

## Allogeneic Stem Cell Transplantation in Therapy-Related Myelodysplasia after Autologous Transplantation for Lymphoma: A Retrospective Study of the Francophone Society of Bone Marrow Transplantation and Cellular Therapy

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#### ▶ To cite this version:

Daniel Jaimes-Albornoz, Lionel Mannone, Stephanie Nguyen-Quoc, Yves Chalandon, Patrice Chevallier, et al.. Allogeneic Stem Cell Transplantation in Therapy-Related Myelodysplasia after Autologous Transplantation for Lymphoma: A Retrospective Study of the Francophone Society of Bone Marrow Transplantation and Cellular Therapy. Biology of Blood and Marrow Transplantation, 2019, 25 (12), pp.2366-2374. 10.1016/j.bbmt.2019.07.013. hal-02414852

### HAL Id: hal-02414852 https://uca.hal.science/hal-02414852v1

Submitted on 21 Jul 2022

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Version of Record: https://www.sciencedirect.com/science/article/pii/S1083879119304458 Manuscript\_b862fe7aabff698b3319771de558212d

# Allogeneic stem cell transplantation in therapy-related myelodysplasia after autologous transplantation for lymphoma: a retrospective study of the SFGM-TC

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33					
34	Running title: Allogeneic transplant in secondary myelodysplasia.				
35					
36	Key words: Allogeneic stem cell transplantation, therapy-related myelodysplasia,				
37	autologous stem cell transplantation, lymphoma.				
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52 Therapy-related myelodysplastic syndrome (t-MDS) after autologous stem-cell 53 transplantation (ASCT) is a rare complication with no curative option. Allogeneic 54 hematopoietic stem cell transplantation (allo-HSCT) may be considered for eligible 55 patients and has been understudied in t-MDS. We report 47 consecutive patients with 56 t-MDS after an ASCT who underwent allo-HSCT with a median age of 58 years 57 (range: 30-71) at transplantation and a median follow-up of 22 months (range: 0.7-58 107). The median overall survival (OS) was 6.9 months (95% confidence interval, 0-59 19). OS rates were 45% (29-60%) and 30% (15-45%) at 1 and 3 years after 60 transplantation, respectively. On univariate analysis prior therapy for t-MDS before 61 allo-HSCT (p=0.02) and mismatched donors (p=0.004) were associated with poor OS. 62 Three-year non-relapse mortality (NRM) and relapse rates were 44% (25-63%) and 63 41% (22-61%), respectively. Mismatched donors (p<0.001) were associated with 64 higher NRM and a high-risk MDS (p=0.008) with a higher relapse risk. On 65 multivariate analysis HLA mismatch was associated with higher NRM (HR 6.21; 95% 66 CI 1.63-23.62; p=0.007).

In conclusion, our results suggest that one third of the patients who develop t-MDS after an ASCT for lymphoma are cured after an allo-HSCT. The use of mismatched donors with standard GVHD prophylaxis should be avoided in such indication for allo-HSCT. It will be worth to see if the implementation of CY post-transplantation will improve the outcome with mismatched donors.

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#### 75 INTRODUCTION

76 Treatment options for lymphoid neoplasms include autologous stem-cell 77 transplantation (ASCT). ASCT may be used for the treatment of relapsed or refractory 78 follicular, diffuse large B-cell, Hodgkin's, or T-cell lymphoma and is associated with improved remission rates and prolonged survival.<sup>1-3</sup> Adequate patient selection and 79 80 advances in supportive care have improved outcomes of intensive chemotherapy in 81 recent years. However, such prolongation of survival is also associated with late 82 complications, such as the development of myeloid neoplasm consecutive to the treatment received, including the conditioning chemotherapy of ASCT.<sup>4-6</sup> 83

The risk of developing therapy-related myelodysplastic syndrome (t-MDS) or acute myeloid leukemia (t-AML) secondary to the use of both alkylating agents and topoisomerase inhibitors ranges from 5 to 7% in most series, although there are studies with very variable and extreme incidences between 1% at 30 months to 11.7% at six years and may continue to increase until 12-15 years after ASCT.<sup>7,8</sup>

Therapy-related MDS is associated with a high incidence of cytogenetic abnormalities, with frequent deletions or monosomies of chromosomes 5 and 7.<sup>8,9</sup> These abnormalities have been described after the use of alkylating agents, confirming their role in the development of myelodysplasia.<sup>7,8</sup> Most cytogenetic alterations and acquired mutations are associated with a poor prognosis linked to a low response to chemotherapy and short duration of remission.<sup>10,11,12</sup>

Although allo-SCT has been widely studied in the context of t-AML studies focusing
on t-MDS are rare. Furthermore, the best time to perform allogeneic allo-HSCT in
this group of patients is unknown as well as predictors that might help patient
selection are lacking.<sup>13,14</sup>

We thus launched a retrospective multicentric study to evaluate the results of all
consecutive allogeneic transplantation in population MDS secondary to autologous
SCT for lymphoma

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#### **103 PATIENTS AND METHODS**

104

#### 105 Data collection

106 The registry coordinated by the Francophone Society of Bone Marrow 107 Transplantation and Cellular Therapy (SFGM-TC) of the European Project Manager 108 Internet Server database (ProMISe) was used as the data source. An electronic letter 109 of authorization for the collection and use of the data for this retrospective study was 110 sent to each center. All patients had given written consent before transplant for data 111 collection in ProMISe for future research, in accordance with the Declaration of 112 Helsinki. The scientific council of the SFGM-TC approved this study on February 2, 113 2017.

114

#### 115 **Patient selection**

116 For this retrospective study, we considered all consecutive adult patients who received

an allogeneic transplant for the treatment of t-MDS to ASCT for lymphoid neoplasms

registered from 2006 to 2016 in the ProMISe SFGM-TC database.

119 Patients who received an ASCT and developed secondary t-AML or those who had t-

120 MDS which progressed to t-AML before allo-HSCT were not included. Patients who

121 received ASCT due to neoplasms other than lymphoid neoplasms were not included.

122

#### 123 **Definitions**

124 Lymphoid neoplasms were categorized according to the WHO 2008 classification. 125 The types of MDS followed the WHO 2008 criteria for MDS and were adapted to the WHO 2016 classification.<sup>15</sup> The cytogenetic classification was assessed according to 126 the IPSS score.<sup>16</sup> Lower-risk MDS comprised MDS with low-risk and intermediate-1 127 128 risk IPSS scores, and higher risk MDS those with intermediate-2 and high-risk IPSS 129 scores. MDS-EB-1 and MDS-EB-2 were defined according to WHO and were 130 analyzed together. The other categories of MDS with <5% blasts in bone marrow 131 (MDS SLD, MDS MLD, MDS-RS-SLD, MDS-RS-MLD, MDS del(5q), MDS-U) 132 were analyzed together. The response criteria in MDS in patients who received some 133 type of treatment after or before allo-HSCT was defined according to the International 134 Working Group response criteria<sup>17</sup>.

135 Acute GvHD was described according to the criteria of the International Bone Marrow Transplantation Registry.<sup>17</sup> For the analysis of CMV serostatus, the most 136 137 hazardous combination, defined as a CMV-seronegative recipient and CMV-138 seropositive donor, was compared with the other possible combinations. Intensity of 139 the allo-HSCT conditioning regimen was analyzed according to previously established working definitions.<sup>18</sup> Only two categories were considered in the 140 141 analysis: myeloablative and reduced-intensity conditioning, due to the multiple 142 schemes and doses used. HLA mismatch was defined as the presence of at least one 143 difference in the HLA-A, -B, -Cw, DR, or DQ loci. For the analysis, no mismatch 144 HLA comprised identical-sibling donor and matched unrelated donor 10/10 (MUD), 145 and HLA mismatched those with mismatched unrelated donor 9/10 (MMUD) and 146 cord blood units 4/6 and 5/6.

147 Overall Survival (OS) was defined as the period from the day of allo-HSCT until the

148 day of death from any cause or date of the last recorded follow-up.

149 Non-relapse mortality (NRM) was defined as death from any cause other than relapse

150 of MDS, including progression to AML, from the day of allo-HSCT.

151 Relapse was defined as a relapse of MDS or progression to AML according to the152 WHO criteria.

153

#### 154 Statistical analysis

The characteristics of the patients and the factors related to lymphoma, ASCT, t-MDS, and allo-HSCT are summarized with descriptive statistics. The primary endpoint of the study was OS, which was calculated using the Kaplan-Meier method.

158 The frequency of NRM and relapse was calculated by cumulative incidences.

The prognostic effects of the factors with respect to OS, NRM, and relapse were analyzed with a log-rank test (Mantel-Cox) by univariate analysis. The multivariate analysis was performed using the potential predicting factors that were significant by univariate analysis with Cox proportional hazards regression models. The calculations were performed with SPSS software version 23.0.

164

#### 165 **RESULTS**

To date, the French ProMISe database includes 74,779 autologous transplant and 38,860 allogeneic transplant observations from 98 centers in France, Belgium, and Switzerland. We searched the French ProMISe database for all registered patients from January 2006 to December 2016. A total of 47 patients who met the inclusion criteria were included.

#### 171 Lymphoma and ASCT

172 The initial neoplasm for eight patients (17.0%) was a Hodgkin Lymphoma, whereas it 173 was a Non-Hodgkin Lymphoma for 37 (78.7%) (Table 1). Fourteen patients (29.8%) 174 received more than two lines of chemotherapy before autologous transplantation 175 (range 0-8). Conditioning consisted in Carmustine - Etoposide - Cytarabine -176 Melphalan (BEAM) for 80.8% of patients and other regimens for only 19.2%. Thirty-177 seven patients (78.7%) achieved complete remission. There were no patients with 178 relapsed lymphoma at the time of allo-HSCT, but it is unknown whether there were 179 patients who relapsed or progressed from lymphoma between ASCT and allo-180 HSCT. There were no patients who relapsed from lymphoma after allo-HSCT.

181

182 MDS

The median time from ASCT to the diagnosis of t-MDS was 74.4 months (range 2.2259). Eleven patients (23.5%) had MDS-EB-1, 12 (25.6%) MDS-EB-2, and 22
(46.7%) other types of MDS. No diagnostic information was obtained for two patients
(4.2%).

187 Cytogenetic data were obtained for 41 patients: 82.9% had at least one cytogenetic 188 abnormality. The most frequent cytogenetic anomalies were on chromosomes 7 (11 189 patients, 23.4%) and 5 (six patients, 12.8%) or both of those (fifteen patients, 31.9%). 190 Among the patients, 23.4% were considered to be at low risk and 59.6% at high-risk. 191 Sixty-eight percent of patients received at least one treatment line before allo-HSCT: 192 16 (34.1%) received hypomethylating agents and 11 (23.4%) AML-like induction 193 treatment. Ten patients (21.3%) achieved complete remission before transplantation 194 and 36 (76.6%) were not in complete remission prior to allo-HCST.

196 HSCT

197 The characteristics of the patients at allo-HSCT are shown in Table 2.

The median age at allo-HSCT was 58 years (range: 30 - 71 years) and most of the patients were male (78.7%). The median time interval from diagnosis of t-MDS to allogeneic transplantation was 7.9 months (range: 2.5 - 16.8).

201 Peripheral blood stem-cells (PBSC) were the source used for 87.4% of patients. 202 Nineteen patients (40.5%) were transplanted from an HLA-identical sibling donor and 203 17 (36.2%) from a matched unrelated donor. 10 patients (21.2%) had an mismatched 204 unrelated donor including three patients (6.3%) received umbilical cord blood 205 transplantation (one with a double cord). None of them received a haploidentical 206 donor. Myeloablative conditioning (MAC) was used in nine patients (19.1%), four 207 received a combination of busulfan/cyclophosphamide regimen (two BuCy and two 208 CyBu), with conventional doses (12.8 mg/kg busulfan IV, 120 mg/kg 209 cyclophosphamide), three a combination of fludarabine/busulfan with ATG (150 mg/m<sup>2</sup> fludarabine, 12.8 mg/kg busulfan and various doses of ATG between 2.5 mg-5 210 mg/kg), one fludarabine/busulfan without ATG (160 mg/m<sup>2</sup> fludarabine, 9.6 mg/kg 211 busulfan), and one fludarabine and TBI (120 mg/m<sup>2</sup> fludarabine, 8 Gy TBI). 212

Thirty-eight patients (80.9%) received reduced intensity conditioning (RIC). 22 patients received a combination of fludarabine-busulfan-based RIC with ATG or ALG (100–150 mg/m<sup>2</sup> fludarabine, 3.2–6.4 mg/kg busulfan, 2.5-5 mg/kg ATG or 5–20 mg/kg ALG). 4 patients received fludarabine-busulfan-based RIC without ATG. Eight patients received fludarabine-TBI-based RIC (2Gy–8 Gy) and four patients received sequential FLAMSA-RIC.

Eighteen patients (38.3%) received prophylaxis with cyclosporine (CsA) and mycophenolate mofetil (MMF) and 11 (23.4%) cyclosporine and methotrexate (MTX). Twelve patients (25.5%) received only cyclosporine and six (12.8%) received other regimens. The most frequent serological status for cytomegalovirus (CMV) was 29.7% R<sup>+</sup>/D<sup>+</sup>. Thirteen transplants (27.1%) were performed with the combination R<sup>+</sup>/D<sup>-</sup>.

- Performance status was assessed using the Karnofsky Performance Score (KPS): 29
  patients (61.8%) had a score of 90-100 before transplantation.
- 227

#### 228 **Response to allo-HSCT and complications.**

The median duration of post-transplant follow-up was 22 months (range 0.7-107) and
median survival 6.9 months (95% CI 0-19).

Thirty-four patients (72.3%) were in complete response after allo-HSCT and nine patients (19.1%) had a relapse/progression post-transplantation. The response was not evaluated for four patients (8.6%) because of the early death of three and lost to follow-up one of them.

Acute GvHD occurred in 20 patients (42.5%), seven (14.8%) had grade I, and 13 (27.7%) had a grade requiring treatment (Grade II-IV). Ten patients (21.3%) developed chronic GvHD, of whom four (8.6%) had extensive and six (12.7%) limited involvement At the time of the analysis in August 2017, there were 15 relapses or progressions of which 13 died: 12 related to relapse and one could not determine the cause. 11 patients in complete remission died due to transplant complications, mainly infectious. 3 patients had early death due to transplant complications without knowing the response obtained to the
transplant. No deaths due to veno-occlusive disease or graft rejection were reported.
The causes of death are listed in Table 3.

245

#### 246 Univariate and Multivariate analysis of patients

247 OS for all patients was 45% (95% confidence interval 29-60) at first year, 39% (95% 248 CI 24-55) at 2 years and 30% (95% CI 15-45) at 3 years (Table 4 and Figure 1). 249 Univariate analysis found that prior therapy of t-MDS with hypomethylating agents 250 before allo-HSCT (p=0.02) and the presence of an HLA mismatch (p=0.004) were 251 associated with poorer OS (Table 5). Multivariate analysis showed only a non-252 statistically significant association with poorer OS for patients receiving a treatment 253 based on hypomethylating agents (HR, 3.55; 95% CI, 0.97 - 12.97; p=0.06), and no 254 other clinically significant factors (Table 6).

NRM was 35% (95% CI 18–51) at first year, 39% (95% CI 21–56) at 2 years, and
44% (95% CI 25–63) at 3 years (Table 4 and Figure 1). Univariate analysis identified
gender (p=0.02), graft type (p=0.02) mismatched unrelated donor type (p<0.001) and</li>
the presence of an HLA mismatch (p=0.001) to be significant risk factors (Figure 2).
Multivariate analysis showed an association between the use of a mismatch unrelated
donor (MMUD) and shorter survival after transplantation, relative to identical siblings
or matched unrelated donors (MUD) (HR, 6.21; 95% CI, 1.63–23.62; p=0.007)

The risk of relapse was 35% (95% CI 18–53) at first year, 41% (95% CI 22–61) at 2 years and 41% (95% CI 22–61) at 3 years (Table 4 and Figure 1). The type of MDS and presence of marrow blasts (p=0.008) were the most significant predictive factor of relapse in univariate analysis, however multivariate analysis did not identify the presence of marrow blasts as a significant factor of relapse.

#### 267 **DISCUSSION**

268 This retrospective study of the SFGM-TC, which examined the experience of allo-269 HSCT in patients, with t-MDS after an ASCT for lymphoid neoplasm, over 10 years, 270 showed donor mismatch to adversely affect OS. This is the first series published in 271 this specific population. In other studies, such patients account for 7 - 32% of the sample.<sup>13,14,19-26</sup> Most of the survival data of these studies show results for a mix of 272 273 patients, including those with t-AML and t-MDS exposed to multiple treatments 274 (chemotherapy and/or radiotherapy, and not necessarily previous ASCT) and various 275 primary diseases (solid organ neoplasms, lymphoid neoplasms, myeloid neoplasms, 276 and congenital anomalies).

Although allo-HSCT has been used as a curative therapeutic modality for eligible patients, data published by several groups have shown poor long-term survival.<sup>13,14,19-</sup> <sup>21</sup> In our series, the median overall survival was 6.9 months with OS at first and third year of 45 and 30%, respectively. The high NRM and relapse rates at three years (44% and 41%, respectively) were similar to those previously reported by other studies, consistent with the poor prognosis of these patients.

283 Relapse or progression of t-MDS was the main cause of mortality (44.4%) in our 284 study. Patients with MDS-EB-1 and MDS-EB-2 had a higher risk of relapse by 285 univariate analysis. The association of these aggressive variants of myelodysplasia 286 with higher relapse rates has been described in other studies on patients with various 287 primary diseases. It is not known whether treating t-MDS before transplantation can 288 decrease the risk of relapse or whether maintenance treatment should be started after 289 transplant to prevent relapse. In our study, the only type of pre-transplant treatment 290 associated with improved survival was AML-like therapy even if they had not

291 achieved CR before transplantation, in univariate analysis (p= 0.02). The multivariate 292 analysis shows a non-statistically significant association of hypomethylating agents 293 before transplantation with a poorer OS (HR 3.55, 95% CI 0.97-12.97, p=0.06) 294 regardless of the response they obtained with this treatment before transplantation, in 295 contrast to patients with de novo MDS and poor risk cytogenetics, who normally benefit from this approach.<sup>28</sup> Our results are not valued for the sample size and the 296 297 retrospective analysis. Prospective studies are necessary to determine the benefit of a type of treatment before transplantation<sup>27</sup> A retrospective analysis that included more 298 299 patients with t-AML than t-MDS, reports relapse rates of 42% and 44% at 5 and 10 300 years with OS of 38% and 24% respectively, when AML-like chemotherapy is used before transplantation.<sup>25</sup> 301

No patient in our study received post-transplant maintenance therapy. The use of maintenance therapy, with low doses of azacytidine after allo-HSCT<sup>28</sup> and azacytidine<sup>29</sup> or decitabine with infusion of donor lymphocytes<sup>30</sup>, has been published recently and may be worth considering. Targeted therapies, directed against mutated oncogenes, such as IDH-1, IDH-2, or FLT-3 genomic alterations, may improve the outcome of specific subsets of patients in the future.

308 Infections and GVHD were the main cause of non relapse mortality.

In our series there were 27 deaths, 12 were due to relapse, 8 due to infections and 3 due to GVHD. The few deaths due to GVHD show the intensity of the immunosuppressive prophylaxis used, limiting the graft-versus-tumor effect, increasing the possibility of relapse and the appearance of infections. Multiple schemes and doses used in conditioning and immunosuppressive prophylaxis, testing the usual drugs, have not allowed to identify whether one scheme is superior to

315 another. A significant factor to improve survival was the absence of a mismatch by 316 using either HLA identical-sibling or matched unrelated donors, as previously reported.<sup>14</sup> A prospective study of the SFGM-TC reported better OS (37% vs 15%, 317 318 p=0.02) of patients with high-risk MDS who had identical HLA donors versus those who did not.<sup>31</sup> In our series, there was a lower frequency of NRM in patients with 319 320 10/10 donors than 9/10 donors (HR, 6.21, 95% CI, 1.63-23.62, p=0.007), although 321 was not significative in OS by multivariate analysis. NRM associated with 322 mismatched HLA donors could be improved with new prophylactic strategies to 323 counter GvHD, such as post-transplant cyclophosphamide (PT-Cy). Reports over the 324 last years on a small number of patients with de novo MDS/AML and t-MN who 325 underwent allo-HSCT show similar results for HLA identical-sibling, MUD, or haploidentical donors.<sup>32</sup> A recent retrospective series of EBMT in patients with MDS 326 327 who underwent haploidentical transplants reported better OS for patients treated with 328 PT-Cy than those who were not (OS at three years of 38% vs 28%), but with high 329 NRM (41% vs 55%).<sup>33</sup> This modality may be an acceptable option, although the risk 330 of relapse remains high and variations in the dose of PT-Cy have even been tested in high-risk patients with refractory MDS/AML.<sup>34</sup> Indeed, older patients were recently 331 reported to have OS of 42%, and a relapse rate of 24% at two years.<sup>35</sup> Strategies to 332 333 improve progression-free survival and decrease NRM using PT-Cy have been 334 implemented. The ALTERGREF trial (NCT03250546), currently in the inclusion 335 phase, will evaluate the effect of PT-Cy for prevention of GvHD in haploidentical and 336 HLA-9/10 mismatched unrelated donors transplant.

The proportion of patients with high risk cytogenetic has been previously reported to be 17 - 49%.<sup>13,14,23,24</sup> In our study, 68% (32 patients) had adverse cytogenetic

alterations, especially abnormalities in chromosomes 5 and 7. However, the resultswere probably not significant for the few patients in the other risk categories.

341 The best time to perform the transplant is unknown. It is possible that rapid 342 transplantation could reduce the risk of t-MDS/t-AML-related deterioration.<sup>25</sup> 343 Although a higher frequency of NRM is possible if the transplant is performed 344 beyond six months (50% vs 12.5%, p=0.03) caused by toxicity and infections due to multiple chemotherapy cycles<sup>23</sup>, we found no difference in OS or NRM. Although 345 346 the median age of 58 years is higher than for other large related series and 60% of 347 patients had a KPS  $\geq$  90, these factors did not play a significant role in survival, as in other series.<sup>12-14,20,23,25</sup> 348

349 Our data also do not support that factors related to the primary disease and its 350 treatment can influence survival. Interestingly, no veno-occlusive disease was 351 observed in this population having received a previous autologous transplantation.

A retrospective report of EBMT found better results in the period between 1998 and 2006 than for transplants performed before 1998 (40% vs 29%, p=0.02).<sup>13</sup> We found no differences between the periods of 2006-2010 and 2011-2016, probably because support care has not changed as much in the last 10 years as during the transition from the 80's to 90's.

357 Treatment options are still limited for patients who are not candidates for allo-HSCT.
358 Other therapeutic strategies have been tested using azacitidine or clofarabine
359 associated with chemotherapy with encouraging results.<sup>36,37</sup>

360 The limitations of our study include the collection of retrospective data and the 361 absence of a historical group for comparisons. Patient selection may have varied

between centers. The small number of patients and the low incidence of secondarymyelodysplasia makes it difficult to perform a prospective study.

364 In summary, although the number of patients in this study was small, the results 365 suggest that patients receiving an ASCT for a lymphoid neoplasm who develop t-MDS have short OS after allo-HSCT, with few long-time survivors. The use of 366 367 MMUD donors with standard GvHD prophylaxis should be avoided in such 368 indications for allo-HSCT. Studies that attempt to determine whether the 369 implementation of Cy post transplantation would improve these outcomes with 370 mismatched donors are still ongoing. It remains necessary to explore more 371 alternatives and transplant strategies in this critical population.

372

#### 373 AUTHOR CONTRIBUTIONS

EG and DJA designed the study, analyzed the data, and wrote the paper. All authors
provided patient data, approved the study design, critically reviewed the manuscript
and approved the final draft.

377

#### 378 ACKNOWLEDGEMENTS

The authors thank Nicole Raus and Réda Chebel from the SFGM-TC for data
collection and management, and all the data managers of the participating centers for
providing supplementary data.

382

#### 383 CONFLICT OF INTEREST

384 The authors declare no conflict of interest.

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Figure 1. Overall survival (OS), Non-relapse mortality (NRM) and Relapse incidence of all patients who underwent <u>allo-HSCT</u> for a t-MDS to ASCT for lymphoid neoplasms.



Figure 2. OS and NRM of patients with or without HLA mismatch donor (A-B). OS in regard to prior therapy of MDS before allo-HSCT (C). Relapse rate by type of MDS at diagnosis (D).

Patient characteristics	Overall population, n (%)
Number of patients	47 (100)
Gender	
Male	37 (78.7)
Female	10 (21.3)
Lymphoid Hematological malignancy	
Follicular lymphoma	12 (25.6)
Diffuse Large B-cell lymphoma	12 (25.6)
Hodgkin lymphoma	8 (17.0)
Mantle cell lymphoma	5 (10.6)
Burkitt lymphoma	2 (4.3)
Nodal marginal zone lymphoma	1 (2.1)
Small lymphocytic lymphoma	1 (2.1)
Lymphoplasmacytic lymphoma	1 (2.1)
Waldenstrom macroglobulinemia	1 (2.1)
Peripheral T-cell lymphoma	1 (2.1)
Angioimmunoblastic T-cell lymphoma	1 (2.1)
Unknown	2 (4.3)
Previous lines of therapy	
1	9 (19.1)
2	24 (51.1)
$\geq 3$	10 (21.3)
Unknown	4 (8.5)
Conditioning ASCT	
BEAM	38 (80.8)
Chemotherapy + TBI	7 (14.9)
Unknown	2 (4.3)
Response to ASCT	
Complete remission	37 (78.7)
Not in Complete remission	4 (8.5)
Unknown	6 (12.8)

#### Table 1. Patient characteristics at autologous stem cell transplantation

ASCT, autologous stem cell transplantation; BEAM, carmustine – etoposide – cytarabine – melphalan; TBI, total body irradiation.

# Table 2. Patient characteristics at allogeneic hematopoietic stem celltransplantation

Patient characteristics	Overall population $n(\%)$		
Number of patients	47 (100)		
Madian aga at allo HSCT, yaara (ranga)	$\frac{47(100)}{58(20-71)}$		
A co	38 (30 - 71)		
	22(48.0)		
$\geq$ 38 years	23(48.9)		
< 38 years	24 (31.1)		
Gender	27(79.7)		
Male	3/(/8./)		
	10 (21.3)		
Myelodysplastic Syndrome diagnosis	5 (10.0)		
MDS-SLD	5 (10.6)		
MDS-MLD	12 (25.6)		
MDS-RS-SLD	1 (2.1)		
MDS-RS-MLD	1 (2.1)		
MDS del(5q)	1 (2.1)		
MDS-EB-1	11 (23.5)		
MDS-EB-2	12 (25.6)		
MDS-U	2 (4.2)		
Unknown	2 (4.2)		
Interval from ASCT to diagnosis of t-MDS			
$\geq$ 74 months	36 (76.5)		
< 74 months	11 (23.5)		
Cytogenetics			
Good/Favorable prognosis	3 (6.4)		
Intermediate prognosis	6 (12.8)		
Poor/Unfavorable prognosis	32 (68.0)		
Unknown	6 (12.8)		
IPSS diagnosis			
Low risk / Intermediate-1	11 (23.4)		
Intermediate-2 / High risk	28 (59.6)		
Unknown	8 (17.0)		
Prior therapy of MDS before allo-HSCT			
AML-like induction treatment	11 (23.4)		
Hypomethylating agents	16 (34.1)		
ESA	4 (8.5)		
Immunosuppressants	1(2.1)		
Nothing	11 (23.4)		
Unknown	4 (8.5)		
Disease status prior to allo-HSCT			
Complete remission	10 (21.3)		
Not in Complete remission	36 (76.6)		
Unknown	1(2.1)		
Interval from diagnosis of MDS to allo-HSCT	. (		
< 6 months	17 (36 2)		
> 6 months	30 (63.8)		
	50 (05.0)		

Year of transplantation	
2006-2010	18 (38.3)
2011-2016	29 (61.7)
Conditioning regimen	
Myeloablative	9 (19.1)
Reduced intensity/non myeloablative	38 (80.9)
Conditioning regimen with TBI	
Yes	9 (19.1)
No	38 (80.9)
Graft type	
PBSC	41 (87.4)
BM	3 (6.3)
СВ	3 (6.3)
Type of donor	
HLA-identical sibling (10/10)	19 (40.5)
Matched unrelated (10/10)	17 (36.2)
Mismatched unrelated (9/10, 4/6, 5/6)	10 (21.2)
Unknown	1 (2.1)
ABO Match	
Major incompatibility	15 (32.0)
Minor incompatibility	8 (17.0)
Compatible	18 (38.3)
Unknown	6 (12.7)
Sex Match	
Male/Male	21 (44.6)
Male/Female	16 (34.0)
Female/Male	4 (8.6)
Female/Female	6 (12.8)
GvHD prophylaxis	
CsA - MTX	11 (23.4)
CsA - MMF	18 (38.3)
CsA	12 (25.5)
Other	6 (12.8)
CMV serostatus	
R-/D-	12 (25.5)
R-/D+	8 (17.7)
R+/D-	13 (27.1)
R+/D+	14 (29.7)
Karnofsky score	
90-100	29 (61.8)
< 90	16 (34.0)
Unknown	2 (4.2)

HSCT, hematopoietic stem cell transplantation; MDS SLD, myelodysplastic syndrome with single lineage dysplasia; MDS MLD, myelodysplastic syndrome with multilineage dysplasia; MDS-RS-SLD, myelodysplastic syndrome with ring sideroblasts and single lineage dysplasia; MDS-RS-MLD, myelodysplastic syndrome with ring sideroblasts and multilineage dysplasia; MDS del(5q), myelodysplastic syndrome with isolated 5q deletion; MDS-EB-1, myelodysplastic syndrome with

excess blasts-1; MDS-EB-2, myelodysplastic syndrome with excess blasts-2; MDS-U, myelodysplastic syndrome unclassifiable; ASCT, autologous stem cell transplantation; IPSS, international prognostic scoring system; AML, acute myeloid leukemia; ESA, erythropoietin stimulating agents; TBI, total body irradiation; PBSC, peripheral blood stem cell; BM, bone marrow; CB, cord blood; HLA, human leukocyte antigen; GvHD, graft versus host disease; CsA, cyclosporine; MTX, methotrexate; MMF, mycophenolate mofetil; CMV, cytomegalovirus.

Cause of death		CR	Not in CR	Unknown
Relapse/Progression	12 (44.4%)	4	8	
NRM causes	14 (51.9%)	11		3
Sepsis / MODS GvHD Pulmonary toxicity Hemorrhage Graft rejection VOD	8 3 1 2 0 0			
Unknown	1 (3.7%)	1		
Total deaths	27 (100 %)	16 (59.3%)	8 (29.6.%)	3 (11.1%)

#### Table 3. Causes of death according to the response at allo-HSCT.

HSCT, hematopoietic stem cell transplantation; NRM, non-relapse mortality; MODS, multiple organ dysfunction syndrome; GvHD, graft versus host disease; VOD, veno-occlusive disease

	Non-relapse mortality	Relapse	Overall survival
At 1 year	35% (18 – 51)	35% (18 - 53)	45% (29 - 60)
At 2 years	39% (21-56)	41% (22 - 61)	39% (24 – 55)
At 3 years	44% (25 - 63)	41% (22 - 61)	30% (15-45)

Table 4. Outcome of patients treated with allo-HSCT for t-MDS to ASCT for lymphoid neoplasms.

HSCT, hematopoietic stem cell transplantation; t-MDS, therapy-related myelodysplastic syndrome; ASCT, autologous stem cell transplantation. All values are represented as % (95% confidence interval, CI).

Table 5. Univariate analysis for probabilities of outcomes of non-relapse mortality, relapse, and overall survival at 3years after HSCT.

	Non-relapse Mortality		Relapse		Overall survival	
	(NRM)		(R)		(OS)	
All probabilities	Cumulative incidence	p-value	Cumulative incidence	p-value	Cumulative incidence	p-value
estimated at 3 years	(95% CI)		(95% CI)		(95% CI)	
Age at HSCT		0.23		0.21		0.69
$\geq$ 58 years	0.43 (0.20 – 0.66)		0.37(0.05 - 0.68)		0.33 (0.09 – 0.57)	
< 58 years	0.45 (0.18 – 0.73)		0.44 (0.20 – 0.68)		0.29(0.09 - 0.48)	
Gender		0.02		0.19		0.002
Male	0.38 (0.17 – 0.59)		0.38 (0.16 – 0.59)		0.37 (0.18 – 0.55)	
Female	0.74(0.34 - 1.14)		0.50(0.15 - 0.85)		0.10 (0 – 0.29)	
Hematological malignancy		0,24		0.65		0.98
NHL	0.50 (0.29 – 0.71)		0.53(0.22 - 0.85)		0.30 (0.13 – 0.47)	
HL	0.25 (0 – 0.68)		0.63 (0.08 – 1.17)		0.21 (0 – 0.56)	
Previous lines of therapy		0.75		0.40		0.34
1-2	0.52 (0.29 – 0.75)		0.41 (0.18 – 0.65)		0.27 (0.10 – 0.44)	
>2	0.37 (0.03 – 0.70)		0.11(0-0.31)		0.48 (0.11 – 0.84)	
Conditioning ASCT		0.45		0.44		0.85
BEAM	0.44 (0.23 – 0.66)		0.39 (0.17 – 0.60)		0.31 (0.14 – 0.48)	
Others	0.57 (0.13 – 1.01)		0.20(0-0.55)		0.34(0-0.73)	
Conditioning ASCT		0.34		0.76		0.91
TBI	0.65 (0.26 – 1.03)		0.44(0-0.93)		0.29 (0 – 0.63)	
Non-TBI	0.39 (0.19 – 0.59)		0.33 (0.13 – 0.52)		0.33 (0.15 – 0.51)	
Interval from ASCT to						0.27
diagnosis of MDS		0.52		0.80		
$\geq$ 74 months	0.41 (0.16 – 0.65)		0.55 (0.20 – 0.89)		0.20 (0 – 0.39)	
< 74 months	0.43 (0.17 – 0.68)		0.36 (0.12 – 0.59)		0.39 (0.17 – 0.60)	
MDS Diagnosis		0.49		0.008		0.08

MDS EB-1/EB-2	0.55 (0.24 - 0.87)		0.65 (0.37 - 0.93)		0.17 (0 – 0.34)	
Other MDS	0.40 (0.14 – 0.65)		0.10(0-0.23)		0.45(0.20 - 0.70)	
IPSS diagnosis		0.25		0.94		0.64
Lower	0.21 (0 – 0.47)		0.45(0.06 - 0.85)		0.34 (0.03 – 0.65)	
Higher	0.62 (0.37 – 0.87)		0.41 (0.11 – 0.70)		0.25 (0.06 – 0.44)	
Cytogenetics		0.40		0.27		0.40
Good prognosis	0(0-0)		0(0-0)		0.67	
Intermediate prognosis	0.5 (0.01 – 0.1)		0.6 (0.02 – 1.18)		0.20	
Poor prognosis	0.57 (0.31 – 0.83)		0.48 (0.21 – 0.75)		0.21	
Prior therapy for MDS		0.12		0.21		0.02
before HSCT						
AML-like therapy	0.1 (0 – 0.29)		0.38 (0.04 - 0.71)		0.56 (0.24 – 0.89)	
Hypomethylating agents	0.52 (0.21 – 0.83)		0(0-0)		0(0-0)	
No chemotherapy	0.51 (0.20 – 0.82)		0.42 (0.08 – 0.76)		0.35 (0.09 – 0.61)	
Year of transplantation		0.72		0.95		0.61
2006-2010	0.39 (0.14 – 0.63)		0.43 (0.15 – 0.70)		0.33 (0.12 – 0.55)	
2011-2016	0.52 (0.22 – 0.81)		0.41 (0.12 – 0.71)		0.28(0.07 - 0.49)	
Duration of MDS from		0.37		0.31		0.13
diagnosis to HSCT						
< 6 m	0.36 (0.06 – 0.65)		0.22(0-0.43)		0.47 (0.20 - 0.74)	
> 6 m	0.50 (0.25 – 0.75)		0.55 (0.28 – 0.81)		0.22 (0.05 – 0.39)	
Disease status prior to		0.67		0.64		0.70
HSCT						
Complete remission	0.25 (0 – 0.55)		0.61 (0.20 – 1.01)		0.25(0-0.55)	
Partial remission	0.53 (0.10 – 0.96)		0.30 (0.01 – 0.59)		0.25(0-0.53)	
Active/Progression	0.48 (0.23 – 0.74)		0.33 (0.07 – 0.58)		0.38 (0.15 – 0.60)	
MDS prior to HSCT		0.10		0.37		0.07
MDS EB-1/EB-2	0.64 (0.30 – 0.98)		0.44 (0.19 – 0.68)		0.14 (0 – 0.32)	
Other MDS	0.22(0-0.45)		0.30(0.03 - 0.56)		0.44(0.17-0.71)	

IPSS prior to HSCT		0.50		0.47		0.70
Lower	0.31 (0.01 – 0.60)		0.40(0.10-0.70)		0.27 (0.01 – 0.54)	
Higher	0.58 (0.25 - 0.91)		0.37(0-0.75)		0.34 (0.12 – 0.58)	
Conditioning regimen		0.49		0.45		0.54
Myeloablative (MAC)	0.61 (0.21 – 1.01)		0.58 (0.18 - 0.98)		0.22(0-0.49)	
Non-myeloablative/RIC	0.37 (0.18 – 0.56)		0.36 (0.15 – 0.57)		0.33 (0.15 – 0.50)	
Graft type		0.02		0.32		0.18
PBSC	0.31 (0.13 – 0.49)		0.37 (0.19 – 0.56)		0.36 (0.19 – 0.53)	
BM	0(0-0)		0(0-0)		0(0-0)	
СВ	0(0-0)		0(0-0)		0(0-0)	
GvHD prophylaxis		0.95		0.24		0.55
CsA - MTX	0.49(0.15 - 0.84)		0.35 (0.01 – 0.70)		0.40(0.10 - 0.71)	
CsA – MMF	0.47 (0.19 – 0.75)		0.49 (0.15 – 0.82)		0.18 (0 – 0.39)	
CsA	0.17(0-0.38)		0.58 (0.17 – 0.98)		0.25(0-0.54)	
Other	0.33 (0 – 0.71)		0(0-0)		0.67 (0.29 – 1.04)	
Type of donor		<0.001		0.90		0.58
HLA-identical sibling	0.30 (0.04 – 0.56)		0.46 (0.19 – 0.71)		0.35 (0.12 - 0.58)	
Matched unrelated	0.36 (0.05 – 0.66)		0.33 (0.04 - 0.62)		0.40 (0.13 – 0.66)	
Mismatched unrelated	0.83 (0.53 – 1.13)		0.25 (0 – 0.55)		0.12 (0 – 0.34)	
Mismatch HLA		0.001		0.86		0.004
No (10/10)	0.34 (0.13 – 0.54)		0.41 (0.21 – 0.61)		0.37 (0.19 – 0.54)	
Yes (No 10/10)	0.83 (0.53 – 1.13)		0.25 (0 – 0.55)		0.12 (0 – 0.34)	
ABO Match		0.58		0.72		0.59
Compatible	0.49 (0.19 – 0.78)		0.46 (0.11 – 0.81)		0.19 (0 – 0.41)	
Major incompatibility	0.59 (0.21 – 0.96)		0.66 (0.30 – 1.02)		0.16 (0 – 0.36)	
Minor incompatibility	0.33 (0 – 0.87)		0.29(0-0.62)		0.40(0-0.83)	
Sex Match		0.50		0.66		0.40
Yes	0.37 (0.13 – 0.61)		0.35 (0.11 – 0.58)		0.36 (0.14 – 0.58)	
No	0.15(0-0.44)		0.50(0.19-0.81)		0.25(0.05 - 0.45)	

CMV serostatus		0.31		0.16		0.07
R+/D-	0.60 (0.19 – 1.01)		0.64 (0.14 – 1.16)		0.12(0-0.34)	
Others	0.41 (0.20 – 0.62)		0.37 (0.16 – 0.57)		0.36 (0.18 – 0.53)	
Karnofsky score		0.50		0.65		0.92
90-100	0.47(0.24 - 0.69)		0.44(0.20-0.68)		0.30 (0.12 – 0.48)	
< 90	0.27(0-0.57)		0.35 (0.05 - 0.64)		0.34 (0.05 – 0.63)	
Acute GvHD		0.29		0.61		0.81
Yes	0.63 (0.34 – 0.93)		0.43 (0.08 – 0.79)		0.24(0.02 - 0.47)	
No	0.29(0.08 - 0.49)		0.42 (0.17 – 0.66)		0.35 (0.15 – 0.54)	

NRM, non-relapse mortality; R, relapse; OS, overall survival; HSCT, hematopoietic stem cell transplantation; NHL, non-hodgkin lymphoma; HL, hodgkin lymphoma; ASCT, autologous stem cell transplantation; BEAM, carmustine – etoposide – cytarabine – melphalan; TBI, total body irradiation; MDS, myelodysplastic syndrome; MDS-EB-1, myelodysplastic syndrome with excess blasts-1; MDS-EB-2, myelodysplastic syndrome with excess blasts-2; IPSS, international prognostic scoring system; AML, acute myeloid leukemia; MAC, myeloablative conditioning; RIC, reduced intensity conditioning; PBSC, peripheral blood stem cell; BM, bone marrow; CB, cord blood; GvHD, graft versus host disease; CsA, cyclosporine; MTX, methotrexate; MMF, mycophenolate mofetil; HLA, human leukocyte antigen; CMV, cytomegalovirus; R: receptor; D, donor; KPS, karnofsky performance status.

Table 6. Multivariate analysis of outcomes of non-relapse mortality, relapse, and overall survival after HSCT.

	Overall survival		Non-relapse mortality		Relapse	
	(OS)		(NRM)		(R)	
	HR (95% CI)	p-value	HR (95% CI)	p-value	HR (95% CI)	p-value
Prior therapy for MDS						
before HSCT						
AML-like induction	1.0	0.13				
Hypomethylating agents	3.55 (0.97 - 12.97)	0.06				
No chemotherapy	1.79 (0.52 - 6.15)	0.35				
HLA Mismatch		0.13		0.007		
No (10/10)	1.0		1.0			
Yes (9/10, 4/6, 5/6)	2.04 (0.80 - 5.22)		6.21 (1.63 – 23.62)			
MDS Diagnosis						0.21
Other MDS					1.0	
MDS EB-1/EB-2					2.63 (0.57 – 12.03)	

NRM, non-relapse mortality; R, relapse; OS, overall survival; MDS, myelodysplastic syndrome; HSCT, hematopoietic stem cell transplantation; AML, acute myeloid leukemia; HLA, human leukocyte antigen; MDS-EB-1, myelodysplastic syndrome with excess blasts-1; MDS-EB-2, myelodysplastic syndrome with excess blasts-2.