



Review Article

Mixed Phenotypic Acute Leukemia

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Introduction

The majority of acute leukemias can be classified by the lineage of origin of blast cells as either myeloid, B Lymphoblastic (B-ALL) or T lymphoblastic. Acute leukemias are categorized by flow cytometry and now, increasingly, molecular results which help to determine whether the leukemia arises from the myeloid or lymphoid series. There does exist, however, an undefined or ambiguous group of leukemias in need of further characterization. Previous World Health Organization (WHO) criteria had defined a population of leukemic cells with no clear lineage restricting markers as Acute Undifferentiated Leukemias (AUL). Mixed Phenotypic Acute Leukemias (MPAL) represent a relatively new diagnostic designation, only being formally recognized and defined by the World Health Organization in 2008 [1]. Given the short time that these formal criteria have been established and the rarity of MPAL, there is a paucity of data regarding the precise biologic characterization of the disease as well as established guidelines on the management and prognosis of these patients. Herein, we review the defining characteristics, cytogenetic and clinical features and potential treatment options available for these patients.

Defining Mixed Phenotypic Leukemias

MPAL are quite rare, comprising roughly 3% of all new acute leukemia diagnoses. In order to establish the diagnosis, a specific set of pathologic criteria must be met to assign a designation of MPAL. Prior to 2008, mixed phenotypic leukemias were classified by a scoring system developed by the European Group for the Immunologic Classification of Leukemias (EGIL) (Table 1). The EGIL scoring system represented a refinement on a previous scoring system developed by Catovsky et al., in 1991 [2]. In this first scoring system, points were assigned to a limited number of Immune-Histochemical (IHC) markers from either a B cell, T cell or myeloid lineage. At the time, the Cavovsky scoring system weighted CD 22, CD3 and MPO

most heavily for defining their respective lineages. A score of greater than or equal to 2 was required to assign a lineage. If leukemia scored positively in more than one lineage, it was given the label of biphenotypic [2].

Points	B Lineage	T Lineage	Myeloid Lineage
2	CD 79a	CD3 (cyt/m)	Anti-MPO
	cyt IgM	Anti-TCR alpha/beta	
	cyt CD22	Anti-TCR gamma/delta	
1	CD 19	CD 2	CD 13
	CD 10	CD 5	CD 33
	CD 20	CD 8	CDw65
		CD 10	CD 117
0.5	TdT	TdT	CD 14
	CD 25	CD 7	CD 15
		CD 1a	CD 64

Table 1: EGIL Classification system.

Abbreviations: cyt: Cytogenetic; m: Monoclonal; MPO: Myeloperoxidase; TCR: T-Cell Receptor; Tdt: Terminal Deoxynucleotidyl Transferase

Note: >2 points required to assign a lineage designation

The EGIL scoring system added new immune-histochemical markers to improve upon the accuracy of diagnosing ambiguous leukemias. For example, CD79a, a marker known to have strong specificity, was added to the B cell lineage [3]. To detect a positive antigen, 20% of cells had to express the IHC markers. According to the EGIL, only 10% of cells had to stain for CD3, MPO and CD79a, owing to their increased specificity. A score of greater than two was required to designate the presence of a particular lineage. A distinction under this system was continued in which leukemias were considered to be bilinear if they had two distinct blast populations each expressing one lineage or biphenotypic if one blast population co-expressed multiple lineage markers (Weinberg and Arber 2010). Both of these groups would often broadly be defined as leukemias of ambiguous origin. There remained drawbacks to this EGIL scoring system, though, that made accurate classification of these leukemias difficult. As an example, AML with t(8;21) often expressed B cell markers including CD 19, CD 20 and CD 79a [4]. Depending on the number of B cell surface markers present, these leukemias could garner anywhere from two to four points on the EGIL system, leading to a misclassified biphenotypic leukemia. Additionally, clearly defined B-cell acute leukemias often co-express MPO in up to 23% of cases [5].

The WHO reclassification in 2008 represents an effort to simplify and clarify the diagnosis of MPAL (Table 2). First, bilinear and biphenotypic leukemias have been brought together under one diagnosis. A clear delineation was made to exclude genetically defined lesions, such as t(15;17), t(8;21) and inv(16). These core binding factors are AML defining and as such, the presence of B cell markers in these cases would not be classified as biphenotypic [1]. Additionally, irrespective of markers leukemias with complex karyotypes or other MDS related cytogenetics without previous evidence of MDS should all be classified as AML with myelodysplastic changes [1]. The most significant changes in the WHO criteria are related to very specific definitions for assigning a particular lineage.

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Lineage	Markers
Myeloid	Myeloperoxidase OR Monocytic differentiation (at least two of the following: NSE, CD11c, CD14, CD64, lysozyme)
T lineage	Cytoplasmic CD3 OR Surface CD3
B Lineage	Strong CD 19 AND at least one of the following: CD79a, cytoplasmic CD22 or CD 10 OR Weak CD19 AND at least two of the following: CD79a, cytoplasmic CD22 or CD 10

Table 2: 2008 WHO Classifications of leukemias of ambiguous lineage.

Myeloid defining markers include demonstration of immune-histochemical, flow cytometry or cytochemistry positivity of Myeloperoxidase (MPO) [4]. Additionally, presence of monocytic maturation is also evidence of myeloid differentiation. B cell lineage is defined by the expression of strong CD19 plus a secondary B-cell marker such as CD79a, cytoplasmic CD22 or CD10. Additionally, B cell lineage can also be defined by weak CD 19 marking with at least two of the three secondary B-cell markers. T cell lineage is defined by CD3 expression either in its cytoplasmic or surface form. A cohort of 100 patients who met the criteria for MPAL revealed that the B cell/myeloid subtype of MPAL made up the majority of cases. 59 of the 100 patients had B cell/myeloid MPAL versus 35 who demonstrated a T-cell/myeloid subtype. The rest were either classified as T/B cell or trilineage populations [4,6]. As we shall discuss later, these changes have led to a change in the perceived incidence of these ambiguous leukemias.

Clinical Features

The majority of presenting symptoms in mixed phenotypic leukemias are not strikingly different from other forms of acute leukemia. The presence of bone marrow failure, as manifested by anemia and thrombocytopenia, is common. This disease affects patients in the pediatric population as well as adults. There is a slight male preponderance with a ratio of 1.6 to 1 in favor of men in a cohort of 100 patients [7]. Additionally, MPAL have been associated with rather high white counts at presentation [6]. The incidence of the disease has been difficult to pinpoint, mainly owing to the relatively new diagnostic category. In 2009, Al-Seraihy et al., [8] reported on a group of 633 new cases of acute leukemia. Of these, 24, or 3.8%, were reported as being biphenotypic based on the EGIL criteria [8]. Of these, only roughly 1.7% would still be classified as MPAL under the WHO criteria. Owaidah et al., [9] in 2006 found 23 cases of biphenotypic leukemia in a sample size of 676 in whom only 7 would have met the current accepted guidelines for diagnosing MPAL [9]. Looking at a collection of studies of 7,627 new acute leukemias, 213 cases were classified as biphenotypic. Of these, roughly 119, or 1.6%, would still be classified as MPAL [6]. The WHO definition represents a refinement of criteria, which is manifested by a much lower reported incidence of the disease.

Cytogenetic and Molecular Features

Within this newly recognized group of mixed phenotype acute leukemias we find further subcategories based on cytogenetic and molecular features. In general, many patients with MPAL have complex cytogenetics with normal karyotypes seen in only 13% of patients in one case series [10]. A report of a group of pediatric patients by Manola in 2008 reported that 29 of 33 patients (88%) had

some form of cytogenetic aberrancies. Overall there exists a broad range in the incidence of cytogenetic abnormalities, ranging from 59%-91% of cases [11], but irrespective of the study, the literature clearly suggests that the vast majority patients had some form of cytogenetic irregularity.

The significance of categorizing chromosomal abnormalities is manifested by the inclusion of two translocations, t(9;22) and t(v;11q23) in the WHO 2008 criteria [1]. The type and degree of cytogenetic alterations appears to be affected by age. t(9;22), resulting in the Philadelphia chromosome, tends to occur in patients of older age and is in fact, the most common chromosomal change seen in this group [11]. One study by Lee et al., [12] published in 2008 found that roughly 32% of patients with a biphenotypic diagnosis were found to be Philadelphia chromosome positive [12] while only 3% were reported to be positive in the pediatric population [8]. A high white count is typical of patients with the Ph chromosome with the median reported to be 41.8x10⁹/L [13]. The majority of these patients tend to express a B cell/myeloid combination of cell surface markers. By contrast, younger patients tend to most frequently have chromosome changes involving (v;11q23) causing a form of MLL gene rearrangement [8,10]. These patients, too, were noted to have an elevated white count at presentation [11]. In the adult population, only roughly 8% of patients have been noted to carry the MLL rearrangement [10]. Identifying some of these changes translate directly to therapeutic decision making. The two aforementioned genetic abnormalities do seem to have some varying partiality for age. However, many of the following changes do not necessarily have such strong evidence of correlation with age, gender or immunophenotype [4]. Various abnormalities, including del (1) (p32) and t(2;5p13;p13~15.3) have been described in case reports in the pediatric biphenotypic population [11]. Del 1p32 has also been described in pediatric ALL and tends to carry a favorable prognosis [14]. Monosomy 5 has been described in patients with expression of B cell/T cell and myeloid/T cell leukemias. Monosomy 5 is often seen in conjunction with a complex karyotype and portends a worse prognosis [11]. Monosomy 7 or del (7q) has also been reported. Both of these changes generally portend a poorer prognosis. These changes in chromosome 5 and 7 are also seen in patients with MDS and AML. One cytogenetic change worth mentioning involves del (6q) which, when found, represents a favorable prognosis [15]. There remain a host of other abnormalities, including t(10;11), trisomy 8, polysomy 8, t(7;12) and t(8;12), all of which have reported in the literature [11].

Treatment

There are no firm, data-driven established guidelines for the treatment of patients with MPAL. Overall the literature is varied and controversial especially regarding the choice of an induction chemotherapy regimen and the role of Hematopoietic Stem Cell Transplantation (HSCT). In childhood MPAL, there are reports of using modified ALL regimens originally designed for high risk lymphoblastic patients. Many of the drugs in these regimens have activity in myeloid leukemias as well. Once consolidated, patients were maintained on weekly chemotherapy for up to 120 weeks [9].

The St. Jude's Research hospital experience in a pediatric population suggests that ALL-type regimens seem to be more efficacious than AML types. 20 of the 35 patients had a T/Myeloid marker pattern and 12 of 35 had B/Myeloid markers. Two patients had T/Myeloid and B cell markers and 1 patient had an undifferentiated leukemia. No major difference was noted between

these two phenotypic groups. 83% of the patients in this study who received an ALL type regimen were able to achieve a complete remission after induction compared with 52% in the AML group. Additionally, 80% of patients in the AML group who did not receive a complete remission were able to when re-treated with an ALL regimen [16]. In this pediatric group, many patients were maintained on a 120 week regimen. Overall, a total of 91% of the 35 patients in the study were able to achieve a CR. 49% of these patients remained leukemia free survivors. The study did not report data on overall survival as defined by initial induction therapy regimen. Based on their findings, the authors suggest that patients who achieve a molecular remission, defined as <0.01% blasts, might not have to move forward with transplant altogether. Patients who had >1% blasts were recommended to undergo transplant evaluation [16].

In 2011, Matutes et al., [7] reported on outcomes in a group of a 67 patients, comprised of both adult and pediatric age patients [7]. In the study, 27 patients received an ALL type regimen, 34 received an AML regimen with 2 receiving additional imatinib, 5 received a combination AML/ALL therapy and 1 patient received only imatinib therapy. Twenty moved forward with hematopoietic stem cell transplantation once in remission. 85% of patients receiving an ALL regimen went into a complete remission versus 41% of patients receiving an AML regimen. Three of the five patients who received a combination regimen achieved a complete remission. Interestingly, when looking at specific characteristics, median survival was 8 months for Ph positive patients compared with 139 months for patients with normal karyotypes compared with 28 months for patients complex cytogenetics. Median overall survival was noted to be only 11 months for patients treated with an AML regimen versus 139 months for patients who received an ALL treatment course.

A 2008 study by Mikulic et al., [17] reviewed the outcomes of 21 patients defined as biphenotypic acute leukemias based on the EGIL scoring system [17]. In their review, 72% of patients receiving high dose chemotherapy achieved a CR. At 5 years, overall survival was quoted at 21%. When broken down, by therapy, ALL regimens appeared to be more effective. All patients receiving an ALL regimen were noted to be in a CR when compared with 60% when treated with an AML regimen. Some authors advocate the use of a combined AML and ALL chemotherapy regimens for remission induction [18,19]. In the analysis by Zhang et al., of 40 patients, the rate of first complete remission after chemotherapy was 71.4% when a combined AML/ALL regimen was used versus 42.9% when either an ALL or AML regimen was selected [18].

The question of transplantation remains controversial. There is a paucity of data regarding outcomes since the WHO reorganization of these leukemias in 2008. There are no hard and fast strategies to guide the clinician in deciding in whom and when in the treatment course transplantation should occur. Unfortunately, in the evaluation by Matute, there was no report on the outcomes for the 20 patients who underwent either an autologous or allogeneic transplant. In a 3 patient case series reported by Aribi et al., in 2007 [20], 3 patients received allogeneic transplantation after having failed a multitude of therapies. After transplantation all were alive and in remission with a median duration of 4 years [20]. In 2010, Gerr et al., reported a poor prognosis in bilineal acute leukemias and stating that only 2 of their 6 patients were still alive at the time of reporting, both of whom had received an allogeneic transplant [10]. The use of haploidentical HSCT has been employed with some suggestion of efficacy. In their retrospective analysis, Zhang et al., looked at the use of haploidentical

hematopoietic stem cell transplantation. Patients who were treated with chemotherapy alone had a one year survival rate of 37.5% as compared with 60% in patients treated with chemotherapy followed by haploidentical HSCT [18]. The use of haploidentical transplantation offers advantages to the patient such as shorter wait times for a potential donor. Given the generally poor prognosis of MPAL, coupled with newer sources of hematopoietic stem cells and improved techniques to minimize and manage graft versus host disease, transplant seems to be an important option in the management of this leukemia.

There are limited studies evaluating the appropriate transplant regimens in patients with biphenotypic acute leukemia [21]. A retrospective analysis of 59 patients examined the use of a standard conditioning regimen consisting of Total Body Irradiation (TBI) with Cyclophosphamide (CY) or CY with busulfan. The intensified regimen consisted of TBI+ CY+ etoposide or fludarabine+cytarabine plus TBI+ CY+ etoposide. The 5 year relapse rate for all patients after allogeneic transplantation was 52.9%. 80.8% of patients in the standard conditioning group had relapsed at 5 years compared with 28.8% of patients in the intensified conditioning group. Additionally, the overall 5 year survival rate was 23.8% and 64% in the standard and intensified groups. Taken together, these data suggest that if the decision is made to move forward with transplantation, for patients who can tolerate, a more aggressive regimen might be in order.

Summary and Conclusions

Mixed lineage leukemias are a rare entity within the spectrum of acute leukemias. The incidence of these leukemias is likely in the 2% - 3% range. The WHO 2008 classification system has led to a simplification of categorizing these conditions. Fewer, but more specific and robust surface markers have been incorporated to help remove ambiguity in diagnosis. Additionally, specific inclusion criteria, such as exclusion of t(8;21) help to ensure that leukemias are not misplaced. Though these have led to a refinement of the diagnosis of MPAL, many questions regarding how to manage these patients remain.

Mixed phenotypic acute leukemias are generally thought to have a generally poor prognosis with overall survival quoted as low as 10 months. Analysis of cytogenetics is becoming increasingly important as potential targets for therapy in the future. In particular the presence of the Philadelphia of chromosome should always be checked by FISH as it is fairly common and would affect the treatment. Patients with complex cytogenetics or Ph chromosome tend to fare even worse than their other MPAL counterparts. A host of other cytogenetic abnormalities have been discovered, though it is not entirely clear what, if any predictive or prognostic value these may have.

There are no agreed upon treatment protocols and standardization has been difficult to come by. In the absence of prospective randomized data to guide treatment decisions, it has been our practice at Cedars-Sinai Medical Center to use an adult ALL regimen such as hyper-CVAD for young fit patients as it is more inclusive than an AML regimen. A tyrosine kinase inhibitor such as imatinib or dasatinib should be employed in conjunction with cytotoxic chemotherapy for patients with the Philadelphia chromosome. The role of transplantation is not clear. Given the bad prognosis we tend to favor an allogeneic transplantation for all medically fit patients in first remission. Future studies regarding the choice of initial regimen and the role of transplant are clearly necessary even in the era of molecularly targeted therapies.

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