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# The Effects of Stem Cell Therapy in the Treatment of Childhood Acute Lymphoblastic Leukemia

Emma Lawrence and Vincent S Gallicchio\*

Department of Biological Sciences, College of Science, Clemson University, Clemson, USA

\***Corresponding author**: Vincent S Gallicchio, Department of Biological Sciences, College of Science, Clemson University, Clemson, USA

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#### Abstract

Acute Lymphoblastic Leukemia (ALL) is a cancerous transformation in the blood and bone marrow beginning with progenitor cells. With developing research into treatment options, there is now a 90% survival rate. However, high relapse rates have become the next challenge to overcome. Relapse rates increase with a variety of markers including age of diagnosis, white blood cell counts, and response to chemotherapy. Chemotherapy and radiation have been the standard of care in the past, but are typically accompanied by secondary malignancies later in life. Researchers believe that stem cell therapies may offer alternative options to high-risk patients.

#### Keywords

Stem cells; Therapy; Childhood ALL

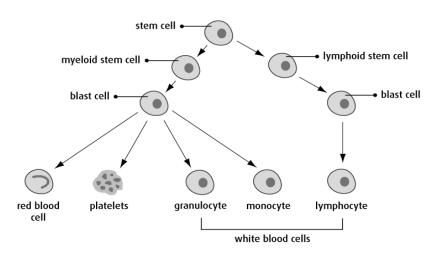
#### **Abbreviations**

**Review Article** | Lawrence E, et al. J Stem Cell Res 2020, 1(2)-8. DOI: <u>https://doi.org/10.52793/JSCR.2020.1(2)-08</u> ALL: Acute Lymphoblastic Leukemia; RBCs: Red blood cells; CBC: Complete Blood Count; HR: High Risk;
VHR: Very High Risk; WBC: White Blood Cell; EFS: Event Free Survival; CAR: Chimeric Antigen Receptor;
MUD: Matched Unrelated Donor; MRD: Matched Related Donor; GvHD: Graft vs. Host Disease; TBI:
Total Body Irradiation; ASC: Adult Stem Cells; ESC: Embryonic Stem Cells; iPSC: Induced Pluripotent Stem
Cells; HSCT: Hematopoietic Stem Cell Transplant; MLL: Mixed-lineage-Leukemia; NK: Natural Killer;
NRM: Non-remission mortality; VHRR: Very High Risk of Relapse

### Introduction

Acute Lymphoblastic Leukemia (ALL) is a rare type of leukemia in adults, but the most common type found in children, peaking between the ages of two and five [1]. It is a rapidly progressing cancer that occurs when errors in DNA cause bone marrow cells to continue to grow and divide when they would normally stop. These abnormal cells are unable to mature into proper white blood cells and eventually crowd out healthy cells. Risk factors of developing ALL include previous cancer treatments, radiation exposure, genetic disorders, and a family history [2].

In its maturation, a blood stem cell will become a myeloid stem cell or a lymphoid stem cell. Myeloid stem cells can then develop into a red blood cell (to carry oxygen), white blood cell (to fight infection), or platelet (to form blood clots). A lymphoid stem cell may become a B lymphocyte (to make antibodies), T lymphocyte (to help make B lymphocytes), or a Natural Killer cell (to attack cancer cells and viruses). ALL is confirmed when too many stem cells becoming lymphoblasts, B lymphocytes, and T lymphocytes. This change is unfavorable because these cells cannot fight infection well. Genetic risk factors include Down syndrome, Neurofibromatosis type 1, Bloom syndrome, Fanconi anemia, Li-Fraumeni syndrome, Ataxia-telangiectasia, and errors in the mismatch repair system [3] (Figure 1).



#### **Development of Blood Cells**

**Figure 1:** Shows the normal development of a stem cell into RBC's, platelets, granulocytes, monocytes, and lymphocytes. In ALL, lymphoid stem cells divide abnormally leading to cancer [4].

Signs of ALL include fever, easy bruising, petechiae, bone and joint pain, lumps in the neck, underarm, stomach and groin, weakness, and loss of appetite. A diagnosis can be confirmed by a complete blood

**Review Article** | Lawrence E, et al. J Stem Cell Res 2020, 1(2)-8. DOI: <u>https://doi.org/10.52793/JSCR.2020.1(2)-08</u> count (CBC), blood chemistry study, bone marrow biopsy, or cytogenic analysis (analysis of chromosome rearrangements). Prognosis factors include age at diagnosis, whether leukemia cells came from B or T lymphocytes, chromosome rearrangements, and how quickly leukemia cell counts drop in the first month. There are three risk groups for childhood ALL. The standard risk group includes children age 1-10 who have a WBC< 50,000 uL. The high-risk (HR) group includes children age 10 and older or a WBC> 50,000 u. The very high-risk (VHR) group includes children less than 1 year old, those with certain gene abnormalities, children who have a slow response to treatment, and those who continue to show signs of leukemia after 4 weeks of chemotherapy [3].

Treatment options vary based on diagnosis factors, but include chemotherapy, radiation therapy, stem cell therapy, and target therapy [3]. Chemotherapy can be given systemically or regionally, depending on the categorized risk group. High-risk children receive higher doses and more aggressive treatments than standard risk patients. Radiation therapy may be administered externally or internally. External radiation is used to treat ALL that has spread, or may spread, to various areas of the body, most notably the brain and spinal cord. It may also be used in preparation for a stem cell transplant. Stem cell transplants may be used to replace healthy blood-forming cells after treatments of chemotherapy and total-body irradiation. Stem cells are typically used only for cases in which there is a high risk of relapse [5-7]. Additionally, many target therapies show promise in the treatment of childhood ALL including tyrosine kinase inhibitors, monoclonal antibody therapy, and proteasome inhibitor therapy [3].

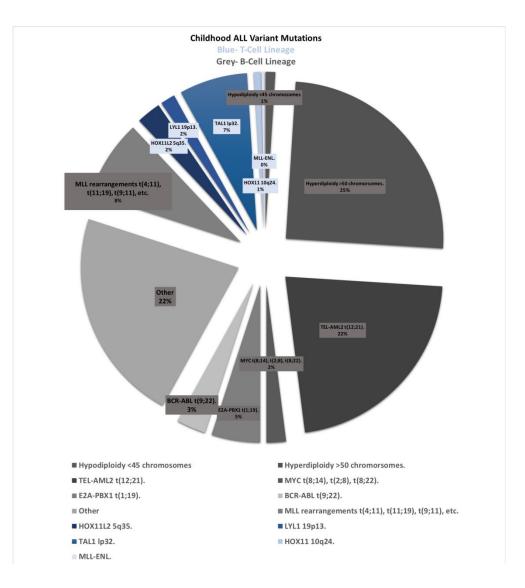
Approximately 3,000 children are diagnosed with ALL every year but due to advancements in treatment options, childhood ALL now has a 5-year survival rate of 90% (compared to 60% for adults) [8,9]. While recovery rates are high, relapse still occurs in 15-20% of children and remains a significant issue of concern. Patients of older age and/or higher WBC counts fare less well than younger patients and/or those with low WBC counts. Additionally, chromosome abnormalities account for heightened risks. A 20-fold increase occurrence of ALL has been associated with Down syndrome, with an additional increase risk of relapse [10]. Aneuploidy, Hyperploidy, and Hypoploidy of chromosomes are associated with higher risk ALL and poor outcomes. 75% of B-ALL cases involve somatic aneuploidy or recurrent chromosomal translocations and are most frequently found in younger children [8] (Table 1).

Variable	Favorable Variable	Adverse Factor	Use in Risk Stratification				
Demographic and clinical							
Demographic and clinical features							
Age	1 to <10 yr	<1 yr or ≥10 yr	This feature is a part of				
			NCI risk group definition				
Sex	Female	Male	No				
Race or ethnic group	White, Asian	Black, Native American,	No				
		Hispanic					
Initial white-cell count	Lower	Higher (≥50,000/mm³)	Part of NCI risk group				
	(<50,000/mm³)		definition				

Immunophenotype	B-cell lineage	T-cell lineage	Often used to select	
			therapy backbone	
Cytogenetic features	ETV6-RUNX1,	BCR-ABL1, MLL	Often used to select	
	hyperdiploidy,	rearrangements,	treatment intensity,	
	favorable	hypodiploidy	assign the patient HSCT,	
	chromosome		or both; some features	
	trisomies		(e.g., BCR-ABL1) can be	
			used to select targeted	
			therapy	
Genomic features	ERG deletions	IKZF1 deletions or	Some research groups	
		mutations; Philadelphia	use IKZF1 deletions to	
		chromosome- like ALL	assign patients to more	
		with kinase gene	intensive therapy; kinas	
		alterations	gene mutations may be	
			used to assign patients	
			to targeted therapy, but	
			this is not yet part of	
			routine care	
Early response to treatme	ent			
Response to 1 week of	Good response	Poor response to	Easy to measure and	
glucocorticoid therapy	to prendisone	prendisone (≥1000	used by many groups;	
	(<1000	blasts/mm <sup>3</sup> )	may be supplanted by	
	blasts/mm³)		MRD	
Marrow blasts after 1-2	M1 marrow	No M1 marrow (≥5%	Easy to measure and	
weeks of multiagent	(<5% blasts) by	blasts) by day 8 or 15	used previously by many	
therapy	day 8 or 15		groups; now being	
			supplanted by MRD	
MRD quantitation during	Reaching low	Persistence of MRD	Most important single	
or at end of induction	(<0.01%) or	≥0.01% at specific time	prognostic factor for	
	undetectable	points; the higher it is,	contemporary therapy;	
	MRD by specific	the worse the prognosis	critical for modern risk	
	time points		stratification	
MRD at 3-4 months	Low (<0.01%);	Persistence of MRD	May help select patients	
	preferably	≥0.01%	for HSCT or new	
	undetectable		therapies in first	
			remission	

**Table 1:** Shows important prognosis factors in Childhood Acute Lymphoblastic Leukemia. These factorsinclude age, sex, ethnicity, initial WBC, genomic factors, and responses to treatment. Determining thesefactors for a patient leads to a risk level analysis [11].

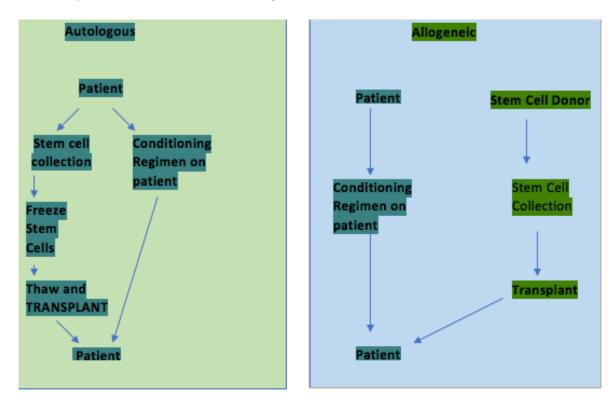
Two subtypes of childhood ALL are B-ALL and T-ALL. B-ALL makes up 85% of pediatric cases, while T-ALL accounts for 10-15% of pediatric cases and results in poor early treatment responses and high rates of relapse [12,13]. Current treatment of T-ALL involves the use of chemotherapy drugs cyclophosphamide, mercaptopurine, and cytarabine, which have shown promising outcomes [13]. However, chemotherapy can lead to drug toxicity and side-effects including glucocorticoid-related osteonecrosis, Asparaginaserelated pancreatitis, and vincristine-related peripheral neuropathy. Relapsed ALL is associated with a greater resistance to chemotherapy [14]. Anthracyclines are a type of antibiotic derived from Streptomyces bacteria that treat cancer by damaging the DNA in cancer cells. They have been shown to be effective against bone marrow relapse, but result in no significant increase in event free survival (EFS) for childhood ALL. Anthracycline can also result in cardiotoxicity and heart failure, which has led to treatment protocols of its use in selected high-risk cases [14-16]. Novel approaches to ALL treatment have led to CAR-T therapy, which involves the infusion of T-cells that have been genetically engineered to express the chimeric antigen receptor (CAR) in cells [17]. These T-cells tailor the body's immune response against specific pathogens. CAR-T therapy could be an option to replace failed treatment for patients with relapsed or refractory B- ALL [17,18]. Melphalan and cyclophosphamide with etoposide have also shown high rates of EFS when paired with hematopoietic stem cell transplants (HSCT) from an alternative donor [19] (Figure 2).

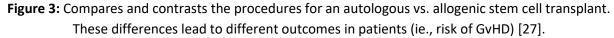


**Figure 2:** Shows the distribution and rates of various genetic alterations leading to childhood ALL. Additionally, it compares the rates between the two subgroups of ALL: B-Cell lineage and T-Cell lineage [20].

HSCT has offered a promising treatment option to induce antileukemic control in children with ALL who don't respond favorably to current chemotherapy approaches [21]. Two types of donor groups include matched unrelated donor (MUD) and matched related donor (MRD). Both MUD and MRD are classified as allogenic, meaning that they are derived from a matched, partially mismatched, or an unrelated donor. Reduced intensity allogenic stem cell treatment uses a lower dose of chemotherapy so the immune system isn't completely inactivated, which can lead to better outcomes for some older and sicker patients [22]. Allogenic transplants however, do lead to a heightened risk of graft-vs-host disease (GvHD), where the donors' immune cells attack the recipients normal tissue. GvHD presents with skin rashes, jaundice, severe diarrhea, fatigue, and muscle aches [22,23]. Immunosuppressant's like cyclosporine are administered following transplant to prevent GvHD, but can leave the body vulnerable to infection24,25. Steroids in combination with calcineurin inhibitor are currently used to treat GvHD.

Conversely, autologous transplants are when a patient receives their own stem cells (previously frozen), but is not a typical option due to its association with a high-relapse rate and the risk that the stem-cells are still contaminated with leukemia cells. In most cases, a matching related donor can't be found, so alternative approaches to allo-HSCT are used, including umbilical cord blood and unrelated/haploidentical allo-HSCT [24,26] (Figure 3).





Total body irradiation has been associated with the late onset of other complications, including cardiac, pulmonary, and renal complications; cataract; and a heightened risk of developing secondary malignancies. These secondary malignancies include organ dysfunction, growth retardation, as well as hormonal imbalances [28]. Cognitive impairments and developmental abnormalities have caused concerns for TBI in pediatric patients. Treatment without TBI has shown comparable results by using busulfan and cyclophosphamide in addition to HSCT, eliminating the adverse effects associated with TBI. This regimen has provided promising results in pediatric and infant ALL [19].

Stem cells are in a broad classification of which all other cells derive from. Once they divide into daughter cells, they can become new stem cells or differentiate into a more specific function (blood cells, brain cells, heart muscle, etc.). Different types of stem cells include adult stem cells (ASCs), embryonic stem cells (ESCs), cord blood/amniotic fluid stem cells, and induced pluripotent stem cells (iPSC's) [29].

Embryonic stem cells are derived from 3-5 day old embryos, and are pluripotent, while making up about 0.01-0.2% of total mononuclear bone marrow cells in humans. They are collected by the process of invitro fertilization, which occurs in a lab. ESC's are grown in test tubes or Petri dishes, typically to later be used during testing of new drugs for safety and effectiveness. These stem cells are defined by their origin in the inner cell mass of the blastocyst [29].

Adult stem cells are derived from a multitude of different tissues, and are more limited in their ability to differentiate into specific tissues. Like all stem cells, they have the ability to make identical copies of themselves in the process of long-term self-renewal. The undifferentiated cell is known as a progenitor cell, or precursor. To be classified as an ASC, the cell must be efficient at self-renewal for an organisms' entire lifespan. It also must be clonogenic, meaning it can give rise to a lineage of genetically identical cells [29].

Recent developments have been able to genetically reprogram adult stem cells to act as if it were an embryonic stem cell. These cells are called iPSC's, and can differentiate into all types of specialized cells to produce new cells, organs, or tissues. iPSC's are derived from a patient's skin or blood before being reprogrammed back into ESC's. These stem cells have offered an option for treatment without the ethical issues surrounding ESC's because the stem cells come from the patient. This option is also safer because it limits the possibility of rejection [30]. There is a preference for the use of embryonic stem cells over adult stem cells due to their versatility, but due to iPSCs, adult stem cells have shown new potential. In HSCT in the treatment of ALL, adult stem cells or umbilical cord blood are used to replace cells damaged by chemotherapy or disease [29].

A stem cell transplant is suggested in rare cases, including those in which ALL shows a likelihood of coming back (high risk), has come back (relapse ALL), or is no longer responding to standard treatment. In preparation for a HSCT, chemotherapy is first administered to kill leukemia cells, killing health cells in bone marrow in the process, making room for stem cells following transplant. Chemotherapy works to dampen the immune system, preventing the host from rejecting the donor stem cells following transplantation6. Long-term side effects of HSCT include the growth of another cancer, lung problems, lack of a menstrual period (damage to ovaries), and vision problems (damage to the lens of the eye) [23].

HSCT has been found to be a valuable course of action for high risk patients and a subgroup of infant mixed-lineage-leukemia (MLL+) ALL that carry poor prognosis factors [28,31]. These prognosis factors included infants being below the age of 6 months, prendiso(lo)ne response (poor predniso(lo)ne response [PPR]:  $\geq$  1.0 g/L blasts at day 8), and WBC counts in peripheral blood at diagnosis ( $\geq$  300 g/L vs < 300 g/L). This study found that 80% of infant cases harbor rearrangements in the MLL gene. Because of the high risks in infant ALL, HSCT survivors can face various other chronic conditions [28].

The information together regarding the MLL arrangement and ALL's peak between ages 2 and 5 supports the theory that additional exogenous exposures play a part in the development of ALL. Endogenous exposures influencing ALL development include inflammation and oxidative stress, while

exogenous exposures include an abnormal response to infection. The influenza virus is a promising candidate as an exogenous exposure. Other chromosomal abnormalities also increase the risk of ALL, including Trisomy 21 (down syndrome), hyperploidy, and translocations. All of these rearrangements are related to a dismal prognosis. Additionally, gene mutations of IKZF1, ARID5B, CEBPE, and CDKN2A have been associated with the occurrence of ALL due to their role as key regulators of blood development, proliferation, and differentiation. Racial and ethnic differences due to socioeconomic factors as well as genetic differences cause a disparity in the occurrence of ALL. Ancestry related genetic variations have led to a heightened occurrence of ALL in individuals with Hispanic ethnicity, followed by those of Asian Pacific and European Descent, and finally those of African ethnicity [31] (Table 2).

Representative Genetic Alterations or Subtypes						
Favorable	-Hyperdiploidy	Diagnosis	Therapy	Remission		
outcomes-	-ETV6-RUNX1		<b></b>			
standard therapy						
Improved	-TCF3-PBX1	Diagnosis	Therapy	Remission		
outcomes-	-BCR-ABL1		<b></b>			
intensified	-T-ALL					
therapy						
Chemoresistance	-Hypodiploidy		Therapy	Chemorefr		
and/or relapse-	-KMT2A	Diagnosis	$\longrightarrow$	actory ALL/		
due to inherent	rearrangement			relapse		
genetic	-iAMP21					
alterations	-Ph-like					
Chemoresistance	-CREBBP	Diagnosis	Therapy	Chemorefr		
leading to	mutations		$\longrightarrow$	actory ALL/		
relapse- due to	-NT5C2			relapse		
acquired genetic	mutations					
alterations	-ABL1					
	"gatekeeper"					
	mutation					

**Table 2:** Genetic alterations lead to variable outcomes based on responses to chemotherapy. The firsttwo boxes represent chemosensitive ALL, while the second two boxes represent chemoresistant ALL.Chemoresistant ALL may develop from drug-resistant subclones, leading to the development of acquiredmutations and later relapse [32].

In animal trials, mouse models have been used due to mice having at disease manifestation of ALL similar to that of humans. Xenograph models have been used on mice to analyze original patient samples for disease characterization and evaluation for potential therapeutics. Limitations to the xenograph model approach include the fact that the immunosuppressants represent a separate environment to that of human. Normal immune cells are usually infused in mice for pre-clinical

immunotherapy studies. Genetic murine models are another approach to ALL in animal studies. Many of these models manifest the symptoms of human ALL including leukemia cell accumulation in the thymus, lymph nodes, bone marrow, spleen, and infiltration in the liver and brain. In ras transgenic mice, ALL developed spontaneously, and half of the mice acquired a NOTCH1 gain of function mutation which led to accelerated T-ALL onset and decreased survival rate. NOTCH proteins are transmembrane receptors involved in cell development and in the transcriptional activation of target gees. This information suggests that NOTCH1 is a driving force in T-cell transformation in ALL. Animal models have shown to be an excellent way to study GvHD in transplants. Canine models have been essential in the testing of immunosuppressant therapies following transplantation. Methotraxate is an example, as it was first tested in dogs and was found to be effective in reducing the occurrence of GvHD33.

In human trials, CD16+ Natural Killer cells (NK) from cord blood stem cells have been used in cancer therapy by using an anti-CD47 antibody. This antibody triggers an antitumor response [34]. With ALL showing a higher level of CD47, the cord blood stem cells eradicated the ALL cells by using this antibody. Additional human trials of HSCT in patients found that transplants before a third or subsequent complete remission had better survival rates. Patients with transplants after a second complete remission had a 2.3 times greater risk of death [35].

HSCT from an allogenic donor is now the standard of care for the high risk subcategory, a qualification applying to only 10% of ALL patients [22]. The option of using HLA-matched unrelated donors (MUDs) has increased the availability to use stem cell transplants. In the Berlin-Frankfurt-Muenster study group, patients received HSCT from either a MUD or matched sibling donor (MSD). They found that engraftment was faster in the MSD-HSCT transplant group, but chronic GvHD also occurred more in this group and was more extensive in patients older than 12. This increase in occurrence could have been from the MUD group receiving more intensive immunosuppressant's. Graft failure was rare, but occurred more often in those receiving transplants from a MUD. Superior outcomes were attributed to the MSD group due to rapid immune reconstruction and fewer infections, but MUD had highly successful outcomes as well. MUD-HSCT transplants could be applied to patients without a suitable related donor [24].

HSCT results have improved through a decrease in nonremission mortality (NRM). Relapse is not uncommon, with rates or relapse reaching 20% following the first complete remission. These high relapse patients are eligible for MSD of MUD HSCT, along with those who are VHRR (very high relapse risk) [7]. These patients receive hyperfractioned TBI followed by immunosuppressant's, which can be ceased if there is no occurrence of GvHD after 100 days [25]. HSCT from both related and unrelated donors has been associated with non-remission mortality and relapse. For this reason, HSCT is limited to patients of high-risk. Relapse has been the most difficult issue to address in childhood ALL due to it commonly being associated with resistance to chemotherapy. Variations of immunomodulation and tailored chemotherapy are plausible ways to decrease the risk of relapse [7].

### Conclusion

Acute Lymphoblastic Leukemia is a cancer of the blood and bone marrow caused by the inability of progenitor cells to replicate properly. A high-risk factor indicative of ALL is the occurrence of chromosomal translocations and replications [36]. While ALL survival rates have reached 90%, relapse rates remain high [37]. Survival depends on a number of factors including the age of diagnosis, white blood cell count, and response to chemotherapy [36].

In attempts to treat patients following relapse, researchers have looked to stem cell use. Both MRD and MUD stem cell transplants have shown to be potential options. HSCT's correlate with a heightened risk of secondary malignancies, leading transplants to be a valuable option only to a specific subgroup of ALL who are at a heightened risk of relapse [28]. While studies have supported the use of HSC's in the treatment of ALL, there is more research to be done in the hopes of reducing both secondary malignancies and relapse.

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