# Original Article Blinatumomab for HLA loss relapse after haploidentical hematopoietic stem cell transplantation

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Received January 27, 2021; Accepted April 1, 2021; Epub June 15, 2021; Published June 30, 2021

Abstract: Loss of patient-specific HLA after haploidentical hematopoietic stem cell transplantation (haplo-HSCT) is considered as a relapse mechanism for lacking the incompatible molecule to elicit alloreactivity, which extensively diminishing graft-versus-leukemia (GVL) effects. Blinatumomab, as a CD3/CD19 bispecific antibody, can yield a profound response via redirecting T cells towards malignant lymphoblasts in B-cell acute lymphoblastic leukemia (B-ALL). We aimed to assess the feasibility of blinatumomab in treating patients with HLA loss relapse after haplo-HSCT. Four eligible patients undergoing HLA loss relapse after haplo-HSCT were enrolled in the study. Four patients achieved a complete remission/complete remission with partial he-matologic recovery (CR/CRh) with three minimal residual disease (MRD)-negative response within the first cycle of treatment. Three of the four met a primary endpoint with CR/CRh and MRD-negative response within 2 cycles of treatment. One patient developed new extramed-ullary sites of skin after the first cycle. Cytokine release syndrome was observed in one patient. Cytopenias, as well as elevated alanine aminotransferase and aspartate aminotransferase, were two common adverse effects during treatment. By redirecting lysis of CD19-positive lymphoblast who losing the incompatible HLA, blinatumomab is a potential strategy to eradicate malignant cells via restoring GVL effects. A randomized clinical trial assessing blinatumomab in patients with HLA loss relapse after HSCT is warranted.

Keywords: B-cell acute lymphoblastic leukemia, HLA loss relapse, blinatumomab

### Introduction

Allogeneic hematopoietic stem cell transplantation (allo-HSCT) can durably control and even cure B-cell acute lymphoblastic leukemia (B-ALL) in the long-term owing to the graft-versusleukemia (GVL) effect of donor T cells [1]. In the case of HLA (human leukocyte antigen) haploidentical HSCT (haplo-HSCT), which provides an accessible option for nearly all patients with allo-HSCT indications lacking fully compatible hematopoietic stem cell resources, the incompatible HLA is an additional target for donor T cells exerting GVL effects compared with HLA identical HSCT, however, relapse is still an unsolved problem [2, 3]. A retrospective study examined patients of B-ALL with recurrence after the first allo-HSCT showed 1- and 2-year overall survival rates of 17% and 10%, respectively, although receiving salvage therapy including a second HSCT, donor lymphocyte infusion (DLI), radiation therapy, mild chemotherapy [4]. Besides, the median survival for relapsed patients is less than a year [4, 5]. Given the relatively high relapse rate and its dismal prognosis following allo-HSCT, further exploration of the underlying mechanism of relapse is warranted as the basis of treatment with pertinence.

Occurring in a proportion of patients who relapse after haplo-HSCT, there is a kind of relapse characterizing in loss of patient-specific HLA genomes in leukemic cells, which facilitates the malignant cells escaping from the GVL effect [6]. A study examined patients with myeloid malignancies found that HLA loss accounted for 33% of relapse in patients undergoing haplo-HSCT [7]. Because of the loss of patient-specific HLA, it is not difficult to explain why DLI as an extension of the GVL effect is redundant in some relapsed patients.

Blinatumomab (Blincyto, Amgen) is the first bispecific T-cell engager (BiTE) consisting of two single-chain variable fragments joined by a linker, bypassing the HLA pathway and dragging T cells closer to malignant B cells by its dual-specificity for CD19 and CD3 [8, 9]. CD3 of T-cell receptor and CD19 expressed by nearly all B-cell linage are connected by blinatumomab, which allows the patient's endogenous T cells to recognize and eliminate CD19-positive B-ALL blasts through several antileukemia processes such as perforin and granzyme release and interferon-y release [10-13]. Because of its HLA-independent feature, it holds a therapeutic promise in HLA loss recurrence. Currently, as there are no clinical data on blinatumomab for posttransplantation HLA loss relapse, we herein report the preliminary results of blinatumomab in treating B-ALL with HLA loss recurrence after haplo-HSCT.

# Method

# Patients

From January 2017 to July 2019, patients relapsed after haplo-HSCT fulling the criteria were enrolled in the study as part of a larger registered study (CTR20170176/NCT034762-39). Eligible patients met the criteria: 1) diagnosed with B cell precursor ALL undergoing haplo-HSCT; 2) achieved complete remission (CR) and full donor engraftment posttransplantation before relapse; 3) had HLA loss relapse after haplo-HSCT; 4) confirmed CD19 expression on blast cells by flow cytometry (FCM). Relapse was defined as ≥5% morphologic lymphoblasts counts in bone marrow (BM) and the reappearance of previously detected clonal cytogenetic abnormalities. The study was conducted in accordance with the Declaration of Helsinki and approved by the ethics review committee of the First Affiliated Hospital of Zhejiang University School of Medicine. Informed consent was obtained from all recruited patients in accordance with the Declaration of Helsinki.

### HLA typing and HLA loss

HLA typing of patients and donors on peripheral blood lymphocytes (PBLs) was performed at diagnosis by the Blood Center of Zhejiang Province or Shanghai Tissuebank Diagnositics Co., Ltd. evaluating 5 loci HLA-A, -B, -C, -DRB1, -DQB1. HLA loss was defined as the no detection of the genome of patient-specific HLA loci in purified CD19<sup>+</sup>/CD34<sup>+</sup> leukemic cells harvested in BM. HLA-KMR, a methodology to detect HLA loss, was based on quantitative polymerase chain reaction (qPCR) by targeting patient-specific HLA markers and non-HLA makers with gene polymorphisms [14]. After haplo-HSCT, the HLA loss relapse or classic relapse (non-HLA loss relapse) pattern was monitored by detection of patient-specific HLA markers and patient-specific non-HLA markers. HLA loss confirmed diagnosis was drawn when patient-specific HLA markers were negative (<3%) and patient-specific non-HLA markers were positive (>3%). For those without ad hoc covered allele groups, next-generation sequencing was applied for chimerism.

### Study design

Blinatumomab was administered via continuous intravenous infusion (CIVI). One cvcle of blinatumomab treatment lasted for 6 weeks, including a 4-week continuous i.v. infusion period and a 2-week treatment-free interval. During the first induction cycle, an initial dose of 9 µg/ day was used for the first 7 days (to reduce the likelihood of cytokine release syndrome [CRS] and immune effector cell-associated neurotoxicity syndrome [ICANS] adverse events associated with blinatumomab), and the dose was increased to 28 µg/day starting on day 8 (week 2) and continuing through day 28 (week 4). The 28 µg/day dose was used for all subsequent cycles. Patients received up to 5 cycles of blinatumomab, with efficacy follow-up at 3, 6, 9, 12, 18, and 24 months after treatment initiation. If a patient experiences a grade 3 central nervous system (CNS)-related adverse event or other clinically relevant grades 3 or 4 adverse event, blinatumomab therapy is discontinued and treatment is restarted when the adverse event is reduced to grade 1 or baseline.

### Outcomes evaluation

The primary endpoint observed complete remission/complete remission with partial hematologic recovery (CR/CRh) rates in the 2 cycles of blinatumomab treatment. The secondary endpoints included minimal residual

disease (MRD) response, relapse-free survival (RFS) rates, overall survival (OS) rates, and adverse events (AEs) during the course of blinatumomab treatment. CR was defined as BM lymphoblasts  $\leq$ 5%, no evidence of active disease, and complete recovery of peripheral blood counts (platelet count >100×109/L, absolute neutrophil count >1×10 $^{9}/L$ ); CRh was defined as BM lymphoblasts ≤5%, no evidence of active disease, and partial recovery of peripheral blood counts (platelet count> 50×10<sup>9</sup>/L and absolute neutrophil count >0.5×10<sup>9</sup>/L). A BM examination was performed on the first 2 cycles after blinatumomab administration assessing the response. RFS and OS were calculated from the start of blinatumomab treatment. AEs with clinical relevance were based on the National Cancer Institute-Common Terminology Criteria for Adverse Events (NCI-CTCAE) v5.0; ICANS and CRS were graded according to American Society for Transplantation and Cellular Therapy (ASTCT) consensus [15].

# Statistical analysis

The patients' characteristics were presented in descriptive form. Survival curves were described using the Kaplan-Meier method. Data were all analyzed using SPSS statistical software version 22.0.01 (IBM, NY, USA).

# Results

# Patients characteristics and transplantation algorithm

Between January 2017 and July 2019, four patients (two females, two males; age range: 22-52 years) with Philadelphia (Ph) chromosome-negative B-ALL receiving haploidentical grafts from their relatives and having posttransplant incipient relapse were enrolled in this study (baseline characteristics and HSCT details were presented in Table 1). Four patients all had CR1 status before HSCT, received cytarabine, busulfan, cyclophosphamide, Me-CCNU and antithymocyte globulin as the conditioning regimen, and peripheral blood stem cell as graft resource. The graft-versushost disease (GVHD) prophylaxis algorithm consisted of cyclosporine, mycophenolate mofetil, and methotrexate [16]. No patient was weighted less than 45 kg.

# HLA loss relapse

Only Patient #4 relapsed in both EM sites of testicles and BM, while other patients had isolated BM relapse. Of note, patient #4 showed an early relapse within 6 months after HSCT. Patients had confirmed HLA loss relapse when enrollment. Patients' and donors' HLA typing outcomes before haplo-HSCT and patients' specific HLA markers showing in red were summarized in Table 2. As shown in Table 2 and Figure 1, when HLA makers and non-HLA markers were discordant, indicating an HLA loss relapse.

# Response and outcomes

Response and outcome data were summarized in **Figure 2**. Two male patients had one course of salvage chemotherapy before blinatumomab treatment. No patient received blinatumomab in remission status. CD19 expression was detected and confirmed by FCM during the course of blinatumomab.

Blinatumomab can exert a potent and prompt efficacy. Four patients (#1, #2, #3, #4) achieved a CR/CRh with three MRD-negative response (#1, #2, #3) within the first 1 cycle of treatment. Three of the four (#1, #2, #3) met a primary endpoint with CR/CRh and MRD-negative response within 2 cycles of treatment. Patient #4 developed new EM sites of skin after the first cycle, for which disrupted the further course of blinatumomab despite remained BM morphological remission and FCM MRD-positive of 0.644%.

Patient #1 completed 5 cycles of blinatumomab, remaining CR since the first cycle. Patient #2 experienced relapse with 19.5% of lymphoblasts in BM after finishing the third cycle. Patient #3 achieved morphological and molecular remission after the first cycle of blinatumomab and kept the leukemia-free for 2 cycles. He developed EM involvement for the first time during the course of the third cycle, manifesting by subcutaneous nodules, while BM kept in remission.

Patient #2 and patient #3 achieved CR/CRh had an RFS of 5.1 and 3.5 months, respectively. The RFS of patient #1 has already sustained for 34.3 months until the latest follow-up (**Figure 3A**). The median overall survival was

Pa- tient	Gen- der	Gene abnor- mities	Age (year)	Status at HSCT	Donor age (year) type, gen- der, HLA match	KIR/ KIR ligand mis- match	Condi- tioning regi- men	CD34+ (×10 <sup>6</sup> / kg re- cipient weight)	Neu- trophil engraft- ment post HSCT	platelet engraft- ment post HSCT	aGVHD	cGVHD		Extra- medullary disease when relapse	BM blasts at relapse (%)	Salvage therapies before blinatu- momab	Time from relapse to the first blinatu- momab (days)	BM blasts at blinatu- momab	Weight at blinatu- momab (kg)
#1	F	None	27	MRD-negative CR1	23, F, 5/10 sister	C2	MAC	8.5	11	12	Liver, GI tract	Mild	8.8	No	38.5	No	13	38.5%	46
#2	F	E2A- PBX1	52	MRD-negative CR1	25, F, 5/10 daughter	No	MAC	6.0	13	13	Skin	Mild	10.7	No	5.5	No	21	37.5%	59
#3	Μ	None	22	MRD-negative CR1	54, M, 5/10 father	C2	MAC	3.1	11	11	Skin	Mild	19.0	No	69.0	VICP	35	61.5%	62
#4	Μ	None	32	MRD-negative CR1	31, M, 5/10 brother	No	MAC	10.6	12	15	No	No	9.5	Testicles	18.9	VMCP	18	6.0%	58

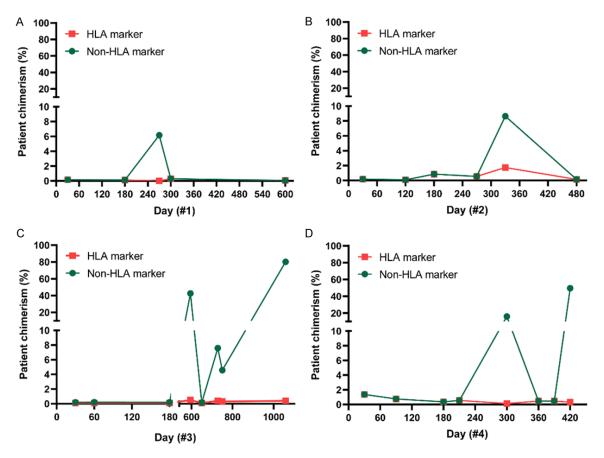
### Table 1. Characteristics of patients and baseline data

aGVHD, acute graft-versus-host disease; BM, bone marrow; cGVHD, chronic graft-versus-host disease; CR, complete remission; F, female; GI, gastrointestinal; HSCT, hematopoietic stem cell transplantation; KIR, killer-cell immunoglobulin-like receptor; M, male; MAC, myeloablative conditioning; MRD, minimal residual disease; VICP, vincristine, idarubicin, cyclophosphamide and prednisone; VMCP, vincristine, melphalan, cyclophosphamide and prednisone.

 Table 2. HLA typing performed on blood samples diagnosis and specific HLA marker used for loss detection

	HLA-A		HLA-B		HL	A-C	HLA-I	DRB1	HLA-DQB1		
#1	A*02:01	A*33:03	B*15:18	B*58:01	C*03:02	C*07:04	DRB1*03:01	DRB1*04:01	DQB1*02:01	DQB1*03:01	
#1 Donor	A*02:01	A*30:01	B*15:18	B*13:02	C*06:02	C*07:04	DRB1*07:01	DRB1*04:01	DQB1*02:02	DQB1*03:01	
#2	A*11:01	A*24:02	B*15:02	B*40:06	C*08:01	C*08:01	DRB1*04:05	DRB1*08:03	DQB1*06:01	DQB1*04:01	
#2 Donor	A*02:01	A*11:01	B*15:02	B*40:01	C*03:04	C*08:01	DRB1*04:05	DRB1*11:01	DQB1*03:01	DQB1*04:01	
#3	A*02:01	A*02:06	B*07:02	B*13:01	C*03:04	C*07:02	DRB1*12:02	DRB1*15:01	DQB1*06:02	DQB1*03:01	
#3 Donor	A*02:06	A*02:07	B*13:01	B*40:01	C*03:04	C*04:82	DRB1*04:03	DRB1*12:02	DQB1*03:01	DQB1*03:02	
#4	A*11:01	A*24:02	B*27:04	B*40:01	C*03:04	C*12:02	DRB1*09:01	DRB1*12:02	DQB1*03:01	DQB1*03:03	
#4 Donor	A*11:01	A*11:01	B*27:04	B*55:12	C*01:02	C*12:02	DRB1*04:05	DRB1*12:02	DQB1*03:01	DQB1*04:01	

Red label indicates the patient-specific HLA loci for HLA-KMR detection.



**Figure 1.** Detection of patient-specific HLA marker (C\*03 in patient #1 [A], A\*24 in patient #2 [B], C\*07 in patient #3 [C], A\*24 and C\*03 in patient #4 [D]) and a patient-specific non-HLA marker after haplo-HSCT.

12.8 months (range, 7.2-34.3), and patient #2, #3, #4 died of disease relapse (**Figure 3B**).

Safety

CRS was observed in patients #1 who had a high fever with a maximum temperature of 39.6°C, requiring medications of temporary dexamethasone, which then pacified in four days. There were no ICNAS observed. Cytopenia, as well as elevated alanine aminotransferase (ALT) and aspartate aminotransferase (AST), were two major AEs during blinatumomab treatment. Cytopenia of grades 1-2 was reported in all patients and no transfusions of blood products required. Patient #1 experienced grade 2 neutropenia; patient #2 experienced grade 2 neutropenia; patient #3 experienced grade 2 pancytopenia; patient #4 experienced

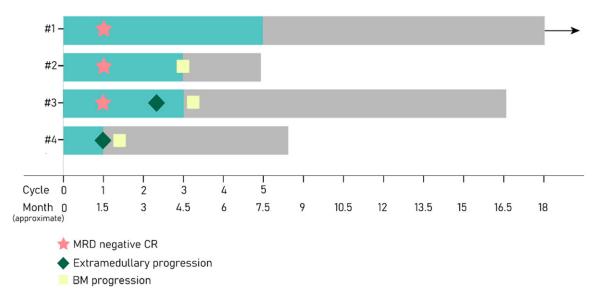


Figure 2. Survival span after blinatumomab. The green bar indicates time treated with blinatumomab.

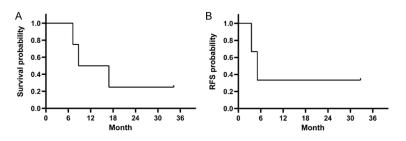


Figure 3. Overall survival (OS) (A) and the incidence of relapse-free survival (RFS) (B) after blinatumomab.

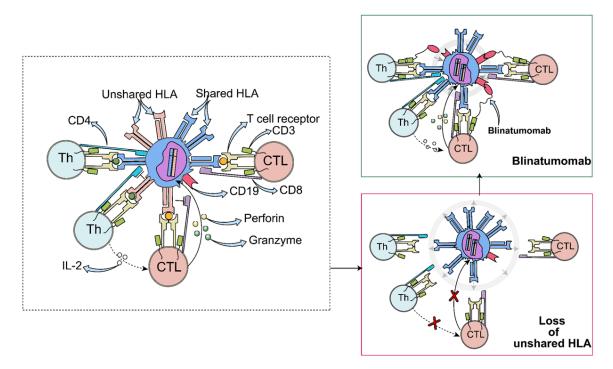
grade 4 elevated ALT (max 699 U/L) and AST (max 595 U/L), thus the study was discontinued for 11 days, thereafter reinitiated when AEs reduced to grade 1 after medication interventions. Chronic GVHD was observed in patient #2 involving mouth and skin (mild) and controlled with prednisone 0.2 mg/kg/d.

### Discussion

Loss of mismatched HLA has been considered as an underlying mechanism for leukemia recurrence after haploidentical HSCT [17]. Because the entire HLA haplotype mismatch between patient and donor, effective alloreactivity could be induced by donor T cells which directly contributes to the GVL effect [18]. Since the relapse after HSCT is characterized by a subset of leukemic cells losing unshared HLA haplotype, it is not difficult to speculate how the relapse occurred: the GVL effect is greatly diminished (Figure 4). For those with HLA loss relapse, selection of accessibly salvage therapeutic options were endowed with more prudent and sensible manner. Since donor T cells acting as an immunological pressure of selection, unlike pan-killing effects of chemotherapies, it may trigger leukemic cells displaying great initiative in removing detect-

able HLA haplotype [19]. Lost patient-specific HLA haplotype partly explains the nonresponsiveness to DLI in HLA-loss patients, for who lacks incompatible HLA to elicit furious alloreactivity, though DLI intends to bring additional donor T cells to magnify GVL effects. Early detection of HLA loss on blasts rather than radically introducing DLI in relapsed patients after allo-HSCT could be more beneficial for saving a lot of troubles to balance the pros and cons of DLI. However, it is crucial to clarify that the seemingly useless DLI might be feasible in some circumstances, which will be discussed later.

Based on the rationale that loss of unshared HLA disguises leukemic cells for escaping T-cell surveillance, to retain antileukemia effects, donor cytotoxic T cells must redirect against those without patient-specific HLA expression once inducing alloreactivity [20]. Hence, as an



**Figure 4.** A proposed mechanism for the escape of malignant lymphoblasts and mode of blinatumomab. The figure illustrates how cytotoxic T cell (CTLs) and T helper (Th) of donors cooperate to kill lymphoblasts via the incompatible HLA molecules, that is alloreactivity. Under the immunological pressure of donor T cells, leukemic cells escape from immunosurveillance by loss of the genome of patient-specific HLA. Blinatumomab can redirect donor T cells to malignant lymphoblasts.

antibody could bind together CD3 of T cells and CD19 of leukemic B cell, blinatumomab is apropos to restore T-cell antileukemia ability by forming a cytolytic synapse.

A profound and rapid response can be induced by blinatumomab. Previous clinical studies showed the CR/CRh rate in relapsed/refractory (r/r) B-ALL was approximately 50% after two cycles [21, 22]. In this study, three patients (75%) achieved CR/CRh after the first cycle, showing a comparable response in the allo-HSCT context, which could be somehow postulated that T cells derived from donors are powerful enough to smash leukemic cells. Patient #1 has been keeping CR after full treatment course and it is vital to note she was the one developed CRS, indicating the furious T cell activation [23].

A study indicated that concomitant or prior EM disease of B-ALL might vitiate blinatumomab's efficiency, and EM relapse or progression was frequently observed in blinatumomab responders [24]. Several cases showed the poor response of blinatumomab in EM disease treat-

ment [25, 26]. In this study, patient (#4) with testicle relapse did show an inferior response than others. One patient (#3) developed EM disease for the first-time during treatment.

This response differentiation between BM and EM sites might lay in the concentration of blinatumomab, quality and capacity of donor T cells, and the level of target antigen expression on lymphoblasts (Figure 5). The traffic incompetence of blinatumomab resulting in inadequate focal concentration, presenting an attachment with BM, is consistent with the clinical evidence that the inferior efficacy in non-Hodgkin lymphomas compared with B-ALL at the same dose while meeting a better response at a higher dose level [27]. Shortcoming of insufficient concentration of blinatumomab outside BM might be overcome by combining with chemotherapy (NCT03023878). Total body irradiation or focal radiation therapy represents an approach to EM sites, for example, a patient with leukemic optic nerve infiltration recovered by combining radiotherapy with blinatumomab [28].

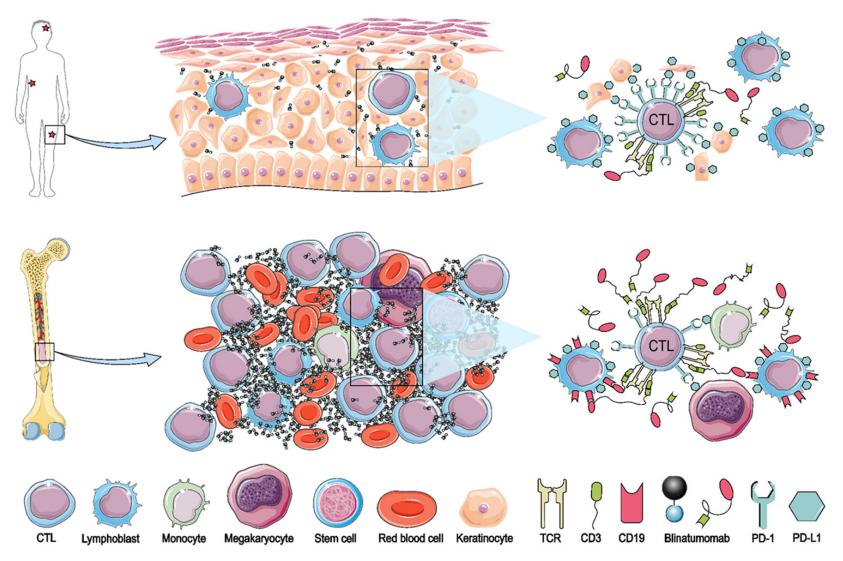


Figure 5. A possible mechanism of response differentiation between BM and EM sites. Compared to EM sites (for example, skin), BM might have a higher concentration of blinatumomab, more cytotoxic T cell (CTLs), and less inhibitory signals (PD-L1/PD-1), as well as a higher level of CD19 in leukemic cells.

Meanwhile, prior publications indicate that GVHD has a positive effect on preventing BM relapse rather than EM relapse [29-31], which can be assumed that the shortage and inefficiency of cytotoxic T cells in EM compartment and these EM sites play as sanctuaries escaping for immunosurveillance. When privileged niches lacking sufficient donor T cells, or in other terms, the spatial heterogeneity of GVL effects in patient's body parts [32-34], for example, T cells with high killing potential accumulating in BM, blinatumomab might increase the anatomic compartmentalization, favoring EM relapse or progression. To date, the mechanisms behind the segregation remain enigmatic, however, interestingly, one of the relevant reasons can be imposed by the varied level of PD-L1 expression in the microenvironment [35, 36]. Furthermore, this mechanism decreases the antineoplastic capacity of donor T cells such as reducing effector cytokines and curtailing proliferation, resulting in exhausted T cells [37, 38]. By artificially fusing extracellular domain of PD-1 to CD3, ex vivo experiment generated highly cytotoxic and targeted T cells [39]. Given the established rationale, inhibiting PD-L1/PD-1 pathway such as bifunctional checkpoint-inhibitory T cell engagers (CITE), nivolumab (NCT04546399) and pembrolizumab (NC-T03512405), might facilitate blinatumomab against EM disease and rescue effector T cells of the donor.

The downregulation or loss of CD19 expression on lymphoblast can explain the EM relapse after blinatumomab to some extent, which is observed in about 40% responders [24]. Conversions of cell lineage such as myeloid shift after blinatumomab blunting treatment efficiency has been reported [40-42]. Combination of immunotherapy with other targets provides a feasible option for eradicating CD19-negative variant (NCT03739814).

On the other hand, in patients with HLA loss relapse, DLI seems to be ineffective since leukemic cells have lost the key incompatible molecular to elicit GVL effects [20]. But DLI provides additional T cells in the patient's body as arsenal despite the possibly disproportionate T-cell distributions, and when combined with blinatumomab, it may arouse robust antileukemia efficacy and overcomes T-cell exhaustion, which is being investigated in a clinical trial (NCT03982992). The AEs observed in these four patients with HLA loss relapse after allo-HSCT were consistent with previous clinical reports in r/r B-ALL, suggesting a well-tolerant feature of blinatumomab. Nealy no AEs greater than grade 3 were reported except patient #4 who developed grade 4 hepatic toxicity showing an elevated ALT and AST after the first dose and then recovered to baseline after herbal medicines treatment. The incidences of GVHD and CRS were reported in prior study approximately 11% and 3%, respectively [43]. No fatal infection event was observed relating to blinatumomab. From the aspect of safety profile, blinatumomab appears safer compared with anti-CD19 chimeric antigen receptor T-cell therapy which has an incidence of CRS approximately 37%-93% and ICANS about 12%-30% [44]. Although the aforementioned idea of combining DLI with blinatumomab in patients with HLA loss is theoretically feasible, it should be interpreted and practiced with caution because the increased risk of severe GVHD and GVHD-related mortality, since GVHD incidence after DLI being about 30-70% [45-47].

In summary, the early detection of patient-specific HLA loss on leukemic cells is of the great importance of selecting salvage treatment and our study was the first one reported the clinical efficacy of blinatumomab in HLA loss occurrence. Blinatumomab can redirect allogeneic T cells to leukemic cells, restoring GVL effects in patients of HLA loss relapse after allo-HSCT. It can exert quick resolution with controllable AEs. Further investigations concerning combination with blockage of PD-1/PD-L1 pathway or DLI are therefore warranted, aiming the goal of sustained remission and EMD clearance.

# Acknowledgements

This work was supported by the National Natural Science Foundation of China (Grant Nos. 81730008) and the Key Research and Development Program of Zhejiang Province (Grant Nos. 2019C03016).

# Disclosure of conflict of interest

### None.

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