Clinical targeting recombinant immunotoxins for cancer therapy

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Abstract: Recombinant immunotoxins (RITs) are proteins that contain a toxin fused to an antibody or small molecules and are constructed by the genetic engineering technique. RITs can bind to and be internalized by cells and kill cancerous or non-cancerous cells by inhibiting protein synthesis. A wide variety of RITs have been tested against different cancers in cell culture, xenograft models, and human patients during the past several decades. RITs have shown activity in therapy of several kinds of cancers, but different levels of side effects, mainly related to vascular leak syndrome, were also observed in the treated patients. High immunogenicity of RITs limited their long-term or repeat applications in clinical cases. Recent advances in the design of immunotoxins, such as humanization of antibody fragment, PEGylation, and modification of human B- and T-cell epitopes, are overcoming the above mentioned problems, which predict the use of these immunotoxins as a potential therapeutic method to treat cancer patients.

Keywords: targeted therapy, hematologic malignancies, solid tumors, vascular leak syndrome, immunogenicity

Introduction
Recombinant immunotoxins (RITs) are chimeric proteins for cancer therapy that contain a toxin fused to a targeting moiety. After the initial success of antibody therapy for cancer, monoclonal antibodies (Mabs) were used to link with the toxin molecules, which have higher specificity in targeting and higher potency in killing cancer cells. Thirty-five years ago, RITs were created by chemically conjugating a whole protein toxin to an Mab or a protein toxin devoid of its natural binding domain.1,2 Other immune proteins such as cytokines and growth factors have also been conjugated and genetically fused to toxins.3 The traditional immunotoxins coupled the toxin and the targeting moiety by chemical method, but the cumbersome steps and the high cost encouraged the development of novel RITs. Nowadays, these RITs are generally synthesized by recombinant DNA techniques through constructing conditional expression plasmid and expressing interest protein in Escherichia coli rapidly and efficiently. RITs combine the cell-killing power of toxin and the specificity of antibody therapies. The careful design of both target moiety and toxin is the key of a successful therapy, because each type of cancer cell expresses different kinds of surface antigens. Compared with other commonly used immunotherapies, RITs’ cytocidal action is not dependent on antibody or complement-dependent cytotoxicity, but it is dependent on the specificity of the targeting moiety and the high activity of the toxin moiety, which will induce development of resistance. RITs directly target the surface of cancer cells, and 1 single internalized toxin molecule could kill the cell. Based on this, patients would not develop myelosuppression dose-limiting toxicity (DLT). Some RITs have been approved by the
US Food and Drug Administration (FDA) for the treatment of cancer, such as denileukin diftitox (ONTAK®). This review considers RITs with different target moieties and toxins directed to cancer cells and focuses on those that have been tested clinically in the recent decades.

**Mechanism of action of bacterial toxins**

The target moiety facilitates the transfer and internalization of RITs into cancer cells. For this purpose, various bacterial toxins have been used in RITs targeting cancer cells, and among these toxins, *Pseudomonas* exotoxin (PE) and *Diphtheria* toxin (DT) are the most commonly used bacterial toxins. Nearly all toxins kill cancer cells by enzymatically inhibiting protein synthesis. Similarly, both PE and DT also inhibit adenosine diphosphate (ADP)-ribosylate elongation factor 2 (EF-2) in the cytosol, which is a critical component of the protein synthesis machinery. Each toxin catalyzes the ADP-ribosylation of his-699 of EF-2, which is posttranslationally modified to a diphthamide residue. Despite their similar action, PE and DT differ greatly in amino acid sequence, and PE’s binding domain is located near to its amino terminal, but DT is located on the opposite terminal.

PE is a single-chain protein, consisting of 613 amino acids in length, which is further composed of 3 functional domains. Domain Ia (amino acids 1–252) is the binding domain, domain II (amino acids 253–364) mediates translocating the toxin to the cytosol, and domain III (amino acids 400–613) contains the ADP-ribosylating enzyme that inactivates EF-2 in the cytosol (Figure 1). The function of domain Ib (amino acids 365–399) is unclear. PE kills the cells following these steps (Figure 2): 1) Lys613 is removed by a carboxypeptidase in the plasma. 2) Domain Ia binds to the α2 macroglobulin receptor on animal cells and internalizes via endosomes to the Golgi. 3) The protease furin cleaves domain II between amino acids 279 and 280. 4) The disulfide bond that joins the 2 fragments generated by proteolysis is reduced. 5) Amino acids 609–612 (arginine–alutamic acid–aspartic acid–leucine, REDL) bind to an intracellular sorting receptor that transports the carboxy terminal fragment from the Golgi to the endoplasmic reticulum (ER). 6) Amino acids 280–313 mediate translocation of the toxin to the cytosol. 7) The ADP-ribosylating enzyme in amino acids 400–602 inactivates EF-2. 8) Though inhibition of protein synthesis is sufficient to induce cell death eventually, cell death from toxins is facilitated by apoptosis.

Full-length 535-amino acid DT is a single-chain protein containing an enzymatic A domain (amino acids 1–193) and a binding B domain (amino acids 482–535). A translocation or transmembrane (T) domain is located in the center of the molecule (Figure 1). DT undergoes the following steps to kill cells (Figure 2): 1) DT is proteolytically cleaved outside the cell, domain II (amino acids 253–364) mediates translocating the toxin to the cytosol, and domain III (amino acids 400–613) contains the ADP-ribosylating enzyme that inactivates EF-2 in the cytosol (Figure 1). The function of domain Ib (amino acids 365–399) is unclear. PE kills the cells following these steps (Figure 2): 1) Lys613 is removed by a carboxypeptidase in the plasma. 2) Domain Ia binds to the α2 macroglobulin receptor on animal cells and internalizes via endosomes to the Golgi. 3) The protease furin cleaves domain II between amino acids 279 and 280. 4) The disulfide bond that joins the 2 fragments generated by proteolysis is reduced. 5) Amino acids 609–612 (arginine–alutamic acid–aspartic acid–leucine, REDL) bind to an intracellular sorting receptor that transports the carboxy terminal fragment from the Golgi to the endoplasmic reticulum (ER). 6) Amino acids 280–313 mediate translocation of the toxin to the cytosol. 7) The ADP-ribosylating enzyme in amino acids 400–602 inactivates EF-2. 8) Though inhibition of protein synthesis is sufficient to induce cell death eventually, cell death from toxins is facilitated by apoptosis.

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the cell between Arg193 and Ser194, which is within a disulfide loop formed by Cys186 and Cys201. 2) DT binds on the cell surface via carboxyl residues 482–535 to a complex of heparin-binding epidermal growth factor (EGF)-like growth factor precursor and CD9. 3) DT internalizes in an endosome and unfolds at low pH, and the disulfide bond between amino acids 186 and 201 is reduced. 4) The TH8 (amino acids 326–347) and TH9 (amino acids 358–376) domains form a hairpin that inserts in the membrane of the endosome and forms a channel through which the enzymatic fragment translocates to the cytosol. 5) In the cytosol, nicotinamide adenine dinucleotide (NAD) binds to the active-site cleft of DT (amino acids 34–52), and the ADP ribose of NAD is transferred to EF-2. 6) Similar to PE, DT also induces cell death by apoptosis.

It has been shown that one or only a few of those toxin molecules delivered to the cytosol are sufficient to kill a target cell. However, the inhibition of protein synthesis by toxins
was thought to be lethal recently.26,27 Because some toxin-treated cancer cells appear to survive from toxin treatment which suggests the existence of resistance to RIT-mediated apoptosis. Therefore, assays that focus more precisely on the mechanisms of cell death have been developed by these toxins later. The activities of bacterial toxin and RITs have been connected to apoptosis in some cell systems, but the mechanisms have not been extensively stated.17,28,29 Keppler et al indicated that a PE-based RIT, B3(Fv)-PE38, kills MCF-7 breast cancer cell line by 2 mechanisms: one is due to the inhibition of protein synthesis caused by inactivation of EF-2 and the other requires caspase activation.30 There was also evidence that PE toxin-induced apoptosis of human mast cells involves downregulation of anti-apoptotic proteins and activation of caspase-8 and -3 pathways.31 When treated with BL22, B cells of chronic lymphocytic leukemia (CLL) underwent programmed cell death that was characterized by caspase-9 and caspase-3 activation, poly(adenosine diphosphate[ADP]-ribose)polymerase (PARP) cleavage, DNA fragmentation, and membrane flipping.32 From the above, cell death mediated by bacterial toxins, such as PE and DT, is often facilitated by caspase-dependent programmed cell death, in addition to the inhibition of protein synthesis.

### Mechanism of action of plant toxins

Plant toxins are classified into 2 classes, holotoxins and hemitoxins. Holotoxins, or class I ribosome-inactivating proteins, include ricin, abrin, modeccin, and mistletoe lectin. Hemitoxins, which are also referred to as class II ribosome-inactivating proteins, include gelonin, saporin, bouganin, and bryodin.33 Plant toxins such as ricin and gelonin also arrest protein synthesis but by inactivating the ribosome instead of EF-2.34,35 As shown in Figure 1, holotoxins such as ricin contain both binding and catalytic domains, whereas hemitoxins contain only catalytic domains. Only the catalytic domains of both holotoxins and hemitoxins translocate to the cytosol, and hence the binding domains of holotoxins would be removed by reduction of the disulfide bond prior to translocation. Plant toxins have been reported to prevent the association of EF-1 and EF-2 with the ribosomal subunit by removing the base A3724 in 28s ribosomal RNA (rRNA).36 Ricin also removes the neighboring base G3723. These toxin-mediated processes stimulate the apoptotic pathway, as well as the bacterial toxins, leading to cell death in the end. How plant toxins translocate to the cytosol from the cell surface is still unknown. The intracellular transport of ricin is dependent on sorting receptors that cycle between the ER and the Golgi.37

### RITs targeting hematologic malignancies

Generally, receptors of cytokines and growth factors are overexpressed in all types of cancer cells. In hematologic malignancies, several receptors are overexpressed compared with normal blood cells and have been targeted successfully in several studies (Table 1). Here, we discuss the recently published RITs targeting hematologic malignancies in clinical application with different surface markers.

#### Interleukin-2 receptor (IL-2R)

Interleukin-2 (IL-2), one of the first lymphokines to be identified, plays a central role in the clonal expansion of activated T cells by interacting with its specific cell surface

### Table 1 RITs targeting hematologic malignancies

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<td>Refractory AML</td>
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**Abbreviations:** RITs, recombinant immunotoxins; IL, interleukin; DT, Diphtheria toxin; CTCL, cutaneous T-cell lymphoma; HCL, hairy cell leukemia; ATL, adult T-cell leukemia; HD, Hodgkin’s disease; NHL, non-Hodgkin’s lymphoma; B-NHL, B-non-Hodgkin’s lymphoma; CLL, chronic lymphocytic leukemia; AML, acute myeloid leukemia; GM-CSFR, granulocyte-macrophage colony-stimulating factor receptor; NA, not applicable.
receptor (IL-2R). IL-2R, which binds the IL-2 with high affinity, is composed of 3 subunits (alpha [CD25], beta [CD122], and gamma [CD132]). IL-2R is overexpressed in hematologic malignancies such as adult T-cell leukemia (ATL), cutaneous T-cell lymphoma (CTCL), Hodgkin’s disease (HD), and other B- and T-cell leukemias and lymphomas. But only a small percentage of normal T cells are IL-2R positive. Thus, IL-2R has been broadly used to target leukemias and lymphomas.

**Denileukin diftitox (ONTAK, DAB$_{389}$IL-2)**

Denileukin diftitox was granted initial approval in 2001 for the treatment of CTCL. Targeting domain of this RIT does not contain an antibody but a recombinant human IL-2 fused to the C-terminus of the DT toxin. The ligand targets those cells that express the IL-2R, which is transiently expressed on activated T cells but constitutively present in a number of hematologic malignancies. It does not cause generalized DT-related toxicity because the binding parts of the DT are replaced by the IL-2 segment targeting only those cells expressing IL-2R.$^{38}$

In a single-arm Phase III trial, patients with IL-2R-positive CTCL received denileukin diftitox (9 or 18 μg/kg/d) for 5 consecutive days every 3 weeks. This study demonstrated 20% partial remission (PR), 10% complete regressions (CRs), and 30% overall response rate (ORR) in 71 patients. The duration of response ranged from 2.7 to >46.1 months with a median of 6.9 months. The side effects included flu-like symptoms, acute infusion-related events, and a vascular leak syndrome (VLS). Through measuring the neutralizing antibody of denileukin diftitox, there was no difference of tolerability at the 2 dose levels.$^{39}$

A later Phase III trial confirmed and improved the response rate of denileukin diftitox with placebo.$^{40}$ A total number of 144 patients with CTCL were assigned to 9 μg/kg/d denileukin diftitox, 18 μg/kg/d denileukin diftitox, or placebo infusion. The agents were administered for 5 consecutive days every 3 weeks. Compared with 15.9% ORR for placebo-treated patients (2% CR and 13.6% PR), 44% of patients (n=100) treated with denileukin diftitox achieved response (10% CR and 34% PR). Higher ORR was observed in the 18 μg/kg/d group than 9 μg/kg/d group (49.1% vs 37.8%, respectively). In addition, progression-free survival was significantly longer for patients treated with denileukin diftitox. This study demonstrated that denileukin diftitox had a significant and durable effect with an acceptable safety profile in patients with CTCL. Based on this study, Prince et al$^{41}$ examined the efficacy and safety of denileukin diftitox in 36 patients with IL-2R low expression skin biopsies. The result demonstrated that the safety profile of denileukin diftitox in IL-2R low expression disease was similar to that in IL-2R-positive disease, which suggests that IL-2R low expression would not preclude clinical response to denileukin diftitox in patients with CTCL.

Denileukin diftitox has also been reported to reduce the percentage of regulatory T cells (Tregs) in the peripheral blood of patients with renal carcinoma, ovarian cancer, and melanoma.$^{42-46}$ It is used infrequently because of poor tolerability and some kinds of side effects, but previously reported adverse effects can be managed effectively by supportive measures without dose reduction.$^{37}$ Hence, denileukin diftitox can be used for the treatment of CTCL currently.

**IL-2Rα (CD25)**

The human IL-2Rα, also described as the Tac antigen or CD25, is a 55-kDa membrane glycoprotein (p55). The deduced amino acid sequence of IL-2Rα predicts a mature protein of 251 amino acids with a signal peptide of 21 amino acids in length. The amino-terminal 219 residues constitute the extracellular region. The next 19 residues constitute the membrane spanning region, and the carboxy-terminal 13 residues, the cytoplasmic region. Mutational analysis showed that the N-terminal 83 residues of the IL-2Rα, especially residues 1–6 and 35–43, were essential for its binding function.$^{48}$

**LMB-2 [anti-Tac(Fv)-PE38]**

LMB-2, also named anti-Tac(Fv)-PE38, consists of a modified PE toxin and an antibody fragment. The antibody fragment, which contains the V$_{H}$ of anti-Tac fused to the V$_{L}$ via a 15-amino acid linker, selectively binds the α-subunits of IL-2R.$^{49}$ LMB-2 has been shown to be cytotoxic to CD25+$^+$ malignant cells that were either established cell lines or directly obtained from patients with hematologic malignancies.$^{50-53}$ LMB-2 produced CR in murine xenografts in vivo.$^{49}$ Blood levels of LMB-2 causing tumor regression in mouse are well tolerated by monkeys.$^{54}$

The Phase I trial of LMB-2 in patients with hematologic malignancies began in 1996.$^{55}$ In this test, 4 patients with CD25+$^+$ hairy cell leukemia (HCL) received LMB-2 at 3 dose levels (30, 40, and 63 μg/kg every other day [QOD] x3). All patients reacted to LMB-2 after a single cycle; 1 patient who received 2 cycles had a CR, marked by regression of HCL cells from blood and marrow, and did not relapse after 11 months. Three additional patients had 98%–99.8% reductions in malignant circulating cells. This study represents evidence that LMB-2 may be an effective new therapy for patients with CD25+$^+$ HCL.
To evaluate the pharmacokinetics, toxicity, immunogenicity, and antitumor activity of LMB-2, more patients with hematologic malignancies received dose levels that ranged from 2 to 63 μg/kg for a total of 59 cycles. Seven PRs were observed in patients with CTCL, HCL, HD, ATL, and CLL. One HCL patient achieved a durable CR for >20 months. All 4 patients with HCL responded to LMB-2. At the maximum tolerated dose (MTD; 40 μg/kg QOD ×3), toxicity was transient and included transaminase elevation and fever. Only 6 of 35 patients developed neutralizing antibodies after the first cycle. This study demonstrated that LMB-2 has clinical antitumor activity in CD25+ hematologic malignancies, especially HCL.

Although LMB-2 showed antitumor activity in Phase I trial, its application was limited by immunogenicity and rapid tumor growth between cycles. Currently, LMB-2 is being investigated in combination with fludarabine and cyclophosphamide for patients with ATL. In the previous report, treatment with fludarabine was associated with lower immunogenicity to murine antibodies, and the fludarabine–cyclophosphamide combination was associated with reductions in normal T and B cells. Patients received fludarabine (25 mg/m²) and cyclophosphamide (250 mg/m²) for 3 consecutive days before cycles began. Two weeks later, patients were treated cyclically every 3 weeks. Fludarabine and cyclophosphamide were administered on days 1, 2, and 3, followed by 30–40 μg/kg LMB-2 on days 3, 5, and 7. An ORR of 50% with 2 CRs and 2 PRs was achieved in 8 evaluable patients, 1 of 2 CRs occurred after 6 months only in a sanctuary site. The toxicity of LMB-2 was not increased by fludarabine and cyclophosphamide, while doses of fludarabine and cyclophosphamide used were also without DLT. With fludarabine and cyclophosphamide, normal T and B cells were reduced to 70% and 96% on average, respectively, which allowed more cycles to result in long-term remission. However, additional patients will be needed to determine if chemotherapy can delay immunogenicity of LMB-2 significantly.

**RFT5-dgA**

RFT5-dgA is an anti-CD25-ricin A chain RIT that is formed by a murine anti-CD25 monoclonal antibody (immunoglobulin G [IgG]) and a deglycosylated ricin A chain (dgA). RFT5-dgA was prepared by using the hindered heterobifunctional cross-linker to dgA. Bell et al evaluated the effect of depleting activated T cells with RFT5-dgA through an in vitro model of acute HIV infection, and the results showed that RFT5-dgA inhibited viral production by activated CD25+ HIV-infected cells and suppressed the breed of infection to uninfected T cells. The antitumor activity of this RIT was evaluated in severe combined immunodeficiency (SCID) mice with disseminated human Hodgkin’s lymphoma (HL) and in nude mice with subcutaneous solid HL. In these mice, 40 μg of RFT5-dgA decreased the diameter of >60% of solid Hodgkin and Reed–Sternberg (H-RS) tumors and inhibited the growth of HL in the majority of the treated mice. And RFT5-dgA was at least 7 times more effective than other ricin A chain-based RITs.

RFT5-dgA was tested in a Phase I trial that involved 15 patients with refractory HL. In this dose escalation trial, all patients received RFT5-dgA intravenously on days 1, 3, 5, and 7 for doses per cycle of 5, 10, 15, or 20 mg/m². After the treatment of 1–4 cycles, 7 of 15 patients made anti-ricin antibodies and 6 of 15 patients developed anti-mouse antibodies. Only 2 PRs were achieved, and 9 of 15 patients got progressive disease. Side effects were mainly related to VLS, which was characterized by decreases in serum albumin, hypotension, edema, myalgia, and tachycardia. In conclusion, RFT5-dgA showed encouraging efficacy at a dose level, and multiple cycles of treatment could be given without cumulative toxicities. In an extension of this Phase I trial, 5 additional patients were treated at the MTD (15 mg/m²). Tumor evaluations were performed after the end of completion of treatment. Overall, 2 patients at the MTD achieved PRs lasting 2 and 21 months, respectively. Of 20 patients, 10 showed progressive disease. Half of the patients developed anti-ricin antibodies or anti-mouse antibodies.

According to the preceding studies, more patients were enrolled in the subsequent trial at the MTD. A total of 27 patients were treated at this level, and all patients had signs of progressive disease before treatment with RFT5-dgA. Of 17 evaluable patients, 2 patients achieved PRs and 1 minor response (MR) and 5 stable diseases. Eleven of 16 patients receiving 2 or more cycles produced anti-ricin antibodies or anti-mouse antibodies, and the side effect in this study remained moderate and related to VLS.

**CD19**

CD19 is the hallmark differentiation antigen of the B lineage and has been proposed to serve as an important co-receptor molecule in conjunction with CD21 and CD81 for modifying signals generated through the B-cell antigen receptor complex. CD19 is a lineage-specific glycoprotein expressed on follicular dendritic cells and B cells. It is present on B cells from the earliest recognizable B lineage cells.
during development to B-cell blasts but is lost on mature plasma cells. CD19 has been used to diagnose cancers that raised from cells – notably B-cell lymphomas. Treatments targeting CD19 have begun to enter trials from 2012. Most experimental anti-CD19 drugs work by exploiting the presence of CD19 to treat B-cell cancers specifically. But another study indicated that CD19 plays an active role in driving the growth of these cancers, which suggests that CD19 and its downstream signaling may be a better therapeutic target.

**Anti-B4-bR**

Anti-B4-bR is an anti-CD19-blocked ricin (bR) RIT that contained an anti-B4 Mab chemically linked to intact ricin (A + B chain). The reactivity of anti-B4-bR with tissues lacking the B4 epitope was diminished by chemically blocking the natural binding site of the B chain. But the B chain of bR RITs can facilitate the efficient internalization of the A chain because both the ricin A and B chains have carbohydrate residues that are recognized by liver cells. Anti-B4-bR has been shown to be highly cytotoxic for B4-positive cells, and the cytotoxicity is restricted to these cells.

Anti-B4-bR induced clinical responses when administered either as continuous infusions or as a daily bolus. Thirty-four patients with relapsed or refractory B-cell neoplasms (twenty-six NHL, four CLL, four ALL) received 7-day continuous infusion of anti-B4-bR at dose levels ranging from 10 to 70 mg/kg/d. Potentially, therapeutically serum levels of anti-B4-bR could be maintained for 4 days in patients treated at the MTD. And there were 2 CRs, 3 PRs, and 11 transient responses (TRs). Another Phase I trial that enrolled 25 patients with refractory B-cell malignancies was conducted to detail the toxicity and clinical response data when anti-B4-bR was administered as daily bolus infusions for 5 consecutive days. One CR, 2 PRs, and 8 mixed responses/ TRs were observed after the treatment of anti-B4-bR. Human anti-mouse antibody and anti-ricin antibody were developed in 9 patients. These studies indicated that anti-B4-bR can be administered safely both by continuous infusion and as a daily bolus infusion. In a Phase I trial using anti-B4-bR immunotoxin, 11 of 12 patients with B-cell non-Hodgkin’s lymphoma (B-NHL) remained in CR to the treatment.

**HD37-dgA**

HD37-dgA was constructed by linking the murine HD37 Mab via a sterically hindered disulfide cross-linker, \( N\text{-succinimidyl-oxycarbonyl-\textalpha-methyl-\textalpha-(2-pyridyldithio)}\text{toluene (SMPT)}\), to dgA. Stone et al used 2 regimens, intermittent bolus infusion and continuous infusion, for the administration of HD37-dgA to patients with NHL in 2 concomitant Phase I trials. In the intermittent bolus regimen, 2, 4, 8, 16, and 24 mg/m\(^2\) of HD37-dgA were divided into 4 equal doses administered every other day. Of 23 evaluable patients, 1 patient achieved CR that persisted >40 months and 1 patient achieved PR (overall 9%). In the continuous infusion regimen, HD37-dgA was administered continuously at 2 dose levels (9.6 and 19.2 mg/m\(^2\)) for 8 days. One of 9 (11%) evaluable patients developed PR on the continuous infusion regimen. The MTD of each regimen was 16 and 19.2 mg/m\(^2\)/8 d, respectively. The DLT mainly consisted of VLS, aphasia, and acrocyanosis on the 2 regimens. Almost 25% of patients on the bolus infusion regimen and 30% on the continuous infusion regimen developed antibody against mouse Ig and/or ricin A chain antibody. This test showed that HD37-dgA can be administered safely and can be used in the later clinical trial. In the following Phase II trial, HD37-dgA was used as adjuvant treatment for patients with relapsed B-NHL combined with anti-B4-bR; 26 of 49 patients remained in CR.

**CD22**

CD22 is a molecule belonging to the sialic acid-recognizing immunoglobulin lectin (SIGLEC) superfamily of lectins. It is a B-lineage-restricted surface molecule that modulates B-cell receptor signaling and mediates cellular adhesion. Mature B cells express CD22 on their surface, while it is found to a lesser extent on some immature B cells. It prevents the overactivation of the immune system and the development of autoimmune diseases. It is also expressed on B cells in most B-cell leukemias and lymphomas; therefore, it is thought to be a potential targeting therapy for B-cell leukemias and lymphomas.

**BL22 [RFB4(dsFv)–PE38/CAT3888]**

BL22, which is also named RFB4(dsFv)–PE38 or CAT3888, was the first RIT designed to kill CD22 overexpressed cells. It contained a single-chain Fv of the anti-CD22 BFR4 antibody fused to a 38 kDa portion of PE toxin. In preclinical studies, BL22 was cytotoxic to a wide variety of CD22 cell lines and fresh malignant cells from patients. And it induced CRs in murine xenograft models at doses tolerated by cynomolgus monkeys. To assess the clinical activity of BL22, a dose escalation trial was approved. Of the 31 patients with B-cell cancers, 16 had HCL and were resistant to cladribine. Between 0.2 and 4.0 mg BL22 diluted in 50 mL of 0.2% albumin in 0.9% sodium chloride was administered intravenously every other
day for a total of 3 doses. Of the 16 patients, 11 had CRs and 2 had PRs, while 6 of them had a CR after only one cycle of BL22. Surprisingly, only 4 of the 16 patients generated neutralizing antibodies against the BL22 after cycles 4, 1, 2, and 4, respectively.

In the following study, 46 patients with B-cell hematologic malignancies were enrolled, of whom 11 patients had CLL, 4 patients had NHL, and 31 patients had HCL. Of the 31 HCL patients, 16 had a high response rate to BL22 in an interim report. BL22 was administered QOD ×3 doses per cycle, dose levels were 3–50 μg/kg. There were no drug-related deaths, and the MTD was determined to be 40 μg/kg. Of the 31 patients with HCL, 19 (61%) CRs and 6 (19%) PRs were achieved, and the ORR was 81%. Only one CLL patient had a PR. The most common DLT was neutropenia with a median IC₅₀ = 9.8 ng/mL to CD22+ fresh bone marrow or blasts from children with acute lymphoblastic leukemia (ALL) and also prolonged leukemia-free survival of xenografts from children with acute lymphoblastic leukemia (median IC₅₀). The results showed that BL22 was cytotoxic (median IC₅₀ = 9.8 ng/mL) to CD22+ fresh bone marrow or blasts from children with acute lymphoblastic leukemia (ALL) and also prolonged leukemia-free survival of xenografts. Although no obvious responses were achieved in patients with CLL, 4 patients had NHL, and 31 patients had HCL.

In this HCL clinical trial, the highest dose for treatment was in an interim report. Remarkable difference had been noted between moxetumomab pasudotox, CAT-8015, or HA22. Moxetumomab pasudotox contains Thr-His-Trp instead of Ser-Ser-Tyr at positions 100, 100A, and 100B in the antigen-binding site of V₃. It was determined to have higher cytotoxicity and a 14-fold improved binding affinity, as a result of lower off-rate.

To determine its safety and efficacy in the treatment of HCL, 28 patients were enrolled in a Phase I trial. Moxetumomab pasudotox was given at 5–50 μg/kg QOD ×3 for 1–16 cycles (median, 4 cycles), including 3 patients each at 5, 10, 20, and 30 μg/kg, 4 patients at 40 μg/kg, and 12 patients at 50 μg/kg. After all 114 cycles of moxetumomab pasudotox were administered, major responses were observed at all dose levels. The ORR ranges from 67% to 100% at each dose level without apparent correlation with dose. The ORR of 86% was achieved in 28 patients, whereas only one patient (5%) made neutralizing antibodies after cycle 1. Neutralizing antibodies were detected in 1 of 20, 1 of 13, and 1 of 9 patients after cycles 2, 3, and 4, respectively, but not in patients receiving 5–16 cycles. Thus, it permitted us to retreat most of the HCL patients to increase the chance and degree of response. Remarkable difference had been noted between moxetumomab pasudotox and BL22 in the results of clinical trials. In this HCL clinical trial, the highest dose for treatment was 50 μg/kg, a level higher than the MTD for BL22, and DLT was not observed. The ORR of moxetumomab pasudotox, 86%, was higher than the 72% ORR of the Phase II trial of BL22, and response rates at all dose levels were high.

Arons et al analyzed the plasma levels of 49 patients who achieved moxetumomab pasudotox to determine the relationship between response and high CD22 density on HCL cells. Those analyzed included the original 28 enrolled prior, an additional 20 at the highest dose level (50 μg/kg), and 1 additional patient enrolled prior. Moxetumomab
pasudotox was administered to 49 patients at different dose levels every other day for 3 doses. There were 28 (57%) CRs and 88% ORR of 49 patients, while CRs at the highest dose level and lower doses were 64% and 44%, respectively. The difference in micro-residual disease (MRD)-free CR rate was significant (39% vs 6%, P=0.02). Correlations between dose level and both peak level and area under the curve were varied in each dose level. At the highest dose level, CR was more likely with lower volume of distribution and clearance. The pharmacokinetics analysis showed that moxetumomab pasudotox can achieve durable CR in relapsed and refractory HCL patients without MRD. This study also indicates that lower HCL tumor burden minimizes the CD22 “sink” effect allowing higher plasma levels and suggests that patients with lower tumor burden may improve the chance of the durable CR. Furthermore, no DLT or HUS cases were observed in any of the cycles administered to 49 patients.99 Moxetumomab pasudotox is the only agent that can eliminate MRD in HCL in a high percentage of patients without causing myelosuppressive toxicities.

RFB4-dgA

CD22 molecules expressing the RFB4 epitope are present in 60%–70% of NHL cells. RFB4-dgA was prepared with the anti-CD22 Mab, RFB4, coupled to chemically dgA via the heterobifunctional cross-linker (SMPT). The antibodies used to construct RITs targeting RFB4 contain a mouse IgG1-k and a Fab’ of RFB4. RITs prepared by these antibodies have had notable potential for patients with NHL. Phase I studies of IgG-RFB4-dgA and Fab’-RFB4-dgA have been completed using an intermittent bolus regimen.96,97 Patients with relapsing NHL were treated with Fab’-RFB4-dgA, via 4-h bolus infusion regimen every second day, 5 of 14 (36%) evaluable patients achieved PRs.96 IgG-RFB4-dgA was tested in a similar NHL patient group treated by the same regimen; 1 CR and 5 PRs (ORR, 25%) were observed in 24 patients,97 and a continuous infusion Phase I study of IgG-RFB4-dgA has also been completed.98 In this subsequent Phase I trial, IgG-RFB4-dgA was administered over 8 days with comparable clinical responses (4 of 18 PRs, 22%).98 In all these trials, 20%–40% of the patients achieved responses and the DLT mainly consisted of VLS.99

Combotox (RFB4-dgA + HD37-dgA)

Combotox is a 1:1 mixture of HD37-dgA and RFB4-dgA, which are RITs that target the CD19 and CD22 antigens, respectively. A previous Phase I trial showed that RFB4-dgA induced 24% PRs and 1 long-lasting CR by continuous infusion regimen.98 A previous Phase I trial using continuous infusion of HD37-dgA also showed evidence of antitumor activity,78 concordant with the less potent action of it in vitro,100 and in SCID mice with Daudi lymphoma.101 Preclinical data showed that combotox is effective in killing cells in the Daudi disseminated lymphoma in SCID mice.101

To determine the MTD, clinical pharmacology, toxicity, and clinical responses of combotox, a Phase I trial was conducted involving 22 patients with refractory B-cell lymphoma.102 All patients expressed CD19 and CD22 on at least 30% of their tumor cells. Patients received a continuous infusion of combotox at 3 dose levels ranging from 10 to 30 mg/m²/192 h. After the treatment, only patients with circulating tumor cells (CTCs) in peripheral blood tolerated all doses without major toxicity, and prior therapies of these patients appeared to have little impact on toxicity. Analysis of the results indicated the fact that both lots of HD37-dgA tested in the trial had a tendency to aggregate after thawing, and the multimerization of HD37-dgA may have contributed to patient toxicity. Toxicities induced by combotox in this trial, including VLS and HUS, were similar to other types of RITs reported in previous trials. However, combotox appeared to be safe in patients with even minimal numbers of CTCs.102

Combotox, the mixture of HD37-dgA and RFB4-dgA, can bind to and kill human precursor-B-ALL blasts in vitro, and the combination was more effective than either RIT alone.103 Encouraging clinical data were observed in pediatric patients with pre-B-ALL.104 Hence, combotox is a new candidate for the treatment of patients with relapsed B-ALL. In the subsequent Phase I trial, combotox was administered to adult patients with refractory or relapsed B-ALL. Seventeen patients received combotox at 5 different dose levels (3, 5, 6, 7, and 8 mg/m²) by intravenous infusion. A cycle of treatment of combotox included 3 doses every other day for 3 doses. There were 28 (57%) CRs and 88% ORR of 49 patients, while CRs at the highest dose level and lower doses were 64% and 44%, respectively. The difference in micro-residual disease (MRD)-free CR rate was significant (39% vs 6%, P=0.02). Correlations between dose level and both peak level and area under the curve were varied in each dose level. At the highest dose level, CR was more likely with lower volume of distribution and clearance. The pharmacokinetics analysis showed that moxetumomab pasudotox can achieve durable CR in relapsed and refractory HCL patients without MRD. This study also indicates that lower HCL tumor burden minimizes the CD22 “sink” effect allowing higher plasma levels and suggests that patients with lower tumor burden may improve the chance of the durable CR. Furthermore, no DLT or HUS cases were observed in any of the cycles administered to 49 patients.99 Moxetumomab pasudotox is the only agent that can eliminate MRD in HCL in a high percentage of patients without causing myelosuppressive toxicities.

CD30

CD30 shows sequence homology to members of the tumor necrosis factor (TNF) receptor superfamily and is expressed only in activated but not resting T and B cells. It was initially described as a surface marker on neoplastic cells of HD.106 Screening for CD30 expression in normal and neoplastic cells led to recognition of CD30 as an activation-induced...
antigen mostly expressed on lymphoid cells and to the identification of a new category of NHL: CD30/Ki-1-positive anaplastic large-cell lymphoma (ALCL) by using monoclonal antibodies. The CD30/CD30 ligand system triggers cytolytic cell death in malignant lymphoma cell line and induces proliferation and cytokine production in T cells or neutrophils.

Ki-4.dgA

Ki-4.dgA was constructed by linking the anti-CD30 Mab (ki-4) via a sterically hindered linker to dgA. Normal human organs revealed no major cross-reactivity of the anti-CD30 Mabs. Ki-4.dgA was 5 times more potent in vitro than other anti-CD30 dgA-RITs tested previously and showed high efficacy in the treatment of human HL in SCID mice. Therefore, ki-4.dgA was selected for a Phase I trial in patients with refractory CD30+ HL and NHL.

In the first Phase I trial that aimed to determine the MTD, DLT, antitumor activity, and pharmacokinetics of ki-4.dgA, 17 patients with relapsed CD30+ lymphoma received escalating doses (5, 7.5, or 10 mg/m²/cycle) QOD ×4 of the RIT for 1–3 cycles. One PR and 1 MR were achieved in 15 evaluable patients. Side effects and DLT were also associated with VLS, which is similar to RIT RFT5-dgA. Seven of 17 (40%) patients developed anti-ricin antibodies, and only one patient made anti-mouse antibodies.

RITs targeting other molecules

CD33 and HUM-195/Rgel

CD33 is a surface protein that is found on hematopoietic colony-forming cells but not in their more primitive precursors. CD33 has structural and binding characteristics that identify it as a member of the sialoadhesin family. Studies using samples from patients indicate that in 90% of patients with AML, leukemic cells express CD33. HUM-195/Rgel was prepared by the humanized anti-CD33 Mab, M195, conjugated to gelonin. HUM-195/Rgel was used in CD33+ cell lines and xenografts in nude mice. In a Phase I trial of refractory myeloid leukemias, HUM-195/Rgel induced a 38%–50% decrease in peripheral blood blasts or bone marrow blasts in 7 of 28 patients, and there was no CR or PR.

CD3 and UCHT1 (A-dmDT390-bisFv)

The CD3 antigen was first identified as a glycoprotein that is present on the surface of all human T lymphocytes. As more diverse labeling and immunoprecipitation methods were used, the structure of the CD3 antigen appeared to be more complex. The CD3 was expressed as a complex composed of 4 glycoproteins (CD3-γ, δ, ε, and ζ). CD3 is overexpressed in T-cell malignancies and is a potential target for the treatment of these diseases. UCHT1 (A-dmDT390-bisFv) is a kind of RIT composed of 2 single-chain Fv fragments of an anti-CD3ε Mab fused to DT390. UCHT1 was administered to 5 patients with CTCL by intravenous infusions. Two of 5 patients had PRs lasting 1 and >6 months. Anti-DT antibodies developed in all patients after 2 weeks, and the side effects of UCHT1 were fever, nausea, chills, hypoalbuminemia, and transaminasemia.

IL-3R and DT388-IL3

IL-3 is a cytokine that supports the proliferation and differentiation of myeloid progenitors, but it is absent from mature myeloid cells. It is overexpressed in myeloid leukemic progenitors and used as a therapeutic target. DT388-IL3, an RIT fused with a Met-His linker, showed antitumor activities in an SCID model of acute myeloid leukemia (AML). A Phase I trial of DT388-IL3 was constructed in patients with chemorefractory AML and myelodysplasia (MDS). One CR and 1 PR were observed out of 40 evaluable patients with AML and 1 PR out of 5 patients with MDS.

GM-CSFR and DT388-GM-CSF

Granulocyte-macrophage colony-stimulating factor (GM-CSF) is a cytokine responsible for the growth and differentiation of granulocytes and macrophages. GM-CSF receptor (GM-CSFR) is overexpressed in leukemic cells and targeted in leukemia. The RIT’s GM-CSF-PE38KDEL and DT388-GM-CSF exhibited specific cytotoxicity in leukemic cell lines and patients, and DT388-GM-CSF was more cytotoxic than GM-CSF-PE38KDEL. Thirty-one patients with refractory AML were treated with DT388-GM-CSF in a Phase I trial, all of those patients were resistant to chemotherapy. One CR and 2 PRs were observed among these patients, and the major DLT was cytokine release syndrome. Neutralizing antibodies against DT were observed in 28 of 31 patients.

RITs targeting solid tumors

RITs have produced many durable CRs in refractory HCL, where patients rarely develop neutralizing antidrug antibodies (ADAs) to the toxin component of the RIT. Targeting solid tumors with RITs is much more difficult than targeting hematologic malignancies. Not only because the cellular junctions between cells are tighter and the tumor was more tightly packed, but also the patients are less immunosuppressed and more likely to develop neutralizing
antibodies to the toxin. Later, we discuss the recently published RITs targeting solid tumors (Table 2).

**Mesothelin (MSLN)**

MSLN is a cell-surface glycoprotein that is normally expressed only in mesothelial cells, and it is also overexpressed in many solid tumors including mesothelioma,28 pancreatic adenocarcinoma,29,30 cholangiocarcinoma,31 nonmucinous ovarian cancer,32 triple-negative-type breast cancer,33 gastric cancer,34,35 cervical cancer,36 and lung adenocarcinoma.37 Otherwise, MSLN is not critical because MSLN knockout mice grow normally and have no discernible phenotype.38 All the above make MSLN one of the few targeting antigens with sufficient differential expression to allow safe treatment of solid tumor.

**SS1P [SS1(dsFv)-PE38]**

Phage display technology was used to generate new Fvs binding to MSLN. SS1P is an RIT obtained that underwent affinity improvement, which consists of the PE38 fragment fused to a murine anti-MSLN variable antibody fragment. SS1P was developed for systemic therapy of patients with MSLN-positive solid tumors. One Phase I, dose escalation study of single-agent SS1P was performed in 34 patients with MSLN-expressing advanced mesothelioma (n=20), ovarian (n=12), and pancreatic cancers (n=2).39 Dose level escalated from 8 to 60 μg/kg, and there were 3 patients enrolled at each dose level. The initial cohort of 17 patients received SS1P QOD ×6 doses, and the MTD was 18 μg/kg. DLT included urticaria (1 patient) and grade 3 VLS (2 patients). To allow further dose escalation, 17 patients were treated QOD ×3 doses and MTD was 45 μg/kg. But SS1P was well tolerated at the highest dose level with pleuritis as the DLT. Following this study, another Phase I trial was performed in patients with chemoresistant solid tumor expression MSLN.40 SS1P was administered by continuous infusion for 10 days and repeated cycles at 4-week intervals until the observation of neutralizing antibodies or progressive disease. Twenty-four patients received 4, 8, 12, 18, and 25 μg/kg/d ×10. One of 6 patients, who received the maximum dose level of SS1P, had DLT because of the reversible VLS. Neutralizing antibodies were observed in 18 (75%) of 24 patients, and 5 (21%) patients received a second cycle. Only one patient had a PR. These 2 Phase I trials showed similar efficacy and toxicity of SS1P by different administration schedule. The majority of patients developed ADAs after their first cycle. The main DLT included on-target pleura. These studies demonstrated the need for an effective means to suppress the host immune reaction to SS1P.

So SS1P was tested in patients with newly diagnosed malignant mesothelioma in combination with standard cisplatin and pemetrexed.41 Pemetrexed (500 mg/m² on day 1) and cisplatin (75 mg/m² on day 1) were administered every 3 weeks for up to 6 cycles with escalating doses of SS1P. SS1P was administered i.v. on days 1, 3, and 5 during this study, the toxicity of the combination was similar to that observed with the single agent. Grade 3 toxicities associated with SS1P included fatigue, hypoalbuminemia, back pain, and hypotension. The grade 3 fatigue was dose limiting in only 1 patient at 55 μg/kg. Of 20 evaluable patients, 12 (60%) had a PR. Of 13 patients who received the MTD (45 μg/kg), 10 (77%) had a PR. This compares favorably to the response rate (41.3%) reported for pemetrexed and cisplatin alone.42 Overall, SS1P given with pemetrexed and cisplatin is safe and exhibits significant antitumor activity in patients with pleural mesothelioma, but the hematologic suppression caused by the chemotherapy failed to delay development of neutralizing ADAs.

### Table 2 RITs targeting solid tumors

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**Abbreviations:** RITs, recombinant immunotoxins; PE, Pseudomonas exotoxin; MSLN, mesothelin; IL, interleukin; EpCAM, epithelial cell adhesion molecule.
In a pilot study, SS1P was tested in combination with the pentostatin plus cyclophosphamide immunosuppressive regimen. After depleting T and B cells through immunosuppressive regimen, antibody formation was largely delayed, thereby allowing multiple cycles of therapy with SS1P that was previously limited to 1 therapeutic cycle. The immune modulation regimen allowed 2 patients to receive 4 or 6 cycles of SS1P. In this study, of 10 patients with advanced, refractory mesothelioma, 3 patients experienced durable response that persisted for >18 months and 2 patients responded to chemotherapy after immunotoxin therapy. As a result, it is essential to reduce the immunogenicity of SS1P to develop better therapeutic effect.

**Ovarian antigen and OVB3-PE**

OVB3-PE is the first PE-based RIT that was tested in a clinical trial. It consisted of a murine antibody that targets an unknown antigen on ovarian cancer cells fused to the entire PE toxin, but it failed to be used in more patients after its first clinical trial. OVB3-PE was administered to 23 patients with refractory ovarian cancer intraperitoneally, but it showed a high level of non-specific toxicity, including central nervous system (CNS) toxicity, transient elevation of liver enzymes, and gastrointestinal (GI) toxicity. Moreover, 100% of the patients made antibodies against the PE toxin 14 days after the first therapy. Human anti-mouse antibodies were also detected in 75% of patients 28 days after therapy.

**CD25 and LMB-2**

LMB-2 is a CD25-directed PE-based RIT that was usually used to treat CD25+ hematologic malignancies. Considering CD25+ CD4+ Treg cells regulate peripheral self-tolerance and possess the ability to suppress antitumor responses, LMB-2 was administered to 8 patients with metastatic melanoma followed by Melanoma Antigen Recognized By T-cells 1 and gp100-specific peptide vaccination. LMB-2 administration resulted in a reduction in Treg numbers and did not augment the immune response to cancer vaccination. This study indicated that LMB-2 can selectively mediate a transient partial reduction in circulating and tumor-infiltrating Treg cells in vivo.

**ErbB2**

The ErbB receptors constitute a group of related transmembrane proteins that belong to the subclass I of the receptor tyrosine kinase superfamily. Four members of this family have been identified: ErbB/epidermal growth factor receptor (EGFR), ErbB2 (HER2/Neu), ErbB3 (HER3), and ErbB4 (HER4). Overexpression of ErbB glycoprotein and its tyrosine kinase activity induces loss of growth control and plays an important role in the development of several human cancers. ErbB2 has been reported to be minimally expressed in normal tissues. Thus, ErbB2 is an attractive target for immunotherapy.

**ERB-38**

ERB-38 is an RIT composed of the Fv portion of Mab e23 that reacts with ErbB2, fused to PE38, a truncated form of PE. The IC$_{50}$ of ERB-38 is 0.2–4 ng/mL on the various ErbB2-positive tumor cell lines. ERB-38 is capable of causing CR in nude mice bearing epidermoid carcinoma and breast cancer. Then ERB-38 was administered in a Phase I trial in patients with advanced carcinoma. In this trial, 5 breast cancer patients and 1 esophageal cancer patient were treated with 3 doses of ERB-38. But hepatotoxicity was observed in all patients. Immunohistochemistry showed the presence of ErbB2 on hepatocytes.

**ScFv(FRP5)-ETA**

ScFv(FRP5)-ETA is a recombinant single-chain antibody toxin that contained a scFv portion of murine Mab FRP5, which recognizes the extracellular domain of ErbB2, linked to a truncated ETA. ScFv(FRP5)-ETA displayed potent antitumoral activity against a wide range of tumor cells including breast and ovarian carcinomas, prostate carcinomas, and squamous cell carcinomas. ScFv(FRP5)-ETA effectively inhibited growth of established murine tumor xenografts. ScFv(FRP5)-ETA was applied first in 11 patients with metastatic breast and colorectal cancers and with malignant melanoma from 4 clinical centers. ScFv(FRP5)-ETA was administered by intratumoral injection into cutaneous lesions for 7–10 days. Of 10 evaluated patients, treatment-induced shrinkage of tumors was observed in 6 patients and 4 CRs and 2 PRs among these 6 patients. Systemic liver toxicity was observed only in 1 patient treated at the highest daily dose levels.

**Lewis Y**

Lewis Y (Le$^+$) is overexpressed as a surface membrane component of many solid tumors, and it is also expressed on gastrointestinal epithelium and in the pancreas.

**LMB-1**

LMB-1 was the first RIT reported to have anti-tumor activity targeting an epithelial tumor. It is composed of a Mab that recognizes Le$^+$, B3, chemically linked to PE38. The clinical test of LMB-1 was conducted in 38 patients.
with LeY-positive solid tumors who had failed conventional therapy. Objective antitumor activity was observed in only 5 patients, and 1 CR was observed in a patient with metastatic breast cancer. A tumor reduction >75% and resolution of clinical symptoms lasting for >6 months were observed in a colon cancer patient. The major toxicity of LMB-1 was VLS, but 33/38 of patients made antibodies against LMB-1 3 weeks after the first cycle of treatment.\textsuperscript{161}

**SGN-10 (BR96 sFv-PE40)**

SGN-10 is an RIT consisting of sFv regions of the murine Mab, BR96, fused to a binding-defective portion of PE toxin (PE40).\textsuperscript{162} BR96 binds to the LeY carbohydrate antigen that is overexpressed on the surface of many human solid tumors.\textsuperscript{160} SGN-10 showed significant antitumor activity in murine xenografts of human breast and lung tumors and made CR in these tumor xenografts.\textsuperscript{163,164} On the basis of these favorable data, SGN-10 was developed for clinical trial in 46 patients with LeY-positive advanced carcinomas.\textsuperscript{165} In this test, cohorts of 3 patients were treated at each dose level on days 1, 4, 8, and 11, followed by a 14-day break. A treatment cycle was 28 days, and patient received 2 or more cycles until there was disease progression or unacceptable toxicity. No CR or PR was observed after 8-week treatment, although 31% patients had stable disease. In this study, DLT was due to GI toxicity rather than to VLS. The immunogenicity of the toxin portion limits the ability of SGN-10 by day 11 of therapy.\textsuperscript{165}

**IL-4R and NBI-3001 [IL-4(38-37)-PE38KDEL]**

IL-4R is overexpressed on many different tumor surfaces. Human malignant glioma cell lines have been shown to express IL-4R,\textsuperscript{166,167} and primary cell lines from glioma also express IL-4R,\textsuperscript{168} but not normal brain cells. To treat these tumors that overexpress IL-4R on their surface, NBI-3001 was composed of circularly permuted IL-4 fused to a truncated portion of PE. NBI-3001 was highly cytotoxic to glioblastoma cell lines\textsuperscript{169} and made CRs in murine xenografts of human glioma in all the animals.\textsuperscript{170} NBI-3001 was first used in 9 patients with malignant high-grade gliomas.\textsuperscript{171} Six of 9 patients showed glioma necrosis, and one of them also showed extensive necrosis of tumor leading to CR. Additional patients were treated to determine the appropriate dose level for patients with malignant glioma.\textsuperscript{172} Thirty-one patients with astrocytoma were enrolled in this following trial, and 25 of them were diagnosed with glioblastoma multiforme (GBM), while the other 6 patients were diagnosed with anaplastic astrocytoma. The results showed decreased signal intensity in the tumor consistent with tumor necrosis after the treatment in most patients.

**EpCAM and VB4-845**

Epithelial cell adhesion molecule (EpCAM) is overexpressed in many carcinomas relative to normal tissue, as in the case of transitional cell carcinoma (TCC).\textsuperscript{173,174} TCC refers to those bladder tumors derived from urothelial tissue. And increase of EpCAM expression was regarded as these cancers progress from lower to higher grades.\textsuperscript{175,176} Thus, EpCAM is a potent clinically relevant antigen for targeted treatment of bladder cancer. VB4-845 is an RIT that targets EpCAM+ cancer cells. It contains an anti-EpCAM humanized scFv fused to a truncated form of PE that lacks the cell-binding domain.\textsuperscript{177} VB4-845 was tested in a clinical Phase I trial to determine its safety, tolerability, immunogenicity, and efficacy.\textsuperscript{178} Sixty bacillus Calmette-Guerin (BCG)-intolerant patients with TCC or in situ carcinoma were enrolled in this test. CRs were observed in 39% of patients after the treatment. Although the majority of patients developed antibodies against the toxin portion of VB4-845, VB4-845 therapy was safe and without the most adverse events.

**Discussion**

The introduction of the “magic bullet” concept by Paul Ehrlich led to the search for agents that can selectively target cancer cells. After the initial success of antibody therapy for cancer, Mabs reacting with cancer cells became widely available. The first RITs that consist of a protein toxin fused to a Mab targeting moiety were constructed in the early 1980s. From then on, toxins from a variety of plants and bacteria were investigated, as well as the continuous optimization of the targeting moiety. To make the RIT more suitable for clinical development, portions of the toxin that were not essential for processing or cytotoxic activity were deleted from the sequence, and point mutations were created in the native toxins to improve activity, limit immunogenicity, or reduce off-target toxicity. Nowadays, the generation of an RIT involves the genetic fusion of a modified form of the toxin and a cell-selective ligand. The ligand can be a recombinant antibody or an antibody fragment, carbohydrate antigen, growth factor, or tumor-associated antigen. To be superior to conventional treatments, the ligand must be directed toward antigens that are exclusively or at least preferentially expressed on tumor cells compared to normal tissues. Meanwhile, one or only a few of toxin molecules delivered to the cytosol is sufficient to kill a target cell. So RITs perfectly combine the high specificity of targeting ligand and the excellent cytotoxic activity of toxins.
As one of the most commonly used antibody drugs, antibody–drug conjugates (ADCs) also have the higher specificity and lower toxicity compared with standard chemotherapy. ADCs consist of an Mab chemically attached to a highly toxic chemotherapeutic agent for use in traditional systemic therapy for cancers. The antibody portion localizes the drug to the tumor but limits its deposition elsewhere, increases antitumor activity, and decreases systemic toxicity of the drug. Although ADCs have so many advantages, RITs have several favorable properties not shared by ADCs. First, the RITs induced the cell death by disrupting the process of protein synthesis (Figure 2). And RITs could bring cell death by activation of caspases, which means that RITs can also be used to treat apoptosis-resisted cancers. Second, the mechanism action of novel RIT allows easy combination with standard agents. Third, unlike chemotherapy agent used in ADCs, RITs can effectively kill nondondiving cells, and RITs have little cross-resistance with other chemotherapy agents. And last, ADCs can cause off-target toxicity due to inappropriate payload, but RITs do not have this problem. In addition, RITs and ADCs share the same principles of selecting targets for therapy. But the requirement of differential expression is more stringent for RITs, so many targets suitable for ADCs are not suitable for RITs.

Because of higher specificity of RITs, selecting differential expressed surface markers is more essential than the category of toxin. Otherwise, RITs will induce some side effects and toxicities. Variety of toxicities has been observed with RITs that have limited the long-term treatment and efficacy in clinical practice. The most common toxicity for these agents is VLS characterized by weight gain, generalized edema, hypoalbuminemia, and orthostatic hypotension. RITs-mediated damage to endothelial cells appears to be responsible for VLS, because the cytotoxic protein must traverse endothelial cells to exit the blood vessels. Studies have shown that PE binds directly to endothelial cells, while truncated PE requires a ligand that reacts with the endothelium and a mutant form of PE showed less VLS. In addition, mutant toxins that lack enzymatic activity do not cause VLS, suggesting that VLS is an off-target effect of RITs. Ricin toxin contains short amino acid motifs that bind endothelial cells. Modification or deletion of these motif sequences reduced toxin-induced VLS. Most recently, Weldon et al deleted nearly entire PE domain II to prevent VLS, while preserving 2 of 3 putative endothelial binding motifs. Clinical factors may also induce the VLS. Clinical factors, such as administration of RITs, vary among patients that may affect the severity of VLS. Ricin-based RITs have reported more severe VLS than PE-based RITs. A retrospective study of HL patients showed that patients with a history of prior radiation therapy will have more frequent and more severe VLS. In addition to mutations of short amino acid motifs of toxin part, patients receiving premedication with dexamethasone have also been shown to have less severe VLS.

As a novel class of immune therapeutics, RITs have been used in the treatment of many kinds of tumors. The clinical responses present a promising approach to fighting cancers. RITs produced limited responses in relapsed and refractory hematologic malignancies mainly because of their side effects and high immunogenicity. Nevertheless, their half-lives were too limited for diffusion to occur in solid tumors. Immunogenicity of protein drugs and antibody drugs suddenly attracts broad attention around the world. For regulatory agencies, immunogenicity assessments are required for the licensure of all biologics to ensure safety and efficacy of the proteins. The causes mainly include 2 aspects: 1) immunogenicity will affect safety and efficacy of the drugs, even life-threatening interaction with endogenous protein and 2) lacking of efficient predictive tools means ADAs can be detected only in late Phase III trials after significant expenses have accrued. Based on a wide range of clinical trials, the incidence of immunogenicity after 1 cycle of RIT ranges from 50% to 100% for solid tumors and from 0% to 40% for hematologic tumors. Patients with solid tumors are more likely to develop neutralizing antibodies to the toxin because of their less immunosuppression. The presence of neutralizing antibodies lowers the level of active RITs and their efficacy. There is a multitude of factors responsible for ADA formation against RITs. Studies on immunogenicity to RITs indicate that ADA had a strong correlation with a decrease in drug serum concentrations and resultant reduced efficacy. ADAs impact the PK of RITs in diverse ways. They can enhance clearance besides sustaining the circulation of RITs. ADA–RIT complexes circulating in the bloodstream trigger regular endogenous elimination processes that are mediated by the reticuloendothelial system, predominantly phagocytic cells in the liver and spleen. The complexes are internalized and undergo subsequent lysosomal degradation. At the same time, ADA–RIT complexes are often clearly slower than the free activated RIT, and compensatory upregulation of shed target may result in concentration increases in total, while “free” concentrations are actually decreasing. Moreover, the mechanism of action of an ADA-induced impact on the PK of RITs still needs further and extensive investigation.
At present, several approaches have been used to prevent the development of neutralizing antibodies of RITs in patients. 1) To diminish the immunogenicity of RITs, the new generation of RITs consists of a humanized Fab or Fv fragment of antibody. It was reported that humanized RIT lost some of its epitopes. But the majority of the antibodies that have been found were against the toxin portion of RITs. Genetic engineered single-chain variable fragments (scFVs) also are fused to toxins instead of full-size antibodies. Compared with full-size Mab therapeutics, low immunogenicity scFV toxin therapeutics has several pros. Antibody fragments, such as scFVs, penetrate tissues and tumors more rapidly and deeply than full-size Mab. In addition, the scFVs have been suggested to permit binding to cryptic epitopes not accessible to full-size Mabs.

2) The most useful method for some biologic agents, such as interferon and L-asparaginase, is PEGylation. Covalent attachment of polyethylene glycol (PEG) to RIT has been found to be useful in “masking” the immunogenic epitopes in the protein. PEGylation also prolongs the circulation of RIT by reducing renal clearance.

3) Domain II of PE appeared to be the most immunogenic portion of the PE molecule. Mazor et al found that domain II of PE was sensitive to protease digestion and that almost all of domain II except the furin cleavage site (amino acids 274–284) could be removed without loss of activity. Liu et al identified the human B-cell epitopes of PE toxin by M13 phage display. Then they constructed a variant RIT with point mutations of the residues that make up the B-cell epitopes. The variant RIT, which with a deletion of domain II and 7-point mutations that modified human B-cell epitopes, had significantly reduced reactivity with human antisera and retained cytotoxic and antitumor activity. Liu et al used this approach to develop a new RIT, RG7787. The cytotoxic activity of RG7787 was significantly improved, but the immunogenicity results are not clear yet.

5) Mazor et al used the similar approach to develop 2 variant RITs that have their T-cell epitopes removed or suppressed. The immunogenicity results suggested that removal of T-cell epitopes is more effective than the removal of B-cell epitopes. All these approaches were widely used in the production of new RITs, especially the deletion of domain II of PE toxin and modification of human B- or T-cell epitopes (Figure 3). The smaller molecular weight, the lower immunogenicity of RITs. So we constructed an RIT with small molecules that consisted of a PE38 toxin and 17 amino acids of amidated gastrin. This RIT, named

![Diagram](image_url)

**Figure 3 Development of the PE-based immunotoxins to increase the toxicity and to decrease the immunogenicity.**

**Notes:** (A) RITs targeting CD22. (B) RITs targeting MSLN. BL22, also named RB4(dsFv)-PE38 and CAT-3888, contained a single-chain Fv of the anti-CD22 antibody fused to truncated PE toxin. HA22 was mutated from BL22 by replacing the residues at positions 100, 100a, and 100b of V, represented here by horizontal red bars. In HA22-LR, the deletion mutant, most of domain II of PE toxin was deleted. HA22-LR-8M, a mutant of HA22-LR, was reported to contain 8 mutations, D406A, R432G, R467A, R490A, R513A, E548A, K590S, and Q592A. SS1P consists of the Fv fragment from anti-MSLN monoclonal antibody coupled to the same PE fragment with BL22. SS1-LR-GGS was developed by deleting the domain II of PE toxin with GGS linker between the antibody and the domain III. RG7787 used a humanized Fab fragment of anti-MSLN antibody and deleted PE domain II. In RG7787, there were 7-point mutations in domain III of PE toxin at B-cell epitopes to eliminate binding to B-cell receptor. LMB-T20 consisted of the Fv fragment coupled to the similar PE fragment with RG7787, but with 6-point mutations in domain III at T-cell epitopes.

**Abbreviations:** PE, Pseudomonas exotoxin A; RIT, recombinant immunotoxins; MSLN, mesothelin.
rG17PE38KDEL, targets the overexpression of cholecystokinin II receptor (CCK2R) on gastric cancer cells, and it has lower immunogenicity in xenograft model, which means it can be used in further development and application.

Recent advances in the design and administration of RITs are overcoming partial challenges. Serial modifications have been used to reduce nonspecific toxicities, to increase stability and to improve targeted cellular killing. To overcome another major challenge, immunogenicity, several approaches were developed in the past years. Onda et al.\textsuperscript{196} developed the less immunogenic PE by engineering variant of the toxin. Based on this strategy, a bispecific ligand-directed toxin EGF4KDEL-7mut was developed, in which human EGF and IL-4 are linked to low immunogenic variant of PE38. It showed the dual benefit by increasing targeting specificity and reducing immunogenicity. Fused micromolecule or ligand also allowed long-term treatment of tumors. An RIT aiming at the treatment of gastric cancer, rG17PE38, was developed by our laboratory, in which amidated gastrin 17 (rG17) was linked to truncated modificatory PE38. It can suppress the growth of tumor and prolong the survival time in murine xenograft models, and the tumor-bearing mouse did not develop the neutralizing antibody against rG17PE38 after continuous infusion.

Several RITs have shown remarkable success against hematologic malignancies. It had been reported that RITs targeting MSLN produced major tumor regression in some patients with advanced mesothelioma.\textsuperscript{179} Although there is a further step to mitigate nonspecific toxicities and to enhance the activity of the toxin, we still can anticipate exciting successes in the future application of RITs based on appropriate combinations of cancers and selective target.

**Conclusion**

The success rate of immunotoxin therapy in clinical trials of leukemia has attracted more attention once again and they could be ideal molecules for combination therapy.

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The authors report no conflicts of interest in this work.

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