REVIEW



Harnessing natural killer cells for refractory/ relapsed non-Hodgkin lymphoma: biological roles, clinical trials, and future prospective



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Abstract

Non-Hodgkin lymphomas (NHLs) are heterogeneous and are among the most common hematological malignancies worldwide. Despite the advances in the treatment of patients with NHLs, relapse or resistance to treatment is anticipated in several patients. Therefore, novel therapeutic approaches are needed. Recently, natural killer (NK) cell-based immunotherapy alone or in combination with monoclonal antibodies, chimeric antigen receptors, or bispecific killer engagers have been applied in many investigations for NHL treatment. The functional defects of NK cells and the ability of cancerous cells to escape NK cell-mediated cytotoxicity within the tumor microenvironment of NHLs, as well as the beneficial results from previous studies in the context of NK cell-based immunotherapy in NHLs, direct our attention to this therapeutic strategy. This review aims to summarize clinical studies focusing on the applications of NK cells in the immunotherapy of patients with NHL.

Keywords Non-Hodgkin lymphoma, Natural killer cells, Immunotherapy, Chimeric antigen receptors, Monoclonal antibodies

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Introduction

B-cell non-Hodgkin lymphoma (NHL) collectively represents the most common type of hematologic malignancy [1]. While advances in chemotherapy, monoclonal antibodies, and stem cell transplantation have improved survival rates, many NHL patients remain resistant to therapy or experience relapse. This highlights the necessity for finding novel curative therapeutic options for these patients [2–4].

Recently, novel therapeutic approaches, such as chimeric antigen receptor (CAR)-T cell therapy, have been utilized in several clinical trials for patients with relapsed/ refractory (R/R) B-cell NHL, resulting in promising clinical responses [5–7]. However, this therapeutic approach is expensive and associated with unique and severe side effects such as cytokine release syndrome (CRS), immune effector cell-associated neurotoxicity syndrome



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(ICANS), and graft-versus-host disease (GVHD) in allogenic settings [8, 9]. Other immune cells, such as natural killer (NK) cells, which exhibit significant cytotoxic activity against cancer cells and possess a safer immune profile, can be used as alternative approaches for immunotherapy of patients with R/R NHL. Moreover, it is also feasible to equip NK cells with a CAR structure [10, 11].

This study will delve into the B-cell NHL tumor microenvironment (TME) and the interaction between NK cells and malignant cells. Additionally, we provide a comprehensive review of clinical trials focused on the utilization of NK cells in patients with R/R NHL. Finally, new approaches recently used to increase NK cells effectiveness for B-cell NHL immunotherapy are summarized.

NK cells: biology, receptors, and functions

NK cells, a specialized subset of innate lymphoid cells (ILCs), can distinguish between self-cells and non-self-cells through the recognition of self-major histocompatibility complex (MHC) I molecules [12, 13]. They constitute approximately 10-15% of the lymphocyte population in peripheral blood, and characterized as large granular lymphocytes with kidney-shaped nuclei, a high cytoplasm-tonucleus ratio, and large azurophilic granules in cytoplasm [14, 15]. NK cell development and maturation primarily occur in the bone marrow, where common lymphoid progenitors (CLPs) differentiate into NK precursors (NKPs), immature NK cells, and finally mature NK cells [12]. Notably, recombinant interleukin (rIL)-15 plays a crucial role in NK cell development from hematopoietic stem cells [16]. In humans, CD122 expression on NKPs is crucial for NK cell lineage commitment, and CD56 expression is a final step in the differentiation of NKPs into NK cells [17]. NK cells are typically identified by the expression of CD56 and CD16, and the absence of CD3 (T cell marker) [18].

Human NK cells can be classified into two main subsets: the CD56^{bright} subset, which is characterized by immaturity, limited cytolytic activity, but high cytokine production; and the CD56^{dim} subset, which is mature, exhibits higher cytolytic activity, but lower cytokine production [19].

NK cells are equipped with various germline-encoded activating and inhibiting receptors [20]. The function of NK cells is delicately regulated by the balance of the activating and inhibitory signals that are transmitted through their receptors [21]. Table 1 provides an overview of NK cell receptors.

NK cells express the HLA-specific activating receptors such as KIRs/CD158 (2DS1–2DS5 and 3DS1), NKG2C, and NKG2E. NKG2C and NKG2E are expressed as heterodimers with CD94. Upon interaction with HLA-E, they transmit activating signals through the DNAX-activation protein (DAP)-12 adaptor molecule [22–24]. The natural cytotoxicity receptors (NCRs) including NKp46, NKp44, and NKp30 are the major non-HLA specific activating NK receptors, which evoke an immune response upon detection of cognate viral and cellular ligands [20, 25]. NKG2D is another non-HLA-specific activating NK cell receptor [26]. The UL16-binding protein (ULBP) and MHC class I chain-related proteins A and B (MICA/B), which are increased in the tumor, stressed, and infected cells, are representative of NKG2D ligands [27]. Additionally, some other molecules, such as 2B4, NTB-A, CD59, NKp80, and DNAX accessory molecule-1 (DNAM-1), are essentially coreceptors; in fact, they can intensify the NK cell triggering induced by NCRs or NKG2D (See Table 1) [28–32]. NK cells are also equipped with the CD16a (FcyRIIIa, a low-affinity Fc), which plays a crucial role in their antibody-dependent cell-mediated cytotoxicity (ADCC) effector function. It is worth noting that CD16 is the only receptor that can activate NK cells without the need for further activation from other receptors [33].

Besides activating receptors, NK cells express inhibitory receptors that modulate the strength of activating receptors and contribute to regulating immune responses and tolerance [34]. The CD94/NKG2A (CD94/ CD159a) heterodimer and members of the KIR/CD158 family are two distinct classes of HLA-specific inhibitory receptors [35, 36]. The LIR-1/ILT2/CD85 is an inhibitory receptor with broad specificity for both classical and non-classical MHC molecules [37-39]. Program-cell death receptor 1 (PD-1), Sialic acid recognizing Immunoglobulin-like Lectins (Siglecs)/p75/AIRM1/CD328, leukocyte-associated immunoglobulin-like receptor-1 (LAIR-1)/p40/CD305, and IRp60/CD300a known as another non-HLA-specific inhibitory receptor that hamper NK cell-mediated antitumor immunity through the recognition of different ligands on the surface of cancerous cells (See Table 1) [40, 41].

NK cells have diverse functions, including their natural antitumor and antiviral activities, as well as their regulatory roles in modulating immune responses and promoting tissue growth. They are abundant in the TME, where they kill cancer cells in a variety of ways [21]. The antitumor functions of NK cells include missing self-mechanisms, direct cytotoxicity, and activation of adaptive immune responses (Fig. 1) [42–47].

TME in NHLs: composition and functions

The development of B-cell lymphoma involves an intricate interplay between tumor cells and the surrounding TME (Fig. 2). The microenvironment in B-cell lymphoma is fascinating because it has crucial functions in regulating the survival and growth of tumor cells, promoting immune evasion, and contributing to the development of resistance to treatment [48–52]. It is worth mentioning

Table 1 Overview of NK cells receptors

Receptors	Molecular structure	CD marker	Ligand (s)
Activating receptors			
HLA-specific activating receptors	Immunoglobin superfamily		
(I) Killer immunoglobin receptors (KIRs)			
• KIR2DS1		CD158h	HLA-C2
• KIR2DS2		CD158j	HLA-C1
• KIR2DL4		CD158d	HLA-G
• KIR2DS5		CD158g	?
• KIR3DS1		CD158e2	HLA-F
(II) CD94/NKG2	C-type lectin family		
NKG2C		CD159c	HLA-E
NKG2E		CD159e	?
Non-HLA-specific activating receptors			
(I) Natural cytotoxicity receptors (NCRs)	Immunoglobin superfamily		
• NKp46 (NCR1)		CD335	Heparin, viral HA and HN
• NKp44 (NCR2)		CD336	viral HA and HN, NKp44L, PCNA
• NKp30 (NCR3)		CD337	B7-H6, BAT3, viral HA
(II) NKG2D	C-type lectin	CD314	MIC-A, MIC-B, ULBP
(III) Coreceptors			
• 2B4	Immunoglobin superfamily	CD244	SLAMF2 (CD48)
• NTB-A	Immunoglobin superfamily	CD352	NTB-A
• DNAM-1	Immunoglobin superfamily	CD226	PVR (CD155), Nectin-2 (CD112)
• NKp80	C- type lectin-like family	-	AICL
(IV) FcγRIII	Immunoglobin superfamily	CD16	lgG1, lgG2, lgG3
Inhibitory receptors			
HLA-specific inhibitory receptors			
(I) Killer immunoglobin receptors (KIRs)	Immunoglobin superfamily		
• KIR2DL1		CD158a	HLA-C2
• KIR2DL2		CD158b	HLA-C1
• KIR2DL3		CD158b2	HLA-C1
• KIR2DL5		CD158F	?
• KIR3DL1		CD158e	HLA-Bw4
• KIR3DL2		CD158k	HLA-A3, A11
(II) CD94/NKG2	C-type lectin family		
• NKG2A		CD158a	HLA-E
(III) Other			
• LIR-1(ILT2)	Immunoglobin superfamily	CD85	?
Non-HLA-specific inhibitory receptors			
(I) PD-1	Immunoglobin superfamily	CD279	PD-L1/PD-L2
(II) Siglec7/p75/AIRM1/	Immunoglobin superfamily	CD328	Sialic acid
(III) LAIR-1/p40	Immunoglobin superfamily	CD305	Collagen, C1q, SP-D, ADP
(IV) IRp60	Immunoglobin superfamily	CD300a	?

HN Hemagglutinin neuraminidases, PVR Poliovirus receptor, AICL Activation-induced C-type, SP-D Surfactant protein D, ADP Adiponectin, PCNA Proliferating cell nuclear antigen, SLAMF2 Signaling lymphocytic activation molecule 2, MIC MHC class I chain-related protein, ULBP UL16 binding protein 1

that different cells within the TME can display protumorigenic or anti-tumorigenic functions, as shown in Table 2.

Intratumoral T lymphocytes constitute 50% of the total cells within the TME and are categorized based on the

expression of CD4 or CD8. Typically, CD4⁺ T lymphocytes support other immune cells, while CD8⁺ cytotoxic T lymphocytes (CTLs) are known to trigger target cell killing by releasing perforin and granzyme B, expression of death ligands and IFN- γ and TNF- α production [53,



Fig. 1 Natural killer (NK) cell function within tumor microenvironment (TME). A) Missing-self recognition against tumor cells lacking MHC class I ligands for inhibitory NK receptors. B) Direct cytotoxicity against tumor cells mediated by releasing cytotoxic granules containing perforin/ granzymes, IFN-γ and TNF-α production, antibody-dependent cell-mediated cytotoxicity (ADCC) via CD16, and induction of apoptosis pathway through death receptor ligands like TRAIL/FasL. C) Triggering the adaptive antitumor immunity by recruiting dendritic cells via chemokines and then amplifying CD4⁺ and CD8⁺ T cells antitumor immune response

54]. CTLs are activated by antigen presentation through MHC-I molecules and the interaction of costimulatory molecules (B7-1 and B7-2) with CD28 [55, 56]. Conversely, inhibitory signals from molecules like CTLA-4, PD-L1/PD-L2, and LAG-3 regulate the activation of CTLs [57, 58]. Clinical studies have shown that higher numbers of intratumoral CD8⁺ T cells are linked to longer overall survival (OS) and disease-specific survival, regardless of other prognostic factors [59]. CD4⁺ T cells play a crucial role in regulating the immune response by boosting Ab production, attracting granulocytes to areas of inflammation, and supporting an efficient immune response by generating cytokines and chemokines [53]. These cells were categorized into effector $CD4^+$ T cells, follicular helper T (TFH) cells, and regulatory T cells (Tregs). Effector CD4⁺ T cells polarized to TH1, TH2, and TH 17 cells based on the patterns of various cytokines (See Table 2) [60, 61]. TH1 cells aid the activation of macrophages, NK cells, and CTL, while TH2 cells facilitate humoral immune responses by promoting B-cell growth and antibody production [62–64]. In B-cell NHL, the expression of both TH1 and TH2 cytokines at high mRNA levels has been reported [65]. Traditionally, the TH1 immune response is considered more effective at promoting antitumor immunity, while the TH2 immune response may support tumor growth by promoting angiogenesis and inhibiting the TH1-mediated immune response [66]. Higher levels of IL-4, indicative of a TH2 response, are associated with longer survival, while increased levels of IL-12, a cytokine involved in TH1 immunity, are linked to poorer prognosis in certain NHL types, suggesting that malignant B cells can modulate the effects of TH1 and TH2 cells in different lymphoma types [52, 67]. Besides, defects in TH17 cells have been observed in B cell NHL [52, 68, 69]. TFH cells play specific roles in B-cell clonal selection, maturation, and differentiation into memory cells or plasma cells within germinal centers [70]. TFH cells also facilitate B-cell activation, prevent malignant B cells from undergoing spontaneous apoptosis, and stimulate the proliferation of lymphoma cells [52, 71]. Tregs have a critical role in cancer by restricting immune activation and specific



Fig. 2 Schematic representation of tumor microenvironment (TME) constituents in non-Hodgkin lymphomas (NHLs). The TME comprises cellular and noncellular components. The cellular microenvironment consists of immune and nonimmune cells that can play pro- or antitumorigenic roles within the NHL milieu (see Table 2 for more details about the pro-tumorigenic and anti-tumorigenic cells in the TME). CTLs, effector CD4.⁺ T cells (TH1, TH2, TH17, TFH, and Treg cells), follicular dendritic cells (FDCs), natural killer (NK) cells, tumor-associated macrophages (TAMs), tumor-associated neutrophils (TANs), and myeloid-derived suppressor cells (MDSCs) constitute the immune microenvironment. On the other hand, mesenchymal stromal cells (MSCs) and cancer-associated fibroblasts (CAFs) are involved in nonimmune microenvironments. The noncellular components include the extracellular matrix (ECM) as well as various cytokines, chemokines, and molecules produced by cancerous and noncancerous cells and executed by these cells to induce their stimulatory or inhibitory effect on bystander cells (see Sect. "TME in NHLs: Composition and Functions" for more details about the interaction between cells inside the TME)

immune responses [72]. In lymphoma biopsy samples, Treg cells are abundant and have been shown to suppress antitumor immunity by inhibiting other intratumoral $CD4^+$ and $CD8^+$ T-cell populations. TGF- β produced by lymphoma cells can stimulate the expression of FoxP3, a specific marker for Tregs, which can result in the conversion of CD4 + /CD25- T cells into Tregs [73].

In addition to T cells, the immune TME in NHLs also contains tumor-associated macrophages (TAMs), tumor-associated neutrophils (TANs), and myeloid-derived suppressor cells (MDSCs). TAMs are classified to anti-tumorigenic (M1 macrophages) and pro- tumo-rigenic (M2 macrophages) [74]. M1 macrophages support the growth and differentiation of TH1 cells and NK cells, trigger CTLs cytotoxicity, and mediate ADCC [75–77]. Conversely, M2 macrophages hinder antitumor immunity and promote TH2 and Treg function, as well as CTL suppression [76, 78]. In B-cell lymphoma, TAMs play an important role in tumor progression, drug resistance, and recurrence via multiple mechanisms [78]. Similarly, in the early stages of tumor formation,

TANs primarily exhibit the N1 phenotype in the presence of INF-β resulting in CTL activation and recruitment and triggering ADCC. However, as the tumor progresses, TGF- β triggers the transition of TANs to the N2 phenotype. Neutrophils with an N2-like phenotype accompanied by tumor proliferation, blood vessel formation, extracellular matrix (ECM) degradation, and hinder T-cell activation. In addition to their functions, the N1 phenotype and N2 phenotype differ from each other in terms of cell surface markers (See Table 2) [50, 78-81]. MDSCs are a diverse group of cells that include monocytic MDSCs (M-MDSCs) and granulocytic MDSCs (PMN-MDSCs). MDSCs suppress CD4+T cells, CD8+T cells, and NK cells through direct cell contact and the production and activation of inhibitory molecules. Furthermore, MDSCs regulate the expansion and activation of Tregs, support tumor angiogenesis and metastasis, and can transform into TAMs at the tumor site. A high prevalence of the M-MDSC subpopulation has been linked to disease progression and decreased OS in B-cell NHL patients [50, 52, 73, 82].

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Table 2 Overview of pro-tumorigenic and anti-tu

Anti-tumorigenic cell	S		
TH1 cell	• Markers: CD4 ⁺ , TCR ⁺ , CXCR3 ⁺ , CCR5 ⁺	TH17 cell	Markers: CD4 ⁺ , TCR ⁺ , CCR6 ⁺
	Polarization cytokines: IL-12, IFN-I		• Polarization cytokines: IL-6, IL-1β, TGF-β
	• Mechanisms: Produce cytokines (IFN-a, TGF-β, IL-2, INF-y), Increase cells function (CTLs and macrophages)		• Mechanisms: Produce cytokines (IL-17, IL-22), Increase inflammation
M1 macrophage	• Markers: CD68 ⁺ , CD80 ⁺ , CD86 ⁺	N1 neutrophil	• Markers: CD16 ⁺ , CD66b ⁺ , CD170 ^{low}
	• Polarization cytokines: IFN-y, TGF-q, GM-CSF		ullet - Polarization cytokines: INF- eta
	Mechanisms: Produce cytokines (IL12, IL-6, IL-1β, IL-23, TNF-a), Produce chemokines (CXCL9, CXCL10), Increase cells' function (TH1 cells, CTLs, and NK cells), Increase inflammation		Mechanisms: Produce cytokines (IL-12, TNF-a, GM-C5F), Produce chemokines (CXCL9, CXCL10, CCL-3), Produce ICAM-1, produce arginase, Increase CTL activation and recruitment, Mediate ADCC
NK cell	• Markers:	СГ	Markers: CD8 ⁺ , TCR ⁺
	Immature NK cells: CD56 ^{bright} , CD16 [–] Mature NK cells: CD56 ^{dim} , CD16+ • Polarization cytokines: -		• Polarization cytokines: -
	Mechanisms: Produce cytokines (IFN-y, TNF-q, GM-CSF, IL-10, IL-5, and IL-13), Pro- duce chemokines (MIP-1d, MIP-16, IL-8, RANTES), Produce perforin and granzymes, Mediate ADCC, Mediate missing-self mechanisms		• Mechanisms: Produce cytokines (IFN-y, TNF-ol, Produce death-induces granules (granzymes, perforin, cathepsin C, and granulysin), Express death ligand (FASL)
Pro-tumorigenic cells			
TH2 cell	• Markers: CD4 ⁺ , TCR ⁺ , CCR34 ⁺ , CCR8 ⁺	Treg	• Markers: CD4 ⁺ , CD25 ⁺ , CD45RB ⁺ , FoxP3 ⁺ , CTLA-4 ⁺ , TIM-3 ⁺
	• Polarization cytokines: IL-4		\cdot Polarization cytokines: $TGF-eta$
	Mechanisms: Produce cytokines (IL-4, IL-5, IL-6), Increasing antibodies production, Inhibit TH1 function, Promote angiogenesis		\bullet Mechanisms: Inhibit cells' function (CD4+ T cells, CD8+ T cells, B cells, NK cells)
M2 macrophage	• Markers: CD163 ⁺ , CD204 ⁺ , CD206 ⁺	N2 neutrophil	• Markers: CD66b ⁺ , CD11b ⁺ , CD170 ^{high}
	\cdot Polarization cytokines: IL-10, TGF- β		\cdot Polarization cytokines: TGF- β
	Mechanisms: Produce cytokines (IL-10, IL-13, IL-4, TNF- β, TGF-B), Produce chemokines (CCL17, CCL18, CCL22, CCL24), Increase TH2 and Tregs function, Inhibit CTLs function		Mechanisms: Produce cytokines (VEGF), Produce chemokines (CXCL1, CXCL2, CXCL8, CXCL8, CXCL16, CXCR4, CCL3, CCL3, CCL4, CCL8, CCL17), Produce MMP-9, Increase tumor proliferation, Increase angiogenesis, Mediate ECM degradation
FDC	• Markers: CD23 ⁺ , CD21 ⁺ , CD35 ⁺	MDSC	• Markers: PMN-MDSCs: CD11b ⁺ , CD14 ⁻ , CD15 ⁺ , CD66 ⁺
			M-MDSCs: Lin ⁻ , CD11b ⁺ , CD14 ⁺ , HLA-DR ^{low}
	 Polarization cytokines: - 		• Polarization cytokines: -
	Mechanisms: Produce mRNA-181a, Decrease Bim levels, Inhibit malignant cell apoptosis		Mechanisms: Produce cytokines (IL-17), Produce inhibitory molecules (Arg1, iNOS, ROS, PGE2), Inhibit cells, function (CD4+T cells, CD8+T cells, and NK cell), Increase Treg function, Increase angiogenesis and metastasis
MSC	• Markers: CD44 ⁺ , CD166+, CD105+, CD90+, CD45 ⁻ , CD34 ⁻	CAF	• Markers: α-SMA ⁺ , S100A4 ⁺ , FAP ⁺ , CD10 ⁺
	• Polarization cytokines: -		Polarization cytokines: -
	Mechanisms: Produce cytokines (IL-8, TNF-a, IL-18, TGF-8), Produce chemokine (CXCL12, CXCL13, CCL19, CCL2), Produce arginase, increase tumor growth, Recruit- ment monocytes, macrophages, and neutrophils to TME, increase malignant cells homing and adhesion, Transformation to CAFs		• Mechanisms: Produce cytokine (IL-8), Produce chemokine (CCL2), Increase tumor growth, increase angiogenesis, Recruitment monocytes, and neutrophils to TME, Decrease neutrophil cel

NK cell Natural killer cell, CTL Cytotoxic T lymphocyte, Treg Regulatory T cell, FDC Follicular dendritic cell, MDSC Myeloid-derived suppressor cell, MSC Mesenchymal stromal cell, CAF Cancer-associated fibroblast

TME also contains stromal cells such as mesenchymal stromal cells (MSCs), which reduce cell death and support tumor growth by secretion of immune molecules. Within the lymphoma TME, MSCs recruit monocytes, macrophages, and neutrophils to the tumor site. Moreover, MSCs can differentiate into cancer-associated fibroblast (CAF)-like cells and secrete diverse chemokines that contribute to the homing and adhesion of lymphoma B cells [52, 83, 84]. CAFs, as another stromal cell, influence a variety of biological processes that advance cancer, including angiogenesis as well as the production and release of growth factors, cytokines, and exosomes. CAFs actively stimulate tumor cell growth, invasion, and inflammation, and contribute to resistance to treatment [50, 52].

Exosomes are an important part of the TME. In lymphomas, exosomes can decrease NK cell-mediated cytotoxicity, trigger immune cell death, and increase treatment resistance through the delivery of various molecules such as interleukins, PGE2, TGF-B, and microR-NAs. Additionally, tumor-derived exosomes expedite the activation and growth of MDSCs [50, 52, 85, 86]. Apart from exosomes, chemokines and cytokines present in the TME can also support tumor growth and development. Many studies have shown that the serum level of soluble IL-2R α , which is produced by CD4⁺ CD25⁺ T cells, is greater in B-cell NHL patients and is associated with a poorer prognosis [73, 87]. The secretion of TGF- β malignant B cells leads to the suppression of TH1 and TH17 cell growth and hinders the proliferation of T cells [73, 88]. Serum IL-10 levels have been demonstrated to be increased in B-cell NHL patients and to be negatively correlated with prognosis [73]. In addition, chemokines released by MSCs in lymphoma, including CXCL13, CCL19/CCL21, and CXCL12, facilitate B-cell adhesion and homing [52].

NK cell defects in NHL

The NK cell defects in the TME of NHL include quantitative deficiency, distribution abnormalities, functional deficiency, the presence of an immunosuppressive TME, and tumor cell escape from NK cell surveillance. Figure 3 summarizes NK cell defects in the TME of NHLs.

Quantitative deficiency

The absolute NK cell count (A-NKC) decreases in (Diffuse large B cell lymphoma) DLBCL patients. According to Plonquet et al. investigation, one-third of DLBCL patients who present with 2 or 3 adverse prognostic factors of aaIPI have low NKC at diagnosis. Furthermore, they established that NKC was associated with a poorer response to treatment and shorter event-free survival (EFS) [89]. Similarly, flow cytometry analysis of CD3⁻CD56⁺ and/or CD16⁺ cells in the peripheral blood of DLBCL and (follicular lymphoma) FL patients demonstrated that both DLBCL and FL patients had low NKC, which was correlated with a reduction in progressionfree survival (PFS) and OS [90]. Furthermore, a deficit in NKC was detected in other NHLs, such as primary central nervous system lymphoma (PCNSL). Lin et al. conducted a study on 161 patients with PCNSL and found that individuals who responded to treatment had a higher median of circulating NKC and NK cell proportion compared to those who did not respond. Their study revealed that PCNSL patients who have a higher baseline NKC display longer OS than those with a low NKC [91]. Consequently, NKC in NHL patients represents a prognostic biomarker for the assessment of clinical outcomes.

Functional deficiency

Cancer cells have been observed to employ a wide range of mechanisms to escape from the innate immune pressure exerted by NK cells, including abnormalities in NK cytotoxicity function [92]. In the context of NHLs, NK cells have impaired degranulation capacity. According to previous studies, NK cells exhibit defects in the production and exocytosis of cytotoxic granules containing perforin and granzyme [93, 94]. In a protein quantification study involving 12 patients with NHL, it was discovered that while the gene expression levels of perform and granzyme B were higher in NHL patients compared to the control group, the intracellular levels of perforin in NK cells were lower in NHL patients [93]. Furthermore, a separate study conducted by Baychelier et al. found that patients who developed NHL after undergoing lung transplantation exhibited an accumulation of NK cells with low expression of perforin and impaired degranulation against NHL target cells [94]. Other aspects of NK cell hypofunction include decreased cytokines production, e.g. IFN- γ and TNF- α , the overexpression of suppressor receptors such as T-cell immunoreceptors with immunoglobulin and ITIM domains (TIGIT), and the decreased expression of activating receptors such as TIM-3 [95]. It should be noted that TIGIT has been associated with NK cell exhaustion [96]. In addition, decreased CD16 expression in NK cells and impaired ADCC activity were observed in newly diagnosed and refractory NHL patients [95, 97].

NK cells in refractory NHL exhibit downregulated expression of activating receptors, including NKp30, NKp46, and NKG2D. Further investigation revealed that de novo NHL development was correlated with increased NKG2A and CD62L expression but reduced inhibitory KIR and CD57 receptor expression [94]. Essa et al. demonstrated that DLBCL patients with advanced stages of the disease have significantly lower NKp44 levels than



Fig. 3 Possible mechanisms of natural killer (NK) cell defects in non-Hodgkin lymphomas (NHLs). The NK cell defects within the TME of NHLs include functional deficiency, the presence of an immunosuppressive TME, and tumor cell escape from NK cell surveillance (as well as quantitative deficiency and distribution abnormalities that are not shown in Fig. 2). See Sect. "NK cell defects in NHL" for more information). A) NK cell functional defects include decreased expression of activating receptors, overexpression of suppressor receptors, decreased ADCC mechanisms, decreased IFN-γ production, and impaired degranulation capacity. B) Immunosuppressive cells such as myeloid-derived suppressor cells (MDSCs), regulatory T cells (Tregs), and M2 macrophages hinder NK cell function through the production of immunosuppressive factors or the expression of inhibitory receptors. C) NHL cells evade NK cell-mediated cytotoxicity via resistance to the perforin/granzyme-mediated apoptosis pathway, resistance to death receptor-mediated apoptosis pathways, and inhibition of NK cell activation

patients with earlier stages of DLBCL. This decrease in NKp44 may be attributed to the high level of IL6 and TGF- β in the advanced stages of the disease, which in turn downregulate NK activating receptors [98]. The expression of CD16 and NKG2D activating receptors on the surface of CD56^{dim} cells was also reported to be decreased after rituximab treatment [99] (Fig. 3A).

Immunosuppressive TME

Successful interaction between NK cells and dendritic cells (DCs) and the production of chemokines are required to induce effective antitumor immunity by NK cells (Fig. 1). This process is negatively affected by TME, especially cellular and soluble components of the TME, which are associated with the escape of cancer cells due to the lack of effective immune responses [100]. Several immune suppressive cells, like MDSC, TAMs, and Tregs negatively interfere with NK cell activation and function. In a phase 2 clinical trial conducted by Bachanova et al.,

the frequencies of MDSCs and Tregs were investigated about adoptive NK cell therapy response in patients with NHL. Results from the trial indicated that patients who exhibited higher frequencies of MDSCs and Tregs, along with the adoptive NK cells, had a poorer response to therapy [95]. Similarly, Sato et al. demonstrated that the accumulation of MDSCs leads to NK cell depletion in NHL patients [101]. Increased numbers of MDSCs were reported in DLBCL, MZL, MCL, high-grade B-cell lymphoma (HGBL), PCNSL, and FL. Interestingly, MDSCs are markedly increased in high-grade NHLs and may be a potential prognostic marker [102]. The inhibitory effect of MDSCs on NKG2D expression and IFN-y production in NK cells was confirmed in both in vivo and in vitro experiments [103]. The downregulation of other types of NK cell activating receptors, such as NKp30 and NKp46, was also detected [104, 105]. Further analysis in a murine lymphoma model revealed that MDSCs, which can secrete IL-10, reduced the frequency of NK cells [101]. Additionally, the coculture of MDSCs with NK cells has been shown to negatively affect the degranulation capacity of these cells through the TIGIT/CD155 pathway [105].

Tregs are another immunosuppressive cell type that limits adoptive NK cell therapy in NHL patients. Increased numbers of Tregs expressing high levels of Foxp3 following high-dose chemotherapy and IL2 administration before adaptive NK cell infusion interfere with NK cell expansion [106]. Treg infiltration in the TME could be justified via Indoleamine-2,3-dioxygenase (IDO). IDO is an immunosuppressive enzyme that catalyzes the conversion of tryptophan to kynurenine [107]. NHL patients who overexpress IDO simultaneously exhibit increased levels of FoxP3, a Treg marker [108]. In addition, IDO not only inhibits NK cell proliferation but also decreases activating receptors [107]. In a study conducted by Ninomiya et al., it was found that 32% of DLBCL patients exhibit overexpression of IDO, which is associated with unfavorable clinical outcomes [109]. Additionally, Yoshikawa et al. reported elevated levels of tryptophan-derived kynurenine in DLBCL patients [110].

Crosstalk between M2 macrophages and NK cells is another barrier to NK cell function in the TME. M2 macrophages limit NK cells' function by triggering the expression of inhibitory receptors immunoglobulinlike transcript 2 (ILT2/ CD85j), an NK cell inhibitory receptor [111]. In NHL, a high density of M2 macrophages in DLBCL of the central nervous system (CNS-DLBCL) has been detected and accounted for poor clinical outcomes [15].

Among other immunosuppressive factors, TGF- β has been investigated in NHL, and previous studies revealed TGF- β signaling Dysregulation in mantle cell lymphoma (MCL), FL, and DLBCL. The TGF- β signaling cascade is dysregulated through various mechanisms, such as altered receptor expression, disrupted SMAD signaling, and disturbances in epigenetic and genetic processes [112]. TGF- β inhibits IFN- γ expression, affects the metabolic pathway of NK cells, and reduces NKG2D and NKp30 expression, which are essential for tumor cell recognition and elimination, as well as for the effective interaction between natural NK cells and DCs [113, 114]. Interestingly, MDSCs and M2 macrophages participate in NK cell exhaustion by producing TGF- β 1 (Fig. 3B) [103, 115].

Evasion mechanism

Resistance to apoptosis

Tumor cells in NHLs may escape from NK cell-mediated cytotoxicity through resistance to perforin/granzymemediated apoptosis. For this purpose, tumor cells may exhibit elevated intrinsic levels of proteinase inhibitor 9 (PI9), which functions to restrict the proteolytic action of granzyme B and secure their survival [116]. In this line, Bladergroen et al. verified that P19 was overexpressed in different types of T/B-NHL, such as extranodal T-cell NHL, enteropathy type T-cell NHL, NK/T-cell nasal-type lymphoma, and DLBCL [116]. Furthermore, cancer cells may escape apoptosis by inactivating apoptotic pathways activated by death receptors. The death ligands FasL/ CD95L and TRAIL, which are members of the TNF family, are expressed in NK cells. These ligands interact with their respective receptors, Fas/CD95 and TRAIL-R, present on the surface of target cells. Upon interaction, the death domain (DD) is activated, initiating the apoptotic signaling cascade and ultimately leading to apoptosis [117]. According to previous studies, loss of Fas/CD95 expression was found in some FL and diffuse B/T-cell lymphomas [118], mucosa-associated lymphoid tissue lymphomas (MALTLs) [119], and cutaneous B-cell lymphomas (CBCLs) [120], which are associated with poor prognosis. In addition, mutations in Fas/CD95 have been reported in GC-derived B-cell lymphomas, such as primary nodal DLBCL, MALT-type lymphomas, FL, and anaplastic large cell lymphoma (ALCL) [121, 122]. A somatic mutation in TRAIL-R, which is correlated with the loss of chromosome 8p21-22, has also been detected in NHLs (Fig. 3C) [123].

Inhibition of NK cell activation

CD58 or lymphocyte-function antigen 3 (LFA-3) is known as an NK cell activator that interacts with CD2 on the NK cell surface [124-126]. Based on this speculation, mutations or deletions in CD58 also prevent NK cell function, which has been reported in DLBCL and FL [125, 126]. HLA-G is an inhibitory molecule in both membrane-bound and soluble isoforms that suppresses NK cells through interaction with its ILT2 [127]. The serum level of soluble HLA-G increased in NLHs such as DLBCL, FL, and peripheral T-cell lymphoma, which may disrupt NK cell function and be involved in lymphoma development [128]. Notably, HLA-G expression in lymphoma is a double-edged sword with protective and destructive effects [129]. To explore the evasion mechanisms, strategies that disturb NK cell receptors are also considered. The investigation by Satwani et al. revealed that, incubation of NHL cells with romidepsin enhanced NK cell cytotoxicity. Subsequently, they reported that romidepsin increases the surface expression of the NKG2D ligands MIC A/B on lymphoma cells. Based on the results of this study, impairment of NK cell function may be related to decreased expression of activating receptor ligands such as MIC A/B [130]. The immune checkpoints PD-1 and PD-L1 also restrict NK cell function, and PD-1/PD-L1 axis blockade unleashes NK cell cytotoxicity [127]. Research conducted by Laurent et al. revealed that DLBCL cells exhibit notably elevated levels of PD-1 and PD-L1/2 compared to FL cells. Notably, some DLBCL tumor cells coexpress both PD-1 and PD-L1/2. Interestingly, there are more PD-L1/2-positive lymphoma cells in the activated B-cell (ABC) subtype of DLBCL (ABC-DLBCL) than in the GC subtype (GC-DLBCL) [129]. Similarly, Kiyasu et al. reported that PL-1 is frequently expressed in tumor cells in DLBCL and is associated with poor prognosis [131].

NK cell immunotherapy in NHLs

In NHL, the A-NKC of the autograft directly influences clinical outcomes of following HSCT [132]. In a randomized, double-blind phase III clinical trial, patients with NHL who received an autograft with an A-NKC $\geq 0.5 \times 10^9$ cells/kg demonstrated 5-year OS and 5-year PFS rates of 87% and 71%, respectively. In contrast, patients infused with an autograft A-NKC $< 0.5 \times 10^9$ cells/kg experienced 5-year OS rates of 55% and 5-year PFS rates of 32% [133]. With a 10.6year median follow-up in the final update, the 13-year OS rates demonstrated a significant difference between groups, with a rate of 46% for the cohort infused with autograft A-NKC $\geq 0.09 \times 10^9$ cells/kg compared to 36% for the group infused with A-NKC $< 0.09 \times 10^9$ cells/kg (P-value < 0.02) [134]. Faster and robust recovery of NK cells following HSCT is another factor that can affect clinical outcomes [135]. Porrata et al. reported that NHL patients with an A-NKC \geq 80 cells/µL on day 15 after autologous HSCT had longer OS and PFS than patients with lower counts (not reached vs 5 months, p0.001; not reached vs 3 months, p0.0001, respectively) [136]. These findings suggest that the early post-HSCT recovery of NK cells may play a crucial antitumor role in the potential graft-versus-tumor (GVT) effect, given that NK cells are the only immune effector cells that reach normal numbers and function post-HSCT [137].

Several early studies have employed the administration of low-dose subcutaneous rlL-2 to promote the recovery and cytotoxic activity of NK cells as an effective approach to eradicate residual disease and prevent relapse following autologous HSCT in NHL patients [138, 139]. In a clinical trial involving patients with R/R high-grade NHL, researchers demonstrated that the administration of a low dose of rlL2 early after autologous HSCT for a duration of one year is well-tolerated and leads to the in vivo expansion of CD16⁺/CD56⁺ NK cells. Significantly, compared to their baseline quantity and function before starting treatment, the expanded CD56^{bright} NK cell subsets exhibited enhanced activity against K562 cells (an NK-sensitive cell line) and CD16-mediated redirected killing activity against P815 target cells (an NK-resistant cell line). All ten patients who participated in the trial remained free from relapse for a period ranging from 5 to 34 months (median 16 months) after initiating rIL2 therapy. Notably, two patients who still had residual disease following HSCT experienced complete disease disappearance after rIL2 treatment [138]. Building upon the encouraging outcomes of these early studies, clinical trials have explored the adoptive transfer of ex vivo activated autologous NK cells or lymphokine-activated killer cells as a therapeutic approach for patients with lymphoma [140, 141]. The adoptive transfer of autologous NK cells was found to be feasible and safe, although only a limited antitumor effect was observed [142]. This limitation primarily stemmed from the matching of inhibitory receptors on autologous NK cells with self-MHC class I present on tumor cells, leading to "self" recognition signals that dampen NK cell activation and subsequent antitumor effects [142]. Furthermore, the adoptive transfer of autologous NK cells is costly and frequently requires multiple apheresis procedures, and the dose of injected NK cells is limited to approximately $10^7/\text{kg}$ [143]. To overcome these limitations, researchers have recently used allogeneic NK cells for lymphoma immunotherapy. In the phase 1 clinical trial conducted by Green Cross LabCell Corporation, the safety and possible efficacy of allogeneic NK cells were assessed in patients with malignant lymphoma or advanced solid tumors. In this study, allogeneic NK cells (namely, MG 4101) were obtained from random healthy unrelated donors and expanded in culture bags supplemented with IL-2, irradiated autologous feeder cells, and OKT3. Multiple doses of MG4101 were administered in the dose range of 1×10^6 cells/kg to 3×10^7 cells/kg without any signs of GVHD or serious toxicity. Among the 17 evaluable patients, only 8 exhibited stable disease (SD), while the disease progressed in the remaining patients. The median PFS for patients with SD was 4 months, ranging from 2 to 18 months [144]. The results of this study indicated that the use of alloreactive NK cells alone was not sufficient to eliminate the disease mass completely. As a result, researchers explored the combination of NK cells with other strategies to enhance their therapeutic effectiveness in subsequent studies [135].

NK cells combined with mAbs

Over the past two decades, the therapeutic effects of at least 570 monoclonal antibodies (mAbs) have been investigated in clinical trials. Among them, 79 therapeutic mAbs, including 30 mAbs for the treatment of hematological malignancies, have received approval from the United States food and drug administration (FDA) and are currently commercially available [145]. When mAbs bind to their targets, they can kill cancer cells through a

variety of mechanisms, including programmed cell death (PCD), complement-dependent cytotoxicity (CDC), and ADCC [146]. Among these mechanisms, ADCC is an effective immune mechanism that is triggered when therapeutic mAbs are employed to eliminate cancer cells [147]. During the ADCC process, the FC region of the antibody is ligated to its corresponding FC receptor (FcR) on the plasma membrane of immune effector cells, while the Fab portion of the antibody attaches to target antigens on the surface of the cancer cell [148]. Human NK cells serve as crucial effector cells in the context of ADCC by expressing CD16A, which is a low-affinity receptor for IgG1 and IgG3 antibodies [149]. Given the likelihood that the efficacy of ADCC-mediated tumor cell elimination relies on the ratio of effector to target cells, the number and function of NK cells have been investigated as potential biomarkers to predict the response to anti-CD20 immunotherapy in NHL patients [90, 150]. Klanova et al. reported that low peripheral blood NKC in FL and DLBCL patients receiving anti-CD20 mAbs (rituximab or obinutuzumab) plus chemotherapy were linked to shorter PFS in both FL and DLBCL patients and diminished OS specifically in FL patients [90] Hence, the number of NK cells in individuals with lymphoma is important for determining their prognosis [151].

The administration of adoptive NK cells to enhance the ADCC capabilities of mAbs is a growing area of intervention that has been explored in recent years [152]. There are several ongoing and completed clinical trials exploring the safety and effectiveness of combining mAbs with infusions of autologous or allogeneic NK cells in patients with NHL (Tables 3 and 4). In a recent phase I study, Tanaka et al. investigated the infusion of ex vivo-expanded autologous NK cells in combination with rituximab-containing chemotherapy in patients with relapsed CD20⁺ malignant lymphoma [153]. Expanded autologous NK cells with high expression of NKp30, NKp44 and CD16 were intravenously infused (up to 10×10^6 cells/kg) into lymphoma patients one day after rituximab-combined salvage chemotherapy. The combination was safe and feasible, and among the nine lymphoma patients, seven achieved complete response (CR), with a median duration of 44 months (range: 6-56 months). However, it is difficult to determine the precise contribution of autologous NK cells to the clinical response, given that chemotherapy was administered to eight of nine patients after NK cell infusion [153]. In another study on chemotherapy-refractory NHL patients, allogeneic NK cell therapy (dose of $0.5-3.27 \times 10^7$ cells/kg) in combination with IL-2 and rituximab was found to be safe and effective in 4 of 15 evaluable patients, with 2 patients achieving CR lasting 3 and 9 months and 2 patients obtaining partial response (PR) [95]. Moreover, in a recent phase I study employing ex vivo-expanded allogeneic NK cells (namely, MG4101) plus rituximab after lymphodepleting chemotherapy for R/R NHL patients, Yoon et al. demonstrated that the treatment was well tolerated and led to a PR in 4 patients and a CR in 1 patient, yielding an overall response rate (ORR) of 55.6% [154]. Notably, one patient achieved a lasting CR that extended beyond 806 days [154].

In addition to autologous or allogeneic peripheral blood (PB)-derived NK cells, an increasing number of clinical trials have scrutinized the safety and efficacy of other NK cell sources, including cord blood (CB) [6, 155, 157], induced pluripotent stem cells (iPSCs) [158], and immortalized NK cell lines [162], for NHL immunotherapy. For example, in a phase 1/2 clinical trial, the safety and clinical activity of AB-101 (an allogeneic, nongenetically modified, CB-NK cell product) has been evaluated as a monotherapy and combined with rituximab for the treatment of R/R NHL patients [155]. The results from this study indicated that the concurrent administration of both agents was safe, resulting in an ORR of 67% in 6 patients (CR observed in 3 patients and PR in 1 patient), in contrast to an ORR of 27% in cohorts receiving AB-101 alone [155]. Another study by Katayoun Rezvani's group assessed the efficacy of ex vivo-expanded CB-NKs in combination with rituximab and high-dose chemotherapy in NHL patients who were candidates for autologous HSCT [157]. Patients received rituximab and high-dose chemotherapy from days 13 through 7, lenalidomide from days 7 through 2, and CB-NK cells (10⁸/kg) on day 5 before to autologous HSCT. CB-NK cells were detectable in vivo for two weeks, regardless of their HLA mismatch status. Importantly, no adverse events attributable to the CB-NK cells were observed. At a median follow-up of 47 months, the rates of relapse free survival (RFS) and OS were 53% and 74%, respectively [157].

NK-92 is an immortalized IL-2-dependent CD16⁻ NK cell line that was isolated and successfully established by Klingman et al. in 1992 from a patient suffering from lymphoma. NK-92 cells exhibit potent cytotoxicity against several cancer cells, a phenomenon primarily ascribed to the overexpression of numerous activating receptors, concurrent downregulation of almost all inhibitory receptors, and heightened expression of perforin and granzyme. Furthermore, NK-92 cells can continuously proliferate with a doubling time of 2-4 days, are easily obtainable, and have a homogeneous phenotype [163, 164]. However, due to their cancerous nature, NK-92 cells must be mitotically inactivated prior to infusion into patients to inhibit undesired clonal proliferation, which restricts their persistence and expansion in vivo, and allogeneic administration demands very high doses of NK-92 cells [165]. In 2008, Arai et al. demonstrated for the first time the feasibility and safety of administering NK-92

Ta	ble 3 Ongoin <u>c</u>	g clinical trials	s of NK cell therap	oies for NHL						
#	NCT Number	Phase/N	Trial status	Study Design	Conditions	Interventions	Doses	Cells source	Outcome Measures	Sponsor/Location
-	NCT03019666	Phase I/39	Completed	Nonrandomized, parallel assign- ment, open label	MM and NHLs	Cyclophospha- mide + fludara- bine + NAM-NK + elo- tuzumab (for MM) or rituximab (for NHL) + IL-2	Not disclosed	8	Primary: safety Secondary: occur- rence of TRM, occurrence of dis- ease response and number of patients alive without progres- sion	Masonic cancer center, university of Minnesota/USA
7	NCT02259348	Phase II/12	Terminated	Single group assignment, open label	ALL, AML, CML, JMML, MDS and NHL	Cyclophospha- mide + fludara- bine + IL-2 + mel- phalan + thi- otepa + rituximab + NK cells + T-cell depleted HPCT + CD45RA- depleted HPCT + G-CSF	Not disclosed	Not disclosed	Primary: percent- age of partici- pants engrafted by day 42 post- transplant Secondary: inci- dence of malig- nant relapse, EFS, OS, incidence and sever- ity of aGVHD and cGVHD and rate of TRM	St. Jude children's research hospital/ USA
m	NCT02280525	Phase //8	Completed	Nonrandomized, single group assignment, open label	Leukemia and lymphoma	Lenalidomide + fludara- bine and cyclophos- phamide for CLL and low-grade lymphoma + rituximab (for patients with B-cell cancer) + NK cells	1×10 ⁷ cells/kg	nG	Primary: MTD of NK cells with lenalidomide and lymphode- pleting chemo- therapy Secondary: response rate of NK cells with lenalido- mide and lym- phodepleting chemotherapy determined by BMA/BMB	M.D. Anderson can- cer center/USA

		(
#	NCT Number	Phase/N	Trial status	Study Design	Conditions	Interventions	Doses	Cells source	Outcome Measures	Sponsor/Location
4	NCT01181258	Phase IV16	Completed	Nonrandomized, single group assignment, open label	NHL and CLL	rituximab, fludarabine, cyclophosphamide, methylpredniso- lone + NK cells + 11.2	1.5 to 8×10 ⁷ cells/kg	8	Primary: number of patients with a PR or CR Secondary: SAEs, time to disease progression and patients with expansion of NK cells	Masonic cancer center, university of Minnesota/USA
Ś	NCT00625729	Phase I, II/6	Terminated	Single group assignment, open label	Relapsed NHL and CLL	Fludarabine + cyclo- phospha- mide + rituximab + NK cells + aldesleukin (rll-2) + G-CSF	1.5 to 8×10 ⁷ cells/kg	8	Primary: NK cell expansion Secondary: num- ber of patients with IL-15 pro- duction and NK cell expansion, OR, number of patients whose disease progressed after treat- ment, number of patients with adequate NK cells infused and OS	Masonic cancer center, university of Minnesota/USA
9	NCT00383994	Phase I/, 6	Completed	Single group assignment, open label	B-cell lymphoid malignancies	GM-CSF + rituxi- mab + NK cell	Not disclosed	Not disclosed	Primary: DLT for NK cells infu- sions	M.D. Anderson can- cer center/USA
\sim	NCT03019640	Phase II/22	Completed	Single group assignment, open label	R/R B-Cell NHL	Carmustine + etopo- side + cytarabine + mel- phalan + lena- lidomide + rituximab (for patients who are CD20 ⁺) + NK cell + HSCT + filgrastim	Not disclosed	UCB	Primary: TRM30 Secondary: num- ber of participants who survived on day 180	M.D. Anderson can- cer center/USA
∞	NCT02843061	Phase I, II/30	Completed	Randomized, par- allel assignment, open label	Recurrent B-cell lymphoma	Rituximab + NK cells	Each time 10 billion cells, 4 times in all	Not disclosed	Primary: relief degree evaluated by the RECIST Secondary: PFS and OS	Fuda cancer hos- pital, Guangzhou/ China

Table 3 (continued)

Ta	ble 3 (continu	(pər								
#	NCT Number	Phase/N	Trial status	Study Design	Conditions	Interventions	Doses	Cells source	Outcome Measures	Sponsor/Location
6	NCT04074746	Phase I, II/30	Active, not recruit- ing	Single group assignment, open label	R/R CD30 ⁺ HL or NHL	Fludarabine + cyclo- phospha- mide + AFM13- NK + AFM13	Not disclosed	UCB	Primary: SAEs Secondary: OS, EFS, ORR, dura- tion of response and immune reconstitution studies	M.D. Anderson Can- cer Center/USA
0	NCT02890758	Phase //14	Completed	Nonrandomized, single group assignment, open label	NHL, AML, MDS, ALL, CLL, CML,	NK cell infu- sion + ALT803	Dose escalation from 1 × 10 ⁶ to 5 × 10 ⁷ cells/kg	Not disclosed	Primary: MTD and number of patients with- out GVHD Secondary: num- ber of patients response cally measurable lesions, patients with malignant lymphoma response, patients response, patients response durar- average durar- of response, and in vivo NK levels and in vivo NK levels	Brenda Cooper / USA
=	NCT01386619	Phase I, II/15	Completed	Single group assignment, open label	AML, MDS, lymphoma, NB and RMS	HSCT + NK DLI	>1×10 ⁷ NK cells/ kg	PB	Primary: feasibility and safety Secondary: efficacy	University hospital, Basel, Switzerland/ Germany and Swit- zerland

Tab	ole 3 (continu	ued)								
#	NCT Number	Phase/N	Trial status	Study Design	Conditions	Interventions	Doses	Cells source	Outcome Measures	Sponsor/Location
12	NCT01287104	Phase 1/34	Completed	Sequential assign- ment, open label	CML and HD	Stem cell + NK cell	1 × 10 ⁵ to 1 × 10 ⁷ cells/kg	8	Primary: number of patients who received 2 doses of NK cell infu- sions and number of patients who received the high- est dose level of NK cells Secondary: number of partici- pants with mild, moderate and/ or severe cGVHD, DFS, OS since date of transplant, occurrences of viral infection and/or reactiva- tion, decline in IL-7 and IL-15, cell numbers postransplant, number of partici- pants with pres- ence of KIR gene mismatch, SAEs and non-SAE	National cancer institute/USA
13	NCT00990717	Phase I/8	Completed	Single group assignment, open label	NHL, HL, leukemia	Irradiated NK-92 cells	Level I: $1 \times 10^{\circ}$ cells/m ² Level II: 3×10^{9} cells/m ² Level III: 5×10^{9} cells/m ²	NK-92	Primary: DLT Secondary: efficacy, immune response directed against the NK-92 cells and kinetics of NK92 cells	University health network, Toronto/ Canada
<u>4</u>	NCT00823524	Phase I, II/47	Completed	Single group assignment, open label	Lymphoma, leukemia, MDS, myeloma, brain and Solid tumors and solid tumors	HSCT + NK cell	Not disclosed	PB	Primary: safety Secondary: efficacy	Asan medical center/ South Korea

	Sponsor/Location	Fred Hutchinson cancer center/USA iD, m- int ed ed	St. Jude children's - research hospital/ ell USA ie- nc-	Tufts medical center/USA ills ce ce ce
	Outcome Measures	Primary: DLT, number of participants with relapsed c ease and aGVH number of non lapse participa who experienc graft failure Secondary: number of sub jects surviving posttransplant and number of participants who experienc cGVHD	Primary: Safety Secondary: effi cacy and NK ce persistence, ph notype and fur tion of NK cells after infusion	Primary: safety Second- ary: duration of donor NK ce in the recipient blood, patient survival at 100 days and at on year post treat- ment, occurren of new cancer during the first. year post infusi and systemic infections dur-
	Cells source	æ	ЪВ	В
	Doses	Not disclosed	Not disclosed	Not disclosed
	Interventions	Fludarabine + cyclo- phospha- mide + TBI + HSCT + NK cells + mycopheno- late + tacrolimus	Clofarabine + cyclo- phosphamide + etopo- side + NK cell + IL-2	AHSCT + NK cell
	Conditions	NHI, ALL, AMI, CLL, CML CML	NHL, MDS, CML, JMML,	NHL, HL, MM, AML
	Study Design	Single group assignment, open label	Nonrandomized, parallel assign- ment, open label	Nonrandomized, single group assignment, open label
	Trial status	Completed	Completed	Completed
led)	Phase/N	Phase I, II/41	Phase I/48	Phase I/13
able 3 (continu	NCT Number	NCT00789776	NCT00697671	NCT00660166

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#	NCT Number	Phase/N	Trial status	Study Design	Conditions	Interventions	Doses	Cells source	Outcome Measures	Sponsor/Location
8	NCT00586703	Phase 1/21	Completed	Single group assignment, open label	Lymphoma	Nonmyeloablative ASCT+NK cell	Up to 1 × 10 ⁷ CD56 ⁺ cells/kg	BL	Primary: safety Secondary: effi- cacy of the regi- men in terms of OS	David Rizzieri, MD/ USA
6	NCT01619761	Phase I/13	Unknown	Nonrandomized, parallel assign- ment, open label	NHL, HL, ALL, AML, MDS, CML, CLL, Myeloma	Lenalidomide + fludara- bine + melpha- lan + rituximab (for CD20 positive patients) + NK cells + UCB-HSC+ tac- rolimus + mycophe- nolate	Not disclosed	nce	Primary: genera- tion of a minimum of 5 × 10 ⁶ NK/kg cells in at least 60% of patients, treatment-related mortality and inci- dence of SAEs Secondary: proportion of patients with aGVHD and cGVHD, OS, DFS, time to initial platelet trecovery and time to initial absolute neutrophil count recovery	M.D. Anderson can- cer center/USA
20	NCT04673617	Phase I, II/108	Recruiting	Nonrandomized, sequential assign- ment, open label	NHL	Cyclophospha- mide + fludara- bine + rituxi- mab + bendamus- tine + NKcell + IL-2	Not disclosed	UCB	Primary: safety and tolerability, ORR, identify the R2PD and effi- cacy	Artiva biotherapeu- tics, Inc/USA
21	NCT03778619	Phase I, II/9	Unknown	Single group assignment, open label	R/R NHL	Fludarabine + cyclo- phosphamide + NK cells (MG4101) + IL-2	Group 1: 1 × 10' cells/Kg Group 2: 3 × 10' cells/Kg Group 3: 9 × 10' cells/Kg	B	Primary: at phase I MTD and at phase II ORR Secondary: at phase I: ORR at phase II: CR, PR, OS, time to pro- gression and time to response	GC cell corporation/ South Korea

#	NCT Number	Phase/N	Trial status	Study Design	Conditions	Interventions	Doses	Cells source	Outcome Measures	Sponsor/Location
22	NCT04023071	Phase 1/72	Terminated	Nonrandomized, parallel assign- ment, open label	B-cell lymphoma and AML	Cyclophospha- mide + fludara- bine + obinutuzumab or rituximab (for B-cell lymphoma) + NK cell (FT516) + IL-2	Not disclosed	iPSC	Primary: DLT and incidence, nature and sever- ity of AEs Secondary: anti-tumor activity of FT516 and FT516 phar- macokinetic data	Fate therapeutics/ USA
NA/ trar Bor rest	M-NK Nicotinamid Isplantation, ASC7 Ie marrow biopsy, Jonse, RFS Relapse eria in solid tumor	e expanded-nai Autologous Stu OR Overall resp : free survival, <i>C</i> s, <i>H</i> R Hematoloo	tural killer, <i>IL</i> Interleuki em Cell Transplant, G-C zonse, <i>ORR</i> Overall rest, <i>J</i> ^F S Disease-free Surviv gical response, <i>WM</i> We	n, PB Peripheral blood, L 2F Granulocyte colony 5D conserate, OS Overall su al, cGVHD Chronic graft aldenstrom's macroolobi	/CB Umbilical cord blo stimulating factor, AE A rvival, EFS Event-Free S versus host disease, DL ulinemia, CL Cutaneou	od, <i>iPSC</i> Induced pluripoten ddverse event, <i>SAE</i> s Serious / burvival, <i>PFS</i> Progression-free <i>IT</i> Dose limiting toxicities, <i>TF</i> s lymbhomas, <i>MM</i> Multiple.	t stem cell, <i>TRM</i> Treatr Adverse Events, <i>MTD</i> N e survival, <i>aGVHD</i> Acu <i>8M30</i> Treatment-relate mveloma, <i>CML</i> Chroni	ment related morta Maximum tolerated ite graft-versus-hos ed mortality within ic mveloid leukemia	lity, <i>H5CT</i> Hematopoie 1 dose, <i>BMA/BMB</i> Bone 1t disease, <i>CR</i> Complete 30 days, <i>RECIST</i> Respoi	tic stem cell marrow aspiration/ s response, <i>PR</i> Partial sse evaluation sstic leukemia, <i>AML</i>

Table 3 (continued)

Acute myeloid leukemia, MDS Myelodysplastic syndrome, JMML Juvenile myelomonocytic leukemia, NHL Non-Hodgkin lymphoma, HL Hodgkin lymphoma, CLL Chronic lymphocytic leukemia, NB Neuroblastoma, RMS Rhabdomyosarcoma, R2PD Recommended phase 2 dos, KR Killer-cell immunoglobulin-like receptors, DLI Donor lymphocyte infusion, CNS Central nervous system

Table 4 Outcomé	e of clinical trials	of NK cells in 	NHL patients							
# Study	Conditions/n	Cells source	Culture methos	Final product Characteristics	Infused dose of NK cells	Chemotherapy regimen before NK cell infusion	Combination therapy	Adverse event/Toxicity	Clinical response	Reference
1 Phase I, II (NCT04673617)	R/R NHL includ- ing DLBCL (n = 5), RL (n = 2), LPL/WM (n = 1)	ncB	Not disclosed	Not disclosed	1 × 10 ⁹ cell/dose or 4 × 10 ⁹ cell/ dose	Cyclophos- phamide and fludarabine	Rituximab	Grade 1 CRS in 2 patients, grade 1 fever in 1 patient, infections in 3 patients, PD in 3 patients	67% (4/6) ORR in rituximab cohort, 27% (3/11) ORR in monother- apy (NK cells) cohort	Khanal et al. [155]
2 Phase II (NCT01181258)	DLBCL (<i>n</i> = 11), MCL (<i>n</i> = 2), FL (<i>n</i> = 1), CLL (1), LPL/WM (<i>n</i> = 1)	8	CD3/CD19- depleted cells were cultured in X-VIVO media sup- plemented with 10% human AB serum and 1000 U/mL IL-2 for 16–18 h	Viability > 70%, NK cell content ≥ 20%, T-cell cont tent ≤ 3.0 × 10 ⁵ /kg, B-cell content < 3% and endo- toxin < 5EU/kg	0.53.2.7 × 10 ⁷ / kg in association with IL-2	Initially, the regi- men consisted of pentostatin, cyclophos- phamide, and denlieukin diffitiox, but it was later revised twas later revised fludarabine, cyclophospha- mide, and meth- ylprednisolone	Rituximab	No symp- toms or signs of GVHD, neurotoxicity, or persistent marrow aplasia were reported	Of 14 (29%) evaluable patients, 4 had (29%) objec- tive responses (including 2 CR and 2 PR) at two months	Bachanova et al. [95]
3 phase 1 (NCT03019666)	R/R NHL includ- ing DLBCL (n = 9), FL (n = 10), MCL (n = 1)	8	CD3 depleted cells were culture for 14 days with IL-15 (20 ng/ml) and NAM (5 mM)	Viability > 97.46%, mean T-cell con- tent 0.7%, mean NK cell content 96.12% and mean B-cell content 0.58%	Three dose level: 2 × 10 ⁷ , 1 × 10 ⁸ or 2 × 10 ⁸ cells/ kg in association with IL-2	Cyclophos- phamide and fludarabine	Rituximab	There was no evi- dence of CRS, neurotoxicity, GVHD, or mar- row aplasia	Of 19 evalu- able patients, 11 (58%) had CR and 3 (16%) had PR for an ORR of 74%. PFS at 1 and 2 years was esti- mated at 50% reand 35%, reand	Cichocki et al. [156]

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	ference	3] 3]	on et al. [154]
	Clinical Re response	Of 9 patients, Tar seven (78%) [1 <u>5</u> patients achieved CR with a median duration of 44 months	Of 9 patients, 4 Yo (44%) patients achieved PR, one (11%) achieved CR and 4 (44%) showed PD. The median duration of response was 45 days
	Adverse event/Toxicity	There was no AEs related to NK cells infusion	CRS in 1 patient. GVHD, neurotoxicity, and DLTs were not observed
	Combination therapy	Rituximab	Rituximab
	Chemotherapy regimen before NK cell infusion	Not dis- closed	Cyclophos- phamide and fludarabine
	Infused dose of NK cells	1 to 10×10°cells/ kg	level 1:1 × 10 ⁷ , level 2: 3 × 10 ⁷ , level 3: 9 × 10 ⁷ cells/kg in asso- ciation with lL-2
	Final product Characteristics	The final prod- ucts contained 80.5% CD56 ⁺ CD3 ⁻ NK cells and 26.2 ± 11.6% of CD56 ⁺ CD3 ⁺ NKT cells	Viability > 90% with CD16 ⁺ CD56 ⁺ NK cells > 95%,
	Culture methos	The PBMC were culture for 3 weeks in SCGM media supplemented with IL-15 (10 ng/mL), IL-2 (5 ng/mL), anti-CD3 mAb (10-1,000 ng/ mL), tacrolimus (0.1 ng/mL) and dalteparin sodium (5-10 U/mL)	CD3 depleted cells were culture in Cell- Gro SCGM media with 1% autoplasma, irradiated autologous PBMCs, anti- CD3 mAb (10 ng/mL) and IL2 (500 IU/mL)
	Cells source	۵	8
(p	Conditions/n	R/R CD20 ⁺ lymphoma includ- ing DLBCL (n = 5), FL (n = 4)	R/R NHL includ- ing DLBCL ($n = 6$), MCL ($n = 1$) ($n = 1$)
Table 4 (continue	# Study	4 phase l UMIN000014072	5 phase l NCT03778619

Table 4 (continue	(pa									
# Study	Conditions/n	Cells source	Culture methos	Final product Characteristics	Infused dose of NK cells	Chemotherapy regimen before NK cell infusion	Combination therapy	Adverse event/Toxicity	Clinical response	Reference
6 phase II NCT03019640	R/R NHL includ- ing DLBCL (n = 16), MCL (n = 1) (n = 1)	B D	CB mono- nuclear cells were plated in RPMI media with irradiated aAPC feeder cells and IL-2 (100IU). On day 7, cultured cells were cells were and remain- ing cells were then culture in the same conditions for an addi- tional 7 days	viability > 96% purity > 98% median CD3 CD16+CD56+ cells 98.91%	10 ⁸ cells/kg	Carmustine, etoposide, cytarabine, mel- phalan and lena- lidomide	Rituximab	GVHD, neurotoxicity, and DLTs were not observed	At median follow-up of 47 months, the RFS and OS rates were 53% and 74%, respectively	Nieto et al. [157]
7 phase I NCT04023071	RVR BCL includ- ing DLBCL (n = 10), FL (n = 1) and MZL (n = 1)	DSd	iPSCs transduced and differenti- ated in CD34 ⁺ cells hnCD16 ⁺ for 18–21 days, then CD34 ⁺ cells culture in B0 medium supplemented with 20% human serum and NK cell initiating cytokines (IL-3, IL-7, IL-15, SCF, FH3L) on EL08- 1D2 stroma (culture days 28–35)	Not disclosed	dose cohort 1: 3 × 10 ⁷ dose cohorts 2: 9 × 10 ⁷ dose cohorts 3: 30 × 10 ⁷ cells/ dose	Cyclophos- phamide and fludarabine	Rituximab or obinutu- zumab	No CRS, ICANS, or GVHD were reported	Of 11 patients, 8 (73%) patients achieved an objective response (including 6 CR and 2 PR) and 3 (27%) showed PD	Patel et al. [158]

Table 4 (continu	ed)									
# Study	Conditions/n	Cells source	Culture methos	Final product Characteristics	Infused dose of NK cells	Chemotherapy regimen before NK cell infusion	Combination therapy	Adverse event/Toxicity	Clinical response	Reference
8 Phase I, II NCT04074746	CD30 ⁺ lympho- mas includ- ing HL ($n = 37$) and NHL ($n = 5$)	B D	CB-derived NK cells were first activated with IL-12/ IL-15/IL-18-and then expanded with K562 feeder cells expressing mblL-21, 4-1BBL and CD48 and CD48 and CD48 and CD48 and CD48 and CD48 with AFM13 complexed with AFM13	Viability > 95% NK cells bound to AFM1 3 > 90% NK cell con- tent > 94% 0.02%	level 1:10 ⁶ level 2:10 ⁷ level 3:10 ⁸ NK cells/Kg	Cyclophos- phamide and fludarabine	AFM13	No cases of CRS, ICANS or GVHD of any grade were reported	ORR 92.8% and CR 66.7% at median follow-dia of 14 months, the EFS/ OS rates are 31%/76%; median EFS/OS were 8 months	Nieto et al. [159]
9 Phase I, II NCT00625729	Relapsed NHL or CLL $(n = 6)$	B	CD3 depleted cells were acti- vated by IL-2 overnight	Not disclosed	1.5–8×10 ⁷ cells/ kg in association with IL-2	Cyclophos- phamide and fludarabine	Rituximab	Not disclosed	Of 6 patients, 4 (67%) patients had OR at 3 months, while 2 (33%) had PD at 6 months. OS at 6 months was 3/6 patients	Mamo et al. [160, 161]
R/R Refractory/relapse Mantle cell lymphomé Progression disease, C free survival, EFS Even FMS-like tvrosine kina	ed, <i>NHL</i> Non-Hodgkir a, <i>WM/LPL</i> Waldenstri <i>RS</i> Cytokine release s t-Free Survival, <i>RFS</i> R se 3 ligand, <i>mb</i> Mem	i lymphoma, HL F öm macroglobulli syndrome, ICANS I elapse free surviv brane bound	lodgkin lymphoma nemia/lymphoplas. Immune associatec 'al, NAM Nicotinami	a, <i>CLL</i> Chronic lymphoo macytic lymphoma, <i>Pl</i> meurological sympto! de, <i>mAb</i> Monoclonal i	-ytic leukemia, <i>DLBCL</i> ¹ 3 Peripheral blood, <i>U</i> Cl m, <i>GVHD</i> Graft-versus- antibody, <i>DL</i> 5 Dose lir	Diffuse large B-cell ly B Umbilical cord bloc host disease, CR Com niting toxicities, aAP	nphoma, <i>FL</i> Follicu d, <i>iPSC</i> Induced plu nplete response, <i>OS</i> C Artificial antigen	ılar Lymphoma, MZ uripotent stem cell, 5 Overall survival, <i>P</i> presenting cells, <i>IL</i>	<i>L</i> Marginal zone ly <i>ORR</i> Overall respo 7 Partial response, nterleukin, <i>SCF</i> Stt	mphomas, <i>MCL</i> nse rate, <i>PD</i> <i>P</i> FS Progression- em cell factor, <i>FLT3L</i>

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cells (up to 3×10^9) to cancer patients [166]. Recently, a phase I dose-escalation study using NK-92 cells (1×10^9) cells/m², 3×10^9 cells/m² and 5×10^9 cells/m²) for refractory hematological malignancies that relapsed after autologous HSCT was conducted by Williams et al. [162]. A total of 12 patients were enrolled in this trial, including 2 patients with HL and 5 patients with NHL. The infusions of irradiated NK-92 cells were well-tolerated even at high doses and resulted in CR in one HL patient and a minor response (defined as 10-30% regression of target tumor lesions without the occurrence of new lesions or progression of nontarget lesions) in 2 NHL patients. Notably, in this study, no NK-92 cells were detected more than 15 min after infusion [162]. As mentioned earlier, NK-92 cells are highly dependent on exogenous IL-2 for survival and lack the CD16 receptor, thus impeding their capacity to mediate ADCC [167]. To address this, NK-92 cells have been modified to internally express IL-2 and the high-affinity CD16 receptor [168, 169]. Currently, this product, designated high-affinity NK (haNK), is being investigated in several clinical trials for solid tumors [170, 171]. Furthermore, preclinical data indicated that the combination of haNK cells withmAbs, such as daratumumab for multiple myeloma (MM) and rituximab for NHL, may have a synergistic effect. However, further clinical investigation is required to validate these approaches for NHL [163].

NK cells derived from iPSCs (iNKs) are another promising avenue for NK cell therapy and have the potential to address challenges commonly encountered with other sources of NK cells (Fig. 4) [172]. To generate iNK cells, somatic cells are first differentiated into iPSCs and then into CD34⁺ hematopoietic stem and progenitor cells (HSPCs). Subsequently, NK cells differentiate from HSPCs using cytokines (IL-3, IL-7, IL-15, SCF, and FLT3L) or stromal-based feeder cell lines and are then cocultured with feeder cells for further expansion [173, 174]. Currently, iPSC-based NK cell platforms have been evaluated in several clinical trials as monotherapies or in combination with mAbs for the treatment of hematological malignancies or solid tumors [158, 175-177]. As an example, FT516 is an iPSC-derived NK cell product modified to express high-affinity, cleavage-resistant Fc receptor (CD16A), with a preliminary report of 18 patients with R/R B-cell lymphoma in combination with rituximab demonstrating safety, with no evidence of GVHD, ICANS or CRS. Patients received two cycles of treatment consisting of a conditioning regimen (fludarabine and cyclophosphamide, each for 3 days), a single dose of rituximab and three weekly cycles of FT516 (four patients received 90 million cells/dose, seven patients

NK cells source	Advantages	Limitations	Key development programme
iPSC derived NK-cells	 ✓ High proliferative potential ✓ Homogeneous cell population ✓ Easy to manipulate and engineer ✓ Offer a robust and reproducible cell source for clinical application ✓ Minimal immune rejection 	 Immature phenotype Requires genetic manipulation for CD16-mediated ADCC Extended in vitro culture condition (weeks to months) 	Fcte
HSPC derived NK-cell	 ✓ Readily available through global cord blood banks ✓ Off-the-shelf source for NK cell-based therapy ✓ High expansion capacity ✓ Homogeneous cell population ✓ Despite a decrease in cytokine release, the high killing capacity is preserved 	 Extended in vitro culture condition (weeks to months) Low ADCC as a consequence of reduced or absent CD16 expression. Immature phenotype Safety concern 	glycostem
CB derived NK-cells	 Readily available through global cord blood banks High proliferative potential Possibility of cryopreservation Minimize GVHD Safety and persistence established in clinical trials 	Requires in vivo expansion (weeks) Low expression of natural cytotoxicity receptors Heterogeneous cell population Low ADCC as a consequence of reduced or absent CD16 expression. Immature phenotype	MDAnderson Cancer Center
PB NK-cells	 ✓ Mature phenotype ✓ Highly functional and cytotoxic ✓ Easy collection ✓ Easily expanded in vivo 	 Heterogeneous cell population Low number of NK and requires ex vivo expansion Low transduction efficiency Needs donors, not readily available During expansion, telomere length is shortened and cytotoxicity is reduced 	nkarta catamaranBio Wugen Kiadiseharma
Cell lines (NK-92,)	 ✓ High proliferative capacity ✓ Relatively easy to manipulate and engineer ✓ Homogeneous cell population ✓ Decreased sensitivity to cryopreservation 	Limited in vivo persistence Low ADCC as a consequence of reduced or absent CD16 expression Required irradiation prior to infusion Safety concern owing to lymphoma origin	o ImmunityBio

Fig. 4 Overview of the advantages and limitations of different sources of NK cells

received 300 million cells/dose, and seven patients received 900 million cells/dose) accompanied by IL-2 (6 MIU after each dose of FT516). Of the 18 patients, 10 patients were naive to treatment with autologous CD19-targeted CAR-T cells, and eight patients were previously treated with autologous CD19-targeted CAR-T-cell therapy. A total of 8/10 naive patients achieved an ORR (including 5 patients who achieved CR), and 3/8 patients previously treated with CD19-targeted CAR-T-cell therapy achieved an OR and CR [158, 177].

NK cells combined with bispecific antibodies

Bispecific killer engagers (BiKEs) were created with the intention of having one "arm" that binds to CD16 on NK cells and the other "arm" that targets a specific antigen on tumor cells [178]. The engager serves as a replacement for traditional antibody-Fc interactions in facilitating the immunological synapse between tumor cells and NK cells, thereby promoting NK activation and the killing of tumor cells [179]. Therefore, the use of BiKEs could enhance the function of NK cells by creating a stronger interaction when binding to anti-CD16 compared to the interaction between CD16 and the natural Fc portion of antibodies [180]. Moreover, BiKEs are nonimmunogenic and have rapid clearance properties, making them easy to engineer to target known tumor antigens. In addition to these advantages, BiKEs may offer advantages over mAbs due to their smaller size, which allows for better distribution in the body. This approach is especially beneficial for treating solid tumors [181–183]. Currently, several clinical trials are being conducted to evaluate the effectiveness of BiKEs in combination with NK cell therapy as a treatment for patients with lymphoma. Some of these trials focused on AFM13 [159, 184]. AFM13 is a tetravalent, bispecific innate cell engager that targets CD16A/ CD30 and activates innate immune cells such as NK cells and macrophages [185]. AFM13 acts as a mediator by binding to CD16A on NK cells and to CD30 on lymphoma cells, which aids in the recruitment and activation of NK cells in proximity to tumor cells [186]. AFM13 was initially tested as a single therapy in a phase 1 clinical study for patients with R/R lymphoma [187]. The study showed that AFM13 treatment was safe and welltolerated and led to positive tumor responses in several patients [187]. CB-NK cells precomplexed with AFM13 were recently tested within an ongoing phase I/II clinical trial for patients with refractory CD30-positive lymphomas. Forty-two patients (37 patients with HL and 5 patients with NHL with a median of seven prior lines of therapy) received fludarabine/cyclophosphamide followed by CB-NK cells precomplexed with AFM13 and three weekly IV infusions of AFM13. The results of this study showed that AFM13 in combination with NK cells was safe for patients with no instances of CRS, ICANS, or GVHD and resulted in an ORR of 92.8% and a CR rate of 66.7%. All four patients who had previously failed CD30 CAR-T-cell therapy achieved a CR [159].

NK cells combined with CAR structure

The CAR construct plays a crucial role in activating cells that have been transduced with CAR. The CARs employed in CAR-NK cells are often analogous to those utilized in CAR-T cells. A CAR consists of four essential components: an extracellular binding domain, a hinge region, a transmembrane domain, and one or more intracellular signaling domains (Fig. 5). Singlechain antibody variable fragments (scFvs) originate from a tumor-specific antibody and have the ability to bind to a particular antigen displayed on the surface of cancer cells. Moreover, the intracellular signaling domains are obtained from the cytoplasmic domains of ITAMs found in TCRs or other stimulating receptors [188]. The extracellular binding domain of CAR-modified effector cells enhances specificity by targeting tumor-associated antigens (TAAs). The hinge region serves as a connection between the extracellular binding domain and the transmembrane domain. The intracellular signaling domains in different generations of CARs possess different compositions, which affects the potency of the activation signal transmitted and consequently influences the cytotoxic capability against tumor cells (Fig. 5) [189]. The first generation of CARs consisted of only the CD3- ζ activation signaling domain. Subsequent generations of CARs incorporated one or two supplementary costimulatory molecules, including CD28, ICOS, 4-1BB, CD27, OX40, and CD40. CD28 and 4-1BB are the predominant molecules utilized among this group of molecules [190, 191]. Researchers have utilized other molecules as activation signaling domains for NK cells, in addition to the commonly used CARs that are applicable for both CAR-T cells and CAR-NK cells. CD244 (2B4), a member of the signaling lymphocyte activation molecule (SLAM) family, can also serve as a costimulatory molecule. The overexpression of 2B4 in NK cells leads to an enhanced ability to amplify signals and increased innate cytotoxicity against tumor cells [192]. DAP-12 is present on NK cells and plays a role in transmitting signals through the NK-activating receptors NKG2C and NKp44. Additionally, DAP-10 is involved in signal transmission through NKG2D [193, 194]. Hence, DAP-12 and DAP-10 can transmit intracellular signals in CAR-NK cells. In addition, NK cells modified with DAP-12-based CARs exhibited superior performance compared to that of NK cells modified with CD3-ζ-based CARs [193]. Recent research has indicated that NKG2D ligands are overexpressed in several hematological malignancies. Hence, the



Fig. 5 An overview of (A) CAR-NK cell therapy workflow in NHLs, (B) CAR structure and generation and (C, D) various methods of delivering CAR into NK cells

NKG2D-DAP-10-CD3- ζ CAR, which specifically targets NKG2D ligands, holds significant promise for the treatment of blood malignancies [195].

Transduction of the CAR gene into NK cells encompasses viral transduction, namely, retrovirus-based and lentivirus-based approaches, as well as transfection techniques such as electroporation, lipofection, and their combination with transposon systems (Fig. 5) [11]. CAR constructs are commonly integrated into a retrovirus or lentivirus-based expression vector. These vectors are then used to transduce primary NK cells or NK cell lines, with NK-92 being the most frequently used. The transduction of retroviral vectors shows a high level of effectiveness (ranging from 43 to 93%) in primary NK cells. However, the occurrence of insertional mutagenesis and its negative consequences significantly limit the use of this method in clinical applications [196]. However, lentivirus-based transduction is considered to be a safer method. Although its transduction efficiency in peripheral blood mononuclear cell (PBMC)-derived NK cells ranges from 8 to 16%, there is still an opportunity for improvement [197]. RNA transfection methods are economical strategies that have greater efficacy in transferring genes. However, the production of CAR constructs using this method is temporary, lasting for approximately 3-5 days. Although the short therapeutic time frame is a limitation, the temporary nature of CAR therapy may lower the occurrence of CAR-associated adverse effects, such as on-target off-tumor effects [195, 197, 198]. The integration of DNA into cells using transposon systems, such as PiggyBac (PB) and sleeping beauty (SB), in combination with transfection methods has emerged as an appealing strategy for generating cells that express transgenes in a safer and more stable manner [199, 200]. The SB transposon vector has proven to be a cost-effective and efficient means of gene transfer. However, its suitability for use with CAR-NK cells has not yet been evaluated [201].

CAR-NK cells are safer than CAR-T cells. The enhanced safety of CAR-NK cells can be attributed to two primary factors. CRS and neurotoxicity are frequently adverse effects of CAR-T-cell therapy [202]. The cytokine storm triggered by CAR-T cells, specifically TNF-α, is primarily facilitated by proinflammatory cytokines such as IL-1 and IL-6 [203]. CAR-NK cells secrete a variety of cytokines, such as IFN-y and GM-CSF, which differ from the cytokines produced by CAR-T cells. Second, CAR-T cells can cause life-threatening GVHD due to HLA limitations. On the other hand, NK cells, which are considered important cells that initiate the GVT response early on, can potentially prevent GVHD by eliminating recipient antigen-presenting cells and CTLs [204]. Furthermore, CAR-NK cells may exhibit superior effectiveness in targeting and destroying cancerous cells compared to CAR-T cells. CAR-NK cells possess the ability to identify and execute their cytotoxic functions via both their designed and innate killing capabilities. By utilizing CARs, effector cells can enhance their ability to selectively target and eliminate a specific antigen with greater efficiency. Unlike CAR-T cells, CAR-NK cells retain the inherent ability of NK cells to destroy target tumor cells even when the expression of specific tumor antigens is reduced [205]. Moreover, the production of CAR-NK cells is more convenient than that of CAR-T cells. Due to the absence of the risk of GVHD, NK cells can be obtained from either a donor who is a match or a donor who has an HLA mismatch, hence expanding the pool of potential donors and enhancing the overall quality of the end products [206].

Recently, CAR-NK cell therapy has been assessed in various clinical trials for the treatment of lymphoma (Tables 5 and 6). CB-CAR-NK cells are presently employed in a clinical trial at MD Anderson Cancer Center, specifically targeting CD19 cells, and yielding highly favorable outcomes. 37 patients with R/R CD19-positive malignancies were enrolled in this trial and treated with CB-CAR-NK cells in two phases: a dose-escalation phase and an expansion phase. In the dose-escalation phase (n=11), patients received a conditioning regimen (fludarabine and cyclophosphamide, each for 3 consecutive days) followed by the infusion of CB-CAR-NK cells (three patients received 10×10^4 cells/kg, four patients received 10×10^5 cells/ kg, and four patients received 10×10^6 cells/kg). In the expansion phase (n=26), patients were first treated with 10×10^6 cells/kg CB-CAR-NK. Then, the trial was amended to include a second expansion cohort in which patients received a single flat dose of 8×10^8 cells/kg CB-CAR-NK. A retroviral vector including an anti-CD19-CD28-CD3-ζ CAR, an IL-15 gene, and a suicide switch was utilized for transduction. None of the patients developed neurotoxicity or GVHD, and only one patient developed mild CRS (grade I). The ORR (including PR and CR) on days 30 and 100 for the 37 patients was 48.6%. The 1-year OS and PFS were 68% and 32%, respectively. Compared with non-responders, patients who achieved OR had higher levels and longer persistence of CB-CAR-NK cells [6, 207].

Goodridge et al. created a CAR-NK product called FT596. This product was derived from iPSCs. The iPSCs were modified to consistently produce anti-CD19 CAR, a high affinity and non-cleavable CD16 Fc receptor, and a combination of a membrane-bound IL-15 and an IL-15R α fusion protein. In a Raji xenograft mouse model, the combination of FT596 with rituximab resulted in a substantial increase in the elimination of Raji tumor cells. In addition, when a mouse model

that had been engrafted with human CD34 cells, FT596 showed enhanced longevity and safety compared to primary CAR19 T cells [211]. This platform has been translated into a multicenter, phase I clinical trial as monotherapy or in combination with rituximab to treat patients with R/R B-cell lymphoma [210]. A total of 20 patients underwent two treatment regimens, including 10 in regimen A (FT596 alone) and 10 in regimen B (FT596 cells combined with rituximab). Among the 17 evaluable patients, clinical response was observed in 9 patients (5 from regimen A and 4 from regimen B), 7 of whom achieved CR. Notably, no dose-limiting toxicity, ICANS, or GVHD of any grade was observed. Interestingly, 2/4 of patients treated with CAR-T-cell therapy at doses $\ge 9 \times 10^7$ cells/kg achieved CR [210]. An extended follow-up period will provide insight into the durability and efficacy of this platform. More recently, similar peripheral blood-derived anti-CD19 CAR-NK cells (named NKX019, a cryopreserved product utilizing OX40/CD3- ζ signaling domains and expressing a membrane-bound form of IL-15 for activation) were investigated in a phase I trial as a monotherapy for 19 patients with R/R B-cell malignancies. Patients received a daily lymphodepletion regimen of fludarabine and cyclophosphamide for 3 days. Next, they received three infusions of NKX019 at 3 dose levels, with doses ranging from 300 million to 1.5 billion cells per infusion. During the follow-up period, no dose-limiting toxicity, neurotoxicity, CRS or GVHD was reported. Among the 14 patients with NHL, 8 achieved CR; however, 3 patients with indolent lymphoma subsequently experienced relapse after a remission period of greater than 6 months [208, 212].

What's Next? CIML NK cells

NK cells following exposure to happens, viral infection or a combination of cytokines achieve memory properties. NK cells preactivated with IL-12/15/18 have been described as cytokine-induced memory-like (CIML) NK cells [213]. CIML NK cells present distinctive characteristics, such as high proliferative capacity, sensitivity to low doses of IL-2, increased IFN- γ production, resistance to TGF- β , elevated glycolysis, and oxidative phosphorylation, which distinguishes them from conventional NK cells (cNK cells) [214–217]. In addition, the long-term life span and adaptive immune features of CIML NK cells have drawn attention to the use of these cells in cancer immunotherapy. Recent findings from preclinical and clinical trials have shown that CIML NK-based immunotherapy has produced promising results and also offers a safe approach to preventing GVHD, CRS, and neurotoxicity [218]. In the context of hematological malignancies, CIML NK cell-based immunotherapy has aided in the discovery of novel treatments for various

	Sponsor/ Location	Zhejiang university/ China	M.D. Ander- son cancer center/USA	Allife medi- cal science and technol- ogy Co,, Ltd./ China	Allife medi- cal science and technol- ogy Co., Ltd./ China	Allife medi- cal science and technol- ogy Co., Ltd./ China
	Outcome Measures	Primary: safety, tolerability and deter- mine the rec- of anti-CD19 of anti-CD19 Secondary: CR, PFS, DOR, OS, PR, ORR and immu- nogenicity of anti-CD19 CAR-NK cell	Primary: safety and efficacy Secondary: Secondary: and PR(CR and PR(S), determining the per- sistence of injected cells and com- cells and com- prehensive immenensi immen	Primary: occurrence of treatment related AEs	Primary: occurrence of treatment related AEs	Primary: occurrence of treatment related AEs
	CAR construct	Not dis- closed	CD19- CD28-zeta- 2A-iCasp9- IL15	CD19/ CD22- CD-244	CD22- CD-244	CD19- CD-244
	Cells source	nce	nce	Not dis- closed	Not dis- closed	NK-92 cell line
	Doses	2.5 × 10 ⁸ cells, 5 × 10 ⁸ cells, cells cells	Not disclosed	50 to 600 × 10 ³ cells/kg	50 to 600 × 10 ³ cells/kg	50 to 600 × 10 ³ cells/kg
	Interventions	Anti-CD19 CAR-NK	Fludarabine + cyclo- phospha- mide + Mesna + CAR- NK Cells + AP1 903 (in case of GVHD or CRS after CAR-NK infusion)	Anti-CD19/CD22 CAR- NK cells	Anti-CD22 CAR-NK cells	Anti-CD19 CAR NK cells
	Vector	vector	Retroviral vector	Not dis- closed	Not dis- closed	Not dis- closed
	Antigen	CD-19	CD-19	Dual-target CD19/22	CD-22	CD-19
	Conditions	RVR B-cell NHL	R/R B-lymphoid malignan- cies (ALL, CLL, NHL) CLL, NHL)	Refractory B-Cell lym- phoma	R/R B-cell lymphoma	R/R B-cell lymphoma
	Study Design	Single group assignment, open label	Single group assignment, open label	Single group assignment, open label	Sequential assignment, open label	Single group assignment, open label
	Trial status	Recruiting	Completed	Unknow	Unknow	Unknown
	Phase/N	Phase 1/48	Phase I, 11/44	Phase I/10	Phase I/9	phase I /9
6	NCT Number	NCT05472558	NCT03056339	NCT03824964	NCT03692767	NCT03690310
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 Table 5
 Ongoing clinical trials of CAR-NK cell therapies for NHL

Tal	ble 5 (continu	led)											
#	NCT Number	Phase/N	Trial status	Study Design	Conditions	Antigen	Vector	Interventions	Doses	Cells source	CAR construct	Outcome Measures	Sponsor/ Location
ο	NCT04639739	phase 1/9	Unknown	Single group assignment, open label	R/R NHL	CD-19	Not dis- closed	Fludarabine + cyclo- phosphamide + anti- CD19 CAR NK cells	2×10 ⁶ /kg 6×10 ⁶ /kg 2×10 ⁷ /kg	Not dis- closed	Not dis- closed	Primary: DLTs and incidence and sever- ity of AEs and SAEs Secondary: ORR, PFS and OS	Xinqiao hospital of Chong- qing/China
\sim	NCT04796675	Phase 1/27	Recruiting	Single group assignment, open label	B Lymphoid Malignancies (ALL, CLL and NHL)	CD-19	Retroviral vector	Fludarabine + cyclo- phosphamide + Anti- CD19 CAR-NK cells	0.01 × 10 ⁷ 0.1 × 10 ⁷ 1.0 × 10 ⁷ /kg	UCB	Not dis- closed	Primary: occurrence of treatment related AEs Secondary: ORR, CR, PFS, DOR and OS	Wuhan union hospital/ China
∞	NCT04887012	Phase //25	Recruiting	Single group assignment, open label	R/R B-cell NHL	CD-19	Lentiviral vector	Anti-CD19 CAR-NK cells	Not disclosed	В	Not dis- closed	Primary: DLTs and ORR Secondary: OS, PFS, pharmacoki- netics of CAR positive cells and pharma- cokinetics of CAR-NK cells	Second affiliated hos- pital, school of medicine, Zhejiang university Hangzhou, Zhejiang/ China
σ	NCT05020678	Phase //150	Recruiting	Single Group assignment, open Label	B-cell cancer (NHL, ALL, CLL)	CD-19	Retroviral vectors	Fludarabine + cyclo- phosphamide + anti- CD19CAR-NK cells (combination cohorts will additionally receive rituximab with each cycle)	3 × 10 ⁶ NK cells (6 × 10 ⁶ /kg kg) kg)	B	CD19- CD134 (0X40)- CD3z-lL-15	Primary: occurrence of TEAEs, proportion of subjects burstencing DLTs and ORR Secondary: assessment of CAR-NK half-life, duration of persistence, host immune response against CAR- NK and ORR	USA USA

	Sponsor/ Location	Zhejiang university/ China	ImmunityBio, Inc/USA	Kunming hope of health hos- pital/China	M.D. Ander- son cancer center/USA
	Outcome Measures	Primary: incidence of TEAEs and pro- portion of subjects experiencing DLTS Secondary: ORR, DOR, PFS, OS, phar- macokinetics of CAR-NK cells and EFS	Primary: safety Secondary: ORR	Primary: inci- dence of AEs Secondary: granulocyte implantation time, platetion time, red blood cell implantation time, DOR, DCR, OS and PFS	Primary: inci- dence of AEs, CR or PR Secondary: PFS, OS, tresponse sta- tus and num- ber of CAR-NK cells in blood
	CAR construct	Not dis- closed	Not dis- closed	Not dis- closed	CD19- CD28-z-2A- iCasp9-IL15
	Cells source	ipSC	NK-92 cell line	Not dis- closed	NCB
	Doses	Not disclosed	Not disclosed	Dose level 1: 5-10×10 ⁶ /kg, Dose level 2: 1-2×10 ⁷ /kg 2-5×10 ⁷ /kg	Not disclosed
	Interventions	Cyclophospha- mide + Fludara- bine + VP-16 + CAR-NK cell as Monotherapy (for patients with r/r B-ALI) or in combina- tion with rituximab (for patients with r/r B-cell Jymphoma)	lymphodepleting chemotherapy +CAR- NK cell in combination with rituximab (cohort A) or in combina- tion with rituximab and N-803 (cohort B)	HSC + CAR-NK cell	Rituximab + car- mustine, etoposide, cytarabine and mel- phalan + CAR-NK cell + ASCT + filgrastim
	Vector	Not dis- closed	Not dis- closed	Not dis- closed	Retroviral vector
	Antigen	CD-19	CD-19	CD-19	CD-19
	Conditions	B-Cell Malig- nancies (r/r ALL and lym- phoma)	R/R NHL	B-Cell malig- nancies	B-cell NHL
	Study Design	Non-ran- domized, parallel assignment, open label	Rand- omized, sin- gle group assignment, pen label	Single group assignment, open label	Single group assignment, open Label
	Trial status	Recruiting	Not yet recruit- ing	Withdrawn (the prin- cipal investigator decides to stop)	Withdrawn (lack of funding)
(pər	Phase/N	Phase I/24	Phase I/20	Phase I, II	Phase I, II
le 5 (continu	NCT Number	NCT05379647	NCT05618925	NCT05570188	NCT03579927
Tab	#	0	=	12	с

Tak	ole 5 (continu	ed)											
#	NCT Number	Phase/N	Trial status	Study Design	Conditions	Antigen	Vector	Interventions	Doses	Cells source	CAR construct	Outcome Measures	Sponsor/ Location
14	NCT02892695	Phase I, II/10	Unknown	Single group assignment, open label	CD19 ⁺ Lymphoma (NHL)	CD-19	Not dis- closed	Anti-CD19 CAR-NK cells	Not disclosed	NK-92	CAR.19- CD28- 41BB-CD3ζ	Primary: AEs Secondary: ORR	Person gen biotherapeu- tics Co., Ltd/ China
15	NCT05092451	Phase I, II/94	Recruiting	Rand- omized, sin- gle group assignment, open label	R/R Hema- tologic Malignances (AML/MDS and B-Cell Iymphoma)	CD-70	Not dis- closed	Cyclophospha- mide + fludarabine phosphate + anti- CD70 CAR-NK	Not disclosed	ncB	CAR.CD70- IL15 (full construct not dis- closed)	Primary: number of participants with TEAEs, CR, PR and number of Participants who are alive and in remis- sion	MD Anderson cancer center/USA
16	NCT05667155	Phase 1/48	Recruiting	Single group assignment, open label	B-cell NHL	Dual-target CD19/70	Lentiviral vector	Anti-CD19/70 CAR-NK	Not disclosed	UCB	CD19/ CD70-IL-15 (full construct not dis- closed)	Primary: incidence of DLTs Secondary: OS, ORR, DOR, CR, PR and PFS	Second affiliated hos- pital, school of medicine, Zhejiang university/ China
1	NCT05654038	Phase I, II/30	Recruiting	Single group assignment, open label	B-Cell hematologic malignan- cies	CD-19	Not dis- closed	HSCT + Anti-CD19 CAR-NK cells	Dose level 1: 5 -10 × 10 ⁶ / kg; dose level 2:1-2 × 10 ⁷ / g; dose level 3:2-5 × 10 ⁷ /kg	Not dis- closed	Not dis- closed	Primary: inci- dence of AEs Secondary: Granulocyte implantation time, platelet implantation time, red blood cell implantation time, DOR, DCR, OS and PFS	920th hos- pital of joint logistics support force of people's libera - tion army of China/ China

Sponsor/	Location Takeda/USA	Aibin Liang/ China	Changhai hospital/ China	Affiliated hospital to academy of military medical sci- ences/China
Outcome	Measures Primary: inci- dence of AEs, number of participants with clinically significant changes in laboratory param eters, number of participants with clinically significant changes in vital signs and ORR, CR, DO, PFS, OS, number of participants with AFS, OS, number of participants with AFS,	Primary: DLTs Secondary: ORR, OS, CR, DOR, PFS	Primary: inci- dence of DLTs and TEAEs Secondary: ORR, OS, CR and DCR	Primary: DLTs, TEAEs and ORR Second- ary. PFS, OS and pro- portion of subjects with MRD negative response
CAR	Construct Not dis- closed	Not dis- closed	Not dis- closed	Not dis- closed
Cells	ource UCB	UCB	Not dis- closed	Not dis- closed
Doses	200 × 10° TAK-007 TAK-007	Not disclosed	6 × 10 ⁸ cells, 1 × 10 ⁹ cells, 1.5 × 10 ⁹ cells	5×10 ⁶ cells/kg, 2×10 ⁷ /kg
Interventions	Fludarabine + cyclo- phosphamide + CD19 CAR-NK cells (TAK-007)	Dual Anti-19/70 CAR- NK cell	Anti-CD19 CAR-NK cells	Fludarabine, + cyclo- phosphamide + CD19- CAR-NK
Vector	vector	Not dis- closed	Not dis- closed	Not dis- closed
Antigen	CD-19	Dual-target CD19/CD70	CD-19	CD-19
Conditions	R/R B-cell NHL	R/R B-cell NHL	R/R DLBCL	R/R B-cell malignan- cies
Study	Design Nonran- domized, parallel assignment, open label	Single group assignment, open label	single group assignment, open label	Single group assignment, open label
Trial status	Recruiting	Recruiting	Recruiting	Recruiting
Phase/N	Phase II/242	Phase I, II/48	Early phase 1/12	Phase I/12
NCT Number	NCT05020015	NCT05842707	NCT05673447	NCT05645601
#	<u></u>	19	20	21

Table 5 (continu	ued)											
# NCT Number	Phase/N	Trial status	Study Design	Conditions	Antigen	Vector	Interventions	Doses	Cells source	CAR construct	Outcome Measures	Sponsor/ Location
22 NCT05410041	Phase I/15	Recruiting	single group assignment, open label	R/R.Bcell malignancies (ALL, CLL and NHL) and NHL)	CD-19	Not dis- closed	Fludarabine + cyclo- phosphamide + CD19- CAR-NK	1-3×10 ⁷ cells/kg	Not dis- closed	Not dis- closed	Primary: safety and ORR Secondary: concentration of CAR ⁺ NK cells in PB pharmacody- namic data in PB Other out- come meas- ures: DOR, PFS and OS	Beijing Boren hospital/ China
23 NCT05336409	Phase I/75	Recruiting	Nonran- domized, sequential assignment, open label	CD19+ B-cell malignan- cies	CD-19	Not dis- closed	Lymphodeplet- ing chemother- apy + CD19-CAR-NK (CNTY-101) + IL-2	Not disclosed	OS di	CD19 CAR-CD28- I-15-express- ing ing	Primary: MTD and RP2R Secondary: CR, OR, DOR, TTR, PFS, OS, plasmition for cnty-101, Tmax, termi- nal disposition phase half-life for 101, AUCO, percentage of participants with at least one TEAE, percentage of participants with clinically significant laboratory abormali- ties and time to treatment initiation	Century therapeutics, Inc/USA
24 NCT05739227	Early phase 1/12	Recruiting	single group assignment, open label	R/R B-cell hematologic malignancies (ALL, CLL and B-cell Lymphoma)	CD-19	Not dis- closed	Fludarabine + cyclo- phosphamide + etopo- side + CD19-CAR-NK (JD001)	1 × 10 ⁶ cells/kg, 5 × 10 ⁶ cells/kg, 2 × 10 ⁷ cells/kg	BB	Not dis- closed	Primary: DLT and ORR Secondary: PFS, OS, MRD- ORR	Xuzhou medical university/ China

Table 5 (continu	(pər											
# NCT Number	Phase/N	Trial status	Study Design	Conditions	Antigen	Vector	Interventions	Doses	Cells source	CAR construct	Outcome Measures	Sponsor/ Location
25 NCT06206902	Phase I/56	Recruiting	Single group assignment, open label	R/R NHL	Not dis- closed	Not dis- closed	Cyclophospha- mide + CAR-NK Cells (F01)	0.5–3×10° CAR- NK Cells	Not dis- closed	Not dis- closed	Primary: safety Secondary: ORR, DOR, PFS, OS, Cmax of F01 cells, Tmax of F01 cells, AUCO- last of F01 cells, Clast of F01 cells, Tlast of F01 cells, Dynamic changes of cytokine levels, Anti- cells, Dynamic changes of cytokine levels, Anti- cells, Dynamic changes of cytokine levels, Anti- cyte subsets, detection rate of replicable virus	Shanghai Simnova Biotechnol- ogy Co. Ltd/ China
<i>CAR</i> Chimeric antiget <i>Tmax</i> Time to reach t Acute myeloid leuker response, <i>RFS</i> Relaps Overall survival, <i>DOR</i> stem cell transplantai	n receptor, <i>RP2H</i> he maximum cc nia, <i>MDS</i> Myelc e free survival, <i>I</i> Duration of res tion, <i>ASCT</i> Auto	Recommended ancentration, AL odysplastic synd DLT Dose limitin ponse, <i>PFS</i> Prog logous Stem cel	d phase 2 regim JCO The area uno rome, DLCBL Dif g toxicities, AEs gression-free sur II transplantatio	en, <i>TEAE</i> Treatin der the drug cu fuse large B-cel Adverse events vival, <i>DCR</i> Dises n, <i>CRS</i> Cytokine	ient emergent rve, <i>Clast</i> Last of Il lymphoma, <i>N</i> , <i>SAEs</i> Serious of ase control rate release syndro	adverse event detectable cor <i>IHL</i> Non-Hodg adverse events <i>pB</i> Periphera Dme, <i>GVHD</i> Gr	, <i>MRD-ORR</i> Minimal-resid centration point, <i>Tlast</i> Tir kin lymphoma, <i>HD</i> Hodgl <i>t</i> , <i>TTR</i> time-to-response, <i>L</i> I blood, <i>UCB</i> Umbilical co aft-versus-host disease	ual disease negative me of the last detect. kin disease, CLL Chro SSR disease control ra rd blood, <i>iPSC</i> Inducc	overall respoi able concentri nic lymphocy ate, <i>DFS</i> Disea: ed pluripotent	rse rate, <i>Cmax</i> G ation, <i>ALL</i> Acute tic leukemia, <i>CR</i> se-free Survival, stem cell, <i>IL</i> Intr	iene copy number lymphoblastic leu Complete respons ORR Overall respo erleukin, HSCT Her	amplification, kemia, <i>AML</i> .e, <i>PR</i> Partial nse rate, <i>OS</i> natopoietic

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Reference	Dickinson e [208]	Ramachanc et al. [209]
Clinical response	Of 14 patients, 8 (57%) achieved CR and 4 (29%) showed PD: Of the 8 patients who achieved CR, 3 patients subsequently relapsed, each relapsed, each than 6 months of remission	Ongoing CR of a duration of 5 months since the first CNTY-101 infu- sion
Adverse event/Toxicity	No patients developed signs of CRS, ICANS and GVHD	No CRS or neu- rotoxicity were detected
Combination therapy	e N	e O
Chemotherapy regimen	Cyclophospha- mide and fludara- bine	Cyclophospha- mide and fludara- bine
Infused dose	Dose level 1: 1.5 × 10° Dose level 2: 1 × 10° Dose level 3: 3 × 10 ⁸ cells	1 × 10 ⁸ cells
Vector/Target	Retroviral vec- tor/CD-19	CD-19 CD-19
CAR-structure	Anti-CD19. OX40.CD3z. IL-15	Anti-CD19.CD28. CD3zeta-IL-15. Additionally, CNTY-101 prod- uct engineered to avoid recogni- tion by patient CD8+T, CD4+T and NK cells. Knockcout of β2 M, designed to prevent CD8+T-cell rec- ognition, knock- out of the CITA, event CD4+T-cell recognition, and knock- out of the CITA, event CD4+T-cell recognition, and knock- out of the ULA-E gence designed to enable higher expression of the HLA-E protein of CNTY-101 cells by host NK cells. Furthermore, to potentially improve safety, the CNTY-101 cells were engineered with an EGFR
Product name	NKX019	CNTY-101
source	8	ip SC
Conditions/n	DLBCL ($n = 6$), FL ($n = 6$), MCL ($n = 1$) and MZL ($n = 1$)	R/R FL
Study	Phase I NCT05020678	Case report NCT05336409

Table 6 Outcome of clinical trials of CAR-NK cells in NHL patients

TheseLing FTSGL (n=26) FTSG (n=26)	3 Phase I R/R BCL (<i>n</i> =26) iPSC FT596 Anti-CD19. NKG2DTM.: NCT04245722 CD32IL-15R CD32IL-15R						•	-	
4 Phase,II DIBCL (n=17), UCB Not applicable Anti-CD19, Retrovial vec. Dose level 1: Cyclophospha- None No GyHD Overall: the day Marin et al. [207] NCT03056333 E. (n=4), MCL CD28+ICCD32, tor/CD-19 1 × 10 ³ mide and fludara- 0 GyHD Overall: the day Marin et al. [207] NCT03056333 E. (n=4), MCL LL CD28+ICCD32, tor/CD-19 1 × 10 ³ mide and fludara- 0 Gs/HD Overall: the day Marin et al. [207] n=0, CLLRT n=0, CLLRT II-15, Casp9 Dose level 2: inde and fludara- 0 Gs/HD Were 48.6% n=0, CLLRT n=0, CLLRT n=0, CLLRT N 1 × 10 ³ csels/kg csels/kg inde and fludara- 0 Gs/HD were 48.6% inde and Gs/HD were 48.6% inde and Gs/HD were 48.6% inde and Gs/HD		nt-CD19. No G2D TM.284. CD 33z.IL-15RF	3. of disclosed/ 3.	 10⁷ cells 10⁷ cells × 10⁷ cells 	Cyclophospha- bine and fludara- bine	Rituximab or obinutu- zumab	No GvHD and ICANS were observed. Two cases of CRS were reported	Overall: 18/26 patients (69%) achieved ORR, including 12/18 patients (46%) on day 29 CR on day 29 fol- lowing a single dose of FT596. Combination therapy (com- prising a total of 12 patients) 9/12 patients (58%) with CR on day patients (58%) with CR on day 29 follow- ing a single dose of FT566 dose of FT566	Bachanova et al. [2 10]
	4 Phase I,II DLBCL ($n = 17$), UCB Not applicable Anti-CD19. NCT03056339 FL ($n = 4$), MCL ($n = 1$), MZL ($n = 1$), MZL ($n = 3$), CLLRT ($n = 2$), CLLRT ($n = 3$), CLLRT ($n = 6$), CLLRT ($n = 1$) and LPL ($n = 1$) and LPL ($n = 1$)	nti-CD19. Rei 228+IC.CD3z. tor 15. Casp9	/CD-19 CC- 13 /CD-19 CC- 12 DC DC 12 13 13 14 14 15 15 15 15 15 15 15 15 15 15 15 15 15	sse level 1: <10 ⁵ lls/kg <10 ⁶ <10 ⁷ lls/kg lls/kg	Cyclophospha- mide and fludara- bine	e None None None None None None None Non	No GvHD and ICANS were observed. one case developed CRS (grade 1)	Overall: the day 30 and day 100 ORR (including PR and CR) were 48.6% for both. The day 30 and day 100 CR rates were 27% were 27% respectively. The 1-year CR rate was 37.8% 1-year CR rates for patients with FL and MZL were 83 (5/6) and for DLBCL was 22% (5/17)	Marin et al. [207]

disease, $\beta 2$ M $\beta 2$ microglobulin, CIITA Class II transactivator, EGFR Epidermal growth factor receptor

Table 6 (continued)

cancers, particularly myeloid disorders. Similarly, adoptively transferred CIML NK cells trigger CR in 44% of R/R acute myeloid leukemia (AML) patients [219].

Another clinical trial by Shapiro et al. revealed that CIML NK cell infusion into an immune-compatible microenvironment in posttransplant relapsed AML, MDS, and MPN patients resulted in satisfactory expansion and persistence [220]. Similarly, CIML NK cells injected into pediatric/young adults with post-HCTrelapsed AML patients significantly expand and persist in a compatible milieu. Furthermore, this clinical trial established that AML patients were treated with donor lymphocyte infusions (DLIs), and CIML NK cells showed promising outcomes [221].

Unlike for myeloid disease, the therapeutic approach involving CIML NK cells in lymphoid malignancies has received less attention. One of these few studies was performed on a rat model of T-ALL, namely, Roser leukemia (RL). In this in vivo experiment, RL was treated with cNK cells, and NK cells were stimulated with IL12/15/18 (CIML NK cells). Based on these results, RL is resistant to cNK cells but not to CIML NK cells. Therefore, CIML NK cells could be introduced as a possibility for immunotherapeutic clinical trials in T-ALL patients [222].

The role of CIML NK cells in lymphoma was studied by Ni et al. in mice injected with RMA-S lymphoma cells. Tumor-bearing mice were treated with IL-12/15/18-preactivated NK cells and IL-15-pretreated NK cells. The results highlighted that compared with IL-15-pretreated NK cells, IL-12/15/18-preactivated NK cells display greater frequency, persistence, proliferation, and functional killing activity at the tumor site [223]. In another study, Gang et al. further investigated CIML NK cell incorporation in lymphoma. They preactivated NK cells with IL-12/15/18 and then developed them to express the anti-CD19 CAR structure. The 19-CAR-CIML NK cells exhibited improved in vitro cytotoxicity against Raji cells and CD19⁺ primary lymphoma cells, as illustrated by elevated IFN-y production and degranulation capacity. In addition, 19-CAR-CIML NK cells exhibited satisfactory durability, expansion, and effector function in a human lymphoma xenograft mouse model [224].

Finally, the challenges in NK cell-based immunotherapy in NHLs, the highly appreciated features of CIML NK cells, and promising results from current preclinical studies have prompted us to develop new therapeutic options based on CIML NK cells. Hopefully, we will witness a fundamental revolution in the management of patients with NHL.

NK cell expansion

NK cells offer significant potential for immunotherapy in NHL treatment, but there are still obstacles to overcome

to harness their full therapeutic benefits. There are still efforts to obtain a considerable number of NK cells for therapeutic purposes and to ensure that the obtained NK cells are fully functional and capable of effectively targeting and killing abnormal cells. This requires careful selection and expansion of NK cells, which can be technically challenging in the laboratory [225, 226]. Most PB-derived NK cell expansion protocols can be categorized into feeder-cell or feeder-free systems [227].

Feeder cells

The production of a significant amount of NK cells from a small initial quantity relies on feeder cells. These feeder cells, whether naturally or through additional modifications, present ligands for NK cell receptors. When combined with cytokines, this interaction drives a substantial expansion of NK cells outside the body, enabling the generation of a large number of NK cells for therapeutic purposes [228]. Various types of cells, such as EBVtransformed lymphoblastoids and genetically engineered HEK293 or K562 cell lines, are utilized as feeder cells. Among these, genetically modified K562 cells are the most commonly employed [227]. For example, when a mixed lymphocyte population is infected with Epstein-Barr virus (EBV) in vitro, it results in an immortalized cell line that exhibits characteristics similar to those of proliferating B cells. With the expression of different ligands (4-1BBL), CD155, CD48, and CD58) that have specific receptors (4-1BB, DNAM-1, 2B4, and CD2, respectively) on activated NK cells, EBV-bearing lymphoblastoid cell lines (LCLs) play essential roles in NK cell expansion and stimulation [228]. Using this method, an average of 1,000-2,000-fold expansion of NK cells was reported to be observed over a period of 14 days [229]. The addition of IL-21 and IL-2 reportedly improved the expansion efficacy [227, 230, 231]. In another method, irradiated feeder cells were employed to amplify NK cells in laboratory settings. The K562 leukemia cell line has been altered to display particular ligands linked to antigen-presenting cells (CD64, CD86, and truncated CD19, CD137L, 4-1BB ligand, and membrane-bound IL-21). The irradiated K562-mbIL21-41BBL cells seemed to be very effective at rapidly increasing the number of NK cells in RPMI media (containing 10% FBS). These modified cells expanded NK cells 47,967-fold in 21 days [232]. Nevertheless, using feeder cells can pose challenges due to licensing intricacies, difficulties in sourcing, and the requirement for their elimination from the culture. Challenges such as incomplete irradiation of feeder cells (which might lead to teratoma) and separating and thoroughly eliminating cancer cells from the culture environment to avoid injection into patients are additional difficulties [233].

Feeder-free expansion methods

Expanding NK cells without the need for feeder cells has benefits compared to traditional methods, especially in terms of lower contamination risks and improved regulatory compliance. Additionally, other benefits, such as reducing costs through a more straightforward process and even decreasing cytotoxicity, have been reported [227]. Feeder-free NK cell expansion systems rely on cytokines and stimulating supplements or antibodies. Ex vivo cultured NK cells treated with IL-15 and nicotinamide exhibited stable CD62L expression, which was linked to increased FOXO1 levels. Nicotinamide enhanced NK cell metabolism, cytotoxicity, and cytokine production, leading to improved outcomes in adoptive transfer experiments. Recently, Cichocki et al. performed a phase 1 clinical trial in patients with relapsed or refractory NHL using rituximab in association with NK cells expanded with IL-15 and nicotinamide. The final result showed a 74% response rate in 19 patients [156]. Gluk et al. conducted two phase I studies to assess the combination therapy of rituximab and IL-2 (4.5-14 million international units) in relapsed or refractory B-cell NHL to boost ADCC through NK cell activation. The results showed that adding IL-2 to rituximab treatment is safe and effective, particularly with thrice-weekly IL-2 dosing, leading to increased NK cell counts and associated with treatment response [234]. In conclusion, obtaining a sufficient number of functional NK cells for therapeutic purposes remains a challenge. Despite the advancements made in feeder-free NK cell expansion, further investigations are needed to optimize this process and ensure its utility in clinical applications.

Conclusion

In this comprehensive review, we first provided an overview of NK cells, including their function, characteristics, development, and maturation. We then delved into the complex tumor microenvironment and the interplay between various presented cells that can either support or hinder the antitumor activity of NK cells in NHL.

Building on these findings, we explored various strategies to enhance the therapeutic potential of NK cells because based on the findings reported in the literature, the function and number of NK cells are defective in NHL patients. Therefore, it would be beneficial to bolster the innate immune response by injecting and activating NK cells. Also, combinations of NK cells with multiplex immunotherapy strategies such as mAbs, BiKEs, and CARs could be effective and have been investigated in numerous clinical trials. The mAbs and BiKEs augmented NK cell-killing activity mediated by ADCC. However, BiKEs simultaneously bind to the tumor antigen and the NK cell surface Fc receptor, potentially creating a bridge between NK cells and tumor cells and allowing them to act more effectively than mAbs. The use of NK cells engineered with a CAR structure is another type of NK cellbased immunotherapy for NHLs. CAR-NK cells, when equipped with cytokine receptors or cytokine genes, have demonstrated enhanced proliferation and prolonged survival in the patient's bloodstream. They can target TAAs with particular specificity and result in improved treatment responses. CIML NK cells with adaptive immune characteristics and long lifespans are also appropriate for this application, but they have not been well assessed in NHLs.

By reviewing the available clinical trial data, we concluded that NK cell-based approaches are generally well-tolerated, with no major safety concerns observed specifically GVHD. Overall, the available clinical trial data provide an encouraging foundation for the continued investigation and development of NK cell-based immunotherapies for the management of NHLs. The safety profile demonstrated in these studies, coupled with the potential for improved clinical outcomes, warrants further exploration of NK cell-based approaches, either as standalone therapies or in combination with other modalities, to improve the treatment landscape for patients with this complex hematological malignancy.

Abbreviations

HL	Hodgkin lymphoma
NHL	Non-Hodgkin lymphoma
mAb	Monoclonal antibody
SCT	Stem cell transplantation
CAR	Chimeric antigen receptor
R/R	Relapsed/refractory
CRS	Cytokine release syndrome
ICANS	Immune effector cell-associated neurotoxicity syndrome
GVHD	Graft-versus-host disease
NK cell	Natural killer cell
TME	Tumor microenvironment
DLBCL	Diffuse large B-cell lymphoma
MCL	Mantle cell lymphoma
BL	Burkitt lymphoma
FL	Follicular lymphoma
MZL	Marginal zone lymphoma
CLL/SLL	Chronic lymphocytic leukemia/small-cell lymphocytic
HSCT	Hematopoietic stem cell transplantation
CTL	Cytotoxic T-lymphocyte
BITE	Bispecific T cell engager
ILC	Innate lymphoid cell
HSC	Hematopoietic stem cell
MHC	Major histocompatibility complex
LGL	Large granular lymphocyte
TCR	T cell receptor
CLP	Common lymphoid progenitor
NKP	NK precursor
rIL	Recombinant interleukin
NKG2	Natural killer group 2
CCR	Chemokine (C–C motif) receptor
KIR	Killer-cell immunoglobulin-like receptor
DAP	DNAX-activation protein
NCR	Natural cytotoxicity receptor
ULBP	UL16-binding protein

MICA/B	MHC class I chain-related protein A and B
IDO	Indoleamine-pyrrole 2,3-dioxygenase
PGE2	Prostaglandin E2
DNAM-1	DNAX accessory molecule-1
GPI	Glycosylphosphatidylinositol
FcγRIII	IgG Fc region receptor III
ADCC	Antibody-dependent cell-mediated cytotoxicity
PD-1	Program-cell death receptor 1
PD-L1/2	PD-1/2 Ligand
Sialecs	Sialic acid recognizing immunoglobulin-like lectins
I AIR-1	Leukocyte-associated immunoglobulin-like recentor-1
TRAIL	TNE-related apontosis-inducing ligand
Fasl	Fas ligand
TRADD	TNE recentor-associated death domain
ТЦ	Helpor T coll
Trog	Pogulaton/T coll
neg	Departitie cell
	Denantic cell
TAIVI	Tumor-associated macrophage
IAN	lumor-associated neutrophil
MDSC	Myeloid-derived suppressor cell
IFH	Follicular helper I cell
MSC	Mesenchymal stromal cell
CAF	Cancer-associated fibroblast
ECM	Extracellular matrix
IFN	Interferon
TNF	Tumor necrosis factor
CTLA-4	Cytotoxic T lymphocyte-associated antigen 4
LAG-3	Lymphocyte activation gene 3
TGF	Tumor growth factor
CXCR	Chemokine (C-X-C motif) receptor
TFR	Follicular regulatory T cell
FoxP3	Forkhead box P3
TIM-3	T-cell immunoglobulin and mucin domain-containing 3
FDC	Follicular dendritic cell
GC	Germinal center
Aq-Ab complex	Antigen-Antibody complex
GM-CSF	Granulocyte-macrophage colony-stimulating factor
CXCI	Chemokine (C-X-C motif) ligand
APRII	A proliferation-inducing ligand
ICAM-1	Intercellular adhesion molecule 1
VEGE	Vascular endothelial growth factor
M-MDSC	Monocytic MDSC
PMN-MDSC	Granulocytic MDSC
Ara1	Arginase1
INOS	Inducible nitric oxide synthese
ROS	Reactive oxygen species
A NKC	Absolute NK coll count
FEC	Event free survival
DES	Drogression free survival
PF5 00	
	Drimary control nonyous system lymphoma
PCINSL	T cell increases and a state with the and ITMA democia
IIGII	I-cell immunoreceptors with ig and Hiwi domain
HGBL	High-grade B-cell lymphoma
	Inhibitory receptors immunoglobulin-like transcript 2
CNS-DLBCL	DLBCL of the central nervous system
	Death domain
MALIL	Mucosa-associated lymphoid tissue lymphoma
CBCL	Cutaneous B-cell lymphoma
ALCL	Anaplastic large cell lymphoma
LFA-3	Lymphocyte-function antigen 3
HLA	Human leukocyte antigen
ABC-DLBCL	Activated B-cell-DLBCL
GC-DLBCL	Germinal center-DLBCL
GVT	Graft-versus-tumor
SD	Stable disease
FDA	Food and drug administration
CR	Complete response
PR	Partial response
ORR	Overall response rate
PB	Peripheral blood
CB	Cord blood

iPSC	Induced pluripotent stem cell
RFS	Relapse free survival
haNk	High affinity NK cell
MM	Multiple myeloma
iNK	NK cells derived from iPSCs
HSPC	Hematopoietic stem and progenitor cell
BiKE	Bispecific killer engager
scFv	Single-chain antibody variable fragment
TAA	Tumor-associated antigen
SLAM	Signaling lymphocyte activation molecule
PBMC	Peripheral blood mononuclear cell
cNK cell	Conventional NK cell
CIML NK cell	Cytokine-induced memory-like NK cells
AML	Acute myeloid leukemia
DLI	Donor lymphocyte infusion
RL	Roser leukemia
ALL	Acute lymphocytic leukemia
LCL	Lymphoblastoid cell line

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Authors' contributions

MB: literature and clinical trial searches, conceptualization, writing – original draft, design of figures, tables preparations and editing. SY: writing – original draft, tables preparations and editing. SK: writing – original draft and editing. FAS, MS, ASJ and FD: writing – original draft. SMS: writing – original draft and editing. AI and SP: Editing of the final manuscript. AG, ER and AH: Supervision and validation. All authors read and approved the final version of the manuscript.

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