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# IL-15 as a potential target in leukemia

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**Abstract:** Leukemia, one of the most aggressive hematopoietic malignancies, is characterized by excessive proliferation, survival, and impaired differentiation of hematopoietic stem cells. Interleukin 15, a proinflammatory cytokine, induces proliferation and promotes cell survival of human T and B lymphocytes, as well as natural killer cells. However, it may also play a detrimental role in the onset of leukemia. This review provided an overview of the aberrant expression of Interleukin 15 and its role in the development and progression of this hematological malignancy. Also, we critically explored the potential therapeutic opportunities involved in targeting the disruption of interleukin-15 signaling as well as in interleukin-15-mediated enhancement of antitumor immunity.

**Keywords:** leukemia, interleukin-15, immunotherapy

## Leukemia

Leukemia is a complex and heterogeneous group of diseases and is characterized as the abnormal proliferation of immature cells of the hematopoietic system. The most widely accepted classification system is the World Health Organization system, which encompasses etiologic, morphologic, immunophenotypic, clinical, genetic, and molecular features and incorporates scientific and clinical information to refine diagnostic criteria. Among the most common forms of leukemia are two acute variants, acute lymphoblastic leukemia (ALL) and acute myeloid leukemia (AML), and two chronic variants, chronic myeloid leukemia (CML) and chronic lymphoblastic leukemia (CLL).<sup>1</sup>

## Current treatment options and patient outcomes

AML is characterized by the inhibition of differentiation and the subsequent accumulation of hematopoietic stem cells at various stages of incomplete maturation, and by reduced production of healthy hematopoietic elements.<sup>2</sup> AML is a heterogeneous disease that can be classified in as many as seven subtypes.<sup>3</sup> These subtypes are characterized by a variety of cytogenetic and cell surface markers. The standard treatment for AML is induction chemotherapy with an anthracycline/cytarabine combination, followed by either consolidation chemotherapy (commonly high-dose cytarabine) or allogeneic stem cell transplantation, depending on the patient's ability to tolerate intensive treatment and the likelihood of cure with chemotherapy alone.<sup>4</sup> The goal of induction chemotherapy is to achieve complete remission (CR). Once this is achieved, consolidation is given to maintain the remission.<sup>5</sup> The duration of initial CR,<sup>6</sup> and cytogenetics and molecular studies in AML,<sup>7</sup> together with antecedent of myelodysplastic

syndrome, multidrug resistance, or other medical comorbidities<sup>8</sup> provide important prognostic information.<sup>7</sup>

ALL, the most common form of cancer in children, is a clonal stem cell malignancy of excessive B- or T-lymphoblast proliferation.<sup>9</sup> The genetics of ALL are quite complex and are comprised of a variety of chromosome fusions, which can be used to distinguish different subtypes of the disease. Treatment for ALL is categorized broadly as induction phase, consolidation therapy, maintenance therapy, central nervous system (CNS) prophylaxis, and allogeneic stem cell transplantation. The goal of induction therapy is to achieve CR. Standard therapy will achieve CR in 80%–90% of patients.<sup>5</sup>

CML is characterized by a balanced genetic translocation, t(9;22)(q34;q11.2), involving a fusion of the *ABL* oncogene from chromosome 9q34 with the *BCR* gene on chromosome 22q11.2. This rearrangement is known as the Philadelphia chromosome. This translocation results in expression of the constitutively active protein tyrosine kinase BCR-ABL.<sup>10</sup> The current standard of care for CML is tyrosine kinase inhibitor (TKI) therapy, a very specific inhibitor of the BCR-ABL fusion protein. Imatinib, a TKI, has revolutionized the treatment of CML by transforming it from an invariably fatal disease to a chronic but manageable condition. Treatment with imatinib results in a 5-year progression-free survival rate of approximately 89%.<sup>11</sup> Resistance to imatinib occurs in certain cases, usually through mutations in the imatinib binding site on BCR-ABL, hence, second generation TKIs (dasatinib and nilotinib) were introduced to the pharmacological armamentarium, successfully rescuing about 50% of imatinib-resistant patients.<sup>12</sup>

CLL, the most common type of adult leukemia,<sup>13</sup> is characterized by the accumulation of mature CD5<sup>+</sup> B cells in the peripheral blood and lymphoid organs. As in the leukemias described above, chromosomal aberrations and gene mutations are common in CLL. Over the past decades, there has been a transition from single-agent alkylator-based therapies to nucleoside analogues, combinations of both alkylators and nucleoside analogues, and most recently, chemoimmunotherapy. CR rates have improved from 7% to a maximum of 70%.<sup>14,15</sup>

## IL-15

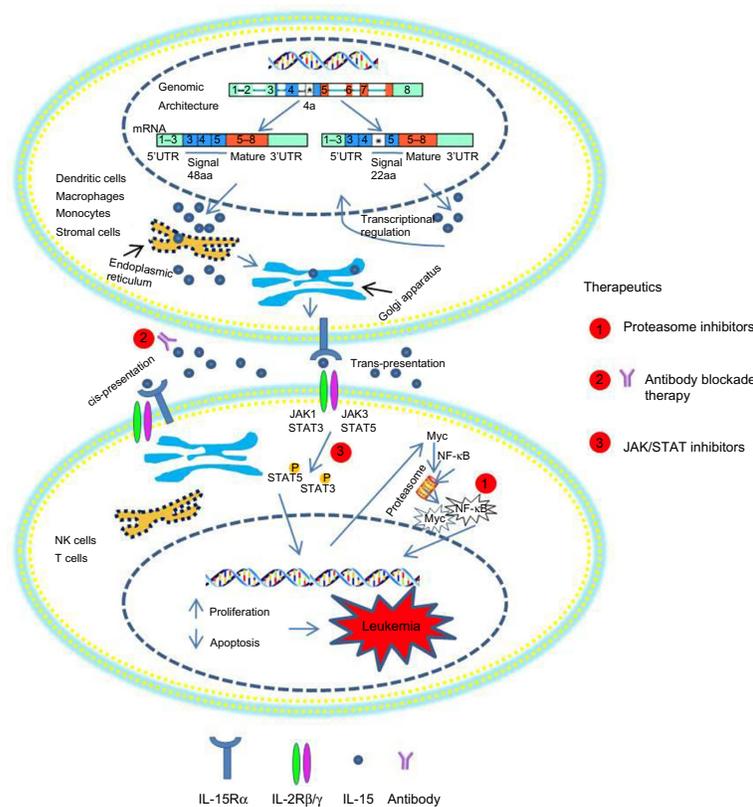
IL-15 was codiscovered in 1994 by two independent groups while studying the human T cell lymphotropic virus-1 (HTLV-1)-infected T-cell line, HUT102, in the absence of IL-2<sup>16</sup> and its ability to stimulate proliferation of the IL-2-dependent CTLL-2 T cell line in the presence of neutralizing anti-IL-2 antibodies.<sup>17</sup> IL-15 is a member of the four

$\alpha$ -helix bundle and shares receptor signaling components (IL-2/15 receptor [R] $\beta\gamma$ ) with IL-2; as a result, the two cytokines have similar biologic properties in vitro. However, specificity for IL-15 versus IL-2 is provided by unique private  $\alpha$ -chain receptors that complete the IL-15R $\alpha\beta\gamma$  and IL-2R $\alpha\beta\gamma$  heterotrimeric high-affinity receptor complexes.<sup>18</sup>

IL-15 is a 14–15 kDa glycoprotein encoded by a 34 kb region on human chromosome 4q31 and the central region of mouse chromosome 8.<sup>19</sup> There are two isoforms of IL-15 messenger(m)RNA that differ in their signal peptides lengths, containing either a short signal peptide or long signal peptide (Figure 1). Both isoforms produce the same mature protein; however, they each have distinct intracellular trafficking, localization, and secretion patterns. The IL-15 long signal peptide is located to the Golgi apparatus, early endosomes, and the endoplasmic reticulum and is often secreted from the cell as a soluble protein. The IL-15 short signal peptide is restricted to the cytoplasm and nucleus and may play a role in its transcriptional regulation.<sup>20</sup>

Although IL-15 mRNA can be found in many tissues and cells, including fibroblasts, muscle cells, keratinocytes, kidney cells, lymphocytes, mast cells, and tumor cells,<sup>21</sup> it is produced as a mature protein mainly by dendritic cells (DCs), monocytes, macrophages, and stromal cells.<sup>22</sup> Translationally inactive IL-15 mRNA is stored in the cell, ready to be rapidly translated upon specific signals.<sup>22</sup> The prevailing mechanism of IL-15 action seems to be trans-presentation (juxtacrine signaling) (Figure 1), although it also includes intracrine, autocrine, paracrine, and endocrine signaling.<sup>20</sup> IL-15 is also capable of efficiently signaling in cis through IL-15 R $\alpha$  and IL2R $\beta/\gamma$ c expressed on the surface of a single cell.

Once IL-15 is secreted out of the cell, it binds to IL-15R $\alpha$  and is presented to IL-2R $\beta\gamma$  complexes expressed on nearby effector cells, leading to a series of signaling events. These include activation of the JAK/STAT proteins for cellular activation.<sup>24</sup> Binding of IL-15 to the IL-2/15R $\beta\gamma$  heterodimer induces JAK1 activation that subsequently phosphorylates STAT3 via the  $\beta$ -chain and JAK3/STAT5 activation via its  $\gamma$ -chain (Figure 1).<sup>24</sup> Additional signaling pathways involve the recruitment of Shc to a phosphorylated site on the IL-2/15R $\beta$  chain, followed by activation of Grb2. From there, Grb2 can proceed down the PI3K-Akt signaling pathway or can bind the guanine nucleotide exchange factor SOS to activate the RAS-RAF-MAPK signaling pathway.<sup>25,26</sup> Collectively, these signaling mechanisms induce expression and activation of downstream effector molecules, such as proto-oncogenes *c-Myc*, *c-Fos*, and *c-Jun*, and antiapoptotic protein Bcl-2 for cellular proliferation.<sup>27</sup>



**Figure 1** IL-15 expression and its potential for targeting in leukemia. The IL-15 gene is made up of nine exons and eight introns, of which four exons code for the mature protein. Long signal peptide is located to the Golgi apparatus, early endosomes, and the endoplasmic reticulum and is often secreted from the cell as a soluble protein, whereas short signal peptide is restricted to the cytoplasm and nucleus. IL-15R $\alpha$  can transpresent IL-15 to IL-2R $\beta$ / $\gamma$ c<sup>+</sup> NK cells or T-cells through cell contact. The possibility of cis-presentation by a IL-15R $\alpha$ /IL-15 membrane complex to an IL-2R $\beta$ / $\gamma$ c heterodimer on the same cell is shown. NK cell or T-cell activation can induce JAK/STAT activation. The phosphorylated STAT proteins form then a heterodimer and traffic to the nucleus, activating transcription factors which contribute to cell survival. Therapeutic interventions for targeting IL-15 include: proteasomal inhibitors, anti-IL-15 receptor antibody, and pharmacologic inhibitors.

**Abbreviations:** IL, interleukin; mRNA, messenger RNA.

IL-15 plays an important role in the development, homeostasis, and function of memory CD8<sup>+</sup> T cells, natural killer (NK) cells, NKT cells, and intestinal intraepithelial T cells.<sup>28,29</sup> This cytokine induces proliferation of naïve CD8<sup>+</sup> and memory CD4<sup>+</sup> and CD8<sup>+</sup> T cells, promoting development of the primary response of effector CD8<sup>+</sup> T cells,<sup>30</sup> and makes both CD4<sup>+</sup> and CD8<sup>+</sup> T cells resistant to the action of regulatory T cells.<sup>31</sup> IL-15, which is itself produced by DCs, participates in their activation, proliferation, and differentiation,<sup>32</sup> promoting expression of major histocompatibility complex (MHC) class II, CD40, and CD80/86 molecules,<sup>33</sup> and making them potent inducers of CD4<sup>+</sup> and CD8<sup>+</sup> T cell responses.<sup>34</sup> More importantly, it inhibits IL-2-induced activation-induced cell death.<sup>35</sup> The antiapoptotic effects of IL-15 in granulocytes and lymphocytes are caused by its regulatory actions on the expression of proteins of the Bcl-2 family and on the activity of NF- $\kappa$ B and caspases.<sup>36</sup> IL-15 induces proliferation and promotes cell survival of human T and B lymphocytes, and NK cells. However, it may also play a role in certain leukemias: human lymphotropic virus type-1 (HTLV-1)-associated adult T cell leukemia (ATL),

large granular lymphocyte (LGL) leukemia, ALL, AML, B-cell CLL, and CML (Figure 1).

## Role of IL-15 in leukemias HTLV-1-associated ATL

Both IL-15 and IL-15R $\alpha$  levels are elevated in ATL patients.<sup>37,38</sup> Using an antibody towards IL-2/IL-15R, Azimi et al<sup>37</sup> demonstrated a role of this cytokine in the proliferation of peripheral mononuclear cells of infected patients. In these patients, an elevation of IL-15 mRNA was also detected. This phenomenon was also confirmed in another study, where an increase of IL-15R $\alpha$ , IL-15 mRNA and protein level was detected in HTLV-1-associated diseases. Furthermore, researchers observed that the viral protein Tax was involved in the activation of IL-15R $\alpha$ . These findings suggest a role for IL-15R $\alpha$  in the aberrant T cell proliferation observed in HTLV-1-associated diseases. Other studies have observed that the mechanism of HTLV-1 oncogenesis relies on a constitutive activation of the JAK/STAT pathway, especially on an association between JAK1 and STAT3, two molecules that are downstream to the IL-2/IL-15 receptor.<sup>39</sup>

However, Chung et al observed that IL-13 mRNA was highly expressed in HTLV-1-infected, IL-2-dependent T-cell lines, but IL-4, IL-10, IL-2, or IL-15 mRNAs were either below detection limits or did not correlate with HTLV-1 infection.<sup>40</sup> These conflicting results may rely on the fact that they were performed on different biological samples. Further studies on the role of IL-15 in ATL patients are warranted to determine definitively whether IL-15 may participate in the initiation, maintenance, or progression of the disease.

## LGL leukemia

LGL leukemia can arise from a CD3<sup>+</sup> T-cell lineage or from a CD3<sup>-</sup> NK-cell lineage, which can be classified into two categories: NK LGL leukemia and chronic NK lymphocytosis.<sup>41</sup> In a study where they deciphered the role of IL-15 in LGL, Zambello et al demonstrated the presence of IL-15 on leukemic LGL blasts, suggested a role of IL-15 in the pathogenesis of this disease, and showed that IL-15 may induce proliferation and cytotoxicity of LGL through the  $\beta$  and  $\gamma$  chains shared with IL-2.<sup>42</sup> The study of Chen et al showed increased expression of IL-15R $\alpha$  in T-cell LGL (T-LGL) leukemia and suggested that higher IL-15R $\alpha$  expression may lower the IL-15 response threshold in vivo and therefore, may contribute to the pathogenesis of the disease.<sup>43</sup> Chronic IL-15-mediated activation via the JAK/STAT pathway, especially STAT3 and STAT5, can be leukemogenic.<sup>44</sup> Somatic mutations in the SH2 domain of *STAT3* have been discovered at the frequency of 40% in T-LGL leukemia and 30% in NK-LGL leukemia patients.<sup>45,46</sup> Mishra et al showed that excessive IL-15 initiates cancer through the induction of *Myc*, *Hdac-1*, and *NF- $\kappa$ Bp65*, which mediate downregulation of microRNA (miR)-29b and consequent overexpression of *Dnmt3b*, thereby hastening the onset of LGL leukemia.<sup>47</sup> Hodge et al demonstrated that IL-15 alters expression of Bcl-2 family members and control of Bid that potentially links this cytokine to leukemogenesis through targeted proteasome degradation of Bid and offers the possibility that proteasome inhibitors may aid in the treatment of LGL leukemia.<sup>36</sup> Yu et al demonstrated T-LGL leukemias in IL-15 transgenic mice that expressed NKp46; NKp46-expression was the hallmark of a minute population of wild-type NKT cells with higher activity and potency to become leukemic.<sup>48</sup> Taken together, all these studies showed a role of IL-15 receptor and its signaling pathway in the pathogenesis of LGL leukemia and suggest that these may be useful targets in the treatment of patients with T-LGL or NKT leukemia.

## AML

Very few studies have been published about the putative role of IL-15 in the development of the disease. Physiologically low amounts of IL-15 can be secreted by bone marrow stromal cells.<sup>49</sup> This suggests that in the pathological condition, IL-15 could play a juxtacrine role, together with other factors, in promoting the survival and proliferation of IL-2R $\beta$ / $\gamma$ <sup>+</sup> neoplastic clones in the bone marrow or lymph node microenvironment. This is likely to occur more frequently in IL-2R $\beta$ / $\gamma$ <sup>+</sup> lymphoproliferative disorders;<sup>50,51</sup> however, it seems that only a fraction of AML express this receptor.<sup>52</sup> Drexler et al showed that only a fraction (3/19) of a panel of AML cell lines examined displayed responsiveness to IL-15, although the possible involvement of IL-2R $\beta$ / $\gamma$  was not studied.<sup>53</sup> These results suggested that IL-15, produced by bone marrow stromal cells, could be an antiapoptotic growth factor for a particular subset of AML cells equipped with a functional IL-2R $\beta$ / $\gamma$  complex.<sup>54</sup>

## ALL

A high expression of IL-15 correlates with CNS disease in childhood ALL<sup>55,56</sup> and neurocognitive impairment.<sup>57</sup> Moreover, several *IL-15* single nucleotide polymorphisms (SNPs) have been associated with minimal residual disease;<sup>58</sup> five SNPs (rs10519612, rs10519613, rs35964658, rs17007695, and rs17015014) located in the *IL-15* locus were associated with childhood ALL treatment response.<sup>59</sup> In adult ALL, the expression of *IL-15* has also been correlated with the immunophenotypes of ALL, therefore Lin et al genotyped the five SNPs of the *IL-15* gene by polymerase chain reaction – restriction fragment length polymorphism assays and observed a excess risk of developing ALL for the rs10519612 CC and rs17007695 TC genotype carriers in the adult Chinese population.<sup>60</sup> Aly et al also observed a higher risk of developing T-cell ALL for rs10519612 CC and an increased risk of developing B-cell ALL for rs17007695 TC and rs17007695 CC genotype carriers in Egypt.<sup>61</sup> These results suggested that several IL-15 SNPs, which were associated with minimal residual disease and correlated with the immunophenotypes of ALL, may provide important prognostic information in ALL patients.

## CML

Autologous transplantation of stem cells could represent an alternative treatment to allogeneic bone marrow transplantation for CML patients, in whom NK cells may constitute the main cytolytic immune cells effect of mediating the graft-versus-leukemia effect.<sup>62</sup> Therefore, Carayol et al investigated

NK cell differentiation of CD34<sup>+</sup> progenitors from CML patients and found that CD34<sup>+</sup> cells from CML patients did not proliferate nor differentiate in the presence of stem cell factor and IL-15. Furthermore, anti-IL-15 monoclonal antibody (mAb) failed to induce the downregulation of the  $\gamma$ c chain or the disruption of its interaction with JAK3 in leukemic cells. Therefore, in leukemic cells, the constitutive expression of the  $\gamma$ c chain, as well as the continuous production of IL-15, would explain the sustained proliferation of leukemic precursors in vivo and their survival. As all the patients included in this study were positive for *BCR/ABL*, the alteration of the IL-15 differentiation pathway could be an indirect consequence of the *BCR/ABL* protein expression and its subsequent enhanced tyrosine kinase activity.<sup>63</sup> In this respect, *BCR/ABL* expression through NF- $\kappa$ B activation<sup>64</sup> may be involved in IL-15 transcription,<sup>65</sup> thus the IL-15 signaling pathway may play a role in the development and progression of CML.

## CLL

Purified CLL cells do not proliferate and tend to undergo spontaneous apoptosis in vitro, suggesting that factors inducing CLL cell survival and proliferation might be present in the microenvironment of lymphoid organs or of the bone marrow.<sup>66</sup> Several cytokines, such as the TNF family members BAFF and APRIL,<sup>67</sup> the chemokine SDF-1,<sup>68</sup> IL-2,<sup>69</sup> and IL-15,<sup>70</sup> stimulate CLL cell proliferation and/or survival in vitro. Bernasconi et al demonstrated that IL-15 and CpG oligonucleotides promote memory B-cell activation and proliferation,<sup>71</sup> and Park et al suggested that IL-15 plays an important role in supporting germinal center B cell proliferation, proposing a new target for treatment of B cell tumors.<sup>72</sup> Importantly, IL-15 has also been described as a growth factor in non-germinal center-derived CLL mature B-cell survival and proliferation, through a strong activation of both STAT5 and ERK1/2 pathways.<sup>73</sup> Interestingly, the mitogenic and antiapoptotic effects of IL-15 were enhanced when CLL B cells were preactivated by CD40. Epron et al found that in the presence of CD40L signaling, IL-15 essentially increased B-cell proliferation.<sup>74</sup> These studies demonstrated that the IL-15 signaling pathway plays a role in CLL cell survival and proliferation and that the cooperation between IL-15 and CD40L increased CLL B cell growth.

## Critical analysis of the potential for targeting IL-15 in cancer

Therapeutic interventions of targeting IL-15 have been found to include: proteasomal inhibitors,<sup>45</sup> anti-IL-15R antibodies and pharmacologic inhibitors (Figure 1).<sup>75</sup>

IL-15 can lower the proapoptotic Bid protein in LGL leukemia, which was shown to result from increased proteasomal degradation, whereas the induction of Bid by the proteasome inhibitor bortezomib increased leukemic cell death, suggesting that proteasome inhibitors could be an effective treatment option for this disease.<sup>36</sup> Furthermore, a formulation of a proteasomal inhibitors provides long-term disease-free survival in leukemic mice,<sup>49</sup> thus offering a new approach to treating patients with aggressive LGL leukemia.

Another therapeutic approach targeting IL-15 in leukemias has been the use of the anti-IL-15R mAb Mik $\beta$ 1 to block the presentation of IL-15 to the IL-2/IL-15R $\beta$ , thereby inhibiting proliferation of IL-15-dependent leukemia cells (Table 1).<sup>76</sup> A Phase I clinical trial of IL-15 blockade in T-cell LGL leukemia was conducted by Waldmann et al using the Hu-Mik $\beta$ 1 mAb that blocks IL-15 transpresentation to cells expressing IL-2/IL-15R $\beta$  and  $\gamma$ c (Table 1).<sup>76</sup> However, Hu-Mik $\beta$ 1 did not block IL-15 action on the cells that express the heterotrimeric receptor in a cis orientation. In this trial, Hu-Mik $\beta$ 1 therapy was not effective in the treatment of patients with monoclonal T-LGL.

Finally, Epling-Burnette et al demonstrated that STAT3 activation contributed to accumulation of the leukemic LGL clones, as a variety of STAT3 inhibitors come forth to the clinic, suggested that investigation could focus on STAT3 inhibitors in the treatment of LGL leukemia.<sup>77</sup>

However, IL-15 can activate many immune antitumor mechanisms and can be considered as a good candidate for application in tumor therapy. The most important cells engaged in IL-15 antitumor activity seem to be T-cells, in particular CD8<sup>+</sup> T cytotoxic cells and NK cells.<sup>22</sup>

IL-15 increases the number of specific CD8 T cells in adoptive T-cell therapy,<sup>78</sup> and in vitro assays have shown that treatment of CML monocytes with IFN- $\alpha$  and GM-CSF resulted in the rapid generation of activated DCs expressing IL-15. An additional activation/maturation step of these DCs can render these cells fully capable of presenting CML-specific antigens inducing tumor-specific CD8<sup>+</sup> effector cells, which represents an important requisite for achieving an immune-mediated control of tumor progression in CML patients.<sup>79–81</sup>

It had been proved that IL-15 increases NK cell cytotoxicity in an NKG2D-dependent fashion,<sup>82</sup> which could be dependent on the presence of the NKG2D ligand ULBP1 on the tumor cell,<sup>83</sup> and that IL-15-expanded NK cells enhanced their expression of KIR and NKG2D.<sup>84</sup> Furthermore, IL-15 may be used in NK cell expansion protocols, in combination with SCF, FLT3-L, and IL-21, to generate NK cells from CD34<sup>+</sup> cells. This protocol was developed by Yoon et al to expanded NK cells into AML or ALL patients shortly after

**Table 1** Clinical trials of targeting IL-15 in cancer

Clinical trial	Trial type	Agents	Disease	Effect of IL-15
Phase I trial of IL-15 trans presentation blockade using humanized Mik $\beta$ -I mAb in patients with T-cell LGL	Cancer immunotherapy	Anti-IL15 antibody Mik $\beta$ -I mAb	LGL	Harmful
Recombinant Interleukin-15 in Treating Patients With Advanced Melanoma, Kidney Cancer, Non-Small Cell Lung Cancer, or Squamous Cell Head and Neck Cancer NCT01727076	Cancer immunotherapy	IL-15	Melanoma, kidney cancer, non-small cell-lung cancer, or squamous cell head and neck cancer	Beneficial
Use of IL-15 After Chemotherapy and Lymphocyte Transfer in Metastatic Melanoma NCT01369888	Cancer immunotherapy (adjuvant)	Tumor-infiltrating lymphocytes, IL-15	Melanoma	Beneficial
A Phase I Study of Intravenous Recombinant Human IL-15 in Adults With Refractory Metastatic Malignant Melanoma and Metastatic Renal Cell Cancer NCT01021059	Cancer immunotherapy	IL-15	Melanoma or renal cell cancer	Beneficial
Continuous Infusion of rhIL-15 for Adults With Advanced Cancer NCT01572493	Cancer immunotherapy	IL-15	Carcinoma or lymphoma	Beneficial
Haploidentical Stem Cell Transplantation and IL-15 NK Cell Infusion for Pediatric Refractory Solid Tumors NCT01337544	Cancer immunotherapy	HSCT, NK cells grown in culture with IL-15 (ancillary agent)	Pediatric solid tumors	Beneficial
IL-15 Super Agonist ALT-803 to Treat Relapse Of Hematologic Malignancy After Allogeneic SCT NCT01885897	Cancer immunotherapy (after SCT)	IL-15 super agonist	Relapse of hematologic malignancy	Beneficial
NK White Blood Cells and Interleukin in Children and Young Adults With Advanced Solid Tumors NCT01875601	Cancer immunotherapy	NK cells, IL-15	Solid tumors, brain tumors, sarcoma, Wilm's tumor, or rhabdomyosarcoma	Beneficial
Haploidentical Donor Natural Killer Cell Infusion With IL-15 in Acute Myelogenous Leukemia (AML) NCT01385423	Cancer immunotherapy	NK cells, IL-15	Acute myelogenous leukemia, myelodysplastic syndrome	Beneficial
A Phase I Study of the Clinical and Immunologic Effects of ALT-803 in Patients With Advanced Melanoma NCT1946789	Cancer immunotherapy	IL-15 super agonist	Melanoma	Beneficial
IL15 Dendritic Cell Vaccine for Patients With Resected Stage III (A, B or C) or Stage IV Melanoma NCT1189383	Cancer vaccine	Dendritic cells grown in culture with IL-15 (ancillary agent)	Melanoma	Beneficial
A Study of Intravesical Bacillus Calmette-Guerin (BCG) in Combination With ALT-803 in Patients With BCG-naive Non-Muscle Invasive Bladder Cancer NCT02138734	Cancer immunotherapy	IL-15 super agonist	Bladder cancer	Beneficial
A Study of ALT-803 in Patients With Relapsed or Refractory Multiple Myeloma NCT02099539	Cancer immunotherapy	IL-15 super agonist	Multiple myeloma	Beneficial
Natural Killer Cells in Acute Leukemia and Myelodysplastic Syndrome NCT02123836	Cancer immunotherapy	NK cells expanded with K562-mb15-41BBL (ancillary agent)	Acute leukemia or myelodysplastic syndrome	Beneficial

**Abbreviations:** HSCT, haploidentical stem cell transplantation; K562-mb15-41BBL, irradiated K562 cell line modified to express membrane-bound IL-15 and 41BB ligand; LGL, large granular lymphocyte leukemia; mAb, monoclonal antibody; NCT, National Clinical Trial; SCT, stem cell transplantation.

haploidentical stem cell transplantation. This feasibility study showed no toxicity of the NK cell infusion.<sup>85</sup>

There are additional ways in which NK cells are important for IL-15 antitumoral therapy. For instance, in vitro assays

have indicated that NK cells in the presence of IL-15 become more efficacious mediators of ADCC against cultures of B-lymphoma cells in the presence of the anti-CD20 mAb rituximab.<sup>86</sup>

The encouraging data of IL-15 immunotherapy in murine models, together with its low toxicity in mice and primates, has led to the design of clinical trials in cancer patients. The first-in-human clinical trial of recombinant human IL-15 (rhIL-15) in patients with cancer, by Conlon et al, demonstrated that IL-15 could be safely administered to patients with metastatic malignancy. IL-15 administration markedly altered homeostasis of NK cells and  $\gamma\delta$  cells in the blood.<sup>87</sup> There were also other clinical trials initiated recently, as shown in Table 1. Six of the trials use the rhIL-15 protein, administered either alone or combined with the administration of patient-derived tumor-infiltrating lymphocyte or combined with NK cells. Three trials used IL-15 as an ex vivo ancillary agent to enhance DC or NK cell expansion and function when used for anticancer vaccination or immunotherapy. The remaining four trials aimed to determine the safety and effectiveness of an IL-15 super agonist complex (ALT-803) administered weekly for 4 or 6 consecutive weeks, alone or mixed together with Calmette-Guerin bacillus.

## Conclusion

The recent published data have highlighted the role of IL-15 and the aberrancies in its regulation in the development and progression of leukemias. The better understanding of the IL-15/IL-15R system has led to the identification of new therapeutic targets, which will open the possibility for the development of new drugs. However, IL-15's ability to stimulate the development and activity of effector and memory CD8<sup>+</sup> T cells, and NK and NKT cells, which all contribute significantly to the antitumor immune response, raises the hope that this cytokine could become a valuable adjuvant in antitumor immunotherapy.

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## Disclosure

The authors report no conflicts of interest in this work.

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