

Excellent prognosis of late relapses of ETV6/RUNX1-positive childhood acute lymphoblastic leukemia: lessons from the FRALLE 93 protocol.

Virginie Gandemer, Sylvie Chevret, Arnaud Petit, Christiane Vermylen, Thierry Leblanc, Gerard Michel, Claudine Schmitt, Odile Lejars, Pascale Schneider, Francois Demeocq, et al.

▶ To cite this version:

Virginie Gandemer, Sylvie Chevret, Arnaud Petit, Christiane Vermylen, Thierry Leblanc, et al.. Excellent prognosis of late relapses of ETV6/RUNX1-positive childhood acute lymphoblastic leukemia: lessons from the FRALLE 93 protocol.. Haematologica, 2012, 97 (11), pp.1743-1750. 10.3324/haematol.2011.059584 . inserm-00700077

HAL Id: inserm-00700077 https://inserm.hal.science/inserm-00700077

Submitted on 22 May 2012 $\,$

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



Early Release Paper

Excellent prognosis of late relapses of ETV6/RUNX1-positive childhood acute lymphoblastic leukemia: lessons from the FRALLE 93 protocol

by Virginie Gandemer, Sylvie Chevret, Arnaud Petit, Christiane Vermylen, Thierry Leblanc, Gerard Michel, Claudine Schmitt, Odile Lejars, Pascale Schneider, Francois Demeocq, Brigitte Bader-Meunier, Francoise Bernaudin, Yves Perel, Marie-Francoise Auclerc, Jean-Michel Cayuela, Guy Leverger, and Andre Baruchel

Haematologica 2012 [Epub ahead of print]

Citation: Gandemer V, Chevret S, Petit A, Vermylen C, Leblanc T, Michel G, Schmitt C, Lejars O, Schneider P, Demeocq F, Bader-Meunier B, Bernaudin F, Perel Y, Auclerc MF, Cayuela JM, Leverger G, and Baruchel A. Excellent prognosis of late relapses of ETV6/RUNX1-positive childhood acute lymphoblastic leukemia: lessons from the FRALLE 93 protocol. Haematologica. 2012; 97:xxx doi:10.3324/haematol.2011.059584

Publisher's Disclaimer.

E-publishing ahead of print is increasingly important for the rapid dissemination of science. Haematologica is, therefore, *E-publishing PDF files of an early version of manuscripts that* have completed a regular peer review and have been accepted for publication. *E-publishing* of this PDF file has been approved by the authors. After having *E-published Ahead of Print*, manuscripts will then undergo technical and English editing, typesetting, proof correction and be presented for the authors' final approval; the final version of the manuscript will then appear in print on a regular issue of the journal. All legal disclaimers that apply to the journal also pertain to this production process.

Haematologica (pISSN: 0390-6078, eISSN: 1592-8721, NLM ID: 0417435, www.haematologica.org) publishes peer-reviewed papers across all areas of experimental and clinical hematology. The journal is owned by the Ferrata Storti Foundation, a non-profit organization, and serves the scientific community with strict adherence to the principles of open access publishing (www.doaj.org). In addition, the journal makes every paper published immediately available in PubMed Central (PMC), the US National Institutes of Health (NIH) free digital archive of biomedical and life sciences journal literature.

Support Haematologica and Open Access Publishing by becoming a member of the Europe Hematology Association (EHA) and enjoying the benefits of this membership, which inc participation in the online CME2program

> Official Organ of the European Hematology Association Published by the Ferrata Storti Foundation, Pavia, Italy www.haematologica.org

Excellent prognosis of late relapses of ETV6/RUNX1-positive childhood

acute lymphoblastic leukemia: lessons from the FRALLE 93 protocol

Running title : Prognosis of ETV6/RUNX1 leukemia relapses

Virginie Gandemer¹, Sylvie Chevret², Arnaud Petit³, Christiane Vermylen⁴, Thierry Leblanc³, Gérard Michel⁵, Claudine Schmitt⁶, Odile Lejars⁷, Pascale Schneider⁸, François Demeocq⁹,

Brigitte Bader-Meunier¹⁰, Françoise Bernaudin¹¹, Yves Perel¹², Marie-Françoise Auclerc²,

Jean-Michel Cayuela¹³, Guy Leverger¹⁴ and André Baruchel³ on behalf of the FRALLE group

¹Department of Pediatric Hematology/Oncology, University Hospital of Rennes, France
²Department of Statistics of St Louis Hospital and Paris Diderot University France;
³Department of Pediatric Hematology of Robert Debre Hospital and Paris Diderot University, France; ⁴Department of Pediatric Hematology of Brussels, Belgium; ⁵Department of Pediatric Hematology, University Hospital of Marseille, France; ⁶Department of Pediatric Hematology/Oncology, University Hospital of Nancy, France; ⁷Department of Pediatric Hematology, University Hospital of Tours, France; ⁸Department of Pediatric Hematology, University Hospital of Rouen, France; ⁹Department of Pediatric Hematology/Oncology, University Hospital of Clermont-Ferrand, France;
¹⁰Department of Pediatric Hematology, University Hospital of Bicêtre, France;
¹¹Department of Pediatric Hematology, University Hospital of Creteil, France;
¹²Department of Pediatric Hematology, University Hospital of St Louis, Paris, France, and ¹⁴Department of Pediatric Hematology/Oncology, University Hospital of St Louis, Paris, France, Paris, France

ABSTRACT

Background.

The prognosis of relapses of *ETV6/RUNX1*-positive acute lymphoblastic leukemia remains to be evaluated, particularly with regards to the frequency of late relapses. We performed a long term follow-up a retrospective study to address the outcome of *ETV6/RUNX1*-positive leukemia relapses.

Design and Methods.

Among the 713 children tested for *ETV6/RUNX1* enrolled into the FRALLE 93 protocol, 43 *ETV6/RUNX1*-positive patients relapsed (19.4%). Most were initially stratified as low or intermediate risk groups. Median follow-up after relapse was 54.2 months. All but three received a second-line salvage therapy and 16 underwent an allogeneic transplantation.

Results.

ETV6/RUNX1 greatly impacted on overall survival after relapse (3 year-survival= 64.7 % for positive *versus* 46.5 % for negative cases) (p= 0.007). The 5-year cumulative incidence of relapse was 19.4% and testes were more frequently involved in *ETV6/RUNX1*-positive relapses (p=0.04). 81.4 % were late relapses, early combined or isolated extramedullar relapses. The 5-year survival rate of *ETV6-RUNX1*-positive acute lymphoblastic leukemia relapses reached 80.8% when relapse occurred after 36 months (*vs* 31.2%). In univariate analysis, female gender was associated with a poor survival, whereas site of relapse, age at diagnosis, leukocytosis and consolidation strategy had no effect. In multivariate analysis, only the duration of first remission remained associated with outcome.

Conclusions.

We found an excellent outcome for *ETV6/RUNX1*-positive leukemia relapses occurring over 36 months post-diagnosis. Duration of first complete remission may thus be a guide to define the treatment strategy of *ETV6/RUNX1*-positive leukemia relapse.

Introduction

Despite significant improvements in treatment, about 20% of children with acute lymphoblastic leukaemia (ALL) still suffer from relapse (1-5). Unfortunately the cure rate remains low with event-free survival (EFS) reaching only 30% at 10 years after relapse (6-9). Few prognostic factors have been identified (10). Blast phenotype (T versus B), site of relapse, and duration of the first remission are the main variables affecting outcome. The current relapse risk group stratifications are based on the system first described by the Berlin-Frankfurt-Munster (BFM) relapse group, which takes into account the time of relapse, the site of relapse (bone marrow and/or extramedullar), and the subsequent immunophenotype to categorize patients into one of four risk categories (S1 to S4) (11). This stratification has been adapted by the Medical Research Council (MRC, UK) and is also used by the Children's Oncology Group which divided marrow relapse into early relapse occurring up to 36 months from diagnosis and late relapse which occurred greater than or equal to 36 months from initial diagnosis (7, 8). The Children's Cancer Group has also divided relapses into early (< 18 months from diagnosis), intermediate (18-35 months) and late relapse (\geq 36 months) (12). In this context, ETV6/RUNX1-positive ALLs have a separate status. Indeed, the ALL subgroup is characterized by a good prognosis which is better than for B-ALL without this

subgroup is characterized by a good prognosis which is better than for B-ALL without this fusion transcript (2-5, 13). Nevertheless, some recent long-term results revealed late relapses in this group of patients (14, 15). It has been hypothesized that a relapse is essentially a *de novo* ALL originating from a preleukemic stem cell (16-19). This notion should be taken into account when considering the treatment options. The increased proportion of extramedullary relapses including testicular or ovarian relapses has also been discussed by a few authors (20, 21). To investigate the outcome of *ETV6/RUNX1*-positive ALL relapses, we report the long-term results of the *ETV6/RUNX1*-positive ALL relapses initially enrolled into the FRALLE93 protocol and we identify some prognostic factors which call into question the current therapeutic approach for treating late relapses.

Design and Methods

The FRALLE (FRench group for childhood ALL) 93 trial was open to children aged 0 to 20 years with untreated ALL, not including those with L3 ALL or Down's syndrome. Between June 1, 1993, and December 31, 1999, 1395 children were enrolled onto the FRALLE 93 trial in 18 French paediatric centers and one Belgian paediatric center. This study was approved by the ethics committee of the Hôpital Saint Louis, France (accepted April 29, 1993). All patients, or their parents, gave their informed consent in accordance with the Declaration of Helsinki. The diagnosis of ALL was based upon morphological, immunophenotypic, and cytogenetic analyses of bone marrow samples. From 1995, children were systematically screened for four fusion transcripts (*ETV6-RUNX1, BCR-ABL, E2A-PBX1, MLL-AF4*). The FRALLE 93 study population was stratified into three groups (low-risk, intermediate-risk, and high-risk) based on the following risk factors: age, white-cell count at diagnosis, haemoglobin level, immunophenotype, karyotype, and response to steroids. Treatment has been described in details by Dufourg *et al*, 2007 (22) (described in supplementary data 1).

Statistical analysis

Summary statistics were calculated, namely median (inter-quartile range, IQR) for continuous variables, and frequency (with percentages) for qualitative variables.

Distribution of qualitative variables was compared with the chisquare test, or —when inappropriate— the Fisher exact test; comparison of continuous variables was based on the nonparametric Wilcoxon rank-sum test.

We first compared the characteristics of tested and non-tested patients, to check for potential selection bias. We then focused on the *ETV6/RUNX1* positive children who relapsed. We compared the cumulative incidence of relapse in positive *versus* negative patients. Relapses were segregated into isolated central nervous system (CNS) relapse, isolated bone-marrow relapse, and others. We also considered bone-marrow relapses (whether isolated or combined), as well as testicular relapses, the latter only assessed in boys.

Finally, we focused on the outcome of *ETV6/RUNX1*-positive compared with negative patients. Relapsed patients were classified according to the risk groups defined in the REZ-BFM 95/96 study (11). This distinguished S1/S2 patients (isolated bone-marrow relapses occurring ≥ 6 months after completion of primary therapy, isolated extra-medullary relapses,

and combined bone marrow relapses occurring ≥ 18 months after completion of primary therapy) from S3/S4 patients (<6 months isolated bone marrow relapses and < 18 months after primary diagnosis combined bone marrow relapses).

Patient follow-up data were updated in October, 2010. The cumulative incidence of relapse was estimated from the date of complete remission, with death prior to relapse considered as a competing risk, and subgroup comparisons based on the Gray test. The Kaplan–Meier method was used to estimate the overall survival (OS) probabilities with 95% confidence intervals (CI) from the date of relapse until death, irrespective of cause. Comparisons of Kaplan–Meier curves were based on the log-rank test. Multivariate Cox stepwise forward-regression analysis (including testicular and non-testicular relapses) was performed to determine the independent set of prognostic variables.

All statistical tests were two-sided, with p-values of 0.05 or less denoting statistical significance. Statistical analysis was performed using the software SAS v9.2 (SAS Inc, Cary, NC), and R 2.10.1 (http://www.R-project.org) software packages.

Results

Total cohort

A flow chart representing the study design is shown in Figure 1. Among the 1,105 B lineage ALL excluding infants and patients with t (9;22) or t (4;11), 724 had a known *ETV6/RUNX1* status (most of the other cases were diagnosed before systematic screening). In terms of age, gender, blood prednisone response, day 21 marrow status, complete remission, and survival, there were no differences between these 724 patients and the 381 who were not tested for *ETV6/RUNX1*. However, the initial median leukocyte count was higher in the tested group $(10.10^9/1 \text{ versus } 6.4.10^9/1 \text{ p}= 0.003)$, and more patients were enrolled in the low-risk group of treatment among those not tested for *ETV6/RUNX1* ALL (22% versus 13% p=0.001).

Tested patients

A total of 186 (97.4%) out the 191 patients with t(12;21) achieved CR1, whereas 6 (1.1%) out of the 533 *ETV6/RUNX1*-negative patients did not achieve CR1 (p= 0.17). Median follow-up after CR was 9.9 years [IQR: 8.3-11.4]. Overall, 162 (45 % of boys) of the 713 children who reached CR1 relapsed, including 43 with t(12;21). The cumulative incidence of relapse did not significantly differ between ETV6/RUNX1positive and negative ALL (p=0.94), with a 5-year estimate at 19.4% and 19.9%, respectively, nor according to gender with 17% of females versus 22% of males who had relapsed at 5 years (p=0.10) (Figures 2A, 2B). There was no difference in the type of relapse according to ETV6/RUNX1 positivity, with 106 (65.4%) isolated bone-marrow relapses (n=27 in ETV6/RUNX1-positive patients and n=79 in ETV6/RUNX1-negative patients), 24 (14.8%) isolated but extra-medullary relapses (n=6 for ETV6/RUNX1-positive patients and n=18 of ETV6/RUNX1-negative patients), and 32 (19.8%) involving combined sites (p=0.80) (Figure 1). Among the 383 boys in CR1, 96 relapsed, with a 5-year cumulative incidence of relapse of 21.6% in ETV6/RUNX1-positive ALL and 22.4% in ETV6/RUNX1-negative ALL (p=0.90). Of note, 11 out of 26 (43%) were testicular relapses in the ETV6/RUNX1-positive ALL males versus 16 out of 70 (23%) in ETV6/RUNX1-negative ALL cases (p= 0.04) (Table 2). As testicular relapses of ETV6/RUNX1-positive ALL occur mostly after 36 months (10 out of 11 cases), we distinguished the 5-year cumulative incidence of relapse in males and females. As depicting in Figure 2B, 5-year cumulative incidence of relapses did not differ significantly neither in males (p=0.90 by the Gray test) nor in females (p=0.81 by the Gray test) between ETV6/RUNX1-positive and ETV6/RUNX1-negative cases of ALL (21.6% versus 22.5% respectively) (Figure 2B). Nevertheless, in males, the cumulative incidence of testicular relapses appeared to be statistically increased in ETV6/RUNX1-positive patients (p= 0.03, by the Gray test), with at 5-year, an estimated cumulative incidence of 11.3% compared to 4.9% in ETV6/RUNX1-negative patients. By contrast, non-testicular relapses were slightly decreased - though not significantly (p= 0.24), in the ETV6/RUNX1-positive patients compared to those with *ETV6/RUNX1*-negativity (Figure 2C).

Median follow-up after relapse was 6.9 years [IQR: 5.7-8.6]. The presence of *ETV6/RUNX1* fusion transcript greatly impacted the outcome of relapses with a 5-year OS rate of 62.1% (95%CI: 49.0-78.7%) in *ETV6/RUNX1*-positive ALL compared with 37.2% (95%CI: 29.3-47.2%) for *ETV6-RUNX1*-negative ALL (p=0.007) (Figure 3A). The difference in survival estimates between *ETV6/RUNX1*-positive and negative ALL persisted though no longer significant in the S1/S2 subgroup with a 5-year OS rate of 67.7% (95%CI: 53.6-85.5%) in *ETV6/RUNX1*-positive and 47.0% (95%CI: 35.6-61.9%) in *ETV6/RUNX1*-negative ALL (p=0.07) (Figure 3B). S3/S4 relapses had a poor outcome, for the whole cohort with a 5-year OS rate of 27.6% (95%CI: 18.6-41.0%), not statistically different according to *ETV6/RUNX1*-positive and 26.2% (95%CI: 16.9-40.6%) for positive and

negative ALL, respectively; p= 0.45 by the log-rank test) (Figure 3B). As the incidence of testicular relapses was high and could contribute to the favourable outcome *ETV6/RUNX1*-positive ALL, we investigated the outcomes according to testicular *versus* other sites of relapse. The improved OS of *ETV6/RUNX1*-positive relapses was observed whatever the site of relapses (but not reaching statistical significance) (Figure 3C).

ETV6/RUNX1-positive ALL relapses

The initial clinical features of the 43 *ETV6/RUNX1*-positive ALL relapses are shown in Table 1 (details are provided in supplementary data 2). There were 26 (60.5%) males, with median age at diagnosis of 4.1 years [IQR: 2.7-5.4], a median leukocyte count of 15.5 x 10^3 /mm³ [IQR: 7.0-35.6] and no cases of CNS involvement. Thirty three (77%) patients were classified as low risk/intermediate risk and displayed good early responses to therapy.

Salvage therapy after relapse

According to the risk groups defined by the REZ-BFM 95/96 study, 35 (81.4%) patients were classified as S1/S2 and 8 (18.6%) patients were classified as S3/S4.

All but three patients received a second-line therapy protocol (three patients were treated according to a first-line protocol after a relapse occurring at 68, 39, and 32 months after diagnosis). Thirty-seven patients were included in the COOPRALL-97 study (23) which recommended a VANDA (VP-16, cytarabine, mitoxantrone, dexamethasone, asparaginase) induction regimen (except for the late extra-medullary relapses) followed by successive blocks (B1, B2, B3). If an HLA identical donor was available, stem cell transplantation (SCT) was performed after 2 or 3 blocks. Otherwise, alternative treatment consisted of 3 successive blocks repeated 3 times followed by autologous transplantation or —for late combined/isolated medullary or isolated extra medullary relapses— maintenance and CNS radiotherapy. Two patients were treated according to a Capizzi scheme (24) and the latest one according to the UKALL R2 (8).

Outcomes

Median follow-up after relapse was 7.7 years [IQR: 6.2-9], with only 1 child lost to follow-up at 12 months after first relapse.

All but one patient (due to early death) achieved CR2 (n=42, 98%). Eighteen children received a consolidation chemotherapy program and 24 children were transplanted (Flowchart of salvage therapy after relapse in supplementary data 3). Transplantations were performed at

 4.2 ± 1.2 months for the 16 allogeneic transplants (10 were classified as S1/S2 and 6 as S3/S4) and 7.6±1.2 months after relapse for the 8 autologous transplants.

A second relapse occurred in 13 patients, including 10 (77%) initially stratified as S1/S2. At 5 year, the cumulative incidence of second relapse was 28.8% (95%CI: 14.8-42.8%), 18.8% in allogeneic transplants, 37.5% in autografted patients, and 36.2% in those who received chemotherapy. Four out of 13 were still alive in CR3, three treated by chemotherapy in CR2 and one allotransplanted.

Finally, a total of 16 deaths were observed. The 5-year OS rate after first relapse was 62.1% (95%CI: 49.0-78.7%), with differences according to the type of relapse: 70.9% (95%CI: 57.3-87.9%) in S1/S2 versus 37.5% (95%CI: 15.3-91.7%) in the S3/S4 risk group (p=0.049).

Prognostic factors

Based on univariate analyses, the overall survival of *ETV6/RUNX1*-positive ALL after relapse was significantly affected by the duration of the first remission with a 5-year OS rate that was significantly better when relapse occurred after 36 months (80.8%, 95%CI:66.9-97.5% *versus* 31.2%, 95%CI:15.1-64.6%, p=0.0008). Female gender was also associated with a poor survival (p= 0.015), whereas the site of relapse (p= 0.13), age at initial diagnosis (p= 0.81), and leukocytosis (p=0.42) were all not found to be of any prognostic value (Table 2). Similarly, the consolidation strategy (allograft or chemotherapy ± autograft) had no affect on survival (p=0.18).

In multivariate Cox-regression analysis, only the duration of first remission remained associated with the outcome (Figure 4A). In particular, the site of relapse (testicular versus non-testicular relapse) did not provide any additional prognostic information (p=0.79). Although timing of relapse of *ETV6/RUNX1*-negative patients impacted also significantly prognosis (Figure 4B), the 5-year OS rate of the 73 tested patients who relapsed after 36 months appeared to be significantly different in the 27 *ETV6/RUNX1*-positive patients (4 extra-medullary, 9 combined and 14 isolated marrow relapses) compared to the 46 *ETV6/RUNX1*-negative ones (81% *versus* 50%, respectively; p= 0.015 by the log-rank test).

Discussion

After a long-term follow-up, we have reported here the results of the ETV6/RUNX1positive ALL relapses which were initially enrolled in the FRALLE 93 protocol. Indeed, long-term results are now mandatory because very late relapses have been reported (20, 21). In agreement with previous studies, we found that relapses occurred in about 20 % of ETV6/RUNX1-positive ALL, and for males after a longer remission period than other ALL of B lineage (13, 15, 25, 26, 27). Indeed, the male cumulative incidence of relapses was impacted by the frequency of testicular relapses which occurred later than in other sites in boys. More recent protocols based on intensified chemotherapy schemes, especially with intensive use of L-asparaginase (28), report lower frequencies of relapse for ETV6/RUNX1positive ALL but should be examined with longer follow-up (3, 5, 13, 29). We also highlight that boys are slightly over-represented which has been rarely reported except by Seeger et al, and more recently by the NOPHO group (15, 27). Moreover, we observed a specific increase of testicular relapses in ETV6/RUNX1-positive ALL without any increase of other extramedullary sites (mainly CNS). Whatever gender, when pooling all sites of relapses, we did not find any difference in cumulative incidence of relapses according to ETV6/RUNX1positivity. To our knowledge, this finding has not been previously reported and highlights the interest of a primary treatment including drugs able to cross the testicular barrier like high dose methotrexate. Most of these testicular relapses occurred late and special attention to the testes, as well as maybe the ovary, should be required in long-term follow-up of ETV6/RUNX1-positive ALL. According to the responsiveness of the disease, ETV6/RUNX1positive relapses are generally considered as well treatable (13, 15), and here we reported a similar good overall outcome of 62.1 % which is significantly better than ETV6/RUNX1negative relapses. We also found that almost 98% of patients achieved a second complete remission which is significantly higher than that reported by the ALL-REZ BFM 90 and the MRC UKALL-R (6, 9) and also higher than FRALLE 93 ETV6/RUNX1-negative relapses. Unfortunately, in our retrospective study, we were unable to explore IgH/TCR rearrangements or sequence the ETV6/RUNX1 genome to better understand treatment failures. The respective prognostic role of age, site, and time of relapse in the outcome of all types of relapses has been often debated (9, 12). In addition, the definition of extramedullary relapses could, in an early date, move considering the prognostic relevance of submicroscopic bone marrow involvement based on *in vitro* amplification sensitive methods (30). Age did not affect prognosis in our exclusively pediatric study but it should be noted that ETV6/RUNX1positive ALL can present a different clinical progress in young adults (14). We showed that only time-point of relapse had prognostic significance in a multivariate analysis for *ETV6/RUNX1* relapses. We thus highlight that the outcome of relapses occurring \geq 36 months reached an OS rate of 80.8% which is excellent and similar to the prognosis of primary ALL. We could not show any contribution of the high incidence of testicular relapse to this very favourable outcome, possibly due to some lack of statistical power. In contrast, the prognosis of early recurrences (stratified as S3/S4 according to ALL REZ-BFM classification) remained poor despite the presence ETV6/RUNX1 positivity. This timing of relapse is also significant for ETV6/RUNX1-negative relapses but not reaching such a high OS for late relapses. This finding may support the hypothesis that the majority of ETV6/RUNX1 "late" relapses could be due to a novel leukemic clone arising from a persistent initial ETV6/RUNX1-positive preleukemic clone (16, 18, 31). Indeed, ETV6/RUNX1 is an early event which is insufficient for leukemic development (32, 33) and requires the occurrence of additional events conferring proliferative advantages for full malignant transformation (19, 34). Bhojwani et al have even suggested that this model could be operating in non-ETV6/RUNX1 subtypes of late relapses (35). Staal *et al*, in a genome-wide expression analysis of paired diagnosis-relapse samples, also found that relapses could result from therapy involving selection of minor clones present at diagnosis and from genetic alteration of the original tumor cells (36).

Although the mechanisms governing relapses remain unclear, the particular natural history of *ETV6/RUNX1*-positive ALL may lead to use intensive but first-line therapeutic approaches for late relapses. The impact of glucocorticoïd resistance at relapse should also probably be explored in the future according to the recent data published by Kuster *et al* who have implicated glucocorticoid signalling in *ETV6/RUNX1* positive relapses (37). In this small study sample we failed to show any impact of the use of allogeneic transplantation, possibly because of an excess of transplantation-related mortality. However, the poor survival rate of early relapses encourages the use of a second-line treatment with cell therapy according to REZ-BFM. In both types of relapse (38), clinical decisions could also be helped by assessing the dynamics of treatment response using minimal residual disease quantification, which is a promising tool being validated in ongoing relapse trials (25, 39). Finally, our data on *ETV6/RUNX1*-positive ALL from the FRALLE 93 study can probably be extended to other first-line protocols because Freyer *et al* have recently shown that post-relapse survival in childhood acute lymphoblastic relapse was independent of initial treatment intensity (38).

In conclusion, relapses of *ETV6/RUNX1*-positive ALL occurring over 36 months postdiagnosis have an excellent prognosis. Testicular relapses are of especially high frequency and suggest using a primary treatment able to cross the testicular barrier and prolonging follow-up with particular interest for screening gonads. If early relapses need common relapse therapeutic approaches, the physiopathology of *ETV6/RUNX1*-positive ALL could allow the postponement of high-dose therapy for late relapses under cover of examining biological markers of responsiveness.

Authorship and Disclosures

All the authors have substantially contributed to the conception and design or acquisition of data or analysis and interpretation of the data in this multicentric study, and participated to drafting and revising the article. SC and MFA performed statistical analysis. GL and AB approved the version to be published. All authors read and approved the final manuscript.

Acknowledgments

We thank all members of the FRALLE group and are grateful to the children and parents who agreed to participate in the study.

Funding

This work was supported by Délégation à la Recherche Clinique (Assistance Publique-Hôpitaux de Paris), a public non profit organization.

Authorship and Disclosures

The information provided by the authors about contributions from persons listed as authors and in acknowledgments is available with the full text of this paper at *www.haematologica.org*. Financial and other disclosures provided by the authors using the ICMJE (*www.icmje.org*) Uniform Format for Disclosure of Competing Interests are also available at *www.haematologica.org*.

References

1. Kamps WA, van der Pal-de Bruin KM, Veerman AJ, Fiocco M, Bierings M, Pieters R. Long-term results of Dutch Childhood Oncology Group studies for children with acute lymphoblastic leukemia from 1984 to 2004. Leukemia. 2010;24(2):309-19.

2. Moricke A, Zimmermann M, Reiter A, Henze G, Schrauder A, Gadner H, et al. Longterm results of five consecutive trials in childhood acute lymphoblastic leukemia performed by the ALL-BFM study group from 1981 to 2000. Leukemia. 2010;24(2):265-84.

3. Pui CH, Pei D, Sandlund JT, Ribeiro RC, Rubnitz JE, Raimondi SC, et al. Long-term results of St Jude Total Therapy Studies 11, 12, 13A, 13B, and 14 for childhood acute lymphoblastic leukemia. Leukemia. 2010;24(2):371-82.

4. Salzer WL, Devidas M, Carroll WL, Winick N, Pullen J, Hunger SP, et al. Long-term results of the pediatric oncology group studies for childhood acute lymphoblastic leukemia 1984-2001: a report from the children's oncology group. Leukemia. 2010;24(2):355-70.

5. Silverman LB, Stevenson KE, O'Brien JE, Asselin BL, Barr RD, Clavell L, et al. Long-term results of Dana-Farber Cancer Institute ALL Consortium protocols for children with newly diagnosed acute lymphoblastic leukemia (1985-2000). Leukemia. 2010;24(2):320-34.

6. Chessells JM, Veys P, Kempski H, Henley P, Leiper A, Webb D, et al. Long-term follow-up of relapsed childhood acute lymphoblastic leukaemia. Br J Haematol. 2003;123(3):396-405.

7. Raetz EA, Borowitz MJ, Devidas M, Linda SB, Hunger SP, Winick NJ, et al. Reinduction platform for children with first marrow relapse of acute lymphoblastic Leukemia: A Children's Oncology Group Study[corrected]. J Clin Oncol. 2008;26(24):3971-8.

8. Roy A, Cargill A, Love S, Moorman AV, Stoneham S, Lim A, et al. Outcome after first relapse in childhood acute lymphoblastic leukaemia - lessons from the United Kingdom R2 trial. Br J Haematol. 2005;130(1):67-75.

9. Tallen G, Ratei R, Mann G, Kaspers G, Niggli F, Karachunsky A, et al. Long-term outcome in children with relapsed acute lymphoblastic leukemia after time-point and site-of-relapse stratification and intensified short-course multidrug chemotherapy: results of trial ALL-REZ BFM 90. J Clin Oncol. 2010;28(14):2339-47.

 Gaynon PS. Childhood acute lymphoblastic leukaemia and relapse. Br J Haematol. 2005;131(5):579-87. 11. Borgmann A, von Stackelberg A, Hartmann R, Ebell W, Klingebiel T, Peters C, et al. Unrelated donor stem cell transplantation compared with chemotherapy for children with acute lymphoblastic leukemia in a second remission: a matched-pair analysis. Blood. 2003;101(10):3835-9.

12. Gaynon PS, Qu RP, Chappell RJ, Willoughby ML, Tubergen DG, Steinherz PG, et al. Survival after relapse in childhood acute lymphoblastic leukemia: impact of site and time to first relapse--the Children's Cancer Group Experience. Cancer. 1998;82(7):1387-95.

13. Loh ML, Goldwasser MA, Silverman LB, Poon WM, Vattikuti S, Cardoso A, et al. Prospective analysis of TEL/AML1-positive patients treated on Dana-Farber Cancer Institute Consortium Protocol 95-01. Blood. 2006;107(11):4508-13.

14. Burmeister T, Gokbuget N, Schwartz S, Fischer L, Hubert D, Sindram A, et al. Clinical features and prognostic implications of TCF3-PBX1 and ETV6-RUNX1 in adult acute lymphoblastic leukemia. Haematologica. 2009;95(2):241-6.

15. Forestier E, Heyman M, Andersen MK, Autio K, Blennow E, Borgstrom G, et al. Outcome of ETV6/RUNX1-positive childhood acute lymphoblastic leukaemia in the NOPHO-ALL-1992 protocol: frequent late relapses but good overall survival. Br J Haematol. 2008;140(6):665-72.

16. Ford AM, Fasching K, Panzer-Grumayer ER, Koenig M, Haas OA, Greaves MF. Origins of "late" relapse in childhood acute lymphoblastic leukemia with TEL-AML1 fusion genes. Blood. 2001;98(3):558-64.

17. Hong D, Gupta R, Ancliff P, Atzberger A, Brown J, Soneji S, et al. Initiating and cancer-propagating cells in TEL-AML1-associated childhood leukemia. Science. 2008;319(5861):336-9.

18. Pine SR, Wiemels JL, Jayabose S, Sandoval C. TEL-AML1 fusion precedes differentiation to pre-B cells in childhood acute lymphoblastic leukemia. Leuk Res. 2003;27(2):155-64.

19. Zuna J, Ford AM, Peham M, Patel N, Saha V, Eckert C, et al. TEL deletion analysis supports a novel view of relapse in childhood acute lymphoblastic leukemia. Clin Cancer Res. 2004;10(16):5355-60.

20. Chow CD, Dalla-Pozza L, Gottlieb DJ, Hertzberg MS. Two cases of very late relapsing ALL carrying the TEL:AML1 fusion gene. Leukemia. 1999;13(11):1893-4.

21. Ly-Sunnaram B, Henry C, Gandemer V, Mee FL, Burtin F, Blayau M, et al. Late ovarian relapse of TEL/AML1 positive ALL confirming that TEL deletion is a secondary event in leukemogenesis. Leuk Res. 2005;29(9):1089-94.

22. Dufourg MN, Landman-Parker J, Auclerc MF, Schmitt C, Perel Y, Michel G, et al. Age and high-dose methotrexate are associated to clinical acute encephalopathy in FRALLE 93 trial for acute lymphoblastic leukemia in children. Leukemia. 2007;21(2):238-47.

23. Domenech C, Mercier M, Plouvier E, Puraveau M, Bordigoni P, Michel G, et al. First isolated extramedullary relapse in children with B-cell precursor acute lymphoblastic leukaemia: results of the Cooprall-97 study. Eur J Cancer. 2008;44(16):2461-9.

24. Capizzi RL, Poole M, Cooper MR, Richards F, 2nd, Stuart JJ, Jackson DV, Jr., et al. Treatment of poor risk acute leukemia with sequential high-dose ARA-C and asparaginase. Blood. 1984;63(3):694-700.

25. Bader P, Kreyenberg H, Henze GH, Eckert C, Reising M, Willasch A, et al. Prognostic value of minimal residual disease quantification before allogeneic stem-cell transplantation in relapsed childhood acute lymphoblastic leukemia: the ALL-REZ BFM Study Group. J Clin Oncol. 2009;27(3):377-84.

26. Harbott J, Viehmann S, Borkhardt A, Henze G, Lampert F. Incidence of TEL/AML1 fusion gene analyzed consecutively in children with acute lymphoblastic leukemia in relapse. Blood. 1997;90(12):4933-7.

27. Seeger K, Buchwald D, Taube T, Peter A, von Stackelberg A, Schmitt G, et al. TEL-AML1 positivity in relapsed B cell precursor acute lymphoblastic leukemia in childhood. Berlin-Frankfurt-Munster Study Group. Leukemia. 1999;13(9):1469-70.

28. Ramakers-van Woerden NL, Pieters R, Loonen AH, Hubeek I, van Drunen E, Beverloo HB, et al. TEL/AML1 gene fusion is related to in vitro drug sensitivity for L-asparaginase in childhood acute lymphoblastic leukemia. Blood. 2000;96(3):1094-9.

29. Bhojwani D, Pei D, Sandlund JT, Jeha S, Ribeiro RC, Rubnitz JE, et al. ETV6-RUNX1-positive childhood acute lymphoblastic leukemia: improved outcome with contemporary therapy. Leukemia. 2011.

30. Hagedorn N, Acquaviva C, Fronkova E, von Stackelberg A, Barth A, zur Stadt U, et al. Submicroscopic bone marrow involvement in isolated extramedullary relapses in childhood acute lymphoblastic leukemia: a more precise definition of "isolated" and its possible clinical implications, a collaborative study of the Resistant Disease Committee of the International BFM study group. Blood. 2007;110(12):4022-9.

31. Konrad M, Metzler M, Panzer S, Ostreicher I, Peham M, Repp R, et al. Late relapses evolve from slow-responding subclones in t(12;21)-positive acute lymphoblastic leukemia: evidence for the persistence of a preleukemic clone. Blood. 2003;101(9):3635-40.

32. Ford AM, Bennett CA, Price CM, Bruin MC, Van Wering ER, Greaves M. Fetal origins of the TEL-AML1 fusion gene in identical twins with leukemia. Proc Natl Acad Sci U S A. 1998 95(8):4584-8.

33. Wiemels JL, Cazzaniga G, Daniotti M, Eden OB, Addison GM, Masera G, et al. Prenatal origin of acute lymphoblastic leukaemia in children. Lancet. 1999;354(9189):1499-503.

34. Mullighan CG, Phillips LA, Su X, Ma J, Miller CB, Shurtleff SA, et al. Genomic analysis of the clonal origins of relapsed acute lymphoblastic leukemia. Science. 2008;322(5906):1377-80.

35. Bhojwani D, Kang H, Moskowitz NP, Min DJ, Lee H, Potter JW, et al. Biologic pathways associated with relapse in childhood acute lymphoblastic leukemia: a Children's Oncology Group study. Blood. 2006;108(2):711-7.

36. Staal FJ, de Ridder D, Szczepanski T, Schonewille T, van der Linden EC, van Wering ER, et al. Genome-wide expression analysis of paired diagnosis-relapse samples in ALL indicates involvement of pathways related to DNA replication, cell cycle and DNA repair, independent of immune phenotype. Leukemia. 2010;24(3):491-9.

37. Kuster L, Grausenburger R, Fuka G, Kaindl U, Krapf G, Inthal A, et al. ETV6/RUNX1-positive relapses evolve from an ancestral clone and frequently acquire deletions of genes implicated in glucocorticoid signaling. Blood. 2011;117(9):2658-67.

38. Freyer DR, Devidas M, La M, Carroll WL, Gaynon PS, Hunger SP, et al. Postrelapse survival in childhood acute lymphoblastic leukemia is independent of initial treatment intensity: a report from the Children's Oncology Group. Blood. 2011;117(11):3010-5.

39. Eckert C, Biondi A, Seeger K, Cazzaniga G, Hartmann R, Beyermann B, et al. Prognostic value of minimal residual disease in relapsed childhood acute lymphoblastic leukaemia. Lancet. 2001;358(9289):1239-41.

Median [IQR]			Stratification according to REZ-BFM				
N (%)		Overall	\$1/\$2	\$3/\$1			
		n=43	n = 35	n-8			
Ducconting alimits	-1 f 4		11-33	li– 0			
Presenting clinica	ai leatures						
Median age [IQR]	at primary dg	4.1 years [2.7-5.4]	4.1 [2.7-5.5]	4.1 [3.0-5.0]			
	< 5 years	n=29	23 (66%)	6 (75%)			
	\geq 5 years	n= 14	12 (34%)	2 (25%)			
Male gender		26 (60%)	21 (60%)	5 (63%)			
CNS involvement		0	0 0				
Median WBC x 10	$0^{3}/\text{ mm}^{3}[\text{IQR}]$	15.5 [7.0-35.6]	12.4 [6.4-28.9]	23.1 [10.7-93.0]			
	<50	34 (79%)	29 (83%)	5 (63%)			
	≥ 50	9 (21%)	6 (17%)	3 (37%)			
Initial risk group	(FRALLE 93)						
	LR	6 (14%)	5 (14%)	1 (12%)			
	IR	27 (63%)	24 (69%)	3 (38%)			
	HR	10 (23%)	6 (17%)	4 (50%)			
Early response to	o therapy						
U I	CR	43 (100%)	35 (100%)	8 (100%)			
	D8 PGR	42 (98%)	34 (97%)	8 (100%)			
	D21 marrow		× ,				
	status						
	M1	32 (74%)	28 (80%)	4 (50%)			
	M2	10 (23%)	7 (20%)	3 (38%)			
	M3	1 (2%)	0	1 (12%)			
Relapse sites		- (-/)	U U	1 (12/0)			
Poo onoo	BM	27 (62%)	19 (54%)	8 (100%)			
	BM+ CNS	2 (5%)	2 (6%)	0			
	RM+testes		7	0			
	Isolated	4 (15% of males)	4	0			
	Testis			U U			
	Isolated CNS	2 (4 6%)	2 (6%)	0			
	Others	2 (1.6%)	2 (6%)	Õ			
Median relance t	ime after da	47 5 [32_55]	48 4 [38 0-58 3]	24 3 [19 2-32 6]			
months	inic alter ug,		-0.0 [JU.0 JU.0]	27.3 [17.2-32.0]			
Second relapse		13 (30%)	10 (29%)	3 (38%)			

Table 1. Initial features of the 43 ETV6/RUNX1 positive ALL relapses.

IQR: interquartile range

PGR: a good prednisone response was defined as a blast count $<1000/\mu$ l blood after the first seven days on prednisone therapy (*i.e.* on day 8) and one triple-drug intrathecal injection.

D21 marrow status: a good early response to chemotherapy was defined by a blast count fewer than 5% in bone marrow smears on day 21 (M1) and a poor early chemotherapy response by a blast count equal to or more than 5% (with two categories: M2 = 5% to 25% and M3 = more than 25%).

CR: complete remission was defined by no physical evidence of disease, no detectable leukemic blasts on blood smears and less than 5% blasts on bone marrow smears, active haematopoiesis, and normal cerebrospinal fluid. WBC: white blood cell count; BM: bone marrow; CNS: central nervous system; LR: low risk group; IR: intermediate risk group; HR: high risk group (details in supplementary data 1)

ALL REZ-BFM S1/S2 classification : isolated bone-marrow relapses ≥ 6 months after completion of primary therapy, isolated extra-medullary relapses, and combined bone-marrow relapses ≥ 18 months after completion of primary therapy). ALL REZ-BFM S3/S4 classification: <6 months isolated bone marrow relapses and < 18 months after primary diagnosis combined bone-marrow relapses.

		Number of		
		patients	3y-OS (%),	P
		(n=43)	95%CI	
Age at diagnosis				
	< 5 years	29	61.3 (45.7-82.2)	0.81
	\geq 5 years	14	71.4 (51.3-95.1)	
WBC (x 10 ³ / mm ³	3)			0.42
	<50	34	61.8 (47.4-80.5)	
	≥ 50	9	77.8 (54.9-100)	
Gender				
	male		80.3 (66.2-97.4)	0.015
	female	17	41.2 (23.3-72.7)	
Duration of first	remission			
< 36 months		16	31.2 (15.1-64.6)	0.0008
	\geq 36 months	27	85.0 (72.5-99.7)	
Classification of	relapse			
	S1-S2	35	70.9 (57.3-87.9)	0.049
	S3-S4	8	37.5 (15.3-91.7)	
Sites of relapse				
	Isolated BM	27	54.7 (38.7-77.5)	0.13
	Isolated extra	6	66.7 (37.9-100)	
	medullary			
	Others	10	90.0 (73.2-100)	
Consolidation tro	eatment*			
Chemotherapy/autologous SCT		26	73.3 (58.2-92.4)	0.18
Allogeneic SCT		16	50.0 (30.6-81.6)	

Table 2. Univariate analyses of factors influencing survival after first relapse of*ETV6/RUNX1*-positive ALL.

WBC: white blood cell count; BM: bone marrow; SCT: stem cell transplantation ALL REZ-BFM S1/S2 classification: isolated bone-marrow relapses ≥ 6 months after completion of primary therapy, isolated extra-medullary relapses, and combined bone-marrow relapses ≥ 18 months after completion of primary therapy). ALL REZ-BFM S3/S4 classification: <6 months isolated bone marrow relapses and < 18 months after primary diagnosis combined bone-marrow relapses.

* 1 patient dead during induction

Figure legends

Figure 1. Flow chart of study design.

43 and 119 relapses respectively occurred in ETV6/RUNX1-positive and -negative cases of ALL. More testicular relapses (p=0.04) characterized ETV6/RUNX1-positive ALL without any increase in other extra-medullary sites. Second complete remission was achieved in 98% of ETV6/RUNX1-positive ALL and in 82% of ETV6/RUNX1-negative ALL.

Figure 2. Cumulative incidence of relapse according to *ETV6/RUNX1* positivity (negative *versus* positive): overall (A), according to gender (B), according to site of relapse (testicular or non testicular) for males(C).

Figure 3. Kaplan-Meier curves of overall survival of FRALLE 93 relapses according to *ETV6/RUNX1* positivity (negative versus positive) overall (A), in REZ-BFM risk group (B), and according to the site of relapse (testicular or non-testicular) (C).

ALL REZ-BFM classification: S1/S2 = isolated bone-marrow relapses occurring ≥ 6 months after completion of primary therapy, isolated extra-medullary relapses, combined bone marrow relapses occurring ≥ 18 months after completion of primary therapy; S3/S4 = <6 months isolated bone marrow relapses or < 18 months after primary diagnosis combined bone marrow relapses

Figure 4. Prognostic value of time elapsed, in months, since diagnosis in terms of survival after relapse according to *ETV6/RUNX1* positivity



DOI: 10.3324/haematol.2011.059584 Figure 2: Cumulative incidence of relapse according to *ETV6/RUNX1* positivity



DOI: 10.3324/haematol.2011.059584





Figure 4: Prognostic value of time elapsed (in months) since diagnosis in terms of survival after relapse according ETV6/RUNX1 positivity



Supplementary data 1. Summary of the FRALLE 93 stratification.

The FRALLE 93 study population was stratified into three groups (low-risk [LR], intermediate-risk [IR], and high-risk [HR]) based on the following risk factors: age, white-cell count at diagnosis, haemoglobin level, immunophenotype, karyotype, and response to steroids.

- LR was defined by age between 1 and 7 years, white-cell count <10.10⁹/l, B lineage c-ALL CD 10 positive, and no adverse cytogenetic findings.
- HR was defined by at least one major criteria: age below 1 year or >15 years, white-cell count>100.10⁹/l, t(9;22) or t(4;11) or hypoploid or tetraploid karyotype, T-cell lineage, or at least 2 minor criteria: age >10 years, tumour syndrome, haemoglobin>10 G/l, white-cell count >50.10⁹/l, expression of two myeloid antigens such as CD13, CD33, or CD14.
- IR was defined by exclusion of LR and HR.

Patients received an initial treatment comprised of a prednisone prophase and a triple-drug intrathecal injection. Induction treatment then included prednisone, vincristine, L-asparaginase, daunorubicin (except for the LR group), and one or two more triple-drug intrathecal injections (TIT). The main treatment features of the SR and IR protocol were induction, consolidation, delayed intensification, and maintenance (total treatment time 26 and 38 months for girls and boys respectively). Treatment of the HR patients consisted of induction, consolidation, two delayed intensifications, and maintenance with a total treatment time of 2 years. LR patients received 16 TIT and IR patients received 18 TIT. HR children over 4 years received 12 TIT before central nervous system (CNS) radiotherapy, whereas younger children received 18 TIT, without radiotherapy. Very high risk patients (steroid-resistant on day 8 or poor bone marrow early response to chemotherapy on day 21) received intensification with six cycles of chemotherapy alternating with methotrexate and high-dose cytarabine, followed by autografting. Allogeneic stem-cell transplantation was recommended for very high risk patients with related matched donors and for all patients with t (9;22) or t (4;11) or hypoploid karyotype.

Patient	Years at dg	Gender	WBC x10 ³ /mm ³	D8 PGR	D21 M status	End of induction	FRALLE 93 treatment risk group	Time of relapse (months)	Site of relapse	REZ-BFM	Salvage therapy	Post CR treatment	2 nd relapse	e status
1	3.7	male	15.5	yes	M1	CR	IR	74	testis	S1-S2	COOPRALL	СТ		alive
2	4.8	male	11	yes	M1	CR	IR	32	BM	S3-S4	NFL	autoSCT	BM	dead
3	4.5	female	4.7	yes	M1	CR	LR	51	BM	S1-S2	COOPRALL	CT		alive
4	5.5	female	27.2	yes	M2	CR	IR	49	BM	S1-S2	COOPRALL	CT		alive
5	2.8	female	35.6	yes	M1	CR	IR	34	BM	S1-S2	COOPRALL	autoSCT		alive
6	7.3	male	10.4	yes	M1	CR	IR	70	BM	S1-S2	COOPRALL	CT	BM+CNS	dead
7	5.3	male	2.3	yes	M1	CR	IR	43	BM+testis	S1-S2	COOPRALL	CT	CNS	alive
8	5.8	male	12.4	yes	M1	CR	IR	55	BM+testis	S1-S2	COOPRALL	CT		alive
9	2.6	male	229	yes	M1	CR	HR	28	BM+CNS	S1-S2	Capizzi	No CR2		dead
10	7.1	female	16.1	yes	M2	CR	IR	60	BM	S1-S2	COOPRALL	alloSCT		dead
11	5.4	male	24.6	yes	M1	CR	IR	47	BM+testis	S1-S2	COOPRALL	autoSCT		alive
12	11.4	male	10.3	yes	M2	CR	HR	35	BM	S3-S4	COOPRALL	alloSCT		alive
13	2.6	female	6.4	yes	M2	CR	LR	64	BM	S1-S2	COOPRALL	alloSCT		dead
14	1.7	female	50	yes	M1	CR	IR	58	BM	S1-S2	COOPRALL	alloSCT		alive
15	4.7	male	80	yes	M3	CR	HR	19	BM	S3-S4	COOPRALL	alloSCT		Alive
16	2.3	female	28.9	yes	M1	CR	IR	32	BM	S1-S2	COOPRALL	alloSCT		Dead
17	1.9	male	5.1	ves	M1	CR	HR	28	testis	S1-S2	COOPRALL	СТ	BM	Dead
18	7.7	male	9.1	yes	M1	CR	IR	51	BM	S1-S2	COOPRALL	СТ	BM	Dead
19	8.6	male	4.2	ves	M2	CR	HR	62	BM	S1-S2	COOPRALL	alloSCT		Alive
20	4.1	female	8.2	ves	M1	CR	IR	57	BM	S1-S2	COOPRALL	СТ		Alive
21	3.9	female	8	ves	M1	CR	IR	20	CNS	S1-S2	COOPRALL	autoSCT	CNS	Dead
22	3.1	female	36.8	ves	M1	CR	IR	32	BM	S1-S2	COOPRALL	alloSCT	BM	Dead
23	4.2	male	19.3	ves	M1	CR	IR	44	BM+testis	S1-S2	COOPRALL	alloSCT	testis	Alive
24	12.3	female	27.8	no	M2	CR	IR	68	other	S1-S2	NFL	СТ		Alive
25	4.5	female	11.7	ves	M2	CR	IR	32	BM	S1-S2	COOPRALL	СТ	BM	Dead
26	4.0	male	6.3	ves	M1	CR	IR	39	CNS	S1-S2	NFL	alloSCT		alive
27	3.4	male	6.1	yes	M2	CR	LR	27	BM	S3-S4	COOPRALL	alloSCT	BM	dead
28	3.7	male	5.2	ves	M1	CR	LR	40	BM+testis	S1-S2	COOPRALL	СТ		alive
29	2.0	male	51.3	ves	M1	CR	HR	40	BM+testis	S1-S2	COOPRALL	autoSCT		alive
30	2.2	male	26.7	ves	M1	CR	IR	66	BM	S1-S2	COOPRALL	СТ	testis	alive
31	2.7	male	61.3	ves	M1	CR	HR	47	BM	S1-S2	COOPRALL	autoSCT		alive
32	5.3	female	223	ves	M1	CR	HR	22	BM	S3-S4	COOPRALL	autoSCT	BM	dead
33	3.8	male	19.8	ves	M1	CR	IR	50	BM	S1-S2	COOPRALL	СТ	BM	alive
34	1.7	male	106	ves	M1	CR	HR	33	BM	S3-S4	COOPRALL	alloSCT		alive
35	3.4	male	7	ves	M2	CR	IR	56	testis	S1-S2	COOPRALL	СТ		Alive
36	2.6	female	24	ves	M1	CR	IR	7	BM	S3-S4	COOPRALL	alloSCT		Dead
37	3.5	female	22.2	ves	M2	CR	IR	20	BM	S3-S4	COOPRALL	alloSCT		Dead
38	5.0	female	6.1	ves	M1	CR	LR	38	BM	S1-S2	COOPRALL	alloSCT		Dead
39	5.8	male	8	ves	M1	CR	IR	52	testis	S1-S2	COOPRALL	CT		Alive
40	5.8	male	7	ves	M1	CR	IR	70	BM+CNS	S1-S2	COOPRALL	CT		Alive
41	3.3	male	181	ves	M1	CR	HR	50	BM	S1-S2	UKALLR2	CT		?
42	2.1	female	3.8	ves	M1	CR	LR	35	BM	S1-S2	COOPRALL	autoSCT		Alive
43	4.8	male	63	ves	M1	CR	IR	43	BM+testis	S1-S2	Canizzi	alloSCT		Alive

Supplementary data 2. Details of characteristics of the 43 ETV6/RUNX1-positive ALL relapses.

D8 PGR: a good prednisone response was defined as a blast count $<1000/\mu$ l blood after the first seven days on 60 mg/m2/d of prednisone (*i.e.* on day 8) and one triple-drug intrathecal injection. D21 M status: a good early response to chemotherapy was defined by a blast count fewer than 5% in bone marrow smears on day 21 (M1) and a poor early chemotherapy response by a blast count $\geq 5\%$ (with two categories: M2 = 5% to 25% and M3 = more than 25%). CR: complete remission was defined by no physical evidence of disease, no detectable leukemic blasts on blood smears and less than 5% blasts on bone marrow smears, active haematopoiesis and normal cerebrospinal fluid.

WBC: white blood cell count; BM: bone marrow; CNS: central nervous system; LR: low risk group; IR: intermediate risk group; HR: high risk group; dg: diagnosis; CT: chemotherapy; auto/allo SCT: stem-cell transplantation; NFL: new first line therapy.

ALL REZ-BFM S1-S2 classification is defined as isolated bone marrow relapses ≥ 6 months after completion of primary therapy, isolated extra medullary relapses, and combined bone marrow relapses ≥ 18 months after completion of primary therapy). ALL REZ-BFM S3-S4 classification is defined as <6 months isolated bone marrow relapses and < 18 months after primary diagnosis combined bone marrow relapses.



Supplementary data 3. Flowchart of salvage therapy after relapse.

SCT: stem cell transplantation.

CR: complete remission.

MRD: matched related donor; MUD: matched unrelated donor; MMUR : mismatched unrelated donor; CB: cord blood.