increased from 60% to approximately 90% for children aged < 15 years and from 28 to 75% for adolescents aged 15 to 19 years between 1975 and 2010 in the United States³.

Much of this improvement is attributable to the tailoring of treatment to the individual characteristics of each patient and of the disease and, more recently, to novel targeted therapies⁴. For decades, groups that study childhood ALL have used risk classification systems to allocate patients to treatment regimens based on their estimated risk of treatment failure. That is, children with favorable clinical and biological features receive less toxic therapy, while more aggressive therapy is reserved for patients who are at higher risk of relapse and less likely to survive in the long term⁵.

This stratification is possible using prognostic factors⁶. Early risk classification systems were based only on clinical factors, such as age and white blood cell count at diagnosis. Current classification systems include the molecular characteristics of leukemic cells at diagnosis and response to treatment. One of the methods used is the detection of minimal residual disease (MRD). Currently, MRD is the most important prognostic factor for

ALL in children⁷. Early response to chemotherapy, with rapid reduction of leukemic cells, especially at the end of remission induction therapy, is an important indicator of a more favorable outcome and a lower risk of relapse⁸.

In Brazil, most centers treating pediatric patients with ALL base their treatment on the Berlin-Frankfurt-Münster (BFM) group protocols or on the Brazilian Cooperative Group for the Treatment of Childhood ALL (Grupo Brasileiro de Tratamento da Leucemia Infantil, GBTLI) protocols. At the hospital, two different protocols were used between 2009 and 2017: treatment suggested by the GBTLI 2009 protocol and treatment regimen proposed by the BFM group.

The present study aimed to describe the clinical and laboratory profile, as well as the outcome of patients with B-cell ALL (B-ALL) treated, and to assess the agreement between risk group classifications in patients classified according to the different childhood ALL treatment protocols or regimens used in the institution.

METHODS

Study population

We conducted a descriptive cross-sectional epidemiological study with retrospective data collection of 75 patients aged 1 to 18 years with B-ALL treated from January 2013 to April 2017. All patients were treated in the same hospital during the study period.

The patients' medical records were reviewed for the following data at the time of diagnosis and after initiation of treatment: age; leukocyte count; central nervous system (CNS) involvement; morphology of leukemic cells; immunophenotyping; cytogenetics; molecular biology; MRD by flow cytometry (FCM) and morphology at day 15 (D15) of induction; MRD by FCM and morphology at the end of induction (D33/D35); risk stratification based on the National Cancer Institute (NCI) criteria and on the criteria of the BFM 95, Intercontinental (IC)-BFM 2002, IC-BFM 2009, GBTLI 99, and GBTLI 2009 treatment regimens/protocols (Table 1)⁹⁻¹⁵; occurrence of relapse or death; current patient status; and date of last update.

Table 1: Risk classification according to the NCI criteria and to the BFM and GBTLI treatment regimens.

NCI criteria and treatment regimens	Risk groups		
	LR	IR	HR
NCI	– Leukocytes < 50,000/mm ³ at diagnosis and age > 1 year and < 10 years		– Leukocytes > 50,000/mm ³ at diagnosis and/or age ≥ 10 years
GBTLI 99	– Age > 1 year and < 9 years; – Leukocytes < 50,000/mm ³ at diagnosis and < 5,000/mm ³ in PB at D8; – Absence of blasts in PB at D14 and M1/M2 BM at D14; – M1 BM at D28; – Patients classified as LR who during treatment meet the requirements for inclusion in the HR group will change groups.		– Age < 1 year and ≥ 1 year and fied as LR who during 3 at diagnosis; – Slow response to treatment (leukocytes ≥ low resp3 at D7 of induction); – Presence of blasts in PB at D14; – M3 BM at D14 and/or M2/M3 BM at D28 and/or evidence of extramedullary leukemic involvement.
GBTLI 2009	– Age ≥ 1 year at 2009 D14 and/or – Leukocytes < 50,000/mm ³ ; – Negative CNS; – No poor-prognosis cytogenetics (BCR-ABL, MLL rearrangement, and hypodiploidy < 46 chromosomes). True low risk: – PB at D8 < 1,000 blasts/μL; – BM at D15 = M1 with MRD-FCM < 0.01% and at D35 = M1 with MRD-PCR < 10 ⁻³ .	PB at D8 < 1,000 blasts/μL; – BM at D15 = M1/M2 with MRD-FCM (0.01 < 10%); – D35 = M1 with MRD-PCR < 10 ⁻³ .	All of the previous criteria. Rapid responders: – D8 < 1,000 blasts/μL; – BM at D15 = M1/M2 with MRD-FCM < 10%; at D35 = M1 with MRD-PCR < 10 ⁻³ . Slow responders: – D8 > 1,000 blasts/μL; – BM at D15 = M2/M3 with MRD-FCM ≥ BM – D35 = M2/M3 with MRD-PCR ≥ 10 ⁻³ .

Table 1: Continuação

NCI criteria and treatment regimens	Risk groups		
	LR	IR	HR
BFM 95	<ul style="list-style-type: none"> – No criteria for high risk; – Leukocytes < 20,000/μL; – Age at diagnosis between 1 and 6 years; 	<ul style="list-style-type: none"> – No criteria for high risk; – And leukocytes < 20,000/μL; – And/or age at diagnosis < 1 year or \geq 6 years 	<ul style="list-style-type: none"> – Poor response to corticosteroids; – And/or no CR at D33 of induction; – And/or evidence of t(9;22) or BCR-ABL; – And/or evidence of t(4;11) or <i>MLL-AF4</i> fusion gene.
IC-BFM 2002	<ul style="list-style-type: none"> – PB at D8 of induction < 1,000 blasts/μL; – And age at diagnosis on < 1,000 blasts/μL; D – Leukocytes at diagnosis < 20,000/μL; – And BM at D15 = M1 or M2; – And BM at D33 = M1. 	<ul style="list-style-type: none"> – PB at D8 of induction < 1,000 blasts/μL; – And age at diagnosis < 1 year or \geq 6 years and/or leukocytes at diagnosis \geq 20,000/μL; – BM at D15 = M1 or M2; – And BM at D33 = M1 or standard risk criteria, but BM at D15 = M3; – BM at D33 = M1. 	<ul style="list-style-type: none"> – At least one of the following criteria: – IR and BM at D15 = M3 (non-standard risk and M3 at D15); – Blasts in PB at D8 \geq Blasts in PB at – BM at D33 = M2 or M3; – Presence of t(9;22) or t(4;11) [<i>MLL-AF4</i>].
IC-BFM 2009	<ul style="list-style-type: none"> – PB at D8: < 1,000 blasts/μL; – And age \geq 1 year and < 6 years; – And initial leukocytes < 20,000/μL; – And if available: MRD-FCM < 0.1% or M1/M2 BM at D15; – And M1 BM at D33. – All criteria must be met. 	<ul style="list-style-type: none"> – All patients who are not stratified as LR or HR. 	<ul style="list-style-type: none"> – IR and, if available: MRD-FCM 10% or M3 BM at D15; – LR and, if available: MRD-FCM > 10%; – PB at D8 \geq 1,000 blasts/μL; – M2 or M3 BM at D33; – Translocation t(9;22) [BCR-ABL] or t(4;11) [<i>MLL-AF4</i>]; – Hypodiploidy \leq ypodiploidy on – Any one of these criteria is sufficient to classify as HR.

BFM: Berlin-Frankfurt-Münster group; BM: bone marrow; CNS: central nervous system; CR: complete remission; D8: day 8 of induction; D15: day 15 of induction; D33: day 33 of induction; FCM: flow cytometry; GBTLI: Brazilian Cooperative Group for the Treatment of Childhood Acute Lymphoblastic Leukemia (Grupo Brasileiro de Tratamento da Leucemia Infantil); HR: high risk; IC: Intercontinental; IR: intermediate risk; LR: low risk; M1: BM blasts < 5%; M2: BM blasts 5 to < 25%; M3: BM blasts \geq 25%; MRD: minimal residual disease; NCI: National Cancer Institute; PB: peripheral blood; PCR: polymerase chain reaction.

Adapted from Smith et al.; Moricke et al.; Stary et al.; Schrappe et al.; Brandalise et al.; Scrideli et al.; Brandalise et al.⁸⁻¹⁴.

Exclusion criteria were patients with missing data for risk stratification or those with non-B-cell ALL.

The study was approved by the Research Ethics Committee (approval number: CAAE 55303916.5.0000.5327).

Data structure and statistical analysis

Sample size was calculated using WinPepi (v11.43). Based on an expected kappa value of 0.8, a maximum range of kappa of 0.28, a significance level of 5%, and a prevalence rate of 47%¹⁶, a sample size of 74 patients was required.

Data were entered into an Excel spreadsheet, version 2016, and then exported to SPSS, version 20.0, for statistical analysis. Data were expressed as absolute or relative frequencies and means, medians, or percentiles. Survival curves were

estimated by the Kaplan-Meier method and compared by the log-rank test. Overall survival was measured from the date of the diagnosis of ALL to the date of death or last contact. Event-free survival was measured from the date of the diagnosis of ALL to the date of relapse, refractoriness to treatment, death, or last contact. Refractoriness was defined as the failure to achieve a complete response after regular chemotherapy.

The risk classifications of each protocol were analyzed for agreement using the kappa coefficient, and the results were interpreted according to the different ranges of kappa values suggested by Landis & Koch¹⁷. Kappa values >0.75 indicate excellent agreement, values <0.40 indicate poor agreement, and values between 0.40 and 0.75 indicate fair to good agreement. To calculate the kappa coefficient,

we excluded the classifications based solely on the NCI criteria and the classifications based on the GBTLI 99 protocol criteria.

All data were updated to October 2017, and a p-value <0.05 was considered significant.

RESULTS

A total of 75 patients aged 1–18 years with a diagnosis of B-ALL were analyzed. Of these, 57.3% were male. Median age was 5 years; 25% of children were younger than 3 years, while 25% were older than 10 years. The main clinical and laboratory characteristics at diagnosis are summarized in Table 2.

Table 2: Clinical and laboratory characteristics and risk group distribution (N = 75).

Characteristics	n	%
Age		
– Median, years (25th/75th percentile)	5	
– Minimum/maximum	(3/10) 1/17	
Male	43	57.3
Initial leukocyte count (/mm ³)		
< 20,000	39	52
20,000 and 50,000	18	24
> 50,000	18	24
CNS		
CNS 1	66	88
CNS 2	4	5.3
CNS 3	5	6.7
ALL subtypes		
Pro-B ALL	3	4
Common-B ALL	70	93.3
Pre-B ALL	2	2.7
Karyotype		
Normal	28	
No metaphases growth	12	
Hyperdiploidy	10	
High hyperdiploidy	3	
Low hypodiploidy	1	
Almost diploid	3	
t(1;19)(q23;p13)	2	
t(4;11)(q21;q23)	1	
t(9;22)(q34;q11)	8	
Trisomy 21	2	
Other changes:	3	
46, XY, – 1, + mar [15]		
46, XX, + 1der(1;16)(q10p10)[20]		
46, XX, dup(1)(q21q32)[26]		
Data not available	2	

Table 2: Continuação

Characteristics	n	%
Molecular biology		
<i>IKZF1</i> deletions	5	
<i>TCF3-PBX1</i>	1	
<i>MLL</i> rearrangement	1	
<i>ETV6-RUNX1</i>	1	
No molecular changes	45	
No data available	22	
Relapse	16	
BM	9	
Testis	2	21.3
CNS	2	
BM + CNS	2	
BM + lymph nodes	1	
Deaths	17	
Disease	9	22.6
In CR after CT	7	
In CR after BMT	1	
NCI criteria		
LR	44	58.7
HR	31	41.3
GBTLI 99 classification		
LR	37	49.3
HR	38	50.7
GBTLI 2009 classification		
LR	10	13.3
IR	24	32
HR	41	54.7
BFM 95 classification		
LR	23	30.7
IR	30	40
HR	22	29.3
IC-BFM 2002 classification		
LR	21	28
IR	32	42.7
HR	22	29.3
IC-BFM 2009 classification		
LR	8	10.7
IR	41	54.7
HR	26	34.7

ALL: acute lymphoblastic leukemia; BFM: Berlin-Frankfurt-Münster group; BM: bone marrow; BMT: bone marrow transplantation; CNS: central nervous system; CR: complete remission; CT: chemotherapy; GBTLI: Brazilian Cooperative Group for the Treatment of Childhood Acute Lymphoblastic Leukemia (Grupo Brasileiro de Tratamento da Leucemia Infantil); HR: high risk; IC: Intercontinental; IR: intermediate risk; LR: low risk; NCI: National Cancer Institute.

Regarding karyotype, 73 (97.3%) patients had available information. Of these, 28 (38.4%) had a normal karyotype and 33 (45.2%) had an altered karyotype; in 12 (16.4%) patients, no growth was observed (no metaphases).

Of the 75 patients, 53 (70.6%) had access to immunomolecular analysis. In these cases, there were five *IKZF1* deletions, one *TCF3-PBX1* fusion, one *MLL* rearrangement, and one *ETV6-RUNX1* fusion.

Sixteen (21.3%) patients relapsed, and 17 (22.6%) patients died.

Risk classifications were compared between protocols, and agreement was assessed by the kappa coefficient. All kappa values were statistically significant.

There was poor agreement between GBTLI 2009 and BFM 95 protocols (kappa = 0.22; p = 0.003), and between GBTLI 2009 and IC-BFM 2002 protocols (kappa = 0.24; p = 0.002). The level of agreement was fair between GBTLI 2009 and IC-BFM 2009 protocols (kappa = 0.44; p < 0.001) (Table 3).

Table 3: Agreement analysis of risk classifications between GBTLI 2009 and BFM 95; GBTLI 2009 and IC-BFM 2002; GBTLI 2009 and IC-BFM 2009.

	GBTLI 2009 classification			Total	
	LR	IR	HR		
BFM 95 classification	LR	7	13	3	23
	IR	3	9	18	30
	HR	0	2	20	22
Total		10	24	41	75
IC-BFM 2002 classification	LR	7	12	2	21
	IR	3	10	19	32
	HR	0	2	20	22
Total		10	24	41	75
IC-BFM 2009 classification	LR	6	2	0	8
	IR	4	20	18	42
	HR	0	2	23	25
Total		10	24	41	75

BFM: Berlin-Frankfurt-Münster group; GBTLI: Brazilian Cooperative Group for the Treatment of Childhood Acute Lymphoblastic Leukemia (Grupo Brasileiro de Tratamento da Leucemia Infantil); HR: high risk; IC: Intercontinental; IR: intermediate risk; LR: low risk.

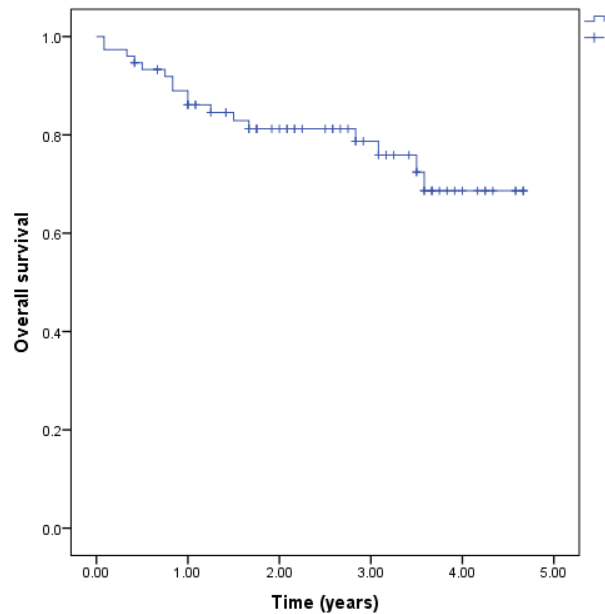
There was better agreement between risk stratification by IC-BFM 2009 and BFM 95 protocols, and between risk stratification by IC-BFM 2009 and IC-BFM 2002 protocols. The kappa coefficient was 0.67 (p < 0.001) and 0.68 (p < 0.001), respectively.

There was an association between risk group and death in cases stratified as high risk, regardless of the treatment protocol used. The percentage of deaths in patients classified as high risk according to the risk classifications of GBTLI 99, GBTLI 2009, BFM 95, IC-BFM 2002, and IC-BFM 2009 protocols was 39.5% (p = 0.003; chi-square test), 36.6% (p = 0.013; chi-square test), 54.5% (p < 0.001; chi-square test), 54.5% (p < 0.001; chi-square test), and 48.0% (p = 0.001; chi-square test), respectively. There was also a significant association between high

risk and relapse: GBTLI 2009, 29.3% (p = 0.03; chi-square test); and IC-BFM 2002, 45.5% (p = 0.003; chi-square test).

The 4-year overall survival was 68.6%, regardless of the treatment protocol used. Mean overall survival was 45.2 months (standard error, 2.2 months). The 4-year event-free survival was 66.6%. Mean event-free survival was 44.7 months (standard error, 2.2 months) (Figure 1).

A



B

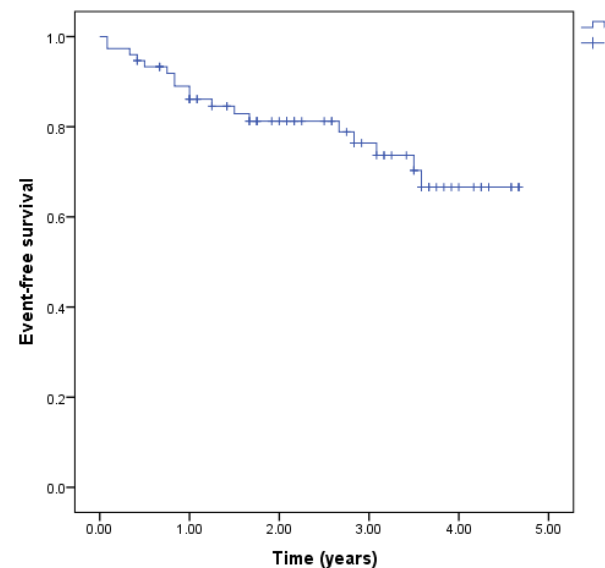


Figure 1: A: Estimated overall survival and; B: estimated event-free survival of the 75 patients, regardless of protocol.

DISCUSSION

Our results show that, as expected, there was better agreement between risk classifications of the BFM group, achieving substantial agreement, or nearly excellent agreement, according to the interpretation proposed by Landis & Koch¹⁷. When risk stratification was compared between BFM and GBTLI protocols, the main differences were age and leukocyte count at diagnosis.

Age ≥ 9 years, leukocyte count $\geq 50,000/\text{mm}^3$, and CNS involvement at diagnosis are considered high-risk features by the Brazilian protocols, regardless of the response to induction therapy¹⁵. In the BFM protocols, however, patients aged ≥ 6 years or with leukocyte count $\geq 20,000/\text{mm}^3$ are initially classified as intermediate risk in the absence of high-risk cytogenetic and molecular abnormalities, and these patients may be transferred to the high-risk group, depending on their response to induction therapy. CNS involvement does not change the initial risk stratification¹¹. This may explain why, as demonstrated in the present study, the Brazilian protocols tend to have a higher percentage of patients classified as high risk (Table 2).

Agreement was only moderate between risk stratification by the GBTLI 2009 and IC-BFM 2009 protocols, although both protocols have already incorporated the concept of MRD into their risk classification systems. A possible explanation is that MRD is assessed somewhat differently in these protocols. While IC-BFM 2009 considers MRD only at D15 of induction by FCM, GBTLI 2009 considers MRD at D15 by FCM and MRD at D35 by polymerase chain reaction (PCR)^{12,15}. However, in the present study, MRD was assessed only by FCM in all patients, because the hospital does not have appropriate technology for detecting MRD by PCR.

When the risk group distribution of our patients, based on the GBTLI 2009, IC-BFM 2002 and IC-BFM 2009 protocols, was compared with the risk group distribution published in the literature, we found a higher percentage of high-risk patients (54.7%, 29.3%, and 34.7%, respectively, vs. 16.7% in the literature). Furthermore, a lower percentage of low-risk patients was observed using the risk classifications of the GBTLI 2009 and IC-BFM 2009 protocols (13.3% and 10.7%, respectively, vs. 30.9% in the literature)^{10,11,18}.

The 4-year overall and event-free survival rates were 68.6% and 66.6%, respectively. These rates are lower than recent international rates from clinical trials conducted in North America and Western Europe, which have reported event-free survival

rates of up to 85% and overall survival rates of up to 90%¹⁹. Some studies have even reported survival rates above 90% for some specific subtypes treated with risk-adapted therapy in B-ALL²⁰.

In sum, the analysis of the present data showed that there was a difference in risk stratification depending on the treatment protocol or regimen used in our institution. This difference, however, had no impact on the overall survival of affected children.

This study provides numerous points to ponder about the treatment of leukemia in our context. The percentage of patients classified as high risk in our sample was higher than that reported in the international literature. In addition, although the patients were treated in a referral hospital in southern Brazil, which has specialized cancer treatment teams and appropriate resources to ensure supportive care for patients, survival rates were lower than those reported in the literature. However, we must consider that this center receives patients with serious clinical presentation and worse prognosis more frequently, due to the high specialized team. This may also have contributed to the finding of a greater number of patients classified as high risk. These data may also be related to the lower survival rate.

Furthermore, the present study raises some questions for future research: would study findings result from a difference in the Brazilian pediatric population with ALL? Are there regional variations in the presentation of childhood leukemia in Brazil? Is there a genetic or molecular alteration more prevalent among Brazilians that should be better studied in a larger sample of patients?

Therefore, further prospective laboratory and clinical studies are needed to explore the reasons underlying these differences, and whether they are sensitive to population or sample size, and to investigate the impact of risk stratification on toxicity and cure rates in our.

New approaches should probably improve diagnosis, prognosis and precision medicine in our population. For example, a study of genomic sequencing would allow the identification of subtypes and genetic alterations with prognostic importance and of their potential role in risk stratification²¹. These questions need to be better explored in our context in larger multicenter studies.

Data statement

All relevant data are within the paper and its Supporting Information files.

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