Original article

Infection-related complications during treatment for childhood acute lymphoblastic leukemia

Short title: Infection in children with ALL

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Background: Comprehensive studies on neutropenia and infection-related complications in patients with acute lymphoblastic leukemia (ALL) are lacking.

Patients and methods: We evaluated infection-related complications that were grade ≥3 on National Cancer Institute's Common Terminology Criteria for Adverse Events (version 3.0) and their risk factors in 409 children with newly diagnosed ALL throughout the treatment period.

Results: Of the 2420 infection episodes, febrile neutropenia and clinically or microbiologically documented infection were seen in 1107 and 1313 episodes, respectively. Among documented infection episodes, upper respiratory was the most common (*n*=389), followed by ear (*n*=151), bloodstream (*n*=147), and gastrointestinal (*n*=145) infections. These episodes were more common during intensified therapy phases such as remission induction and reinduction, but respiratory and ear infections, presumably viral in origin, also occurred during continuation phases. The 3-year cumulative incidence of infection-related death was low (1.0%±0.9%, n=4), including 2 from Bacillus cereus bacteremia. There was no fungal infection-related mortality. Age 1–9.9 years at diagnosis was associated with febrile neutropenia (P=0.002) during induction and febrile neutropenia and documented infection (both P<0.001) during later continuation. White race was associated with documented infection (P=0.034) during induction. Compared with low-risk patients, standard- and high-risk patients received more intensive therapy during early continuation and had higher incidences of febrile neutropenia (P<0.001) and documented infections (P=0.043). Furthermore, poor neutrophil surge after dexamethasone pulses during continuation, which can reflect the poor bone marrow reserve, was associated with infections (P<0.001).

Conclusions: The incidence of infection-related death was low. However, young age, white race, intensive chemotherapy, and lack of neutrophil surge after dexamethasone treatment were associated with infection-related complications. Close monitoring for prompt administration of antibiotics and modification of chemotherapy should be considered in these patients.

Clinical trials number: NCT00137111

Key words: acute lymphoblastic leukemia, children, infection

Key message: This study describes all infection-related complications throughout the treatment phases in children with ALL. Infection often occurs during intensive therapy phases such as induction and reinduction, but respiratory and ear infections occur throughout treatment. Younger age, white race, and lack of neutrophil surge after dexamethasone during continuation are risk factors for infection.

Introduction

The long-term survival of patients with acute lymphocytic leukemia (ALL) has increased to approximately 90% with risk-directed therapy and improved supportive care.[1] However, intensification and prolonged use of chemotherapeutic drugs are associated with the increased risk of infections.[2, 3] The frequency of treatment-related mortality in contemporary ALL trials is reported to be 2%–4%, mostly due to infections.[3, 4] Chemotherapy intensity, neutropenia, Down syndrome patients, and female gender are associated with a higher risk of infection-related deaths.[2-4] A minor modification in an intensive conventional induction regimen can lead to higher infection-related morbidity and mortality. For example, the substitution of prednisone (40 mg/m², daily) with dexamethasone (6 mg/m², daily) in an induction regimen resulted in a high incidence of sepsis (16 of 38 children with ALL, 42.1%) and in 4 toxic deaths (10.5%).[5] To prevent infection-related death, it is essential to evaluate the incidence and pattern of infection-related complications and associated risk factors in patients with ALL.

Therapy-induced infection-related complications in pediatric acute myeloid leukemia (AML) patients are well characterized.[6, 7] However, similar data on ALL patients are limited to those during induction therapy, and data throughout all treatment phases are lacking. We therefore evaluated the spectrum of and risk factors for infection-related complications throughout the entire treatment period in ALL patients treated with a contemporary regimen in a single protocol.

Methods and Patients

patients, treatment, and supportive care

This retrospective study was approved by the institutional review board and included ALL patients (n=409) treated in the Total XV study from June 2000 to October 2010 at St. Jude Children's Research Hospital (St. Jude).[8] Risk classification and treatment regimen are described elsewhere.[8] Patients received oral trimethoprimsulfamethoxazole 3 days per week starting at day 15 of remission induction until 2 months off therapy for *Pneumocystis jiroveci* pneumonia prophylaxis. Apart from this, routine prophylactic antibiotics or antifungals or colony-stimulating factors were not administered. Patients were advised to wear a particulate filtration mask during the induction and reinduction phases or whenever ANC was <0.5×10⁹/L. All patients had a Port-a-Cath or central venous catheter, which was aseptically accessed as necessary for blood draw or drug infusion. Cefepime or ceftadizime was immediately administered for neutropenic patients with fever and suspected infections, and meropenem was given for those with abdominal symptoms. Vancomycin and tobramycin were added for grampositive and gram-negative infections, respectively. Empiric antifungal agents (i.e., voriconazole, liposomal amphotericin B, caspofungin, or micafungin) were recommended for neutropenic patients with persistent fever of unclear etiology for 3 to 5 days.

infection and febrile neutropenia episodes

The National Cancer Institute's Common Terminology Criteria for Adverse Events (version 3.0) was used to define infection episodes, including febrile neutropenia and clinically or microbiologically documented infections, and episodes grade ≥3 were recorded prospectively. Data were discussed in biweekly multidisciplinary meetings to

confirm accuracy. Fever was defined as an oral temperature of 38.0° C for at least 1 h or a single oral temperature of 38.3° C. Neutropenia and profound neutropenia were defined as absolute neutrophil counts (ANCs) of $<0.5 \times 10^{9}$ /L and $<0.1 \times 10^{9}$ /L, respectively.[9] The duration of neutropenia was calculated as the percentage of actual neutropenia period (days) in the treatment duration (days). ANC responses were evaluated 1 week after dexamethasone and vincristine pulses. Responses were defined as poor when there was less than a 2-fold increase of ANC from a pre-dexamethasone ANC of $0.5-1.2 \times 10^{9}$ /L, no increase in ANC from a pre-dexamethasone ANC $>1.2 \times 10^{9}$ /L, or ANC $<1.0 \times 10^{9}$ /L for a pre-dexamethasone ANC $<0.5 \times 10^{9}$ /L. Definitions of the documented infections are given in the supplementary document.

statistical analysis

The treatment period was divided by chemotherapy intensity as follows: remission induction (6 weeks); consolidation (8 weeks); continuation weeks 1–6; reinduction I (weeks 7–9); continuation weeks 10–16; reinduction II (weeks 17–20); and continuation weeks 21–47, 48–71, 72–103, 104–120, and 121–146 (boys only). Incidences (events/100 patient-days) of febrile neutropenia and documented infections were calculated by the number of episodes divided by treatment duration and patient number during the phase.

Associations among clinical factors (age, sex, race, the presence of Down syndrome, white blood cell count at diagnosis, immunophenotype, treatment risk group, CNS status, and body mass index category at diagnosis), ANC, and infection events were evaluated. The Wilcoxon signed-rank test was used to compare the duration of

neutropenia. The Poisson regression model was applied to identify risk factors for infection events. Multivariate regression analysis was performed to identify multiple independent risk factors. Infection events during the 4-week block after dexamethasone and vincristine pulses were analyzed for their association with ANC response by the generalized estimation equation model. Adjustment for multiple hypothesis tests was not applied. All analyses were performed using SAS (r) Proprietary Software Version 9.3.

Results

infection-related complications

Supplementary Table S1 shows patient demographics and clinical characteristics (*n*=409). Supplementary Table S2 lists 2420 episodes of infection-related complications by treatment phase. Febrile neutropenia was the most common infection-related complication (*n*=1107 episodes), followed by documented infections of the upper respiratory tract (*n*=389), ear (*n*=151), bloodstream (*n*=147), and gastrointestinal tract (*n*=145). Incidences of febrile neutropenia and documented infections were higher during induction and the 2 phases of reinduction (Figure 1A). Lip/perioral, gastrointestinal, urinary tract, and fungal infections were also seen in these phases (Figure 1B). Bloodstream infections were mostly seen during induction, followed by reinduction II in which standard- and high-risk patients received high-dose cytarabine therapy. Skin infections were often seen in induction and between week 10 and reinduction II, and pneumonia was frequently seen in reinduction II (Figure 1C). Upper respiratory tract, ear, and catheter infections were seen throughout the treatment course

(Figure 1D), and upper respiratory tract and ear infections tended to increase toward the end of therapy.

Details of blood stream bacterial infections, fungal infections, and respiratory viral isolates are shown in supplementary Tables S3, S4, and S5, respectively. Of the 409 patients, 4 (0.98%) died of infection: 2 patients died due to *Bacillus cereus* bacteremia during induction therapy on days 12 and 14, respectively, and 2 patients died of presumed septic shock. Postmortem culture in a patient showed *Clostridium* species and *Bacteroides caccae* in the cerebrospinal fluid after the second dose of high-dose methotrexate during consolidation. The other patient with Down syndrome had a presumed infection during continuation week 12, but the causative organism was not identified. The 3-year cumulative incidence of infection-related death was 1.0%±0.9%. No patient died from fungal infection.

absolute neutrophil counts during induction and continuation phases and

infection

Because neutropenia is associated with the occurrence of infections, we evaluated the percentage of days with ANC < 0.1×10^{9} /L and ANC < 0.5×10^{9} /L during induction and continuation therapy (Table 1). The median duration of neutropenia (ANC < 0.5×10^{9} /L) was longer during induction (52.0%) than during continuation weeks 1–20 (16.0%) and 21–120 (11.0%). During induction, the longer neutropenia period (for both ANC < 0.1×10^{9} /L and < 0.5×10^{9} /L) was significantly associated with younger age (1–9.9 years) at diagnosis, low-risk subgroup, white race, and B-lineage ALL (all *P*≤0.002). Female

gender was significantly associated with a longer duration of ANC $<0.5 \times 10^{9}$ /L (*P*=0.001).

For continuation weeks 1–20, age 1–9.9 years at diagnosis (P=0.012) and standard- and high-risk subgroups (P=0.007) were significantly associated with longer duration of ANC <0.5×10⁹/L (Table 1). For continuation weeks 21–120, age 1–9.9 years at diagnosis was significantly associated with a longer duration of ANC <0.5×10⁹/L (P=0.010), and white race was significantly associated with longer duration of ANC <0.5×10⁹/L (P=0.010), and white race was significantly associated with longer duration of ANC <0.5×10⁹/L (P=0.010), and white race was significantly associated with longer duration of ANC <0.1×10⁹/L (P=0.044).

risk factors associated with infections

Table 2 lists the clinical factors significantly associated with infection-related complications. During induction, age 1–9.9 years at diagnosis was significantly associated with frequent febrile neutropenia (P=0.002) and white race was significantly associated with documented infection (P=0.034). Female gender was associated with higher frequencies of bloodstream infections (P=0.041). During continuation weeks 1–20, standard- and high-risk subgroups were associated with a higher incidences of febrile neutropenia (P<0.001) and documented infections (P=0.043) than the low-risk subgroup. For continuation weeks 21–120, age 1–9.9 years at diagnosis was associated with febrile neutropenia (P<0.001), documented infection (P<0.001), and upper respiratory infection (P=0.006). B-ALL (P=0.033) and underweight category (P=0.037) were also associated with febrile neutropenia.

Longer duration of ANC < 0.5×10^9 /L was associated with documented infections during induction (*P*<0.001), weeks 1–20 (*P*=0.009), and weeks 21–120 (*P*=0.050) as

well as bloodstream (*P*=0.003) and upper respiratory (*P*=0.022) infections during induction.

absolute neutrophil counts after dexamethasone and vincristine pulses and infection

Between continuation weeks 24 and 103, patients received dexamethasone and vincristine pulses every 4 weeks. ANC was significantly higher 1 week after the start of pulses (median 1.9×10^{9} /L, range 0–30.2 ×10⁹/L) compared to the pre-treatment value (median 1.5×10^{9} /L, range 0–13.5 ×10⁹/L) (*P*<0.001). Among the 6990 total treatment periods, patients had poor ANC responses in 3694 (52.8%) and good responses in 3296 periods (Table 3). Poor ANC response was significantly associated with a high frequency of infection. Among 984 infection episodes during this time period, 702 (71.3%) occurred in patients with poor ANC responses as compared to 282 (28.7%) in patients with good ANC responses (*P*<0.001), with an odds ratio of 2.47 (95% confidence interval: 2.12–2.88). Febrile neutropenia, upper respiratory tract infections, and ear infections were common (supplementary Table S6). When ANC responses were individually evaluated within each 4-week block between pulses, patients with poor responses had significantly higher risk of infections than those with good responses in 14 of 20 of the 4-week blocks (*P*<0.05) (supplementary Table S7).

Discussion

This is the first detailed report of all infection-related complications occurring throughout the entire treatment course of a uniformly treated cohort of patients with ALL. As

expected, febrile neutropenia and documented infections were common during the intensified chemotherapy phases (remission induction and reinduction phases). Intensive chemotherapy increases the frequency and duration of neutropenia, which is a major risk factor for infections.[2, 3] Bloodstream, lip/perioral, gastrointestinal, urinary tract, and fungal infections occurred during these phases of treatment, but upper respiratory tract and ear infections, presumably of viral origin because bacterial pathogens were rarely isolated, occurred throughout the treatment course and increased toward the end of therapy. Febrile neutropenia and documented infections, especially severe bacterial and invasive fungal infections, interfere with the uninterrupted administration of chemotherapy; therefore, reducing these events could further improve outcomes of leukemia treatment.

Compared with low-risk patients, standard- and high-risk patients received more intensive chemotherapy with asparaginase, anthracycline, and high-dose dexamethasone, in addition to 2 intensified reinduction phases during continuation weeks 1–20. Thus, they had significantly longer neutropenia periods and a higher frequency of infection episodes. Interestingly, age 1–9.9 years at diagnosis, compared to age \geq 10 years at diagnosis, was associated with a significantly longer duration of neutropenia in all phases of treatment and with higher incidences of infections during induction and continuation weeks 21–120 when the chemotherapy intensity was not remarkably different between low-risk and standard- or high-risk patients. ALL therapy might induce a more profound immunocompromised status in younger patients, who not only have lower neutrophil counts but also impaired immunoglobulin production and less immunologic memory. White and female patients had a longer duration of neutropenia Downloaded from http://annonc.oxfordjournals.org/ at University of Western Ontario on October 27, 2016

during induction and a higher frequency of documented infections and bloodstream infections, respectively. It is possible that because of pseudo-neutropenia due to a minor reduction in hematopoietic myeloid progenitors at steady state,[10] African-American patients received less intensity of chemotherapy. ALL groups in the United Kingdom and Nordic countries reported increased treatment-related mortality in females.[3, 4] The Nordic group speculated gender differences in immunologic response to infections or in toxicity after cytotoxic chemotherapy,[4] which might explain the longer duration of neutropenia during induction therapy in our study. Underweight patients had a higher incidence of febrile neutropenia in later continuation. Their immune responses can be impaired because of decreased production of complement, cytokines, and immunoglobulins due to malnutrition.[11] Whether a reduction of chemotherapy leads to a decrease in infection-related complications without compromising overall treatment outcome merits further study.[1]

The cumulative risk of infection-related mortality in our study (1.0%) is lower than that in other trials (1.7%–2.4%), although different treatment regimens make direct comparison difficult. Patients were instructed to call our medical staff first if a febrile episode occurred, and IV antibiotics were available immediately upon arrival. However, there were 2 cases of fatal *B. cereus* bacteremia during induction therapy, which can cause fatal fulminant infection with a short interval between onset and irreversible injury.[12] Induction chemotherapy, presence of neutropenia, administration of glucocorticoids or third-generation cephalosporins, lumbar puncture with intrathecal chemotherapy, and tea consumption are risk factors associated with this severe

infection-related complication.[12] It is unlikely that third- and fourth-generation cephalosporin monotherapy (e.g., ceftazidime and cefepime), which is commonly used as empiric therapy for patients with febrile neutropenia, can eliminate B. cereus bacteremia; vancomycin and/or meropenem need to be given.[12] We also instruct patients not to consume tea made from a ball or bag, but allow bottled pasteurized tea. One of our patients with Down syndrome also had a fatal infection-related episode. Increased risk for infection-related complications in patients with Down syndrome is well recognized, and they require special attention.[3] We did not observe fungal infectionrelated mortality, although it has accounted for as many as 20% of cases of infectionrelated mortality in some studies.[3] We also captured all possible invasive fungal infection episodes, not limited to the revised definitions of invasive fungal disease from the European Organization for Research and Treatment of Cancer/Invasive Fungal Infections Cooperative Group and the National Institute of Allergy and Infectious Diseases Mycoses Study Group (EORTC/MSG), and identified these in 31 of 409 (7.6%) patients (supplementary Table S4). This rate is much lower than that from another study reporting proven or probable invasive fungal infection in 30 of 125 (24.0%) patients by using EORTC/MSG criteria.[13] This difference might be due to close monitoring of patients especially during induction and reinduction phases, use of particle filtration masks, early initiation of empiric broad-spectrum antifungal therapy, or different disease epidemiology in our study. Patients receiving prolonged continuation chemotherapy are likely to have frequent viral infections (upper respiratory tract or ear infections), especially in the winter months, as shown in our patient cohort. ALL therapy also induces profound B-cell lymphopenia with abnormally low IgG and IgM levels

during continuation therapy and impairs seroconversion after influenza virus vaccinations.[14] Patients were also allowed to attend schools and other social activities after week 21 of the continuation phase, which increased their susceptibility to community-acquired pathogens. As previously suggested,[3] antibiotics and antifungal prophylaxis could be considered in patients with prolonged neutropenia during intensive phases of treatment (i.e., induction and reinduction phases) and in those at risk such as those with Down syndrome. Several studies have shown a potential reduction in infection with fluoroquinolone prophylaxis without an increase in the incidence of fungal infections.[15, 16] However, further evaluation is required to assess the efficacy and potential harms of prophylaxis in this population. For patients with hypogammaglobulinemia or frequent infection episodes, intravenous immunoglobulin replacement can be considered.

Our continuation therapy included monthly pulses of dexamethasone and vincristine. Dexamethasone can induce marrow release, reduce egress into tissue, delay apoptosis of polymorphonuclear leukocytes, and promote demargination of granulocytes by downregulating L-selectin expression.[17, 18] Thus, the finding of a significantly increased risk of infection in the absence of a granulocyte surge after the dexamethasone block is likely related to decreased bone marrow reserve. Therefore, in the current Total Therapy study for childhood ALL, we decrease the doses or hold subsequent myelosuppressive chemotherapy if ANCs do not increase after dexamethasone pulses. Prolonged use of dexamethasone during the continuation phase is also associated with an increased risk of severe infection and death.[19] As randomized studies with or without dexamethasone and vincristine pulses during

continuation therapy did not show survival benefit for intermediate-risk ALL patients,[20] it is important to evaluate the effects of pulses on survival and infection in all risk groups of ALL.

In conclusion, infection-related complications are associated with young age, white race, intensive chemotherapy, and lack of neutrophil surge after dexamethasone treatment. These findings can be useful to devise future therapeutic interventions, such as close monitoring of patients, use of prophylactic antibiotics, administration of immunoglobulins, modifications of chemotherapy dosing based on bone marrow reserve, or the rational reduction of intensity of chemotherapy regimen stratified by leukemia risk factors.

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disclosure

The authors declare no competing interests.

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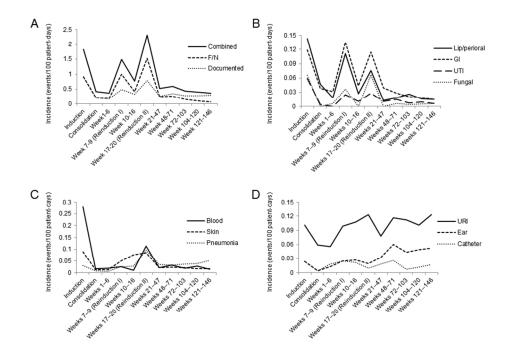
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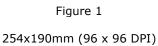
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figure legend

Figure 1. Incidences of infection-related complications during therapy in children with acute lymphoblastic leukemia. (A) All infections (combined), febrile neutropenia (FN), and documented infections (documented). (B) Lip/perioral, gastrointestinal (GI), urinary tract (UTI), and fungal infections. (C) Bloodstream infections, skin infections, and pneumonia. (D) Upper respiratory (URI), ear, and catheter infections. Incidence (events/100 patient-days) was calculated by the number of episodes divided by treatment duration and patient number during the phase.





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Female 173 8.0 0.048.0 0.231 57.0 0.0116.0 0.000 Race Male 212 8.0 0.046.0 47.5 0.096.0 Race White 304 10.0 0.045.0 <0.001 55.5 0.0100.0 <0.002 Other 81 0.0 0.045.0 <0.002 55.5 0.0100.0 <0.002 Immurphenotype 0 0.045.0 0.002 56.0 0.0100.0 <0.002 All patterns 365 0.0 0.050.0 32.0 0.072.0 0.0 Age group 2 0.0 0.0-52.0 0.088 19.0 0.0-72.0 0.0 Risk group 83 0.0 0.0-22.0 11.0 0.0-72.0 0.0 Standard or high risk 196 0.0 0.0-22.0 0.0 0.0-72.0 0.0 Gender 76 0.0 0.0-22.0 0.0 0.0-72.0 0.0 B-ineage 314 0.0 0.0-32.0	Cond	•	190	3.0	0.0-00.0		45.0	0.0-90.0		
Male 212 8.0 0.0-86.0 47.5 0.0-96.0 Race Immume 304 10.0 0.0-86.0 <0.01 55.5 0.0-100.0 <0.0 Other 81 0.0 0.0-45.0 Mail 0.0 0.0-86.0 0.001 55.5 0.0-100.0 <0.0 B-lineage 328 9.5 0.0-66.0 0.02 56.0 0.0-100.0 <0.0 Other 81 0.0 0.0-52.0 16.0 0.0-72.0 0.001 Alg group 7 0.0 0.0-32.0 11.0 0.0-72.0 0.01 Alg group 83 0.0 0.0-25.0 0.068 19.0 0.0-72.0 0.01 Sindard or high risk 196 0.0 0.0-25.0 0.068 19.0 0.0-72.0 0.00 Sindard or high risk 196 0.0 0.0-25.0 0.941 19.0 0.0-72.0 0.00 Male 200 0.0 0.0-25.0 0.941 19.0 0.0-72.0	Gende		170		0.0.00		57.0	0.0.110.0	0.001	
Race Immute and the set of the set			-			0.231			0.001	
White 304 10.0 0.048.0 <0.011 55.5 0.0100.0 <0.00 Dher 81 0.0 0.045.0 40.0 0.093.0 <0.00	_	Male	212	8.0	0.0-64.0		47.5	0.0-96.0		
Other 81 0.0 0.045.0 40.0 0.093.0 Immurophentype 328 9.5 0.068.0 0.002 56.0 0.010.00 <0.000 Tilneage 328 9.5 0.068.0 0.002 56.0 0.010.00 <0.000 All patients 365 0.0 0.050.0 32.0 0.072.0 0.0110 All patients 365 0.0 0.052.0 0.068 19.0 0.072.0 0.011 Immurophentype 282 0.0 0.025.0 0.068 19.0 0.072.0 0.01 Kisk group 7 <th7< th=""> 7 7 <</th7<>	Race	1.0.0.1		10.0						
Immunophenotype Immunophenotype <t< td=""><td></td><td></td><td></td><td></td><td></td><td><0.001</td><td></td><td></td><td><0.00</td></t<>						<0.001			<0.00	
B-lineage 328 9.5 0.0680 0.002 56.0 0.01000 <0.00 T-lineage 57 0.0 0.0500 32.0 0.093.0 All patients 365 0.0 0.052.0 16.0 0.072.0 All patients 365 0.0 0.022.0 16.0 0.072.0 All patients 383 0.0 0.022.0 0.068 19.0 0.072.0 0.01 All patients 83 0.0 0.022.0 0.068 19.0 0.072.0 0.01 Risk group Image 0.0 0.024.0 0.299 15.0 0.072.0 0.00 Standard or high risk 169 0.0 0.025.0 0.941 19.0 0.072.0 0.09 Male 200 0.0 0.024.0 0.931 15.0 0.0-72.0 0.09 Male 289 0.0 0.024.0 0.318 17.0 0.0-72.0 0.19 Male 289			81	0.0	0.0-45.0		40.0	0.0-93.0		
T-lineage 57 0.0 0.0-50.0 32.0 0.0-33.0 All patients 365 0.0 0.0-32.0 16.0 0.0-72.0 Age group	lmmu									
All patients 365 0.0 0.0-32.0 16.0 0.0-72.0 Age group 1 -9.9 years 282 0.0 0.0-32.0 16.0 0.0-72.0 0.01 ≥ 10 years 282 0.0 0.0-25.0 0.068 19.0 0.0-72.0 0.01 ≥ 10 years 282 0.0 0.0-25.0 0.068 19.0 0.0-72.0 0.01 Risk group 11.0 0.0-72.0 0.00 0.0-32.0 28.0 0.0 0.0-32.0 0.0 0.0-72.0 0.00 Standard or high risk 196 0.0 0.0-25.0 0.941 19.0 0.0-72.0 0.00 Gender 165 0.0 0.0-25.0 0.941 19.0 0.0-72.0 0.09 Male 200 0.0 0.0-32.0 0.318 17.0 0.0-72.0 0.09 Male 289 0.0 0.0-32.0 0.204 16.5 0.065.0 11.0 Immurphenotype 7 0.0 0.0-24.0 15.0						0.002	-		<0.00	
All patients 385 0.0 0.0-32.0 16.0 0.0-72.0 Age group 1 1 1 1 1 1 1 1 1 1 1-9.9 years 282 0.0 0.0-25.0 0.068 19.0 0.0-72.0 0.01 Risk group 1 0.0 0.0-32.0 0.068 19.0 0.0-72.0 0.00 Risk group 1 0.0 0.0-25.0 0.068 19.0 0.0-72.0 0.00 Standard or high risk 196 0.0 0.0-25.0 0.941 19.0 0.0-72.0 0.00 Gender 1 0.0 0.0-25.0 0.941 19.0 0.0-72.0 0.09 Male 200 0.0 0.0-32.0 0.941 19.0 0.0-72.0 0.09 Male 200 0.0 0.0-32.0 0.941 19.0 0.0-72.0 0.19 Other 76 0.0 0.0-32.0 0.204 15.0 0.0-72.0 0.19		, ,	57	0.0	0.0-50.0		32.0	0.0-93.0		
Age group Image of the second s	ontinuation	weeks 1-20								
I=9.9 years 282 0.0 0.0-25.0 0.068 19.0 0.0-72.0 0.01 ≥10 years 83 0.0 0.0-32.0 11.0 0.0-60.0 11.0 0.0-60.0 Risk group	All pa	tients	365	0.0	0.0-32.0		16.0	0.0-72.0		
≥ 10 years 83 0.0 0.0-32.0 11.0 0.0-60.0 Risk gr-up Image: Component of the prime	Age g	roup								
Risk group Low risk 196 0.0 0.0-24.0 0.299 15.0 0.0-72.0 0.00 Standard or high risk 169 0.0 0.0-32.0 20.0 0.0-67.0 0.00 Gender Image: Standard or high risk 165 0.0 0.0-25.0 0.941 19.0 0.0-72.0 0.09 Male 200 0.0 0.0-32.0 0.941 19.0 0.0-72.0 0.09 Male 200 0.0 0.0-32.0 0.941 15.0 0.0-72.0 0.09 Male 200 0.0 0.0-32.0 0.318 17.0 0.0-72.0 0.19 Other 76 0.0 0.0-32.0 0.318 17.0 0.0-72.0 0.19 Immurophenotype 15.5 0.0-59.0 15.5 0.0-59.0 15.0 0.0-72.0 0.78 T-lineage 314 0.0 0.0-32.0 0.204 16.0 0.0-72.0 0.78 Immurophenotype 11.0 0.0-41.0 0.178 11.0<		1–9.9 years	282	0.0	0.0-25.0	0.068	19.0	0.0-72.0	0.012	
Low risk 196 0.0 0.0-24.0 0.299 15.0 0.0-72.0 0.00 Standard or high risk 169 0.0 0.0-32.0 20.0 0.0-72.0 0.09 Gender Female 165 0.0 0.0-25.0 0.941 19.0 0.0-72.0 0.09 Male 200 0.0 0.0-32.0 0.941 19.0 0.0-72.0 0.09 Race Male 200 0.0 0.0-32.0 0.318 17.0 0.0-72.0 0.19 Other 76 0.0 0.0-32.0 0.318 17.0 0.0-72.0 0.19 Other 76 0.0 0.0-32.0 0.318 17.0 0.0-72.0 0.19 B-lineage 314 0.0 0.0-32.0 0.204 16.0 0.0-72.0 0.78 T-lineage 51 0.0 0.0-26.0 11.0 0.0-41.0 0.0 All patients 350 2.0 0.0-10.0 0.127 12.0 0.0-41.0 0.0<		≥10 years	83	0.0	0.0-32.0		11.0	0.0-60.0		
Standard or high risk 169 0.0 0.0-32.0 10.0 0.0-67.0 0.0-67.0 Gender Female 165 0.0 0.0-32.0 0.941 19.0 0.0-67.0 0.0 Male 200 0.0 0.0-32.0 0.941 19.0 0.0-67.0 0.09 Male 200 0.0 0.0-32.0 0.941 19.0 0.0-72.0 0.09 Male 200 0.0 0.0-32.0 0.318 17.0 0.0-72.0 0.19 White 289 0.0 0.0-32.0 0.318 17.0 0.0-72.0 0.19 Immurphenotype 76 0.0 0.0-32.0 0.244 16.0 0.0-72.0 0.19 Immurphenotype 79 0.0 0.0-32.0 0.204 16.0 0.0-72.0 0.78 All patients 350 2.0 0.0-10.0 11.0 0.0-41.0 0.01 Alge group 79.0 0.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01	Risk g	group								
Gender Image <		Low risk	196	0.0	0.0-24.0	0.299	15.0	0.0-72.0	0.007	
Gender Image <		Standard or high risk	169	0.0	0.0-32.0		20.0	0.0-67.0		
$\begin{tabular}{ c c c c c c } \hline Male & 200 & 0.0 & 0.0-32.0 & 15.0 & 0.0-60.0 \\ \hline Race & & & & & & & & & & & & & & & & & & &$	Gende	ər								
Male 200 0.0 $0.0-32.0$ 15.0 $0.0-60.0$ Race White 289 0.0 $0.0-32.0$ 0.318 17.0 $0.0-72.0$ 0.19 Other 76 0.0 $0.0-32.0$ 0.318 17.0 $0.0-72.0$ 0.19 Immurplenotype 76 0.0 $0.0-24.0$ 15.5 $0.0-59.0$ 0.78 B-lineage 314 0.0 $0.0-32.0$ 0.204 16.0 $0.0-72.0$ 0.78 Onti-mage 51 0.0 $0.0-32.0$ 0.204 16.0 $0.0-72.0$ 0.78 All patienage 51 0.0 $0.0-32.0$ 0.204 16.0 $0.0-72.0$ 0.78 All patients 3500 2.0 $0.0-10.0$ 0.204 $0.0-35.0$ 0.78 All patients 3500 2.0 $0.0-10.0$ 0.127 12.0 $0.0-41.0$ 0.01 All patients 186 2.0 $0.0-10.0$ 0.595		Female	165	0.0	0.0-25.0	0.941	19.0	0.0-72.0	0.091	
Race White 289 0.0 0.0-32.0 0.318 17.0 0.0-72.0 0.19 Immurphenotype 76 0.0 0.0-24.0 15.5 0.0-59.0 0.19 Immurphenotype 76 0.0 0.0-24.0 15.5 0.0-59.0 0.19 B-lineage 314 0.0 0.0-32.0 0.204 16.0 0.0-72.0 0.78 T-lineage 51 0.0 0.0-26.0 15.0 0.0-56.0 0.78 All patients 350 2.0 0.0-10.0 11.0 0.0-41.0 0.0 Age group 79 0.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01 210 years 79 0.0 0.0-10.0 0.595 11.0 0.0-33.0 0.36 Standard or high risk 164 2.0 0.0-10.0 0.595 11.0 0.0-41.0 0.75 Male 161 2.0 0.0-10.0 0.055 11.0 0.0-35.0 0.75 Kemale							-			
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$	Race									
	1.000	White	289	0.0	0.0-32.0	0 318	17.0	0.0-72.0	0 197	
Immunophenotype Immunoph						0.010			0.107	
B-lineage 314 0.0 0.0-32.0 0.204 16.0 0.0-72.0 0.78 T-lineage 51 0.0 0.0-26.0 15.0 0.0-56.0 0.0 ontimution weeks 21-120 350 2.0 0.0-10.0 11.0 0.0-41.0 0.0-41.0 All patients 350 2.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01 Age group 271 2.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01 1-9.9 years 271 2.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01 Risk group 79 0.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01 Risk group 186 2.0 0.0-10.0 0.595 11.0 0.0-33.0 0.36 Gender 186 2.0 0.0-10.0 0.595 11.0 0.0-41.0 0.75 Male 189 1.0 0.0-10.0 0.063 11.0 0.0-35.0 0.16 Male 189 1.0 0.0-10.0 0.044 12.0 0.0-35.0	Immu		70	0.0	0.0-24.0		15.5	0.0-39.0		
T-lineage 51 0.0 0.0-26.0 15.0 0.0-56.0 All patients 350 2.0 0.0-10.0 11.0 0.0-41.0 Alge grup	IIIIIu		24.4	0.0	0.0.22.0	0.004	10.0	00700	0 705	
All patients 350 2.0 0.0-10.0 11.0 0.0-41.0 Age group 271 2.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01 ≥10 years 271 2.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01 Risk group 200 0.0-10.0 0.127 12.0 0.0-41.0 0.01 Bisk group 200 0.0-10.0 0.127 12.0 0.0-41.0 0.01 Cow risk 186 2.0 0.0-10.0 0.595 11.0 0.0-33.0 0.36 Gender Female 161 2.0 0.0-10.0 0.063 11.0 0.0-41.0 0.75 Kace Female 161 2.0 0.0-10.0 0.063 11.0 0.0-41.0 0.75 Male 189 1.0 0.0-10.0 0.063 11.0 0.0-35.0 0.16 Male 278 2.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Other						0.204	-		0.785	
All patients 350 2.0 0.0-10.0 11.0 0.0-41.0 Age group 1 1.9.9 years 271 2.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01 ≥10 years 79 0.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01 Risk group Image 186 2.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01 Risk group Image 186 2.0 0.0-10.0 0.127 11.0 0.0-35.0 0.36 Risk group Image 186 2.0 0.0-10.0 0.595 11.0 0.0-33.0 0.36 Standard or high risk 186 2.0 0.0-10.0 0.633 11.0 0.0-41.0 0.75 Gender Female 161 2.0 0.0-10.0 0.063 11.0 0.0-41.0 0.75 Male 189 1.0 0.0-10.0 0.063 11.0 0.0-35.0 0.16 Male 278 2.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 0.16			51	0.0	0.0-26.0		15.0	0.0-56.0		
Age group Image: Standard or high risk 271 2.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01 ≥10 years 79 0.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01 Risk group 79 0.0 0.0-10.0 9.0 0.0-35.0 0.0 Risk group Image: Standard or high risk 186 2.0 0.0-10.0 0.595 11.0 0.0-33.0 0.36 Gender Image: Standard or high risk 164 2.0 0.0-10.0 11.0 0.0-41.0 0.0-41.0 Gender Female 161 2.0 0.0-10.0 0.063 11.0 0.0-41.0 0.75 Male 189 1.0 0.0-10.0 0.063 11.0 0.0-41.0 0.75 Male 189 1.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Other 72 0.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Immutrephenotype Immutrephenotype					0.0.40.0		11.0	0.0.44.0		
1-9.9 years 271 2.0 0.0-10.0 0.127 12.0 0.0-41.0 0.01 ≥10 years 79 0.0 0.0-10.0 9.0 0.0-35.0 9.0 0.0-35.0 9.0 0.0-35.0 9.0 0.0-35.0 9.0 0.0-35.0 9.0 0.0-35.0 9.0 0.0-35.0 9.0 0.0-35.0 0.0-41.0 0.0-41.0 0.0-41.0 0.0-35.0 0.0-41.0 0.0-35.0 0.0-41.0 0.0-41.0 0.0-35.0 0.0-41.0 <td< td=""><td></td><td></td><td>350</td><td>2.0</td><td>0.0-10.0</td><td></td><td>11.0</td><td>0.0-41.0</td><td></td></td<>			350	2.0	0.0-10.0		11.0	0.0-41.0		
×10 years 79 0.0 0.0-10.0 9.0 0.0-35.0 Risk gr-up Image: Constraint of the second sec	Age g									
Risk group Image Image <thimage< th=""></thimage<>		-				0.127			0.010	
Low risk 186 2.0 0.0-10.0 0.595 11.0 0.0-33.0 0.36 Standard or high risk 164 2.0 0.0-10.0 11.0 0.0-41.0 0.36 Gender Female 161 2.0 0.0-10.0 0.063 11.0 0.0-41.0 0.75 Male 189 1.0 0.0-10.0 0.063 11.0 0.0-41.0 0.75 Male 189 1.0 0.0-10.0 0.063 11.0 0.0-35.0 0.75 Male 189 1.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Other 72 0.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Immurphenotype Immurphen		,	79	0.0	0.0-10.0		9.0	0.0-35.0		
Standard or high risk 164 2.0 0.0-10.0 11.0 0.0-41.0 0.0-41.0 Gender Female 161 2.0 0.0-10.0 0.063 11.0 0.0-41.0 0.75 Male 161 2.0 0.0-10.0 0.063 11.0 0.0-41.0 0.75 Male 189 1.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 White 278 2.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Other 72 0.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Immurphenotype Immurphenotype <td>Risk g</td> <td>group</td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>	Risk g	group								
Gender Info One find Info One find One f				2.0	0.0-10.0	0.595	11.0	0.0-33.0	0.363	
Female 161 2.0 0.0-10.0 0.063 11.0 0.0-41.0 0.75 Male 189 1.0 0.0-10.0 0.063 11.0 0.0-41.0 0.75 Race V 278 2.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Other 72 0.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Immurphenotype		Standard or high risk	164	2.0	0.0-10.0		11.0	0.0-41.0		
Male 189 1.0 0.0-10.0 11.0 0.0-35.0 Race Male 189 1.0 0.0-10.0 11.0 0.0-35.0 0.0-35.0 White 278 2.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Other 72 0.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Immurphenotype 72 0.0 0.0-10.0 0.620 11.0 0.0-35.0 0.79 B-lineage 303 2.0 0.0-10.0 0.620 11.0 0.0-35.0 0.79 T-lineage 47 2.0 0.0-8.0 10.0 0.0-41.0 0.0-41.0	Gende	ər								
Race Mite 278 2.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Other 72 0.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Immurphenotype 72 0.0 0.0-10.0 0.620 11.0 0.0-35.0 0.79 B-lineage 303 2.0 0.0-10.0 0.620 11.0 0.0-35.0 0.79 T-lineage 47 2.0 0.0-8.0 10.0 0.0-41.0 10.0 0.0-41.0		Female	161	2.0	0.0-10.0	0.063	11.0	0.0-41.0	0.757	
White 278 2.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Other 72 0.0 0.0-10.0 0.044 12.0 0.0-35.0 0.16 Immurphenotype B-lineage 303 2.0 0.0-10.0 0.620 11.0 0.0-35.0 0.79 T-lineage 47 2.0 0.0-8.0 10.0 0.0-41.0 0.0-41.0		Male	189	1.0	0.0-10.0		11.0	0.0-35.0		
Other 72 0.0 0.0-10.0 10.0 0.0-41.0 Immurphenotype 303 2.0 0.0-10.0 0.620 11.0 0.0-35.0 0.79 Flineage 47 2.0 0.0-8.0 10.0 0.0-41.0 0.0-35.0 0.79	Race									
Other 72 0.0 0.0-10.0 10.0 0.0-41.0 Immurphenotype 303 2.0 0.0-10.0 0.620 11.0 0.0-35.0 0.79 Flineage 47 2.0 0.0-8.0 10.0 0.0-41.0 0.0-35.0 0.79		White	278	2.0	0.0-10.0	0.044	12.0	0.0-35.0	0.164	
Immunophenotype Immunophen							-			
B-lineage 303 2.0 0.0-10.0 0.620 11.0 0.0-35.0 0.79 T-lineage 47 2.0 0.0-8.0 10.0 0.0-41.0	Immu									
T-lineage 47 2.0 0.0-8.0 10.0 0.0-41.0			303	2.0	0.0-10.0	0.620	11.0	0.0-35.0	0.796	
		, , , , , , , , , , , , , , , , , , ,					-		2 00	
	hbroviations			2.0	0.0 0.0		10.0	0.0 11.0		

¹Duration of neutropenia was calculated as the percentages of actual neutropenia period (days) among treatment duration (days). [#]No significant differences were seen among groups based on the presence of Down syndrome, white blood cell count at diagnosis, central nervous system status, or body mass index category at diagnosis.

Infection event		Clinical factors	Poisson regression analysis		Multiple Poisson regression analysis	
mecuones	/ent	Cillical lactors	Estimates	Р	Estimates	Р
nduction						
Febrile neutr	openia					
		Age 1–9.9 years vs ≥10 years at diagnosis	2.03	0.002	2.04	0.002
		White vs. others	1.61	0.039	1.52	0.071
		Obese vs. healthy weight	1.63	0.027	1.34	0.107
Documented	l infection					
		White vs. others	1.60	0.034	1.61	0.034
		Obese vs. healthy weight	1.55	0.035	1.27	0.181
		Duration of ANC < 0.5x10 ⁹ /L	1.02	<0.001		
Bloodstream	infection					
		Female vs. male	1.84	0.041		
		Duration of ANC < 0.5x10 ⁹ /L	1.02	0.003		
Upper respir	atory infection					
		Duration of ANC < 0.5x10 ⁹ /L	1.03	0.022		
Continuation w	eeks 1 to 20					
Febrile neutr	ropenia					
		Low vs. standard or high risk	0.62	<0.001	0.64	<0.001
		B-lineage vs. T-lineage	0.74	0.021	0.93	0.628
Documented	l infection					
		Age 1–9.9 years vs ≥10 years at diagnosis	0.73	0.026	0.84	0.268
		Low vs. standard/high risk	0.69	0.006	0.74	0.043
		Duration of ANC < 0.5x10 ⁹ /L	1.01	0.009		
Continuation w	eeks 21 to 120					
Febrile neutr	openia					
	•	Age 1–9.9 years vs ≥10 years at diagnosis	1.79	<0.001	1.73	<0.001
		Low vs. standard/high risk	1.25	0.011	0.91	0.370
		White vs. others	1.26	0.048	1.20	0.123
		B- vs. T-lineage	1.65	0.001	1.44	0.033
		Initial WBC count <100x10 ⁹ /L vs. ≥100x10 ⁹ /L	1.63	0.002	1.33	0.121
		Underweight vs. healthy weight	1.51	0.002	1.29	0.037
Documented	l infection					
		Age 1–9.9 years vs ≥10 years at diagnosis	1.57	< 0.001	1.57	< 0.001
		Low vs. standard or high risk	1.15	0.044	0.93	0.416
		Female vs. male	1.18	0.016	1.15	0.056
		B- vs. T-lineage	1.23	0.042	1.05	0.672
		Initial WBC count <100x10 ⁹ /L vs. ≥100x10 ⁹ /L	1.34	0.010	1.29	0.058
		Underweight vs. healthy weight	1.49	< 0.001	1.01	0.882
		Duration of ANC <0.5x10 ⁹ /L	1.01	0.050		
Upper respir	atory infection					
	-	Age 1–9.9 years vs ≥10 years at diagnosis	1.59	0.007	1.61	0.006
		Underweight vs. healthy weight	1.51	0.047	0.90	0.507

*Multiple analysis was performed when there were 2 or more significant clinical presenting factors by univariate analysis. Duration of ANC <0.5x10⁹/L was not included in the model. Two-sided *P* values <0.05 are listed.

ANC response	ANC pre/post pulse	Infection	No infection	Total	P	OR (95%CI)
Poor		702	2992	3694	<0.001	2.47 (2.12-2.88)
	(500 to 1200/no doubling)	(234)	(1045)	(1279)		
	(>1200/no increase)	(386)	(1712)	(2098)		
	(<500/<1000)	(82)	(235)	(317)		
Good		282	3014	3296		
Total		984	6006	6990		

Table 3. Infection events based on changes in absolute neutrophil counts after dexamethasone and vincristine pulses

Abbreviations: ANC, absolute neutrophil counts; OR, odds radio; CI, confidence interval.