

## LUND UNIVERSITY

#### The European LeukemiaNet AML Working Party consensus statement on allogeneic HSCT for patients with AML in remission: an integrated-risk adapted approach

Cornelissen, Jan J.; Gratwohl, Alois; Schlenk, Richard F.; Sierra, Jorge; Bornhaeuser, Martin; Juliusson, Gunnar; Racil, Zdenek; Rowe, Jacob M.; Russell, Nigel; Mohty, Mohamad; Lowenberg, Bob; Socie, Gerard; Niederwieser, Dietger; Ossenkoppele, Gert J.

Published in: Nature Reviews Clinical Oncology

DOI: 10.1038/nrclinonc.2012.150

2012

#### Link to publication

#### Citation for published version (APA):

Cornelissen, J. J., Gratwohl, A., Schlenk, R. F., Sierra, J., Bornhaeuser, M., Juliusson, G., Racil, Z., Rowe, J. M., Russell, N., Mohty, M., Lowenberg, B., Socie, G., Niederwieser, D., & Ossenkoppele, G. J. (2012). The European LeukemiaNet AML Working Party consensus statement on allogeneic HSCT for patients with AML in remission: an integrated-risk adapted approach. Nature Reviews Clinical Oncology, 9(10), 579-590. https://doi.org/10.1038/nrclinonc.2012.150

Total number of authors:

14

#### **General rights**

Unless other specific re-use rights are stated the following general rights apply:

Copyright and moral rights for the publications made accessible in the public portal are retained by the authors and/or other copyright owners and it is a condition of accessing publications that users recognise and abide by the

legal requirements associated with these rights. • Users may download and print one copy of any publication from the public portal for the purpose of private study or research.

· You may not further distribute the material or use it for any profit-making activity or commercial gain

· You may freely distribute the URL identifying the publication in the public portal

Read more about Creative commons licenses: https://creativecommons.org/licenses/

Take down policy If you believe that this document breaches copyright please contact us providing details, and we will remove access to the work immediately and investigate your claim.

LUND UNIVERSITY

**PO Box 117** 221 00 Lund +46 46-222 00 00

# Allogeneic stem cell transplantation for patients with AML in remission: an integrated risk adapted approach

A position paper from the European Leukemia Net (ELN) AML Working Party

Jan J. Cornelissen<sup>1</sup>, Alois Gratwohl<sup>2</sup>, Richard F. Schlenk<sup>3</sup>, Jorge Sierra<sup>4</sup>, Martin Bornhäuser<sup>5</sup>, Gunnar Juliusson<sup>6</sup>, Zdenek Råcil<sup>7</sup>, Jacob M. Rowe<sup>8</sup>, Nigel Russell<sup>9</sup>, Mohamad Mohty<sup>13</sup>, Bob Löwenberg<sup>1</sup>, Gerard Socié<sup>10</sup>, Dietger Niederwieser<sup>11</sup>, Gert J. Ossenkoppele<sup>12</sup>

Running title: Risk adapted transplantation strategy.

- <sup>1</sup> Department of Hematology, Erasmus University Medical Center, Rotterdam, The Netherlands
- <sup>2</sup> University Hospital, Basel, Switzerland
- <sup>3</sup> Universitätsklinikum, Ulm, Germany
- <sup>4</sup> Hospital Clinic, Barcelona, Spain
- <sup>5</sup> Department of Internal Medicine, University Hospital, Dresden, Germany
- <sup>6</sup> Department of Hematology, Lund University, Lund, Sweden
- <sup>7</sup> Department of Internal Medicine , Masaryk University and University Hospital , Brno, Czech Republic
- <sup>8</sup> Rambam Medical Center, Haifa, Israel
- <sup>9</sup> Department of Hematology, University of , Nottingham, UK
- <sup>10</sup> Service d'Hematologie, Hopital Saint Louis, Paris, France
- <sup>11</sup> Department of Hematology/Oncology, University of Leipzig, Leipzig, Germany
- <sup>12</sup> Department of Hematology, VU Medical Center, Amsterdam, The Netherlands

13 Centre Hospitalier et Universitaire de Nantes, Service d'Hématologie Clinique, Nantes, France; Université de Nantes, Faculté de Médecine, Nantes, France

#### **Reprints/Address correspondence to:**

Jan J. Cornelissen,

Erasmus MC - Daniel den Hoed Cancer Center

Department of Hematology

Groene Hilledijk 301

3075 EA Rotterdam

The Netherlands

Email: j.cornelissen@erasmusmc.nl

Telnr: +31 (0)10 7041153

Faxnr: +31 (0)10 7041004

#### Abstract

Allogeneic hematopoietic stem cell transplantation (alloHSCT) is frequently applied as part of treatment in acute myeloid leukemia (AML) in first or subsequent remission. It reduces relapse, but non-relapse mortality (NRM) and morbidity may counterbalance that beneficial effect. Here we review recent studies reporting new disease specific prognostic markers as well as alloHSCT related risk factors to be identified at specific time points during treatment. We propose risk assessment as a dynamic process during treatment, incorporating both disease and transplant related factors for the decision to proceed either to alloHSCT or with a non transplant strategy, whereby alloHSCT may be favored if projected disease free survival can be expected to be improved by at least 10%, based on individual risk assessment. Pivotal for such an approach are initial disease risk assessment, search for a sibling or unrelated donor early after diagnosis, and the incorporation of time dependent risk factors, all within the context of an integrated therapeutic management approach.

#### Introduction

Myeloablative allogeneic hematopoietic stem cell transplantation (alloHSCT) from an HLAidentical sibling donor is generally recommended for patients with acute myeloid leukemia (AML) in first complete remission (CR) without a favorable risk genetic profile (1-9). Although it offers a strong anti-leukemic effect, the benefit of alloHSCT in terms of overall survival is compromised by non-relapse mortality (NRM). As a consequence, alloHSCT may result in a substantial gain in disease free survival (DFS) and overall survival (OS) in a particular group of patients, but also in a loss of survival in other patients, despite significantly reducing relapse. Therefore, it is important to assess the most significant variables that affect the risk of relapse and variables predicting NRM at diagnosis, but also at later time-points during the course of treatment, as illustrated in Figure 1. The main questions to be addressed repeatedly for the individual patient during that process include:

1. Having obtained remission, to what extent does alloHSCT reduce relapse as compared to an alternative consolidation strategy in this particular type of leukemia?

2. How do alloHSCT and the non-transplant consolidation strategy compare with respect to NRM and also morbidities?

3. Combining the risks of NRM and relapse, what percentage of longterm DFS can be projected for the individual patient?

3. What are the prospects after relapse?

Combining the answers to these questions may yield an estimate to what extent the composite endpoint DFS may be improved and whether quality of life may be compromised. Here we review recent studies reporting new disease specific prognostic markers as well as alloHSCT related risk factors to be identified at specific time points during treatment. We propose risk assessment as a dynamic process during treatment, integrating both disease and transplant related risks for the final decision in the individual patient to proceed either to alloHSCT or with a non transplant approach, aiming for a significant benefit in DFS of at least 10% by alloHSCT, which order of magnitude is based on earlier recommendations following large meta-analyses studies (4-6). Such an integrated approach would deviate from a "one size fits all" strategy and result in a more tailored approach for the individual patient.

#### **Risk of the disease**

Cytogenetic analysis has allowed for distinguishing categories of AML with widely different prognosis and risk of relapse. Three cytogenetic prognostic categories (favorable, intermediate, poor) have long been used. However, cytogenetic risk classification is continuously being refined (12), incorporating new categories such as e.g. the so-called monosomal karyotype (MK) category, which is associated with a very poor outcome and which is already used by several cooperative AML study groups (13). As a detailed review of cytogenetic abnormalities has been performed before, current cytogenetic abnormalities to be taken into account are summarized in Table 1 and presented according to prognosis. The listing is largely based on a recent summary by Grimwade et al (12) and the ELN-recommendations reported earlier (14).

Several larger donor versus no donor studies and their meta-analysis have shown that alloHSCT results in superior DFS and OS in patients with poor-risk AML in CR1. A meta-analysis of 5 earlier studies by Yanada et al had clearly shown improved DFS for patients with poor-risk cytogenetics, but the role of alloHSCT in intermediate risk AML proved less clear (4). The study performed by the HOVON/SAKK consortium, which also included a limited meta-analysis of the combined dataset of the HOVON/SAKK, MRC, EORTC and BGMT studies, showed

improved DFS in both intermediate and poor-risk patients (5). The reduction of relapse was estimated at approximately 50% (hazard ratio (HR) 0.4-0.5), as was derived from an intention to treat analysis. Although relapse was also significantly reduced in favorable risk patients with a risk of relapse below 35%, those patients did not benefit from myeloablative alloHSCT in terms of overall survival as a NRM of approximately 20% attenuated the beneficial effect of alloHSCT in those patients. These results were confirmed and extended in a larger meta-analysis by Koreth et al, including 18 prospective studies in AML CR1 (6). Although these older studies were confined to patients receiving myeloablative alloHSCT using sibling donors, the studies suggested that myeloablative alloHSCT may more generally be recommended for younger patients in first CR with intermediate or poor-risk cytogentic subtypes of AML, but not for patients with cytogenetic favorable subtypes of AML where the relapse probability is 35% or less. The latter applies to most patients with the so-called core binding factor leukemias—AML t(8;21), and AML inv(16)/t(16;16). Meanwhile, continued study of alloHSCT in intermediate risk AML is warranted, because of a continuous trend of progessiviely improved survival following autologous HSCT and/or chemotherapy as consolidation therapy (15-18).

The role of alloHSCT in the new very poor-risk MK subcategory has recently also been addressed (19-22). Although relapse after alloHSCT appeared high, 20% long term survival was reported and virtually no surviving patients were noted among CR patients receiving chemotherapy only or autologous transplantation. Strikingly, the relative reduction of relapse may not differ from what can be observed in other cytogenetic subtypes of AML (20), indicating that the immunotherapeutic effect of alloHSCT is exerted similarly among different AML categories and rather depends on alloreactive minor and major HLA-differences than on leukemia subcategory.

#### **Molecular markers**

The majority of patients with AML in first CR harbor an intermediate risk profile. While most of these leukemias lack a specific, prognostically relevant, karyotypic abnormality, molecular genetic markers such as gene mutations and deregulated gene expression can be identified in the majority and may be associated with a more specific prognosis (23, 24). Approximately 50% of cytogenetically normal AML may carry a mutation in the nucleophosmin gene (NPM1) (25). The prognostic value of the presence of the NPM1 mutation appeared to depend on the additional presence of the internal tandem duplication (ITD) in the FLT3 tyrosine kinase receptor (FLT3/ITD) (26-29). Myeloid leukemia's characterized by the NPM1 mutation but without *FLT3*/ITD, appeared to exhibit a more favorable prognosis with relapse rates less than 30%. Very similar to what was observed in cytogenetic favorable subgroups of AML, characterized by a relapse risk of less than 35%, a German study evaluating alloHSCT in molecularly defined subgroups of cytgenetically normal AML patients showed that patients with NPM1 mutation but without FLT3/ITD did not benefit from alloHSCT due to enhanced NRM, while alloHSCT appeared associated with better survival in patients with the *FLT3*/ITD mutation in their series, although the relapse rate after alloHSCT may be higher as can be observed in intermediate risk AML patients (9-11). Also, the molecular subtype of AML based on the mutation of CEBPA appeared associated with a more favorable prognosis, whereby especially the subtype of AML characterized by a bi-allelic mutation appeared associated with a low risk of relapse (30-34). Therefore, it seems reasonable to withhold myeloablative allogeneic HSCT also in that category of AML patients. Although new molecular abnormalities associated with a better prognosis have been put forward, validation in independent cohorts of patients with mature follow-up are

required before incorporating these abnormalities in our decision making as regards allogeneic HSCT in CR1. Also new molecular markers have been identified that specifically relate to poor or very poor-risk AML, characterized by a very high risk relapse after attainment of first CR (Table 1). These categories for instance include AML with over expression of EVI-1 (35-37). Outcome of younger patients with EVI-1 AML appeared to be dismal, but recipients in CR1, who proceeded to allogeneic HSCT were suggested to benefit (37).

#### **Response and residual disease**

Apart from cytogenetic and molecular prognostic markers that are identified at diagnosis, a number of variables to be monitored during induction and consolidation therapy may offer additional prognostic information, which may affect the decision whether or not to proceed to alloHSCT. Such variables include time to CR, number of blasts early after induction and quantified minimal residual disease (MRD) after induction or consolidation (Table 1) (38). Different groups have shown that quantified levels of MRD relate to outcome and risk of relapse in first CR patients, although prospective validation studies are largely lacking. A study from Italy addressed the question whether multicolor flow cytometry (MFC) applied after induction and consolidation would allow to identify patients with a low risk of relapse < 30-35%, similar to the favorable risk AML subtypes as can be determined prior to the start of treatment by cytogenetics and/or molecular techniques (39). By combining the results of MFC obtained after induction and consolidation, a new subgroup of patients was identified with a favorable prognosis, for whom one may prefer to postpone alloHSCT until eventual relapse. One important caveat in those studies, however, is the effect of alloHSCT itself in that "new" good-risk group. If the majority of patients in that new subgroup actually received an alloHSCT and benefitted

from that modality, it may be hazardous to omit that treatment in the future without having shown favorable outcome of a substantial number of patients, who did not proceed to alloHSCT. In principle, that caveat applies to all studies claiming to identify a new subgroup of good-risk patients, which necessitates the prospective evaluation of risk-adapted treatment, including decisions based on MRD. Furthermore, the presence or absence of MRD before transplantation may provide important prognostic information (40). It may be argued that the presence of MRD identifies a subgroup of patients with a particularly high risk of relapse, thereby qualifying for alloHSCT even in prognostic favorable AML. Minimal residual disease may be quantified by MCF but also by PCR, including quantitative assays. PCR-based monitoring in cytogenetically defined low risk AML, in particular those exhibiting an inv(16) may identify patients with an increased risk of relapse, who then qualify for alloHSCT (41). In addition, the most frequent genetic aberration in AML, the *NPM1* mutation, can also be excellently monitored in a quantitative way, allowing to identify patients with a higher risk of relapse, for whom an alloHSCT may be considered (42, 43).

#### Predicting counterbalancing non relapse mortality (NRM)

As outlined above, there seems to emerge general agreement that NRM associated with myeloablative alloHSCT may outweigh a beneficial effect on relapse in patients with a cytogenetic favorable-risk profile, but with the advent of reduced intensity conditioning regimen as well the identification of the most important parameters predicting for NRM, a careful assessment of NRM-risk should complement the cytogenetic and molecular inventory of the risks associated with the leukemia in each patient. The HOVON analysis of sibling alloHSCT in 4 larger AML-studies (5) showed that age significantly predicted outcome, which effect was

mainly exerted by higher NRM in patients older than 40 years. Apart from age, other variables (Table 2) such as general performance, CMV serostatus, cytokine polymorphism, donor/recipient gender-combination, and comorbidities significantly predict for NRM (44). Taking important individual risk factors into account, composite risk scores were developed, including the European Group for Blood and Marrow Transplantation (EBMT) risk score and the Seattle hematopoietic cell transplantation comorbidity index (HCT-CI). The EBMT risk score is based on 5 criteria: disease stage, patient age, donor type, time interval from diagnosis to transplantation, and donor-recipient gender combination (45). The score was validated in several independent patient cohorts and confirmed over time. Recently, the score was also tested and validated in other hematological disorders, including AML (46). It was shown that AML patients in CR1 receiving myeloablative alloHSCT with a low risk-score ranging in between 0 and 1 point experienced a NRM of less than 15%, patients with scores 2-3 experienced a NRM of approximately 20-25%, and patients with higher risk scores (>4) showed enhanced NRM of approximately 35% (Figure 2, panel A). The risk score was initially conceived to assess NRM and survival. It had, however, also an impact on death from relapse. Loss of overall survival in patients with higher risk scores appeared to be due to both enhanced NRM and a higher relapse rate. Despite that limitation, its application in risk-assessment prior to transplantation appears clinically useful and is now quite widely accepted (47). Risk assessment for an individual patient is complicated as apart from pre-transplant parameters (44-48), also peri-transplant and posttransplant factors influence outcome (Table 2). Peri-transplant factors include the transplant techniques, conditioning regimen, GVHD prevention, and stem cell source (49). Post-transplant risk factors include the predominating factor GVHD, which needs to be assessed itself in terms of predisposing risks (50, see below). Pre-transplant risk factors generally exert additive effects,

but their impact may vary and depend on the sum of the risks. Survival is generally 3-5% worse for CMV seropositive patients compared to CMV seronegative patients, which effect is especially evident in EBMT low risk patients (46). In contrast, the role of Karnofsky performance score may become increasingly important with increasing EBMT risk score, and act independent from comorbidities present (51). Cytokine polymorphisms and single nucleotide changes within HLA locus have recently been described as factors associated with outcome, and may be integrated in risk assessment in the future (52).

Another composite risk-score is the HCT-CI, which was developed in Seattle. In earlier studies, the adapted Charlson comorbidity index predicted NRM, but that index lacked sensitivity (53, 54). Therefore, the hematopoietic cell transplantation (HCT) comorbidity index (CI), based on a number of comorbidities was developed (51, 53-57). A HCT-CI of, respectively, 0, 1 or 2 points resulted in a 2 yr NRM rate of approximately 10%, 15-20% and 25%. A higher CI score of 3 or  $\geq$  4 resulted in NRM rates of 35-40% (Table 3). A validation study showed the impact of the HCT-CI score on OS, NRM and RFS in both AML and MDS (57) patients. The HCT-CI was subsequently also confirmed in other institutions, different diagnoses, and in recipients of reduced intensity conditioning alloHSCT (58-61). A recent study in Spain also confirmed the score, but added more detail in patients with higher scores, enabling to identify subgroups in patients with a score exceeding 3 (62). Collectively, these studies have suggested that acceptable rates of NRM following alloHSCT can be expected in patients with a low EBMT score or with a low comorbidity score. Accordingly, by combining the risk of relapse and NRM (Table 4, Figure 1), it may be argued that patients, whose AML is characterized by a relapse risk > 50%, which may be reduced to less than 25% after alloHSCT, and for whom NRM can be estimated < 25%, those patients may be expected to benefit from alloHSCT by a difference in DFS of at least 10%

and therefore may qualify for alloHSCT. Likewise, a NRM of approximately 30% may still be acceptable in patients, whose leukemia is characterized by a very high risk of relapse (> 80%) (Table 4).

Meanwhile we would like to stress that transplant outcome continues to improve as a result of a number of developments (63-66), including better supportive care, quality management systems, more efficacious infection prophylaxis, and high resolution HLA-typing, which necessitates repeated validation and refinement of NRM risk scores. While it is beyond doubt that high resolution HLA typing has considerably improved matching between donor and recipient and thereby outcome (66), NRM following myeloablative unrelated donor alloHSCT may still be somewhat higher as compared to sibling alloHSCT. The risk of NRM progressively increases with the number of HLA disparities, emphasizing the importance of high-resolution HLA typing and the selection of donors with, preferably, no more than one mismatched allele out of 8 (67, 68). Currently, a number of cooperative groups have incorporated unrelated donor alloHSCT in their protocols for upfront treatment of AML patients, as multiple retrospective and prospective studies have shown that "well matched" unrelated donor grafts may be associated with acceptable NRM and strong reduction of relapse in AML CR1 patients (69-76). In the recent prospective study of the German Austrian AML Study Group, equivalent efficacy and NRM was shown in a head to head comparison of matched related and unrelated donor alloHSCT in adult high-risk patients (76). Collectively, these studies suggest that an unrelated donor alloHSCT is justified if the a priori risk of relapse is sufficiently high and the counterbalancing NRM following unrelated donor alloHSCT can be estimated as moderate. Given the time needed to identify and prepare for an unrelated donor alloHSCT, it implies that the search for a sibling and the subsequent search for an unrelated donor should be performed as soon as possible after diagnosis and initial risk assessment. While the probability of identifying an adult unrelated donor may be as high as 60% for Caucasian patients, still a considerable number of patients with a diverse ethnic background lack a suitable donor. Alternative donors and/or stem cell sources include unrelated cord blood and haploidentical family donors (77-82). Currently, in many centers, such transplants are not routinely performed in patients with AML in first CR given the higher NRM associated with these donors/stem cell sources. However, first CR patients with a very high risk of relapse (> 80%) and lacking a sibling or unrelated donor may qualify for an alternative donor if the risk of NRM can be estimated not to exceed approximately 35%. Preferably, such transplants should be performed by experienced centers, that have validated these transplants with respect to NRM and anti-leukemic efficacy.

### Non-myeloablative (MA) or reduced intensity conditioning (RIC) using sibling or matched unrelated donors in older AML patients or patients with comorbidities.

While alloHSCT has predominantly been studied in younger AML patients, AML predominantly affects older individuals (median age at diagnosis of 71 years (8)). Non-MA or RIC-regimen have been developed in order to reduce NRM in older or medically less fit patients, who do not tolerate a myeloablative conditioning regimen. Several studies have indeed shown that the morbidity and mortality following RIC alloHSCT are less than after MA-conditioning and that encouraging graft versus leukemia effects (GVL) are exerted (reviewed in 83-85, Figure 2 panel B). For the time being, no mature results from prospective, randomized studies comparing these two modalities have been reported in literature sofar. Most comparative studies reported were performed retrospectively and concerned patients with AML/MDS in CR1, CR2, or with advanced disease (86-95). During the prospective German-Austrian study (76), a growing

number of patients received RIC-regimens. Although not randomized, that study prospectively suggested equivalent results in patients receiving RIC or MA regimen in terms of relapse, NRM, and survival. However, several retrospective studies have meanwhile suggested a somewhat higher relapse rate in recipients of RIC alloHSCT and especially in recipients of non-MA alloHSCT (87, 95-97), although these studies relate to patients receiving different types of preceding induction and consolidation therapy, which may impact also on outcome after alloHSCT. Therefore, prospective comparative studies in similarly pretreated patients are highly needed and participation should be encouraged. Nevertheless, a recent prospective comparison of older AML patients by sibling donor availability suggested improved disease free survival for patients with a donor (98, 99). In addition, 2 recent large retrospective studies in older cohorts of AML patients also suggested improved outcome in recipients of RIC alloHSCT as compared to conventional chemotherapy (100, 101). Preferably, a prospective randomized comparison of alloHSCT from sibling or unrelated donors with chemotherapy as consolidation therapy should establish the long-term value of this approach, especially in older patients (Figure 1). Currently, such a prospective randomized study is being performed in Europe by the EBMT and several cooperative groups (ClinicalTrials.gov Identifier: NCT00766779).

#### Adverse effects beyond 2 years after transplantation and quality of life.

So far the decision whether or not to advice alloHSCT as consolidation therapy was discussed by weighing the counterbalancing risks of relapse at one hand and NRM at the other hand. While that discussion is underpinned with mature survival data from a number of studies, late, persistent morbidity and quality of life were not taken into account. However, this issue cannot be disregarded especially when chemotherapeutic approaches and/or autologous HSCT, which

are associated with less morbidity continue to improve outcome in intermediate risk AML (15-18). Several studies in recipients of alloHSCT have addressed the issue of late morbidity and late mortality occurring in patients, who were alive and well at 2 years after alloHSCT (102-106). A relative increase of 20% mortality, gradually occurring during the ensuing 2 decades, has been reported when comparing alloHSCT recipients with age matched controls. Late morbidities include: a long lasting immune deficiency, endocrine dysfunction; skeletal disorders; ocular problems, respiratory tract problems; salivary function and dental problems; liver complications; vascular complications; chronic kidney disease; sexuality; and secondary cancers (107-111), all adversely impacting on quality of life compared to conventional chemotherapy (111). While a detailed discussion of late morbidities and reduced quality of life goes beyond the scope of this review, it should be stressed that the major risk factor for most if not all these morbidities is chronic GVHD and the long lasting immunosuppressive therapy needed for its treatment. Probabilities to encounter each specific morbidity is still relatively low, but the incidence of chronic GVHD following RIC alloHSCT is considerable and a reliable risk estimate that predicts for chronic GVHD is strongly needed (112). A large study from Seattle estimated the risk of persisting GVHD with higher age, peripheral blood as a stem cell source, unrelated donor, and gender combination as the most important risk factors (113). While a reduction of quality of life may be incorporated quantitatively in the so-called "quality of life adjusted life years saved", as was recently reported (114), the complicated quantifications are extremely difficult to convey and to discuss with individual patients, who may value the quality of his/her life on a very individual, personal basis.

#### Transplantation in second remission

Due to its potent anti-leukemic effects, alloHSCT has consistently been considered as the treatment of choice for most relapsed patients. Outcome of allografts beyond first remission, however, is inferior to that in first-remission patients, owing to an increase in both treatmentrelated mortality (25%-35%) and relapse (40%-45%). Breems et al reported a prognostic index for adult patients with relapsed AML based on a cohort of 667 AML patients in first relapse (115). Four relevant parameters significantly predicted for outcome, including length of relapse free interval, age at relapse, cytogenetics at diagnosis, and whether or not patients had received a previous transplantation. Based on these parameters, 3 risk groups could be defined. In all 3 groups, recipients of an alloHSCT faired better than patients treated with chemotherapy or an autograft. However, only 249 out of 667 (37%) relapsing patients entered second CR and less than 50% of those CR2 patients proceeded to allografting (n=109), indicating that ultimately only 15% of relapsed patients received the preferred treatment option. More recent observations were reported by Kurosawa et al (116) and by Armistead (117), who detailed results from 599 relapsing patients treated at MD Anderson. Given the restricted application of alloHSCT in CR2 and in view of the dismal outcome in patients, who did not proceed to alloHSCT, allografting would therefore preferentially need to be considered and weighed in patients in first CR (Figure 1).

#### Conclusion

It has become clear that the decision whether or not to apply alloHSCT in AML patients no longer depends alone on the risk-profile of the leukemia and the availability of a donor. The decision making process has become more complex by also taking into account patient specific parameters that predict for NRM. Table 4 shows a condensed proposal how to integrate the most important risk factors for decision making, whereby we would suggest to favor alloHSCT as consolidation therapy in AML CR1 only if an advantage in DFS of at least 10% may be expected, as can be deduced from the individual risks of relapse and NRM. A continuous risk assessment of the disease in parallel to risk assessment of the transplant procedure is considered pivotal as well as an early search for a donor and stringent cooperation between the leukemia care providers and the transplant teams and unrelated donor registries in the respective countries. While such an integrated system may provide a more tailored approach for the individual patient,, the strategy as such should be subject of ongoing prospective study.. Acknowledgements

Myriam Labopin, statistician from the EBMT acute leukemia working party (ALWP),

is gratefully acknowledged for analyzing NRM in AML CR1 patients, having received myeloablative alloHSCT in Europe between 2000 and 2010 (Figure 2).

#### References

(1) Slovak ML, Kopecky KJ, Cassileth PA, et al: Karyotypic analysis predicts outcome of preremission and postremission therapy in adult acute myeloid leukemia: a Southwest Oncology Group/Eastern Cooperative Oncology Group Study. Blood 2000;96:4075-4083

(2) Burnett AK, Wheatley AH, Goldstone RF, et al: The value of allogeneic bone marrow transplant in patients with acute myeloid leukaemia at differing risk of relapse: results of the UK MRC AML10 Trial. Br J Haematol 2002;118:385-400

(3) Suciu S, Mandelli F, De Witte, T, et al: Allogeneic compared to autologous stem cell transplantation in the treatment of patients < 46 years old with acute myeloid leukemia (AML) in first complete remission (CR1): an intention to treat analysis of the EORTC/GIMEMA AML-10 trial. Blood 2003;102:1232-1240

(4) Yanada M, Matsuo K, Emi N, et al: Efficacy of allogeneic hematopoietic stem cell transplantation depends on cytogenetic risk for acute myeloid leukemia in first disease remission: a metaanalysis. Cancer 2005;103:1652-1658

(5) Cornelissen JJ, van Putten WLJ, Verdonck LF, et al: Results of a HOVON/SAKK donor versus no-donor analysis of myeloablative HLA-identical sibling stem cell transplantation in first remission acute myeloid leukemia in young and middle-aged adults: benefits for whom? Blood 2007;109:3658-3666

(6) Koreth J, Schlenk R, Kopecky KJ, et al: Allogeneic stem cell transplantation for acute myeloid leukemia in first complete remission. Systematic review and meta-analysis of prospective clinical trials. Jama 2009;301:2349-2361

(7) Gupta V, Tallman MS, Weisdorf DJ: Allogeneic hematopoietic cell transplantation for adults with acute myeloid leukemia: myths, controversies, and unknowns. Blood 2011;117:2307-2318

(8) Juliusson G, Karlsson K, Lazarevic, VLj, et al: Hematopoietic stem cell transplantation rates and long-term survival in acute myeloid and lymphoblastic leukemia. Cancer 2011;117(18):4238-4246

(9) Schlenk RF, Döhner K, Krauter J, et al: Mutations and treatment outcome in cytogenetically normal acute myeloid leukemia. N Engl J Med 2008;358:1909-1918

(10) Gale RE, Hills R, Kottridis PD, et al: No evidence that FLT3 status should be considered as an indicator for transplantation in acute myeloid leukemia (AML): an analysis of 1135 patients, excluding acute myeloid leukemia, from the UK MRC AML10 and 12 trials. Blood 2005;106:3658-3665

(11) Brunet S, Labopin M, Esteve J, et al: Impact of FLT3 internal tandem duplication on the outcome of related and unrelated hematopoietic transplantation for adult acute myeloid leukemia in first remission: a retrospective analysis. J Clin Oncol 2012, January 30 [Epub ahead of print]

(12) Smith ML, Hills RK, Grimwade D: Independent prognostic variables in acute myeloid leukemia. Blood reviews 2011;25:39-51, 2011

(13) Breems DA, Van Putten WL, De Greef GE, et al: Monosomal karyotype in acute myeloid leukemia: a better indicator of poor prognosis than a complex karyotype. J Clin Oncol 2008;26:4791-4797

(14) Dohner H, Estey EH, Amadori S, et al: Diagnosis and management of acute myeloid leukemia in adults: recommendations from an international expert panel, on behalf of the European LeukemiaNet. Blood 2010;115:453-474

(15) Vellenga E, Van Putten W, Ossenkoppele GJ, et al: Autologous perpheral blood stem cell transplantation for acute myeloid leukemia. Blood 2011;118:6037-6042

(16) Pfirrmann M, Ehringer G, Thiede Ch, et al: Prediction of post-remission survival in acute myeloid leukaemia: a post-hoc analysis of the AML96 trial. Lancet Oncol 2012;13:207-214

(17) Burnett AK, Hills RK, Milligan DW, et al: Attempts to optimize induction and consolidation treatment in acute myeloid leukemia: results of the MRC AML12 trial. J Clin Oncol 2010;28:586-595

(18) Burnett AK, Hills RK, Milligan DW, et al: Identification of patients with acute myeloblastic leukemia who benefit from the addition of gemtuzumab ozogamcin: results of the MRC AML15 trial. J Clin Oncol 2011;29:369-377

(19) Fang M., Storer B, Estey E, et al:. Outcome of AML patients with monosomal karyotype who undergo hematopoietic cell transplantation. Blood 2011, June 16 [Epub ahead of print]

(20) Cornelissen JJ, Breems D, Van Putten W, et al: A comparative analysis of the value of allogeneic hematopoietic stem cell transplantation in acute myeloid leukemia with monosomal karyotype versus other cytogenetic risk categories. Accepted for publication JCO 2012.

(21) Kayser S, Zucknick M, Döhner K, et al: Monosomal karyotype in adult acute myeloid leukemia: prognostic impact and outcome after different treatment strategies. Blood 2012;119:551-558

(22) Stelljes M, Beelen DW, Braess J, et al: Allogeneic transplantation as post-remission therapy for cytogenetically high-risk acute myeloid leukemia: landmark analysis from a single prospective multicenter trial. Haematologica 2011;96(7):972-979

(23) Marcucci G, Haferlach T, Döhner H: Molecular genetics of adult acute myeloid leukemia: prognostic and therapeutic implications. J Clin Oncol 2009;29:475-486

(24) Mrózek K, Radmacher MD, Bloomfield CD, Marcucci G: Molecular signatures in acute myeloid leukemia. Curr Opin Hematol 2009;16:64-69

(25) Falini B, Mecucci C, Tiacci E, et al: Cytoplasmic nucleophosmin in acute myelogenous leukemia with a normal karyotype (Erratum in: N Engl J Med. 352: 740, 2005). N Engl J Med 2005;352:254-266

(26) Döhner K, Schlenk RF, Habdank M, et al: Mutant nucleophosmin (NPM1) predicts favorable prognosis in younger adults with acute myeloid leukemia and normal cytogenetics: interaction with other gene mutations. Blood 2005;106:3740-3746

(27) Schnittger S, Schoch C, Kern W, et al: Nucleophosmin gene mutations are predictors of favorable prognosis in acute myelogenous leukemia with a normal karyotype. Blood 2005;106:3733-3739

(28) Thiede C, Koch S, Creutzig E, et al: Prevalence and prognostic impact of NPM1 mutations in 1485 adult patients with acute myeloid leukemia (AML). Blood 2006;107:4011-4020

(29) Verhaak RGW, Goudswaard CS, van Putten W, et al: Mutations in nucleophosmin (NPM1) in acute myeloid leukemia (AML): association with other gene abnormalities and previously established gene expression signatures and their favorable prognostic significance. Blood 2005;106:3747-3754

(30) Frohling S, Schlenk RF, Stolze I, et al: CEBPA mutations in younger adults with acute myeloid leukemia and normal cytogenetics: prognostic relevance and analysis of cooperating mutations. J Clin Oncol 2004;22:624-633

(31) Wouters BJ, Löwenberg B, Erpelick-Verschueren CAJ, et al: Double CEBPA mutations, but not single CEBPA mutations, define a subgroup of acute myeloid leukemia with a distinctive gene expression profile that is uniquely associated with a favorable outcome. Blood 2009;113:3088-3091

(32) Dufour A, Schneider F, Metzeler KH, et al: AML with biallelic CEBPA gene mutations and normal karyotype represent a distinct genetic entitiy associated with a favorable clinical outcome. J Clin Oncol 2010;28: 570-577

(33) Green CL, Roo KK, Hills RK, et al: Prognostic significance of CEBPA mutrations in a large cohort of younger adult patients with AML.

J Clin Oncol 2010;28:2739-2747

(34) Taskesen E, Bullinger L, Corbacioglu A, et al: Prognostic impact, concurrent genetic mutations, and gene expression features of AML with CEBPA mutations in a cohort of 1182 cytogenetically normal AML patients: further evidence for CEBPA double mutant AML as a distinctive disease entity. Blood 2011;117:2469-2475

(35) Barjesteh van Waalwijk van Doorn-Khosrovani S, Erpelinck C, Van Putten WLJ, et al: High *EVI1* expression predicts poor survival in acute myeloid leukemia: a study of 319 de novo AML patients. Blood 2003;101:837-845

(36) Lugthart S, van Drunen E, van Norden Y, et al: High EVI1 levels predict adverse outcome in acute myeloid leukemia: prevalence of EVI1 overexpression and chromosome 3q26 abnormalities underestimated. Blood 2008;111:4329-4337

(37) Gröschel S, Lugthart S, Schlenk RF, et al: High Evi-1 expression predicts outcome in younger adult patients with acute myeloid leukemia and is associated with distinct cytogenetic abnormalities. J Clin Oncol 2010;28:2101-2107

(38) Kern W, Schoch C, Haferlach T, et al: Monitoring of minimal residual disease in acute myeloid leukemia. Crit Rev Oncol Hematol 2005;56:283-309

(39) Maurillo L, Buccisano F, Del Principe MI, et al: Toward Optimization of Postremission Therapy for Residual Disease–Positive Patients With Acute Myeloid Leukemia. J Clin Oncol 2008;26:4944-4951

(40) Walter RB, Gooley TA, Wood BL, et al: Impact of pretransplantation minimal residual disease, as detected by multiparametric flow cytometry, on outcome of myeloablative hematopoietic cell transplantation for acute myeloid leukemia. J Clin Oncol 2011;29:1190-1197

(41) Corbacioglu A, Scholl C, Schlenk RF, et al: Prognostic impact of minimal residual disease in CBF-MYH11 positive AML. J Clin Oncol 2010;28: 3742-3749

(42) Schnitger S, Kern W, Tschulik C, et al: Minimal residual disease levels assessed by NPM1 mutation specific RQ-PCR provide important prognostic information in AML. Blood 2009;114: 2220-2231

(43) Krönke J, Schlenk R, Jensen K, et al: Monitoring of minimal residual disease in NNPM1 mutated acute myeloid leukemia: A study of the German-Austrian AML Study Group (AMLSG). J Clin Onc 2011;29:2709-2716.

(44) Anasetti C: What are the most important donor and recipient factors affecting the outcome of related and unrelated allogeneic transplantation? Best Pract & Res Clin Haematology 2008;21:691-697

(45) Gratwohl A, Hermans J, Goldman JM, et al: Risk assessment for patients with chronic myeloid leukaemia before allogeneic blood or marrow transplantation. Chronic Leukemia Working Party of the European Group for Blood and Marrow Transplantation. Lancet 1998;352:1087-1092

(46) Gratwohl A, Stern M, Brand R, et al: Risk score for outcome after allogeneic hematopoietic stem cell transplantation. Cancer 2009;115:4715-4726

(47) Gratwohl A: The EBMT risk score. Bone Marrow Transplant 2011, june 6 [Epub ahead of print]

(48) Passweg JR, Zhang M, Rocha V, et al: Donor characteristics affecting graft failure, graftversus-host disease, and survival after unrelated donor transplantation with reduced-intensity conditioning for hematologic malignancies. Biol Blood Marrow Transplant 2011 Jul 20 [Epub ahead of print]

(49) Körbling M, Freireich EJ: Twenty-five years of peripheral blood stem cell transplantation.Blood 2011;117:6411-6416

(50) Flowers MED, Inamoto Y, Carpenter PA, et al: Comparative analysis of risk factors for acute graft-versus-host disease and for chronic graft-versus-host disease according to National Institutes of Health consensus criteria. Blood 2011;117:3214-3219

(51) Sorror M, Storer B, Sandmaier BM, et al: Hematopoietic cell transplantation-comorbidity index and Karnofski performance status are independent predictors of morbidity and mortality after allogeneic nonmyeloablative hematopoietic cell transplantation. Cancer 2008;112:1992-2001

(52) Martín-Antonio B, Granell M, Urbano-Ispizua A: Genomic polymorphisms of the innate immune system and allogeneic stem cell transplantation. Expert Rev Hematol 2010;3:411-427

27

(53) Diaconescu R, Flowers CR, Storer B, et al: Morbidity and mortality with nonmyeloablative compared to myeloablative conditioning before hematopoietic cell transplantation from HLA matched related donors. Blood 2004;104:1550-1558

(54) Sorror ML, Maris MB, Storer B, et al: Comparing morbidity and mortality of HLA-matched unrelated donor hematopoietic cell transplantation after nonmyeloablative and myeloablative conditioning: influence of pretransplantation comorbidities. Blood 2004;104:961-968

(55) Sorror ML, Maris MB, Storb R, et al: Hematopoietic cell transplantation (HCT)-specific comorbidity index: a new tool for riks assessment before allogeneic HCT. Blood 2005;106:2912-2919

(56) Sorror ML, Giralt S, Sandmaier BM, et al: Hematopoietic cell transplantation-specific comorbidity index as an outcome predictor for patients with acute myeloid leukemia in first remission: combined FHCRC and MDACC experiences. Blood 2007;110:4606-4613

(57) Sorror ML, Sandmaier BM, Storer BE, et al: Comorbidity and disease status based risk stratification of outcomes among patients with acute myeloid leukemia or myelodysplasia receiving allogeneic hematopoietic cell transplantation. J Clin Oncol 2007;25:4246-4254

(58) Sorror ML, Storer B, Storb RF: Validation of the hematopoietic cell transplantation-specific comorbidity index (HCT-CI) in single and multiple institutions: limitations and inferences. Biol Blood Marrow Transplant 2009;15(6):757-8

28

(59) Mohty M, Labopin M, Basara N, et al: Association between the hematopoietic cell transplantation-specific comorbidity index (CI) and non-relapse mortality (NRM) after reduced intensity conditioning (RIC) allogeneic stem cell transplantation (allo-SCT) for acute myeloid leukemia (AML) in first complete remission (CR1). Blood 2009;114 (abstract 650)

(60) Bokhari SW, Watson L, Nagra S, et al: Role of HCT-comorbidity index, age and disease status at transplantation in predicting survival and non-relapse mortality in patients with myelodysplasia and leukemia undergoing reduced-intensity-conditioning hemopoietic progenitor cell transplantation. Bone Marrow Transplantation 2011 Jul 11. doi: 10.1038/bmt.2011.138. [Epub ahead of print]

(61) Eissa H, Gooley TA, Sorror ML, et al: Allogeneic hematopoietic cell transplantation for chronic myelomonocytic leukemia: relapse-free survival is determined by karyotype and comorbidities. Biol Blood Marrow Transplant 2011;17:908-915

(62) Barba P, Pinana JL, Martino R, et al: Comparison of two pretransplant predictive models and a flexible HCT-CI using different cut off points to determine low-, intermediate-, and highrisk groups: the flexible HCT-CI is the best predictor of NRM and OS in a population of patients undergoing allo-RIC. Biol Blood Marrow Transplant 2010;16:413-420

(63) Gooley TA, Chien JW, Pergam SA, et al: Reduced mortality after allogeneic hematopoieticcell transplantation. N Engl J Med 2010;363:2091-2101 (64) Gratwohl A, Brand R, Niederwieser D, et al: Introduction of a quality management system and outcome after hematopoietic stem-cell transplantation. J Clin Oncol 2011;29:1980-1986

(65) Appelbaum FR: Hematopoietic cell transplantation from unrelated donors for treatment of patients with acute myeloid leukemia in first complete remission. Best Practice & Research Clinical Haematology 2007;20:67-75

(66) Petersdorf EW: HLA matching in allogeneic stem cell transplantation. Curr Opin Hematol 2004;11:386-391

(67) Lee SJ, Klein J, Haagenson M, et al: High resolution donor-recipient HLA matching contributes to the success of unrelated donor marrow transplantation. Blood 2007;110: 4576-4583

(68) Lodewyck T, Oudshoorn M, van der Holt B, et al: Predictive impact of allele-matching and EBMT risk score for outcome after T-cell depleted unrelated donor transplantation in poor-risk acute leukemia and myelodysplasia. Leukemia 2011;Oct 25(10):1548-54

(69) Moore J, Nivison-Smith I, Goh K, et al: Equivalent survival for sibling and unrelated donor allogeneic stem cell transplantation for acute myelogenous leukemia. Biol Blood Marrow Transplant 2007;13:601-607

(70) Yakoub-Agha I, Mesnil F, Kuentz M, et al: Allogeneic marrow stem-cell transplantation from human leukocyte antigen-identical siblings versus human leukocyte antigen-allelic-matched unrelated donors (10/10) in patients with standard-risk hematologic malignancy: a prospective study from the French Society of Bone Marrow Transplantation and Cell Therapy. J Clin Oncol 2006;24:5695-5702

(71) Cutler C, Li S, Ho VT, et al: Extended follow-up of methotrexate-free immunosuppression using sirolimus and tacrolimus in related and unrelated donor peripheral blood stem cell transplantation. Blood 2007;109: 3108-3114

(72) Schetelig J, Bornhäuser M, Schmid Ch, et al: Matched unrelated or matched sibling donors result in comparable survival after allogeneic stem-cell transplantation in elderly patients with acute myeloid leukemia: a report from the cooperative german transplant study group.

J Clin Oncol 2009;26:5183-5191

(73) Basara N, Schulze A, Wedding U, et al: Early related or unrelated haematopoietic cell transplantation results in higher overall survival and leukaemia-free survival compared with conventional chemotherapy in high-risk acute myeloid leukaemia patients in first complete remission. Leukemia 2009;23:635-640

(74) Gupta V, Tallman MS, He W, et al: Comparable disease-free and overall survival after Well-Matched unrelated donor and matched sibling donor transplantation in acute myeloid leukemia with adverse risk karyotype in first complete remission. Blood 2010;116:1839-1848

31

(75) Federmann B, Faul Ch, Vogel W, et al: Allogeneic hematopoietic cell transplantation in AML: Comparable results after matched or mismatched unrelated versus matched related transplantation. Blood 2009;114, (abstract 1199)

(76) Schlenk RF, Döhner K, Mack S, et al: Prospective evaluation of allogeneic hematopoietic stem-cell transplantation from matched related and matched unrelated donors in younger adults with high-risk acute myeloid leukemia: German-Austrian trial AMLHD98A. J Clin Oncol 2010;28:4642-4648

(77) Rocha V, Labopin M, Sanz G, et al: Transplants of umbilical-cord blood or bone marrow from unrelated donors in adult with acute leukemia. N Engl J Med 2004;351:2276-2285

(78) Laughlin MJ, Eapen M, Rubinstein P, et al: Outcomes after transplantation of cord blood or bone marrow from unrelated donors in adults with leukemia. N Engl J Med 2004;351:2265-2275

(79) Brunstein CG, Aker KS, Wagner JE: Umbilical cord blood transplantation for myeloid malignancies. Curr Opin Hematol 2007;14:162-169

(80) Brunstein CG, Barker NJ, Weisdorf DJ, et al: Umbilical cord blood transplantation after nonmyeloablative conditioning: impact on transplantation outcomes in 110 adults with hematologic disease. Blood 2007;110:3064-3070

(81) Chen Y-B, Spitzer TR: Current status of reduced-intensity allogeneic stem cell transplantation using alternative donors. Leukemia 2008;22:31-41

(82) Aversa F, Terenzi A, Tabilio A, et al: Full haplotype-mismatched hematopoietic stem-cell transplantation: a phase II study in patients with acute leukemia at high risk of relapse. J Clin Oncol 2005;23:1-8

(83) Baron F, Storb R: Hematopoietic cell transplantation after reduced-intensity conditioning for older adults with acute myeloid leukemia in complete remission. Curr Opin Hematol 2007;14:145-151

(84) Blaise D, Vey N, Faucher C, et al: Current status of reduced-intensity-conditioning allogeneic stem cell transplantation for acute myeloid leukemia. Haematologica 2007;92:533-541

(85) Horwitz ME: Reduced intensity versus myeloablative allogeneic stem cell transplantation for the treatment of acute myeloid leukemia, myelodysplastic syndrome and acute lymphoid leukemia. Curr Opin Oncol 2011;23:197-202

(86) Martino R, Caballero MD, Perez Simon JA, et al, for the AML and alloPBSCT subcommittees of the Spanish Group for Hematopoietic Transplantation (GETH): Evidence for a graft-versus-leukemia effect after allogeneic peripheral blood stem cell transplantation with

reduced-intensity conditioning in acute myelogenous leukemia and myelodysplastic syndromes. Blood 2002;100:2243-2245

(87) Aoudjhane M, Labopin M, Gorin NC, et al: Comparative outcome of reduced intensity and myeloablative conditioning regimen in HLA identical sibling allogeneic haematopoietic stem cell transplantation for patients older than 50 years of age with acute myeloblastic leukaemia: a retrospective survey from the Acute Keukemia Working Party (ALWP) of the European group for Blood and Marrow Transplantation (EBMT). Leukemia 2005;19:2304-2312

(88) Scott BL, Sandmaier BM, Storer B, et al: Myeloablative vs nonmyeloablative allogeneic transplantation for patients with myelodysplastic syndrome or acute myelogenous leukemia with multilineage dysplasia: a retrospective analysis. Leukemia 2006;20:128-135

(89) Shimoni A, Hardan I, Shem-Tov N, et al: Allogeneic hematopoietic stem-cell transplantation in AML and MDS using myeloablative versus reduced-intensity conditioning: the role of dose intensity. Leukemia 2006;20:322-328

(90) Flynn CM, Hirsch B, Defor T, et al: Reduced intensity compared with high dose conditioning for allotransplantation in acute myeloid leukemia and myelodysplastic syndrome: a comparative clinical analysis. Am.J Hematol 2007;82:867-872

(91) Martino R, Iacobelli S, Brand R, et al: Retrospective comparison of reduced-intensity conditioning and conventional high-dose conditioning for allogeneic hematopoietic stem cell

transplantation using HLA-identical sibling donors in myelodysplastic syndromes. Blood 2006;108:836-846

(92) Alyea EP, Kim HT, Ho V, et al: Impact of Conditioning Regimen Intensity on Outcome of Allogeneic Hematopoietic Cell Transplantation for Advanced Acute Myelogenous Leukemia and Myelodysplastic Syndrome. Biology of Blood and Marrow Transplantation 12:1047-1055, 2006

(93) Martino R, Valcarcel D, Brunet S, et al: Comparable non-relapse mortality and survival after HLA-identical sibling blood stem cell transplantation with reduced or conventionalintensity preparative regimens for high-risk myelodysplasia or acute myeloid leukemia in first remission. Bone Marrow Transplant 2008;41:33-38

(94) Luger SM, Ringdén O, Zhang M-J, et al: Similar outcomes using myeloablative vs reducedintensity allogeneic transplant preparative regimens for AML or MDS. Bone Marrow Transplant 2011, Mar 28 [Epub ahead of print]

(95) Ringdén O, Labopin M, Ehninger G, et al: Reduced intensity conditioning compared with myeloablative conditioning using unrelated donor transplants in patients with acute myeloid leukemia. J Clin Oncol 2009;27:4570-4577

(96) Valcárcel D, Martino R, Caballero D, et al: Sustained remissions of high-risk acute myeloid leukemia and myelodysplastic syndrome after reduced-intensity conditioning allogeneic hematopoietic transplantation: chronic graft-versus-host disease is the strongest factor improving survival. J Clin Oncol 2008;26:577-584

(97) Hegenbart U, Niederwieser D, Sandmaier BM, et al: Treatment for acute myelogenous leukemia by low-dose, total-body, irradiation-based conditioning and hematopoietic cell transplantation from related and unrelated donors. J Clin Oncol 2006;24:444-453

(98) Mohty M, De Lavallade H, Ladaique P, et al: The role of reduced intensity conditioning allogeneic stem cell transplantation in patients with acute myeloid leukemia: a donor vs no donor comparison. Leukemia 2005;19:916-920

(99) Mohty M, de Lavallade H, El-Cheikh J, et al: Reduced intensity conditioning allogeneic stem cell transplantation for patients with acute myeloid leukemia: long term results of a donor versus no donor comparison. Leukemia 2009;23:194-196

(100) Farag SS, Maharry K, Zhang M, et al: Comparison of reduced-intensity hematopoietic cell transplantation with chemotherapy in patients age 60-70 years with acute myelogenous leukemia in first remission. Biol Blood Marrow Transplant: 2011, Jun 21 [Epub ahead of print]

(101) Kurosawa S, Yamaguchi T, Uchida N, et al: Comparison of allogeneic hematopoietic cell transplantation and chemotherapy as post-remission strategy in elderly patients with non-M3 AML in CR1: Retrospective analysis with 1036 patients. Biol Blood Marrow Transplantation 2011;17:401-411

(102) Socié G, Veum Stone J, Wingard JR, et al: Long-term survival and late deaths after allogeneic bone marrow transplantation. N Engl J Med 1999;341:14-21

(103) Bhatia S, Francisco L, Carter A, et al: Late mortality after allogeneic hematopoietic cell transplantation and functional status of long-term survivors: report from the Bone Marrow Transplant Survivor Study. Blood 2007;110:3784-3792

(104) Wingard JR, Majhail NS, Brazauskas R, et al: Long-term survival and late deaths after allogeneic hematopoietic cell transplantation. J Clin Oncol 2011;29:2230-2239

(105) Shimoni A, Rand A, Shem-Tov N, et al: Long-term survival and late events after allogeneic stem-cell transplantation with reduced-intensity conditioning (RIC) for AML and MDS; comparable results with myeloablative conditioning. Blood 2009;114 (abstract 518)

(106) Scott Baker K, Ness KK, Weisdorf D, et al: Late effects in survivors of acute leukemia treated with hematopoietic cell transplantation: a report from the bone marrow transplant survivor study. Leukemia 2010;24(12):2039-2047

(107) Robin M, Porcher R, De Castro Araujo R, et al: Risk factors for late infections after allogeneic hematopoietic stem cell transplantation from a matched related donor. Biol Blood Marrow Transplant 2007;13:1304-1312

(108) Socié G, Salooja N, Cohen A, et al: Nonmalignant late effects after allogeneic stem cell transplantation. Blood 2003;101:3373-3385

(109) Curtis RE, Rowlings PhA, Deeg HJ, et al: Solid cancers after bone marrow transplantation.N Engl J Med 1997;336:897-904

(110) Rizzo JD, Wingard JR, Tichelli A, et al: Recommended screening and preventive practices for long-term survivors after hematopoietic cell transplantation: Joint recommendations of the European Group for Blood and Marrow Transplantation, the center for International Blood and Marrow Transplant Research, and the American Society of Blood and Marrow Transplantation. Biol Blood Marrow Transplant 2006;12:138-151

(111) Messerer D, Engel J, Hasford J, et al: Impact of different post-remission strategies on quality of life in patients with acute myeloid leukemia. Haematologica 2008;93: 826-833

(112) Hansen JA, Chien JW, Warren EH, et al: Defining genetic risk for graft-versus-host disease and mortality following allogeneic hematopoietic stem cell transplantation. Curr Opin Hematol 2010;17:483-492

(113) Flowers MED, Storer BE, Lee SJ, et al: Risk factors for the development of acute and National Institute of Health (NIH) chronic Graft-Versus-Host Disease (GVHD). Blood 2009;114 (abstract 345)

(114) Kurosawa S, Takuhiro Y, Uchida N, et al: A Markov decision analysis of post-remission strategies in 2029 patients with AML in first remission (CR1): Should we perform allogeneic hematopoietic cell transplantation CR1? Blood 2011;117:2113-2120

(115) Breems DA, Van Putten WLJ, Huijgens PC, et al: Prognostic index for adult patients with acute myeloid leukemia in first relapse. J Clin Oncol 2005;23:1969-1978

(116) Kurosawa S, Yamaguchi T, Miyawaki S, et al. Prognostic factors and outcomes of adult patients with acute myeloid leukemia after first relapse. Haematologica 2010;95(11):1857-1864

(117) Armistead PM, de Lima M, Pierce S, et al: Quantifying the survival benefit for allogeneic hematopoietic stem cell transplantation in relapsed acute myelogenous leukemia. Biol Blood Marrow Transplant 2009;15:1431-1438

 Table 1. Prognostic factors in AML (leukemia related)

Effect	cytogenetic	molecular	clinical
Favorable	t(8;21) inv(16)/t(16;16) t(15;17)	mutated <i>CEBPA</i> (double) mutated <i>NPM1</i> (without <i>FLT3</i> /ITD)	MRD negative
Adverse	inv(3)/t(3;3) t(9;22) t(9;11) t(6;9) -5 or del(5q) -7 abn(17p) complex karyotype monosomal karyotype	enhanced Evi-1 expression <i>MLL</i> -rearranged <i>FLT3</i> -ITD mutation mutated <i>DNMT3A</i> expression of BAALC expression of ERG expression of MN1 <i>WT1</i> polymorphism <i>BCR/ABL positive</i>	increasing age high WBC extramedullary disease no early CR persistent MRD CD34 positive t-AML

type of parameter

Abbreviations: AML, acute myeloid leukemia; CEBPA, CCAAT/enhancer binding protein; NPM1, nucleophosmin; FLT3-ITD, fms-like tyrosine kinase receptor-3 - internal tandem duplications; Evi-1, ecotropic viral integration site 1; WBC, white blood cells; MLL, mized lineage

leukemia; CR, complete remission; DNMT3A, DNA methyltransferase 3A; MRD, minimal residual disease; BAALC, gene for Brain and acute leukemia, cytoplasmic. ; ERG, Ets-related gene; MN1, meningioma-1; WT1, Wilms tumor 1; t-AML, treatment related AML

 Table 2.
 Prognostic factors for NRM

		AlloHSC1-parameter assessed at timepoint		
Effect	pre-transplantation	peri-transplantation	post-transplantation	
Favorable	sibling donor (HLA-matched) shorter time from diagnosis to transplant* Caucasian race	non-myeloablative conditioning stem cell source (BM/PB) (T-cell depletion)	early immune recovery	
Adverse	higher recipient age* recipient/donor sex* co-morbidities (HCT-CI) CMV serostatus cytokine polymorphism unrelated donor HLA-mismatch performance score refractory leukemia t-AML	myeloablative conditioning regimen alternative stem cell source (UCB)	graft-versus-host disease severity	

#### 

\*factors incorporated in EBMT-risk scrore

Abbreviations: NRM, non-relapse mortality; alloHSCT, allogeneic hematopoietic stem cell transplantation; HLA, human leukocyte antigen; PB, peripheral blood; BM, bone marrow; UCB, umbilical cord blood; HCT-CI, hematopoietic cell transplantation comorbidity index; CMV, cytomegalovirus; t-AML, therapy related AML

Study	Non-relapse mortality (%) by HCT-CI			
	0	1-2	≥ 3	
Sorror et al. (55)				
training set (n=708)	9	14-27	41-43	
validation set (n=346)	14	19-22	40-41	
Sorror et al. (56)	7	19-21	27-37	
(FHCRC n=177, MDACC n=67)				

Table 3. Non-relapse mortality at 2 yrs according to HCT-Comorbidity Index (55, 56)\*

\*The studies included both recipients of matched sibling or matched unrelated donor grafts following either myeloablative or NMA conditioning;

Abbreviations: HCT-CI, hematopoietic cell transplantation-comorbidity index; FHCRC, Fred Hutchinson Cancer Research Center; MDACC, MD Anderson Cancer Center

AML-risk Group**	AML Risk assessment, including respons to induction-I	Risk of relapse following consolidation by		Consider alloHSCT <b>if</b> continued CR after consolidation and <b>if</b> NRM-prognostic scores indicate:		
		Chemo/ autoPBSCT	alloHSCT	EBMT-score	HCT-CI score	NRM
Good	t(8;21), WBC $\leq 20$ Inv(16)/t(16;16) <i>CEBPA</i> + <i>FLT3</i> -ITD-/ <i>NMP1</i> +, and CR1, and no MRD	35-40%	15-20%	NA (≤ 1)	NA (<1)	10-15%
IM	T(8;21), WBC $\leq 20$ CN - X - Y, WBC $\leq 100$ , and CR	50-55%	20-25%	≤2	≤ 2	<20-25%
Poor	Good/IM, but no CR CN –X – Y, WBC >100 CA	70-80%	30-40%	$\leq 3/4$	$\leq 3/4$	<30%
Very poor	MK+ Abn3q26 EVI1+	>90%	40-50%	≤5	≤ 5	<40%

Table 4. Patient specific	, integrated risl	k-based application	of alloHSCT i	in AML CR1*
---------------------------	-------------------	---------------------	---------------	-------------

Abbreviations: alloHSCT, allogeneic hematopoietic stem cell transplantation; AML, acute myeloid leukemia; CR, complete remission; NRM, non-relapse mortality; EBMT, European group for blood and marrow transplantation; HCT-CI, hematopoietic cell transplantation comorbidity index; NA, not advocated; CEBPA, CCAAT enhancer-binding protein alpha; FLT3ITD, fms-like tyrosine kinase receptor-3 - internal tandem duplications; NMP1, nucleair matrix protein; MRD, minimal residual disease; IM, intermediate; WBC, white blood cells; CBF, core binding factor; MK, monosomal karyotype, Evi-1, ecotropic viral integration site .

\*The proposed patient specific application of alloHSCT in AML CR1 integrates the individual risks for relapse and NRM and aims for a DFS benefit of at least 10% for the individual patient as compared to consolidation by a non-alloHSCT approach

\*\*The categorization of AML based on cytogenetic, molecular, and clinical parameters (including WBC) into good, intermediate, and (very) poor subcategories is subject of continuing study and debate. Here, categories are, arbitrarily, presented according to the latest HOVON/SAKK policy (www.HOVON.nl).

#### **Legends to Figures**

#### Figure 1.

Diagnostic and time-dependent parameters predicting for outcome after allogeneic stem cell transplantation versus alternative consolidation therapy in patients with acute myeloid leukemia in first complete remission, who undergo upfront treatment by induction and consolidation therapies.

#### Figure 2.

Cumulative incidence of non relapse mortality, with relapse as competing risk, in AML CR1 patients having received myeloablative alloHSCT (panel A) or reduced intensity conditioning RIC) alloHSCT (panel B) in Europe between 2000 and 2010, as determined by the EBMT risk score, including the parameters patient age, donor type, time interval from diagnosis to transplantation, and donor-recipient gender combination (according to reference 46 and by courtesy of the Acute Leukemia Working Party (ALWP) of the EBMT). Patients receiving RIC alloHSCT were significantly older then patients receiving myeloablative alloHSCT (median age (range): 38 years (35-77) versus 56 (54-77), p,0.0001).



Figure 1

Cumulative Incidence of NRM in AML CR1 (n=8658) after myeloablative alloHSCT: EBMT-ALWP results from transplants performed in 2000-2010.



Figure 2 panel A



Figure 2 panel B

Cumulative Incidence of NRM in AML CR1 (n=3226) after RIC alloHSCT: