

Allogeneic Haematopoietic Cell Transplantation for Myelofibrosis: Proposed Definitions and Management Strategies for Graft Failure, Poor Graft Function and Relapse: Best Practice Recommendations of the EBMT Chronic Malignancies Working Party

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Abstract

Allogeneic Haematopoietic Cell Transplantation (allo-HCT) remains the only curative approach in Myelofibrosis (MF). Despite advances over recent decades, relapse and non-relapse mortality rates remain significant. Relapse rates vary between 15% and 25% across retrospective studies and management strategies vary widely, ranging from palliation to adoptive immunotherapy and, in some cases, a second allo-HCT. Moreover, in allo-HCT, there is a higher incidence of poor graft function and graft failure due to splenomegaly and a hostile 'pro-inflammatory' marrow niche. The Practice Harmonisation and Guidelines subcommittee of the Chronic Malignancies Working Party (CMWP) of EBMT convened an international panel consisting of transplant haematologists, histopathologists and molecular biologists to propose practical, clinically relevant definitions of graft failure, poor graft function and relapse as well as management strategies following allo-HCT. A systematic approach to molecular monitoring, histopathological assessment and chimerism testing is proposed. These proposed recommendations aim to increase the accuracy and uniformity of reporting and to thereby facilitate the development of more consistent approaches to these challenging issues. In addition, we propose management strategies for these complications.

Introduction

Despite many therapeutic advances in the field of Myelofibrosis (MF) over the last decade, allogeneic haematopoietic cell transplantation (allo-HCT) remains the only potentially curative option(1). Registry data reveal an increase in MF allo-HCT activity, with an increasing number of older recipients, the use of reduced intensity conditioning (RIC) and more mismatched donors over time. Current EBMT-European LeukemiaNet (ELN) criteria, last revised in 2015, suggest that 'All patients with intermediate-2 or high-risk disease according to IPSS, DIPSS or DIPSS+, and age <70 years, should be considered potential candidates for allo-SCT'(2). Moreover, patients less than 65 years of age with intermediate risk-1 disease 'should be considered candidates for allo-HCT if they present with either refractory, transfusion-dependent anaemia or peripheral blood blasts >2%, or adverse cytogenetics (as defined by the DIPSS+ classification).' The post allo-HCT course can be complex due to a 'hostile', fibrotic, pro-inflammatory marrow niche, bulky splenomegaly and a higher incidence of poor graft function (PGF)(1). Moreover, despite improving outcomes over time, relapse and non-relapse mortality remain significant. Relapse rates can range from 15-25% and approaches to relapse management vary greatly (3–5).

In MF allo-HCT, clearly defining relapse is made difficult by the dynamics of donor: recipient chimerism, variable rates of molecular disease clearance (when a marker of measurable residual disease (MRD) is available) and widely varying rates of resolution of marrow fibrosis and splenomegaly. We recently performed a survey of 36 international allo-HCT centres with experience in this area in order to get a 'snapshot' of current practice in MF allo-HCT (4). This revealed marked variation in practice relating to disease assessment both pre- and post- allo-HCT, monitoring (bone marrow trephines, chimerism, molecular and cytogenetic annotation) and in the definitions and management of relapse. It highlighted the need for best practice guidelines for the transplant community focused on a working definition of relapse following allo-HCT and on clarifying the difference between graft failure (GF), PGF and relapse. Moreover, there is a need for practical

management recommendations. These guidelines are the first in a series focused on MF allo-HCT with the aim of providing practical, clinically relevant consensus criteria for defining and managing GF, PGF and relapse.

Methodology

The Practice Harmonisation and Guidelines subcommittee of the Chronic Malignancies Working party of EBMT proposed this workshop in January 2020. A survey of 36 international allo-HCT centres with experience in this area was performed, highlighting wide variation in clinical practice (4). A panel of international experts in the field of MF allo-HCT, molecular biology and histopathology was convened. The objective of the panel was to identify the key practical issues relating to GF, PGF and relapse following MF allo-HCT and to produce best practice consensus guidelines. Relevant literature in the MEDLINE and PubMed databases was reviewed by pairs of experts who generated summative conclusions for review by the group. Two teleconferences were held in May 2020 to discuss and refine the suggested recommendations. Given the absence of evidence from randomised trials, the decision was made not to grade these recommendations. They therefore represent the consensus views of the authors.

Definitions

Engraftment

Engraftment after allo-HCT is defined as an absolute neutrophil count (ANC) greater than $0.5 \times 10^9/L$ for three consecutive days. Platelet recovery by definition requires a platelet count $> 20 \times 10^9/L$ for seven consecutive days free of transfusion support. A haemoglobin (Hb) $> 80g/L$, without support, is also necessary to meet the criteria for complete engraftment (6,7).

Post-transplant cytopenias

Mono-, bi- or pancytopenia can occur after allo-HCT in MF patients and can be mild, moderate or severe (i.e. ANC $< 0.5 \times 10^9/l$, anaemia requiring transfusion and/or platelet count $< 20 \times 10^9/L$). In contrast to cytopenias that are related to GF, PGF or MF relapse, cytopenias related to other causes (drug toxicities, vitamin deficiency, viral infections, GVHD, macrophage activating syndrome, hormonal disorders) are usually transient.

Graft Failure

Post-transplant GF can lead to significant morbidity and mortality, and early recognition is key (8). In a recent EBMT study, the five-year survival rate in patients who developed graft failure was 14% (9). In addition to recognised donor-, conditioning- and cell-dose-related risk factors for GF in allo-HCT, there are particular risks associated with MF allo-HCT including the presence of often bulky splenomegaly, an adverse marrow 'microenvironment' due to fibrosis, chronic inflammation, iron overload and potential HLA-sensitisation if the recipient has been heavily transfused (1,10). Of note, Keyzner *et al* found a higher risk of GF in patients undergoing allo-HCT utilizing HLA mismatched donors compared to matched donors (60% versus 13%, $p=0.04$), whereas the bone marrow (BM) fibrosis grade and degree of baseline splenomegaly were not found to affect GF rates (11). GF can be divided into primary GF or secondary GF based on current EBMT definitions (10). Primary GF is defined by an ANC $< 0.5 \times 10^9/L$ by day +28 following stem cell infusion, haemoglobin $< 80 g/L$ and platelets $< 20 \times 10^9/L$. The EBMT guidelines recommend that, in the context of a RIC

allo-HCT, evaluation of donor cell chimerism is required. In cord blood transplantation, where there is a higher incidence of delayed engraftment/ recovery kinetics, count recovery up to day +42 is permitted before the criteria for primary GF apply.

Secondary GF frequently represents a heterogeneous group in 'real world' practice. It is currently defined by EBMT criteria as the presence of an ANC $<0.5 \times 10^9/L$ occurring after initial engraftment and not related to relapse, infection, or drug toxicity. In the RIC setting, loss of donor haematopoiesis (frequently defined by % donor CD15/CD33 population) to below 5% is required, although the use of this threshold is controversial.

Poor graft function

There is no universally accepted definition of PGF. The current EBMT definition requires the occurrence of two or three cytopenias lasting for more than two weeks, after day +28 in the presence of donor chimerism $>5\%$ (10). However, this definition is difficult to apply in clinical practice given chimerism kinetics and potential overlap with evolving graft failure. It is our view that the definitions suggested by both Stasia *et al* and Klyuchnikov *et al* offer a more practical and discriminating set of criteria (12,13). Here, PGF is defined by the presence of mild/moderate cytopenias in at least two haematopoietic lines (ANC $\leq 1.5 \times 10^9/L$, platelet count $\leq 30 \times 10^9/L$, Hb ≤ 85 g/L) lasting for more than two consecutive weeks following documented engraftment, beyond day +14 and in the presence of full donor chimerism. Cytopenias are frequently accompanied by a hypocellular bone marrow although this is not always the case. This definition mandates the absence of severe acute or chronic GVHD, relapse, and either drug- or CMV reactivation-related myelosuppression. Suggested risk factors include underlying host (age)- and disease-related characteristics (higher risk with MF), the use of unrelated donors, major ABO incompatibility, prior HLA sensitisation and low CD34+ cell doses (14–16). The Hamburg group reported on the incidence and risk factors for PGF in a single centre study of 100 patients with MF who underwent RIC allo-HCT, predominantly conditioned with either busulphan- or treosulphan- based regimens and in the pre-ruxolitinib era (14). The cumulative incidence of PGF

was 17% with a median time of onset of 49 (range, 4-99) days. Risk factors within this cohort included older age, particularly in males, and the persistence of bulky splenomegaly at day +30.

MF Relapse following allo-HCT: suggested categorisation

The current International Working Group for MPN Research and Treatment (IWG MRT) criteria for response in MF in the non-allo-HCT setting were last revised in 2013 (17). These encompass not only morphological and haematological criteria for complete remission (CR) and partial remission (PR) but also clinical improvements, anaemia, spleen and symptom responses, where relevant, alongside cytogenetic and molecular evaluation (**Table 1**). These criteria have not been validated in the post allo-HCT setting; direct applicability would be difficult and clearly would not take account of chimerism and variable resolution of fibrosis and splenomegaly.

In the non-transplant setting, the definition of CR in MF includes normal age-adjusted marrow cellularity on the trephine biopsy, <5% blasts and \leq grade 1 MF according to the European grading system (currently the WHO Fibrosis score)(18). Accompanying peripheral blood count criteria include Hb \geq 100 g/L as well as being below the defined local upper normal limit (UNL), ANC $\geq 1.0 \times 10^9$ /L and $<$ UNL, and platelets $\geq 100 \times 10^9$ /L and $<$ UNL with $<$ 2% immature myeloid cells on the blood smear ($<$ 5% immature myeloid cells in splenectomised patients). Finally, clinical criteria include resolution of clinical symptoms, no palpable spleen or liver and no evidence of extramedullary haematopoiesis (EMH), albeit in real world practice this is difficult to ascertain. One potentially controversial point included in the non-transplant setting discussed above is resolution of the fibrosis to \leq grade 1 MF . The dynamics of fibrosis resolution is often very heterogeneous and only occurs in a proportion of patients undergoing drug therapy where duration of response in this regard, if any, remains unpredictable.

A practical approach to defining relapse following allo-HCT would be to further categorise it as one of the following: (1) molecular relapse only, (2) cytogenetic relapse only (rarely reported), (3) molecular AND cytogenetic relapse only, and (4) morphological/clinical relapse. Changes in chimerism would also clearly need to be considered.

Defining Morphological Relapse

In our view, the presence of stable marrow fibrosis, at least within the first twelve months, although of potential concern with increasing time from transplant, is in itself insufficient to diagnose MF relapse. Its significance requires correlation with concurrent clinical findings as well as haematological, MRD and chimerism results. With emerging or established relapse, 'typical' MF-related findings such as age-adjusted hypercellularity, megakaryocytic abnormalities and sustained increases or persistence of marrow fibrosis become more prominent. For the purposes of a new definition of relapse, two post-transplant morphological scenarios should be considered: (1) relapsed disease (recurrence of myelofibrotic morphology and an increased grade of fibrosis following confirmed normalization of marrow / regression of fibrosis), and (2) progressive/accelerated disease (development of myelodysplasia, monocytosis or increased blasts).

The morphologic criteria for post-transplant relapse require that regression of myelofibrotic features had been confirmed previously. In the presence of such normalization of morphology and an associated documented reduction in fibrosis post-allo-HCT, the criteria for relapse include:

1. Increase in age-adjusted cellularity and abnormal M: E ratio
2. Megakaryocytic abnormalities typical of MF (pleomorphism, hyperchromasia, cloud-like nuclei, and megakaryocytic clusters)
3. Increase in grade of reticulin/collagen fibrosis (it should be borne in mind that the previously formed new bone usually takes a long time to be resorbed/resolved and that this should not therefore be used in grading post-transplant unless its density is significantly

greater than the pre-transplant biopsy or there is active evidence of continuing new bone deposition)

The scenarios of progressive or accelerated disease post-transplant can be considered under the following categories:

1. Evidence of myelodysplasia (usual criteria for erythroid & myeloid lineages ensuring that therapy-related changes are excluded; presence of small hypolobated megakaryocytes / micromegakaryocytes for megakaryocytic lineage)
2. MF-related monocytosis that is persistent and not attributable to other causes
3. Increasing peripheral blood / bone marrow blast counts
4. Acute myeloid leukaemia

Defining Molecular Persistence and Relapse

The definition of molecular persistence and relapse in MF allo-HCT is complicated by the variable kinetics of clearance of detectable MRD. It is not feasible to assign an arbitrary time frame to define molecular persistence. However, persistent detectable disease at 3-6 months post-transplant suggests that the patient is at a higher risk of frank clinical relapse. We suggest that molecular relapse could be defined as the re-appearance of the established MRD marker (on two consecutive peripheral blood (PB) samples at least 28 days apart) after documented clearance which had also been confirmed by two consecutive negative PB samples collected at least 28 days apart.

Defining Cytogenetic relapse

Cytogenetic abnormalities are detected in up to 45% of patients with MF, dependent on the risk group and the testing methodology(19). Cytogenetic monitoring following MF allo-HCT is frequently performed using either standard Giemsa-banded (G-banded) karyotyping, FISH for any previously

detected anomaly or SNP Arrays. Cytogenetic relapse can be defined as detection of an informative, previously detected chromosomal abnormality on G-banded, FISH or SNP Array analysis in a patient not meeting the criteria for morphological relapse, as described below. Of note the prognostic significance of cytogenetic evolution post MF- allo-HCT has not been defined.

Defining relapse by Chimerism Analysis after MF allo-HCT

It is well established in allo-HCT practice that rising mixed chimerism (MC) in myeloid malignancies is, in general, associated with an increased risk of relapse. Recent studies have evaluated the effects of mixed chimerism on relapse risk (20,21). Srour *et al* reported on the value of mixed myeloid chimerism (MMC) in predicting early relapse, defined as <95% donor myeloid chimerism at any stage after day +30(21). A total of 35/82 MF allo-HCT patients had MMC, 29 of whom had previous full donor chimerism. Only one patient who maintained full donor chimerism experienced disease progression. In contrast, all apart from two patients with MMC had evidence of morphological or molecular relapse at the time of detection of MMC or soon after. Of note, the majority had detectable molecular relapse concurrent with MMC and, in a few patients, MCC preceded molecular relapse. The four-year progression-free survival (PFS) in the MMC cohort was 4% versus 60% in those with full donor myeloid chimerism ($p < 0.0001$). More recently, Deeg *et al* found that mixed CD33+ chimerism was associated with worse OS and RFS and higher rates of relapse, irrespective of the CD34+ dose infused. Interestingly, in this series, mixed CD3 chimerism was associated with improved RFS and lower rates of GVHD. There was no association between mixed CD3 chimerism and graft failure (22). Although the kinetics of chimerism following the varying range of conditioning platforms used in MF allo-HCT are not fully understood, serial analysis at fixed time points is pivotal for monitoring engraftment, and better defining graft failure facilitates a clearer definition of relapse. Some patients with apparent mixed chimerism remain in long term 'remission' following allo-HCT for MF.

Post-transplant monitoring

Morphological Assessment

Comprehensive, large scale reports of serial marrow morphology and fibrosis remodelling after allo-HCT in MF are lacking, as a result of which it is difficult to make conclusive recommendations regarding the differences between response and relapse. Markedly varied dynamics in the resolution of typical MF features following allo-HCT are likely dependent on disease stage, conditioning intensity, the achievement of sustained engraftment and the extent of donor chimerism. In our experience, post MF allo-HCT marrow samples frequently demonstrate reduced age-adjusted cellularity, particularly in older recipients. Discrepant aspirate and trephine findings are not uncommon, particularly early post allo-HCT. A further challenge is that the distribution and grade of marrow fibrosis can be heterogeneous, making direct comparisons with the pre-allo-HCT trephines challenging (23).

Tamari and colleagues elegantly summarised findings from various studies comparing pre- and post-allo-HCT marrow characteristics (24). MF-associated hypercellularity resolves in the vast majority of cases and this occurs on engraftment or soon thereafter. With regard to MF-related megakaryocytic anomalies (hyperchromasia, cloud-like nuclei, clustering) and increased M:E ratios, resolution was variable. These features were seldom estimated in the early post allo-HCT period but normalisation could be seen early following sustained engraftment. The resolution of bone marrow fibrosis was highly variable and, in some cases, takes over a year (6-23 months), as does resolution of microvasculature anomalies such as increased vessel density. Complete resolution of reticulin fibrosis has been reported in >90% patients at twelve months following myeloablative (high intensity) conditioning (MAC) or RIC transplants. The dynamics of marrow fibrosis resolution with regard to clinical phenotype and outcomes have been reported previously (25,26). Kroger et al evaluated the degree of resolution of fibrosis at day +30 and day +100 in 57 patients with PMF or Post Essential Thrombocythaemia (PET-) or Post Polycythaemia Vera (PPV-) MF (26). By day +100, >50% of patients

had either had no change in fibrosis grade or a 'one grade reduction' only. Those with ongoing grade 2-3 MF (as per European consensus / WHO grading) at day +100 had higher transfusion burdens. There was discordance between the rate of resolution of fibrosis and clearance of MRD despite correlation of fibrosis resolution with degree of donor chimerism. Early clearance (a 2 or 3 grade reduction) by day +100 is associated with improved graft function, a reduced incidence of relapse and good long-term outcome (Five-year OS based on fibrosis score at day +100 was 96% for MF0/1 vs. 57% for MF2/3). Complete resolution after one year occurred in the vast majority of successful allo-HCT. Of note, there appear to be variable responses with respect to modulation of new bone deposition (osteosclerosis) post allo-HC. As a result, its presence cannot be used, in our opinion, to either confirm CR or define relapse (25).

Suggested Marrow Evaluation following MF allo-HCT

In our opinion, it is essential to have a baseline bone marrow trephine sample to allow comparison with the post allo-HCT samples. Bone marrow aspirate evaluation is usually uninformative due to low cell yields. While decalcification protocols vary between institutions, EDTA decalcification yields optimal morphology. Trephine biopsy frequency is often centre- and patient- dependent, but we would suggest a day+100 and a +12 months trephine as a minimum. Additional biopsies at +24 and +36 months, if feasible, would be useful to understand the kinetics of marrow and fibrosis response, as well as aiding in the evaluation of late relapse. As a practical recommendation, when relapse is identified using clinical, haematological, cytogenetic / molecular or chimerism criteria, then a bone marrow evaluation should also be performed.

Reports should encompass the following: (1) assessment of cellularity, (2) the myeloid: erythroid (ME) ratio, (3) megakaryocyte morphology and (4) general comments on engraftment and regeneration. The evaluation of reticulin and collagen fibrosis should be performed using the Gomori technique and Masson's trichrome stain, respectively. Fibrosis grading should utilise the WHO fibrosis score or the

European consensus scoring system(18,27). New bone formation should be assessed (osteosclerosis scoring system, assessed on haematoxylin and eosin stains)(28). Evidence of dysplasia should be evaluated and blast proportion estimated via immunohistochemistry (IHC; minimum CD34/ CD117 and if possible, an extended panel of TdT, HLA-DR, CD56 and CD123). Monocytic components can be assessed via CD14 (preferable) or CD68. One suggested optimal IHC panel for assessment of MF trephines hence incorporates: erythroid markers (one or more of CD71 / e-cadherin / glycophorin A / alpha-Hb), myeloid markers (myeloperoxidase (MPO) +/- CD15), megakaryocyte markers (CD61/CD42b), monocyte/macrophage markers (CD68/CD14) and blast / stem cell markers (CD34 & CD117 at a minimum, optimal panel would also include TdT, HLA-DR, CD56, and CD123).

Detection of driver mutations in MF

Screening investigations for the so-called 'driver mutations' (*JAK2*, *MPL*, *CALR*) performed on peripheral blood DNA will detect a mutation in 80-90% of patients with MF (29). It is recommended that diagnostic testing for *JAK2* V617F should use a test with a limit of detection of 1-3%. Although there are at present no consensus criteria for diagnostic testing limit of detection for either *CALR* exon 9 and *MPL* exon 10 mutations, an assay sensitivity of at least 5% is suggested (29). Low-level *CALR* or *MPL* mutations at diagnosis are thought to be uncommon. A small minority of all cases tests positive for more than one MPN driver gene, particularly when using sensitive detection assays. These are currently considered as being of no specific clinical significance and one clone usually predominates. All such diagnostic or follow-up testing, whether driver mutation testing, MRD analysis or next generation sequencing (NGS, discussed below) should ideally be performed in an ISO 15189 accredited laboratory.

Use of Next Generation Sequencing Panels in MF at diagnosis or prior to allo-HCT

The availability of NGS varies widely. Contrasting data exist regarding the impact of additional somatic mutations on relapse risk, although the number of patients harbouring such mutations has been

limited in the majority of studies. For instance, a higher risk of relapse has been observed in *ASXL1*-mutated cases of Primary MF (PMF) undergoing allo-HCT in some studies (30,31) though not in others(32–34). The Fred Hutchinson group reported a higher risk of relapse in patients with ≥ 3 mutations in addition to *JAK2* or *CALR*, independent of DIPSS-plus risk (33). NGS panels can also be helpful in identifying a clonal marker in cases of ‘triple negative’ MF. We recommend that, at a minimum, MF allo-HCT candidates are screened using an NGS panel for the gene sets included in the mutation-enhanced international prognostic scoring systems for transplant-age patients -MIPPS70+ v2 scoring system -(which includes the genetically inspired prognostic scoring system (GIPSS)) namely; *ASXL1*, *SRSF2*, *EZH2*, *IDH1*, *IDH2*, *U2AF1Q157* (and *CALR* status), although we recognise that wider genomic analysis provides additional prognostic value (35,36). There is a paucity of clinical data to define the appropriate level of detection (LOD). The majority of laboratories use a LOD of 5% but this requires validation by each laboratory.

MRD monitoring in MF allo-HCT- *JAK2/CALR/MPL*

The detection of *JAK2/CALR/MPL* mutations pre-transplant facilitates the assessment of measurable residual disease (MRD) dynamics post-allo-HCT by allowing for the detection of ‘molecular relapse’. Accumulating data support the prognostic relevance of MRD monitoring in MF allo-HCT. For example, in a cohort of 30 MF allo-HCT patients, the detection of a *JAK2* V617F allele burden $>1\%$ on day +28 was associated with a higher risk of relapse and worse overall outcomes (37). Shah *et al* suggested that detection of a *JAK2* V617F allele burden of $>0.05\%$ by allele specific (AS) qPCR within the first 180 days post-transplant would be the optimal cut-off point to identify patients at risk of relapse (38). The largest study, reported from the Hamburg group, investigated the value of MRD analysis in 136 patients undergoing RIC allo-HCT. The rates of molecular clearance varied according to mutation type; Day +100 clearance was higher for *CALR* (92%) than for *JAK2* V617F (67%) and *MPL* (75%)(39). However, the sensitivity of the technique for the detection of *CALR* ranged from 0.02% to 1%, lower than that for *JAK2* or *MPL* (0.01%). This may explain these different rates of molecular response as

opposed to *CALR+* patients having better responses than other genotypes. Patients with detectable molecular disease at day +100 or at day +180 had a significantly higher risk of clinical relapse within five years compared to patients with undetectable disease (62% vs 10%, $P < 0.001$ and 70% vs 10%, $P < 0.001$, respectively), irrespective of the underlying mutation. These figures illustrate the prognostic value of residual minimal disease. Multivariate analysis revealed that detectable MRD at day+ 180 (HR 8.36, 95% CI: 2.76-25.30, $P < 0.001$) was a significant risk factor for subsequent relapse, highlighting the importance of MRD monitoring. MRD can guide utilisation of adoptive immunotherapy with DLI. However, despite these studies, our survey of 'real world' practice revealed wide variations in practice with regard to monitoring with 25% of centres evaluating MRD status only when triggered by haematological or clinical findings suggesting relapse. Furthermore, additional work is required to standardise technical aspects of MRD tests.

The EBMT recommendations for chimerism and measurable residual disease (MRD) assessment following MF allo-HCT are shown in **Table 2**.

Differential diagnoses in MF patients with cytopenias

Given the varying rates of resolution of fibrosis and splenomegaly in parallel with the individual dynamics of MRD clearance and achievement of full donor chimerism, defining relapse, particularly within the first six months following MF allo-HCT, is challenging. While the diagnosis of post-transplant relapse is relatively easy in patients with myeloproliferative features, it can be much more difficult in cytopenic patients. Indeed, a comprehensive laboratory work-up and other tests are required in order to rule out other causes in patients developing post-transplant cytopenias. **Figure 1** provides a suggested algorithm for the evaluation of post-transplant cytopenias in MF patients.

It is of pivotal importance to be able to differentiate between GF, PGF and relapse in MF allo-HCT as both the prognostic implications and the management strategies differ. Most frequently, clinicians

appear to use a combination of clinical findings: failure of resolution or re-emergence of splenomegaly; changes in blood counts/ blood smear; and molecular, cytogenetic and chimerism findings to decide if relapse has occurred following MF allo-HCT, leading to marked variation in relapse criteria and management approaches. There is no uniformly accepted scoring system that can accurately predict the risk of relapse. There is also the possibility of overlapping situations. In **Table 3**, we have summarised the characteristic findings seen in GF, PGF and relapse. Use of these criteria may improve the accuracy and uniformity of reporting across centres and to guide interventions such as DLI and referral for a second allo-HCT. More systematic use of these definitions will aid in the collection of more robust clinical data and contribute to their future refinement.

Management Strategies

Graft failure

Strategies to address risks of graft failure include treatments to reduce splenomegaly (JAK inhibitors), selection of a CMV-seronegative donor for a CMV-seronegative recipient, the evaluation of donor-specific antibodies in HLA haplo-identical or mismatched donor allo-HCT, and patient-stratified conditioning intensity and T cell depletion approaches. Recommendations on the management of GF in MF allo-HCT are shown in **Table 4**.

Poor graft function

As introduced above in the cohort reported from the Hamburg group, PGF patients were found to have similar outcomes at three years when compared to those who did not have PGF. However, this most likely reflects intervention with a CD34+ selected stem cell boost (14). Whether the pre-allo-HCT use of JAK inhibitors will affect the incidence of poor graft function in MF allo-HCT remains unknown. Klyuchnikov *et al* reported on outcomes following CD34+ selected stem cell boost without further conditioning in 32 patients with poor graft function, 14 of whom had MF, with a median interval of five (range, 2-228) months between allo-HCT and infusion of the CD34+ SCB (median CD34+ cell dose:

3.4 x 10⁶/kg). Haematological improvement was observed in 81% of patients, occurring at a median of 30 (range, 14-120) days. The cumulative incidence of grades II-IV acute GVHD was 17% and chronic GVHD 26%(13). The use of a CD34+ stem cell boost in this setting has been reported by other groups (12,16). There is insufficient evidence, in our view, to suggest the routine use of either thrombopoietin agonists or mesenchymal stem cell infusions in the management of poor graft function.

There is no consensus on management strategies to either reduce the risk or treat established poor graft function in MF allo-HCT. The degree of splenomegaly pre-allo-HCT, HLA-sensitisation, iron overload, conditioning platform choice/intensity and persistence of bulky splenomegaly following allo-HCT all contribute to the cumulative risk and hence treatment approach. There is a lack of randomised trial evidence to guide such approaches. The recommendations in **Table 5** therefore represent an expert consensus view.

Post-transplant relapse

Donor Lymphocyte Infusions (DLI)

The first reports demonstrating evidence of a graft-versus-MF effect and highlighting the potential efficacy of DLI were published over two decades ago (40–42).The potential use of DLI should be considered in three discrete settings: prophylactic, pre-emptive and salvage.

Prophylactic DLI

There is no evidence to date to support the routine use of prophylactic DLI in MF allo-HCT. Controlled studies to investigate such an approach would be of interest in high risk and accelerated phase disease.

Pre-emptive DLI

At present, pre-emptive MRD-triggered approaches are the preferred setting for adoptive DLI use. Persistent MRD, increasing mixed patient chimerism and molecular relapse, as defined above, are the

most frequent indications. The weaning and cessation of immunosuppression is generally required prior to pre-emptive DLI. Pre-emptive approaches have demonstrated higher efficacy compared to salvage approaches after frank haematological/ clinical relapse has occurred (43,44). By way of example, Kroger *et al* reported on 17 MF RIC allo-HCT cases who received DLI; in eight patients, the use of pre-emptive DLI was triggered by MRD detection of the JAK2 V617F mutation and nine cases were salvaged by DLI. The median time from allo-HCT was 269 (range, 127-1570) days. The median time to achieve complete remission was 79 (range, 31-495) days. The overall rate of complete molecular remission was 68%, higher in the pre-emptive group than in the salvage group (100% vs 44%; $p = .04$). Our preference is for an escalating dose regimen, where feasible, as opposed to high dose 'bulk DLI', to reduce the risk of severe GVHD (44–46). The dosing regimen of choice is dependent on the transplant platform, the degree of T cell depletion and the donor source. Suggested approaches are shown in **Table 6**. The goal of DLI-based therapy should be clear and may include achievement of CR, ideally MRD negative if there is a suitable marker, or the achievement of full donor chimerism. The continued administration of escalating dose DLI using the end-point of clinically significant GVHD is no longer recommended. It is not currently possible to provide more specific recommendations regarding the timing of DLI or the duration of an escalating dose regimen-based approach given the lack of trial data. The decisions should be based on ongoing clinical assessment of the competing risks of relapse and GVHD. Donor chimerism and MRD, where applicable, should be evaluated after each course. Patients who achieve the targeted endpoints should continue to undergo serial monitoring.

Salvage DLI

Pre-emptive approaches take precedence over salvage DLI approaches. Lower rates of response are observed when compared to pre-emptive approaches(43,44).

EBMT CMWP recommendations on DLI approaches in MF Allo-HCT are shown in **Table 6**.

Dosing of DLI

The dose of DLI depends on the type of donor and the degree of HLA-matching. In addition, subsequent DLI can be given every six-to-eight weeks in the absence of GVHD grade > 1 and if no significant response has been observed.

Recommendations on DLI approaches including dosing and scheduling based on donor type are shown in **Table 7**.

Timing of DLI

The practice survey introduced above revealed marked variation between allo-HCT centres with regard to DLI timing, with some using DLI as early as day +30 for persistent MRD and differing chimerism thresholds between institutions (4). As stated in the EBMT handbook, it is not possible to give precise advice on the timing and dosing schedules for DLI in MF allo-HCT and there is a focus on the collection of more such data in the cellular therapy registries (10). The detection of MRD or falling donor myeloid chimerism between day +100 and day +180 frequently prompts the weaning of immunosuppressive therapy with a view to escalating dose regimen DLI, if required thereafter, in the absence of GVHD. Routine practice is to administer DLI every eight-to-twelve weeks until the end point is achieved. No further DLI is given following the development of GVHD. Improving uniformity in practice may aid in the development of evidence-based dosing schedules.

Combination therapy with DLI

There is no evidence to date to support the routine use of hypomethylating agents, JAK inhibitors, interferon-based therapies or intensive chemotherapy approaches in combination with DLI. Such approaches remain experimental and require further evaluation in a clinical trial setting.

Failure to respond to DLI with progressive disease

If patients do not respond to either escalating dose regimen or salvage DLI and relapse has occurred, then patient suitability for a second allo-HCT should be considered. However, this is often reserved for fitter, younger individuals with a good performance status who are felt to have sufficient reserve to undergo such an approach. Otherwise, the choice between novel agents in clinical trials, the use of JAKi to address constitutional and splenic symptoms or palliative approaches will depend on patient status, clinical context and availability.

JAK inhibitor use for emerging or established relapse

To date, there is no evidence to support the use of JAKi post MF allo-HCT in emerging or frank relapse. The Hamburg group reported on 10 MF allo-HCT cases who relapsed and had mixed chimerism (47). All received ruxolitinib at a median dose of 10 mg twice a day (range, 5-20mg), but all progressed. Further evaluation is required in this setting, in particular in combination with DLI. It is therefore our view that routine use cannot be recommended. There is, however, a possible role post-relapse for the control of splenomegaly, MF-related symptoms and potentially as 'a bridge' to a second allo-HCT in suitable individuals.

Role of second allo-HCT for Relapse or Graft Failure

Data in the literature concerning the role of a second allo-HCT for MF remains sparse, although there have been some comprehensive recent publications (43,48,49). Most patients are not suitable so patient selection is paramount. An adequate Karnofsky performance Score (KPS) in the absence of significant co-morbidities appears to be essential. The main indication is relapsed disease with no response to DLI (or lack of available DLI) or primary/ secondary graft failure. Furthermore, in the context of persistent poor graft function, as discussed above, options include a selected CD34+ stem cell boost or escalation to a conditioned second allo-HCT. Bridging strategies are patient-specific, ranging from proceeding straight to allo-HCT, the use of cytoreductive therapies or JAK inhibitors. The

CMWP recently reported on 216 MF patients undergoing second allo-HCT. The vast majority were RIC allo-HCT, for either relapsed disease (56%) or graft failure (31%), at a median interval from the first procedure of eight months(48). The reported three-year OS estimate was 42% (95% confidence interval [CI]: 34-49), with a median OS of 23 months (95% CI: 10.9–35.7), and 3-year RFS estimate was 39% (95% CI: 31–48). For both OS and RFS, univariate analysis revealed improved outcomes when transplant occurred after the first twelve months, when the indication was relapse rather than graft failure (due to increased NRM) and when the recipient had a KPS >90. There was no benefit in choosing an alternative donor over the original donor unless there was a contraindication or there were donor-directed antibodies. Reports on the use of haploidentical donors in second MF allo-HCT are very limited. The optimal conditioning regimen for a second transplant remains unclear; however, it should ideally be of low toxicity and exert a strong ‘anti-MF’ effect. The Hamburg group reported a single centre experience (n=33) of second allo-HCT for MF using treosulphan (36-42g/m²)-based conditioning in combination with fludarabine and ATG (49). Timely engraftment occurred in all but one patient and the rates of full donor chimerism were satisfactory. The TRM was 16% at day +100, and the five-year disease-free survival and OS were 45% and 47%, respectively. Recommendations regarding a second allo-HCT in MF are shown in **Table 8**.

Conclusions

We acknowledge that these proposed new working definitions of relapse remain unvalidated and will not cover all possible clinical scenarios. They undoubtedly oversimplify the dynamic interplay between blood counts, marrow morphology, the disease clone, and donor-recipient chimerism following an allo-HCT. There is, in addition, the highly variable rate of modulation of marrow topography and fibrosis resolution over time. However, there is an evident need within the field to develop practical empiric definitions of the different kinds of relapse post MF allo-HCT and our suggested guidelines may provide the basis for more consistent reporting in clinical studies and, hence, more informed decisions in routine clinical practice.

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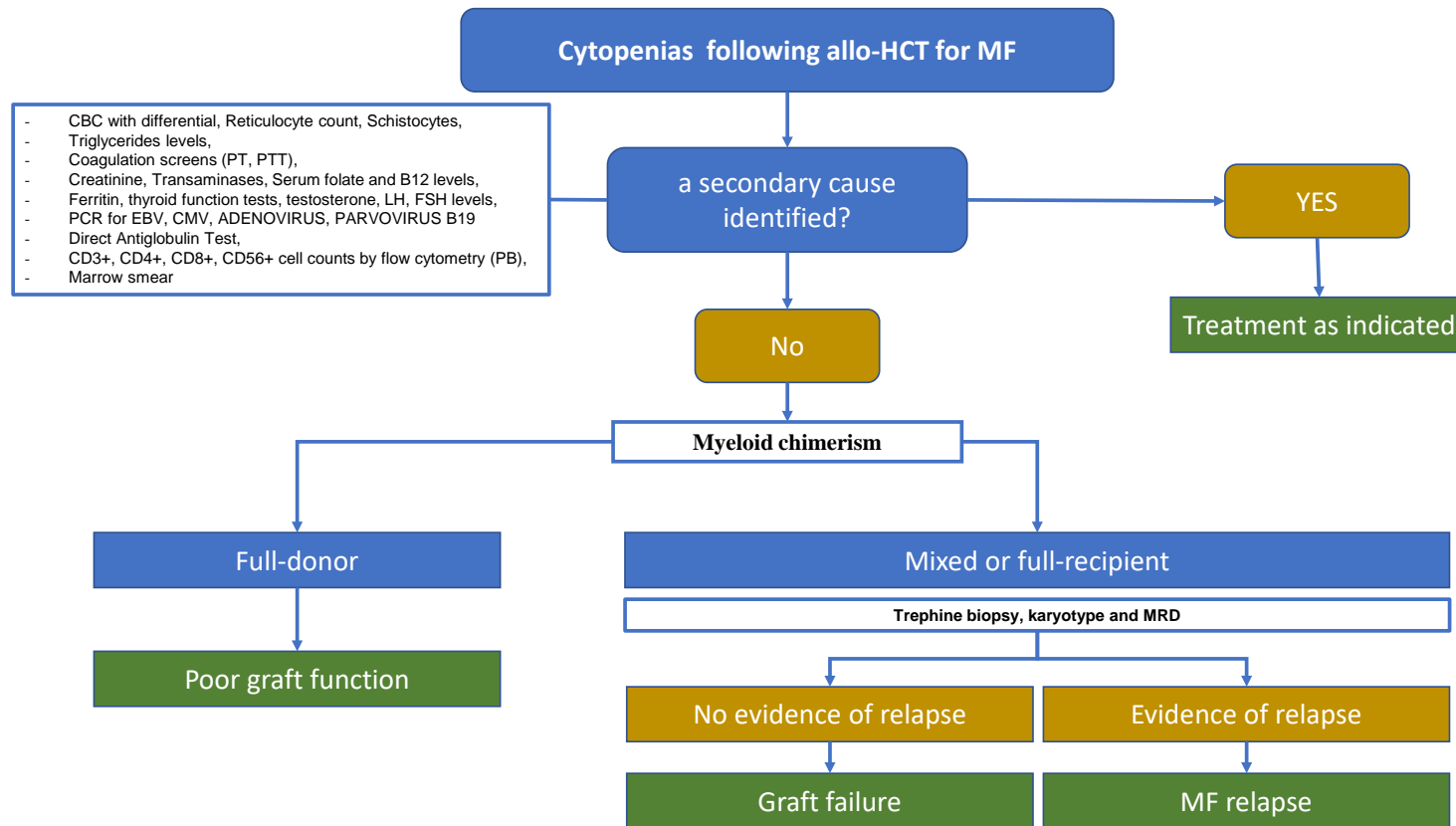
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Figure Legends

Figure 1: Differential diagnosis in MF patients with post-transplant cytopaenias



	Required criteria (duration >12 weeks)	Notes
Complete remission	<p>Bone marrow: Age-adjusted normocellularity; <5% blasts; ≤grade 1 MF*</p> <p>and</p> <p>Peripheral blood: Haemoglobin ≥100 g/L and <UNL; neutrophil count ≥ 1.0 × 10⁹/L and <UNL; Platelet count ≥100 × 10⁹/L and <UNL; <2% immature myeloid cells**</p> <p>and</p> <p>Clinical: Resolution of disease-related symptoms; spleen and liver not palpable; no evidence of extramedullary haematopoiesis</p>	<p>* Grading of MF according to the European classification (WHO Fibrosis score)(18)</p> <p>**<5% immature myeloid cells in splenectomised patients. Immature myeloid cells constitute blasts + promyelocytes + myelocytes + metamyelocytes + nucleated red blood cells</p>
Cytogenetic Remission	<p>CR: eradication of a pre-existing abnormality (N.B. the limit of detection may differ substantially for different methods)</p>	
Molecular Remission	<p>CR: Pre-existing abnormality no longer detectable</p>	<p>Molecular response evaluation must be analysed in PB granulocytes and requires confirmation by repeat testing within six months</p>
Cytogenetic/ Molecular Relapse	<p>Re-emergence of a pre-existing cytogenetic or molecular abnormality that is confirmed on repeat testing</p>	

Table 1: Definition of complete remission of MF in the non-transplant setting, Adapted from the IWG-MPN criteria (2013)(17). CR=Complete remission; PB= peripheral blood

	Chimerism	MRD
Assessment recommended time points	<p>Day +30, Day +100, Day +180, Day +270 and Day +360, or as clinically required</p> <p>As late relapses can occur, there is some evidence to suggest that longer term chimerism testing may be useful(50)</p>	
Technique and interpretation	<p>Chimerism assessment is generally performed on PB but bone marrow-based analysis, if performed, should also be assessed. PB lineage-specific chimerism is recommended</p> <p>Complete chimerism—when all hematopoietic cells are of donor origin, dependent on technique, but frequently defined as >95% of cells analysed as being of donor origin</p> <p>Increasing mixed myeloid chimerism (>5% in the tested lineage when compared to the previous sample of same type) is associated with a higher risk of relapse</p>	<p>Sensitive laboratory techniques are required for MRD assessment, ideally with a sensitivity of 0.01-1%. For <i>JAK2 V617F</i> monitoring, laboratories should ideally use a quantitative PCR, digital PCR or other sensitive methodologies</p> <p>Test accuracy may be validated by use of the WHO Reference Panel (NIBSC (www.nibsc.org) panel code 16/120)</p> <p>Both <i>CALR</i> and <i>MPL</i> MRD monitoring have been used for assessment of MRD</p> <p>There is still a lack of standardisation of quantitative results for <i>CALR</i> and <i>MPL</i>, hence quantitative results should be interpreted with caution and considered within the clinical context</p>
Special cases	<p>Role of CD34+ specific chimerism in MF requires further evaluation</p>	<p>The use of extended NGS panels in MRD monitoring is currently unstandardised and hence remains a research tool. Clinical utility should be evaluated is required in the context of a clinical trial</p>
Role in the diagnosis of relapse	<p>Chimerism assessment is often paired with MRD assessment when there is a suitable marker and this is recommended to increase the predictive value</p> <p>Utilising chimerism to predict and define relapse requires careful individualised interpretation of chimerism kinetics</p>	
Other roles	<p>Chimerism assessment can guide assessment of response to DLI or other relapse-directed therapies</p> <p>Accurate determination of chimerism is technically challenging and testing laboratories should participate in regular external quality assurance schemes e.g., see www.uknegasli.co.uk</p>	

Table 2: EBMT suggested recommendations for chimerism and minimal residual disease (MRD) assessment following MF allo-HCT

Predisposing factors		Primary Graft Failure	Secondary Graft Failure	Poor Graft Function	Disease Relapse
Disease-related	Clinical Factors	Older Age? Bulky splenomegaly Iron Overload? HLA-sensitisation DSA in haploidentical allo-HCT	Older Age? Bulky splenomegaly Iron Overload? HLA-sensitisation DSA in haploidentical allo-HCT	Older age? M>F Bulky splenomegaly Prior HLA-sensitisation DSA in haploidentical allo-HCT	Splenectomy status(51)
	Mutational Profile effect	Lack of evidence	Lack of evidence	Lack of evidence	Controversial, higher risk has been suggested with <i>ASXL1</i>
Transplant-related	Donor	Mismatched/ Haploidentical higher rates Low yield of CD34+ Cells	Mismatched/ Haploidentical higher rates Low yield of CD34+ Cells	Unrelated donors Major ABO incompatibility Low yield of CD34+ Cells	Controversial, ? mismatched unrelated donors/ haplo-identical
	Conditioning	Effect of Intensity undetermined	Effect of Intensity undetermined	Effect of Intensity undetermined	Trends for RIC>MAC
Timing post allo-HCT		Defined by day+28 (not cord where it is day+42)	Occurs after demonstration of initial engraftment	Lasting for > 2 consecutive weeks following documented engraftment, beyond day+14	Most frequent within first 12-months after allo-HCT but late relapses can occur.
Evaluation					
Peripheral blood criteria		<0.5X10 ⁹ /L by day+28, haemoglobin <80 g/L, platelet count <20 × 10 ⁹ /L	ANC <0.5 × 10 ⁹ /L occurring after initial engraftment	Neutrophils ≤1.5 × 10 ⁹ /L, platelet count ≤30 × 10 ⁹ /L, Hb ≤85 g/L	No set criteria. Evaluate for persistent or emerging cytopaenia, or proliferative features.
Marrow smear		Frequently aparticulate and hypocellular	Frequently aparticulate and hypocellular	Frequently aparticulate and hypocellular	Frequently aparticulate and hypocellular. Assess for features suggestive of MF such as megakaryocyte abnormalities
Trephine biopsy		Frequently hypocellular and fibrosis persists	Frequently hypocellular and fibrosis may persist	Frequently hypocellular but may be normo- or even hypercellular.	See Morphological Relapse definition
Karyotype		Not required for definition	Not required for definition	Not required for definition	Detection of previously detected chromosomal abnormality not meeting the criteria for morphological relapse
MRD		Not required for definition	Not required for definition	Not required for definition	Reappearance of established MRD marker after documented clearance confirmed by 2 PB samples collected 28 days apart
Chimerism Analysis		Predominant recipient chimerism	Predominant recipient chimerism In RIC setting, EBMT criteria suggest <5% donor chimerism but controversial	Full donor chimerism	Sensitive techniques will detect early increases in host chimerism suggestive of emerging or established relapse.

Table 3: Characteristics of primary and secondary GF, PGF and disease relapse in MF

Prevention by minimising risk factors where possible and early detection is paramount

A bone marrow aspirate and trephine biopsy should be performed as soon as graft failure is expected

Cytogenetic analysis / Fluorescence in situ hybridisation (FISH; particularly if sex-mismatched allo-HCT) should be performed where appropriate

Both peripheral blood and marrow aspirate sample (where available) should be sent for chimerism analysis following local policies. Depending on practice, this could be whole blood or lineage-specific chimerism and whole marrow or CD34-selected chimerism

Consider myelosuppressive drugs, treatment of viral infections (particularly CMV), and treatment of GVHD

If suspected GF, changes in immunosuppressive therapy are dependent on timing post allo-HCT and the results of the above investigations

Growth factors are often administered though there is little supporting evidence. Frequently they do not improve counts in primary GF and are not a long term-solution

Donor Lymphocyte Infusions (DLI) may have a role in the context of falling donor chimerism and can convert some patients to full donor chimerism and remedy secondary GF; however, this is case-dependent and there are no reports specific to MF(52)

Recipients with primary GF with no functional haematopoiesis and pancytopenia should be considered for a second allo-HCT using either the same or an alternative stem cell donor

In secondary GF, assess for and treat contributing factors such as CMV reactivation, uncontrolled GVHD and myelosuppressive drugs, as indicated. Individuals with true secondary GF should be considered for a second allo-HCT if feasible

Table 4: Recommendations for the investigation and management of Graft Failure in MF allo-HCT

Management of PGF in MF allo-HCT

Infection prophylaxis based on time post allo-HCT, neutrophil count and lymphocyte subset recovery should be based on local institutional policies

Growth factor support with recombinant human EPO (RHuEPO) and GCSF can be considered based on local policies although this is not a long term solution.(14) There is insufficient evidence at present to support the routine use of thrombopoietin (TPO) agonists such as Eltrombopag. Their use remains experimental and more robust evidence is required

Consider a CD34+ selected stem cell boost (SCB), either fresh or cryopreserved, in the presence of full donor chimerism. This can successfully treat poor graft function and does not require further conditioning. The optimal timing of this approach, however, requires further evaluation, as does the associated risk of inducing GVHD(13,14,16)

For some patients with persistent, bulky splenomegaly, there are reports of successful resolution following post allo-HCT splenectomy (14,53)

For eligible patients, in the presence of significant, severe and unresponsive poor graft function, consider a second allo-HCT

Insufficient evidence at present for the routine use of Mesenchymal Stem Cell infusions. Use remains experimental and more robust evidence is required

Table 5: Management of Poor Graft Function in MF allo-HCT

Modality of DLI	Indication for use	Comments	Supporting Evidence
Prophylactic	Not established	No evidence to support use in higher risk or accelerated phase disease to date	Lack of supportive evidence Not recommended in EBMT/ELN 2015 consensus(2) Requires evaluation in a clinical trial setting
Pre-emptive	Mixed chimerism	More common indication in RIC setting PB mixed myeloid chimerism predicts molecular and morphologic relapse better than CD3+ chimerism and overall chimerism status Consider DLI, preferably escalating dose regimen, in absence of achievement of full myeloid chimerism or trend to loss of full myeloid chimerism	(20,21,54)
	Presence of MRD	Directed by MRD or cytogenetic abnormalities as discussed above	(20,39,43,44)
Salvage	Overt morphological relapse	Lower rate of CR compared with pre-emptive strategy	(43,44)

Table 6: Suggested Indications for DLI approaches in MF Allo-HCT

	Donor	HLA-matched Sibling/unrelated (10/10)	HLA-mismatched unrelated (9/10)	Haploidentical (T-cell replete haplo-HCT with PTCy)
1 st DLI, (starting dose *)	Pre-emptive	5 x 10 ⁶ CD3 ⁺ /Kg	1 x 10 ⁶ CD3 ⁺ /Kg	1 x 10 ⁵ CD3 ⁺ /Kg
	Salvage	1 x 10 ⁷ CD3 ⁺ /Kg	1-5 x 10 ⁶ CD3 ⁺ /Kg	1 x 10 ⁶ CD3 ⁺ /Kg
Second and subsequent	Sequential 'Half-log-increase' DLI dose should be given (repeat every 6-8 weeks)			

Table 7: Suggested DLI approaches: dosing and scheduling as per donor type, adapted from De Vos *et al*, 2019 (ref. 55)

Summary of Recommendations for MF second allo-HCT
<p>The main indication for second allo-HCT is relapsed disease with no response to DLI (or lack of available DLI) or primary/ secondary graft failure. Recipient should have a good performance status to be suitable for second allo-HCT with no major co-morbidities.</p> <p>Access to funding for second transplants varies between countries</p>
<p>Bridging therapies to second allo-HCT are very patient-specific</p> <p>No general recommendations can be given</p>
<p>In the absence of a contraindication or donor-specific antibodies, either the original or an alternative donor can be used as outcomes do not appear to differ</p>
<p>Conditioning intensity and the platform of choice depends on recipient age, co-morbidities, the prior conditioning platform and choice of donor</p> <p>Normally these are RIC-based, frequently incorporate ATG, and should be of low toxicity but with a potent 'anti-MF' effect</p>
<p>There is emerging evidence supporting the use of Fludarabine, treosulphan (36-42g/m²) and ATG although further evaluation is required(49)</p>

Table 8: Summary of EBMT Recommendations for MF second allo-HCT

