



# Mainly Post-Transplant Factors Are Associated with Invasive Aspergillosis after Allogeneic Stem Cell Transplantation: A Study from the Surveillance des Aspergilloses Invasives en France and Société Francophone de Greffe de Moelle et de Thérapie Cellulaire

Christine Robin, Catherine Cordonnier, Karine Sitbon, Nicole Raus, Olivier Lortholary, Sébastien Maury, Régis Peffault de La Tour, Stéphane Bretagne, Sylvie Bastuji-Garin

## ► To cite this version:

Christine Robin, Catherine Cordonnier, Karine Sitbon, Nicole Raus, Olivier Lortholary, et al.. Mainly Post-Transplant Factors Are Associated with Invasive Aspergillosis after Allogeneic Stem Cell Transplantation: A Study from the Surveillance des Aspergilloses Invasives en France and Société Francophone de Greffe de Moelle et de Thérapie Cellulaire. *Biology of Blood and Marrow Transplantation*, 2019, 25 (2), pp.354-361. 10.1016/j.bbmt.2018.09.028 . pasteur-02191094

**HAL Id: pasteur-02191094**

**<https://pasteur.hal.science/pasteur-02191094>**

Submitted on 22 Oct 2021

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



Distributed under a Creative Commons Attribution - NonCommercial 4.0 International License

Mainly post-transplant factors are associated with invasive aspergillosis after allogeneic stem cell transplantation. A study from the SAIF and SFGM-TC

Christine Robin<sup>1,2</sup>, Catherine Cordonnier<sup>1,2</sup>, Karine Sitbon<sup>3</sup>, Nicole Raus<sup>4</sup>, Olivier Lortholary<sup>3,5,6</sup>, Sébastien Maury<sup>1,2,7</sup>, Régis Peffault de la Tour<sup>8,9</sup>, Stéphane Bretagne\*<sup>3,8,10</sup>, and Sylvie Bastuji-Garin\*<sup>11,12</sup>, for the SAIF (Surveillance des Aspergilloses Invasives en France) and the SFGM-TC (Société Francophone de Greffe de Moelle et de Thérapie Cellulaire) Aspergillus Group

*\*SB and SBG contributed equally to this work*

1 Assistance Publique-Hôpitaux de Paris (APHP), Henri Mondor hospital, Hematology Department, Créteil,

2 Paris-Est Créteil University (UPEC), Créteil,

3 Institut Pasteur, CNRS, molecular Mycology unit, French National Reference Center for Invasive Mycoses & antifungals, UMR2000, Paris,

4 Société Française de Greffe de Moelle et de Thérapie cellulaire, Saint-Denis,

5 APHP, Necker-Enfant malades hospital, Centre d'Infectiologie Necker Pasteur, and IHU Imagine, Paris,

6 Paris Descartes University, Paris,

7 INSERM U955 Henri Mondor hospital, Créteil,

8 Paris-Diderot, Sorbonne Paris Cité University, Paris,

9 AP-HP, Lariboisière Saint-Louis Fernand Widal hospital, Stem Cell Transplant Unit, Paris,

10 AP-HP, Lariboisière Saint-Louis Fernand Widal hospital, Parasitology-Mycology Laboratory, Paris.

11 APHP, Henri Mondor hospital, Public Health Department, Créteil,

12 UPEC, IMRB, CEpiA (Clinical Epidemiology and Ageing Unit, EA EA7376), Créteil, France.

**Key words:** allogeneic stem cell transplantation, aspergillosis, invasive fungal infection

**Running heads:** Invasive aspergillosis after allogeneic HSCT

**Corresponding author:** Christine Robin, MD  
Department of Hematology  
Henri Mondor University Hospital  
94000 Créteil, France  
Tel: +33 149 812 057 / Fax: +33 149 812 067  
christine.robin@aphp.fr

**Abstract word count:** 250

**Text word count:** 3506

**Number of references:** 37

**Number of tables:** 6

**Number of figures:** 1

This study has been presented at the EBMT Annual Meeting on March 19, 2018, Lisbon (poster no. B121)

## ABSTRACT

Invasive aspergillosis (IA) occurs in up to 23% of allogeneic hematopoietic stem cell transplantation (HSCT) patients. While transplant procedures have changed over time, more late cases of IA are being observed. The objective of this study was to identify the pre- and post-transplant factors of IA in a large cohort of HSCT patients mainly transplanted with reduced-intensity conditioning.

This multicenter, case-control study was carried out using data collected between 2005–2010 by the Surveillance des Aspergilloses Invasives en France (SAIF) program (Institut Pasteur, Paris) and the EBMT ProMISe registry. Four controls without IA were individually matched to each case based on the center, patient's age, and year of the transplant.

We identified 185 cases of probable and proven IA and 651 controls. The median date of IA after the transplant was 133 days, with 35 (19%) cases of early IA (before day 40), 33 (18%) cases of late IA (day 40 to day 100), and 117 (63%) cases of very late IA (after day 100). In the multivariate analysis, early IA was significantly associated with a lack of engraftment, while late and very late IA were significantly associated with > grade 2 acute graft-versus-host disease (GvHD); very late IA was also significantly associated with relapse and secondary neutropenia.

Two thirds of IA cases occurred more than 100 days after HSCT with different risk factors than those occurring earlier. Prophylactic strategies should consider the specific risk factors for late and very late IA, especially GvHD, relapse after transplant, and secondary neutropenia.

## Introduction

Invasive aspergillosis (IA) is a common complication after allogeneic hematopoietic stem cell transplantation (HSCT); it occurs in 2.7–23% of HSCT recipients.<sup>1-4</sup> IA is the most common invasive fungal infection (IFI), representing about two thirds of IFI cases after allogeneic HSCT.<sup>2, 5</sup> Despite improvements in antifungal prophylaxis and treatment, IA remains a life-threatening complication in allogeneic HSCT recipients and has a mortality rate of 20–75%.<sup>2, 6-8</sup>

Several risk factors of IA after allogeneic HSCT have been identified.<sup>9, 10</sup> Pre-transplant factors that increase the risk of IA include age, bone marrow transplant rather than peripheral blood stem cells (PBSC), transplantation of cord blood units (CBU), an active hematological malignancy, administration of antithymoglobulin (ATG), and a pre-transplant iron overload.<sup>7, 9-12</sup> Geoclimatic conditions, such as temperature and precipitation, correlate with spore counts in the air and may also influence the incidence of IA.<sup>13</sup> Post-transplant factors include cytomegalovirus (CMV) infection and disease, delayed neutrophil and lymphocyte engraftment, secondary neutropenia, relapse of the underlying disease, administration of corticosteroids, and graft-versus-host disease (GvHD).<sup>8-11, 14-21</sup>

Two decades ago, IA was mainly considered to be an early complication of HSCT. However, more late cases of IA are being reported and, thus, the role of the risk factors may vary according to the timing of the onset of IA.<sup>9, 10</sup> In addition, transplant procedures have greatly evolved over the last 15 years. While more than 80% of the patients of previous large series of IA were transplanted using myeloablative regimens from HLA-identical donors,<sup>10, 11</sup> there has been a switch to reduced-intensity conditioning (RIC) and non-myeloablative (NMA) regimes and more alternative stem cell sources.<sup>22-24</sup>

The goal of this study was to determine if the risk factors for IA have changed with the evolution of conditioning. We identified the pre- and post-transplant risk factors that were independently associated with IA in a large cohort of allogeneic HSCT recipients transplanted in France. Of the cohort, 95% of the recipients were transplanted using RIC or NMA conditioning. We then assessed

whether the impact of the factors varied according to the timing of the onset of IA after the transplant.

## **Patients and Methods**

### **Design**

This multicenter, case-control study used the data from two prospective programs: the Surveillance des Aspergilloses Invasives en France (SAIF) program from the Institut Pasteur in Paris, France, and the European ProMISe registry. SAIF anonymously collected consecutive cases of IA diagnosed in university hospitals from three regions of France (Paris-Ile de France, Grand Ouest, and Rhône-Alpes) between 2005 and 2010 through a web-based file.<sup>8</sup> The ProMISe registry prospectively collects data on all consecutive HSCTs within the European Group for Blood and Marrow Transplantation (EBMT). These two databases are complementary: the SAIF database contains mainly clinical, imaging, and microbiological data focused on the definition criteria of IA and the ProMISe database mainly contains transplant-related data. This study was registered with the Commission Nationale de l'Informatique et des Libertés (CNIL) (number 1882064 v 0).

### **Cases of IA**

Consecutive cases of IA were selected from the SAIF registry from January 1, 2005 to December 31, 2010. The IA cases were prospectively recorded from the laboratory of each participating center; the local microbiologist completed a standardized questionnaire about the demographics, underlying conditions, diagnostic tools used (imaging, direct examination and mycological culture, histology, and serum or broncho-alveolar galactomannan), dates of hospitalization, and first-line antifungal therapy. A local committee involving the mycologists and the hematologists in charge of the patient reviewed every recorded questionnaire, checked the diagnosis, and classified each episode according to the European Organization for Research and Treatment of Cancer (EORTC)/Mycoses Study Group (MSG)

2002 criteria<sup>25</sup> for patients included before 2008 and the EORTC/MSG 2008 criteria<sup>26</sup> for patients included from 2008 to 2010. The date of IA was set as the date the criteria were fulfilled. Patients were managed in HSCT centers according to the local practices for the diagnosis and treatment of IA. The timing of IA after transplant was classified as follows: *early* IA when diagnosed before day 40, *late* IA when diagnosed between day 40 and day 100, and *very late* IA when diagnosed after day 100.

9

All proven and probable cases of IA diagnosed in patients aged 15 years or over after an allogeneic HSCT were selected from the SAIF database. Only the first episodes of proven or probable IA after HSCT were considered. Patients with previous IA before transplantation were not excluded from the study. Of the 20 French HSCT centers that participated in the SAIF registry, 19 participated in the present study; 3 881 allogeneic HSCT were performed during the period from 2005 to 2010.

### **Control patients**

Our objective was to individually match 4 control patients to each IA case in terms of HSCT center, age of patient ( $\pm 5$  years), and year of transplant. Control patients were recruited from the ProMISe registry among patients who did not develop IA after HSCT.

Most patients (IA cases and controls) were housed in laminar air flow rooms at the initial phase of transplant, except those who received NMA conditioning with fludarabine and 2 Gray total body irradiation (TBI).

### **Data collection and definitions**

The pre- and post-transplant clinical and biological data of the case and control patients were collected from the ProMISe registry. The EMBT definitions were used for the terms. For example, the day of engraftment was defined as the first day of 3 consecutive days  $> 0.5 \times 10^9/L$ . Acute GvHD was reported according to the staging by Glücksberg et al.<sup>27</sup> Secondary neutropenia was defined as the occurrence of a neutrophil count below 0.5 G/L after engraftment. According to the EBMT

recommendations, all patients signed a consent form before their transplant for their data to be entered into the ProMISe registry and used for retrospective studies.

### Statistical analysis

For the present study, the alpha risk was set at 0.05. Thus, 175 cases were needed to provide 80% power for detecting an odds ratio (OR) > 1.65 related to the factors present in 35% of the control population and an OR > 5.21 related to the factors present in 1% of the control population. The factors present in 1% of the population corresponded to the prevalence of secondary neutropenia, which was a less frequent, but well known, risk factor.

Table 1 displays the pre- and post-transplant factors of the cases and controls with IA. The qualitative variables were given as numbers (%) and compared using the chi-squared test or the Fisher exact test. The quantitative variables were given as the median (interquartile range: Q1–Q3) and compared using the Kruskal-Wallis test.

Data were analyzed using the standard methods for estimating ORs and 95% confidence intervals (Cis) in case-control studies. For the univariate analyses, the adjusted ORs (aOR) were estimated separately for each potential risk factor yielding a *P* value < 0.20 using unconditional logistic regression models forcing the matching variables, such as HSCT center, age of patient, and year of transplant, into all models. The quantitative variables were converted into categorical variables using a cut-off according to the literature when available or a median split. The whole population was studied for pre-transplant factors. However, only the controls and their corresponding cases were studied for post-transplant factors; the controls were followed for the same duration as their cases. For the post-transplant risk factors, we only analyzed the events that occurred before IA. As aspergillosis prophylaxis with posaconazole has been recommended to be administered to patients with GvHD since 2008,<sup>28</sup> ORs were also adjusted for the periods 2005–2007 and 2008–2010.

Variables yielding *P* values < 0.10 in the univariate analyses were considered for the multivariate analyses. First-order interactions and confounding factors were investigated using multiple two-by-



two analyses. In the first multivariate logistic regression model, we considered pre-transplant factors and added post-transplant variables. The discrimination of the model was assessed using the C-index (area under the curve (AUC) of receiver operating characteristics (ROC), while the Hosmer–Lemeshow calibration test was used to assess the goodness of fit.

Finally, we conducted similar analyses using a multinomial logistic regression model to compare early IA, late IA, and very late IA cases with the control population. Due to the small number of patients in some groups, we pooled some modalities of the following variables: disease status, stem cell source, and donor (D)/recipient (R) CMV serology.

All tests were two-tailed and a  $P$  value  $\leq 0.05$  was considered statistically significant.  $P$  values for comparisons between each group of IA (early, late and very late) and the control group were corrected using the false discovery (FDR) method for multiple comparisons<sup>29</sup>

Data were analyzed using Stata Statistical Software 12.0 (Stata Corp., College Station, TX, USA). This observational study was reported according to the Strengthening the Reporting of Observational Studies in Epidemiology statement.<sup>30</sup>

## Results

This study included 185 IA cases: 166 (90%) probable IA cases, and 19 (10%) proven IA cases. The time from transplant to IA ranged from 3 days to 4 308 days with a median of 133 days (64–233 days). Thirty-five (19%) cases were classified as early IA, 33 (18%) cases were late IA, and 117 (63%) cases were very late IA. The median number of cases per HSCT center was 7 (ranged from 1 to 34).

We were able to identify 651 controls (Figure 1). A total of 135 cases had 4 matched controls, 24 cases had 3 controls, 15 cases had 2 controls, 9 cases had 1 control, and 2 cases had no controls. For the post-transplant factor analysis, 526 controls had a follow-up that matched the timing of IA in their corresponding case patient. The median follow-up was 1904 days (1448–2505 days). The control patients were well matched with the case patients for age ( $P = 0.85$ ), HSCT center ( $P = 1.00$ ), and year of transplant ( $P = 1.00$ ).

The characteristics of the patients are summarized in Table 1. Briefly, the median age at transplant was 49.6 years. More than two thirds of the patients had acute leukemia or myelodysplastic syndrome. Two thirds of the patients were in complete remission at transplant. Only 4.4% of the patients received a myeloablative (MA) regimen; all others received RIC or NMA conditioning. PBSCs were the main stem cell source. More than half of the patients were transplanted from an unrelated donor.

The median overall survival was 0.70 years (Q1-Q3: 0.60–0.90 years) for the cases and 4.94 years (3.61–7.06 years) for the controls ( $P < 0.001$ ). There was no difference in survival when the period before the introduction of posaconazole prophylaxis (2005–2007) was compared with the period after its introduction (2008–2010).

### **Pre- and post-transplant factors associated with the risk of IA**

#### *Univariate analysis*

The univariate analysis of the pre-transplant factors adjusted for matching variables (Table 1) showed that IA was significantly associated with TBI, D and R CMV serology (D-/R+ versus others), and the absence of the use of ATG-based conditioning. A trend for an association with IA was observed for the use of clofarabine.

Among the post-transplant factors (Table 1), IA occurrence was significantly associated with grade 3-4 acute GvHD, a relapse after transplant, and secondary neutropenia. An adjustment for the period before 2008 did not affect the association between acute GvHD and IA (aOR: 2.65 [1.68-4.17];  $P < 0.001$ ). The time between acute GvHD and IA was 113 days (56-259).

#### *Multivariate analysis*

The multivariate analysis (Table 2) showed that TBI, clofarabine use, and D-/R+ CMV serology were independently associated with IA. When the pre- and post-transplant factors were taken into account, the analysis showed that TBI, grade  $\geq 2$  acute GvHD, and relapse after transplant were independently associated with IA. A trend for an association was observed for secondary

neutropenia. The D-/R+ CMV serology and clofarabine use were not significant because of an inverse association with GvHD and relapse after transplant, respectively. The final model had a good calibration ( $P$  value of the goodness-of-fit test  $> 0.20$ ) and a moderate discrimination (AUC under ROC curve was 0.77 [0.73-0.82]).

### **Pre- and post-transplant factors associated with the risk of early, late, or very late IA**

#### *Univariate analysis*

The univariate analysis (Table 3) showed that the occurrence of early IA was significantly associated with the use of CBU as the stem cell source, clofarabine use, absence of acute GvHD, lack of engraftment, and relapse after transplant. A trend for a high risk of early IA was observed for the absence of complete remission at transplant, an unrelated donor, transplantation during summer and the absence of CMV infection. The occurrence of late IA was associated with, grade 3-4 acute GvHD regardless of the period (before or after 2008; aOR: 3.88 [1.58-9.57];  $P = 0.003$ ), and relapse after transplant. A trend for a high risk of late IA was observed for the absence of the use of ATG-based conditioning. The time between acute GvHD and late IA was 39 days (22-57). The occurrence of very late IA was associated with D-/R+ CMV serology,  $\geq$  grade 2 acute GvHD regardless of the period (before or after 2008; aOR: 1.66 [0.98-2.79];  $P = 0.057$  for grade 2 GvHD and aOR: 4.12 [2.21-7.68];  $P < 0.001$  for grade 3-4 GvHD), relapse after transplant, and secondary neutropenia. A trend for a high risk of very late IA was observed for the absence of the use of ATG-based conditioning and TBI. The time between acute GvHD and very late IA was 158 days (98-401).

#### *Multivariate analysis*

The multivariate analysis showed that, among the pre-transplant factors, only the absence of a complete remission at transplant and an unrelated donor tended to be associated with early IA (Table 4). Clofarabine use was not associated with early IA in the multivariate analysis because of its association with the absence of a complete remission. When the pre- and post-transplant factors were taken into account, the analysis showed that only the absence of engraftment remained

significantly associated with early IA. The absence of a complete remission tended to be associated with early IA. The final model had a good calibration ( $P$  value of the goodness-of-fit test= 0.32) and a moderate discrimination (AUC under ROC curve was 0.72 [0.60-0.84]).

The multivariate analysis (Table 4) showed that, among the pre-transplant factors, only the absence of ATG-based conditioning tended to be associated with late IA. When the pre- and post-transplant factors were taken into account, only the grade 3-4 acute GvHD remained associated with late IA. There was a trend for relapse after transplant to be associated with late IA. The final model had good calibration ( $P$  value of the goodness-of-fit test 0.33) and a moderate discrimination (AUC under ROC curve was 0.69 [0.58-0.80]).

Among the pre-transplant factors, the multivariate analysis (Table 4) showed that D-/R+ CMV serology was associated with very late IA. When the pre- and post-transplant factors were taken into consideration, the analysis showed that grade  $\geq 2$  acute GvHD, relapse after transplant, and secondary neutropenia were independently associated with very late IA. CMV infections were not associated with very late IA in the multivariate analysis because of their association with GvHD. The final model had good calibration ( $P$  value of the goodness-of-fit test 0.54) and discrimination (AUC under ROC curve was 0.80 [0.74-0.85]).

## Discussion

Our study showed that the changes in allogeneic HSCT procedures over time did not change the impact that GvHD has on the onset of IA except for early IA cases, which occur before day 40. It also showed that currently, 63% of IA cases occur after day 100. These findings were drawn from robust data from two prospective registries and the use of a 1:4 ratio of case patients to control patients from the HSCT centers of 19 university hospitals over 6 years. This is the largest study on the occurrence of IA after allogeneic HSCT in Europe.

Our study also showed that the risk of IA after allogeneic HSCT cannot be reliably predicted from pre-transplant characteristics and, thus, should be regularly assessed after transplant. As previously

reported from a cohort with a majority of MA-conditioned patients,<sup>9</sup> the risk factors of IA in our study varied according to the timing of the onset of IA. The factors intervened after transplant; for example, the occurrence of early IA was associated with the absence of engraftment, late IA was associated with grade 3-4 acute GvHD, and very late IA with  $\geq$  grade 2 acute GvHD, post-transplant relapse, and secondary neutropenia. Contrary to previous studies,<sup>1, 9, 10</sup> we found that the pre-transplant factors, such as sex, donor type, stem cell source, and underlying disease, were not associated with the occurrence of IA. As age was one of the matching criteria for choosing the controls, it was not studied as a risk factor. In our study, the pre-transplant factors that showed a trend for an association with early IA were supplanted by the lack of engraftment. The pre-transplant factors had less impact on early IA, late IA, and very late IA when the post-transplant factors were added to the model, despite a trend for an association between absence of complete remission and early IA.<sup>9, 10</sup>

Two thirds of the cases of IA occurred after day 100 independently of the type of conditioning, which confirmed the increasing trend in delayed onset of IA over the last decades.<sup>9, 31</sup> This result suggests that it may be possible to reduce the risk of IA during the first months after transplant, but that the risk is underestimated later on. The later occurrence of IA raises major concerns for the transplant community. First, because the clinical and imaging presentation may be different, the diagnosis of IA may be more difficult several months after transplant than during neutropenia.<sup>32</sup> The serum galactomannan may be less often positive in these patients than in neutropenic patients<sup>33</sup> and, thus, the requirement for bronchoalveolar lavage (BAL) may increase. However, neutropenic patients and patients with GvHD share the same consensus diagnostic criteria<sup>26</sup>. Moreover, the presence of several types of mold in the BAL fluid may be more difficult to interpret in highly immunodepressed patients suffering from severe chronic GvHD than in standard patients.<sup>34</sup> Thus, the current definitions of probable IA, which have been mainly tailored for neutropenic patients, should be refined for patients with GvHD. Second, the late occurrence of IA should prolong the duration of antifungal prophylaxis for months or even years, which raises potential concerns about resistance,

prolonged toxicity, and drug interferences. Studies should now focus on diagnostic criteria of IA in this very specific population and also on prophylaxis approach for very late IA.

Finally, despite the progress in the management of fungal infections over the last two decades, IA remains associated with a very poor outcome in cases when compared with controls. It can be hypothesized that IA acts both as a deadly infection and as a marker of deep immunosuppression associated with GvHD and other infectious complications. It should be noticed that in our study, the poor outcome was not modified by the introduction of posaconazole in 2008.

We acknowledge the following limitations in this study. First, due to its design, some data were not available, including ferritin levels, neutrophil, lymphocyte, or monocyte counts, and post-transplant factors, such as chronic GvHD and steroid use. However, all IA data were prospectively collected in the HSCT centers and the transplant data were exhaustively registered in the ProMISe registry, which ensured the consecutive patterns of the collected data available for study. Second, we were unable to obtain individual data about antimold prophylaxis. However, posaconazole prophylaxis had become routine in most French HSCT centers for patients with GvHD in the second semester of 2007 following the publication of the study by Ullmann et al.<sup>28</sup> The ORs adjusted for the period before or after 2008 for IA and the overall survival were unchanged according to this study, and GvHD remained the main risk factor regardless of the time period. Third, we did not explore the genetic risk factors that have been shown to increase the risk of fungal infection after HSCT.<sup>35-37</sup> Other main post-transplant risk factors, such as relapse of the underlying disease or secondary neutropenia, should be taken into account for prophylaxis indication and be considered in future prospective studies. Last but not least, we did not have any correction for multiple comparisons as it would have eliminated most of the risk factors identified.

In conclusion, in this comparative study, we found that in a population of patients mainly conditioned with NMA or RIC, GvHD remained the main risk factor for IA. Among pre-transplant factors, absence of complete remission tended to be associated with early IA. Clearly, IA cannot be reliably predicted before HSCT. Post-transplant factors have a major impact on IA, especially the lack

of engraftment for early IA cases, and acute GvHD, relapse, and secondary neutropenia for late and very late cases of IA. Although the presence of GvHD in transplant centers should indicate the need to start antimold prophylaxis, relapse, and secondary neutropenia that is often consecutive to the use of ganciclovir should also be considered as major indicators of prophylaxis and the targets of future studies.

### **Conflicts of interest**

The authors certify that there are no potential conflicts of interest.

### **Author contributions**

CR, CC, SB, and SBG conceived and designed the study. SB, OL, and KB generated and provided the SAIF data. NR, RPDLT, and SM provided the ProMISe registry data. CR provided additional material on patients, assembled the data, and ran the analysis. CR, SBG, SB and CC analyzed and interpreted the data and drafted the manuscript. All authors approved the final version.

**Funding source:** The SAIF program was supported by a specific grant from Institut de Veille Sanitaire, 94410, Saint-Maurice, France.

### **Acknowledgments**

The authors are grateful to the microbiology and clinical staff from the French Mycosis Study Group who actively participated in the SAIF network. They are also grateful to the clinicians and data managers of the Société Française de Greffe de Moelle et de Thérapie Cellulaire (SFGM-TC) centers who provided data for this study.

### **Investigators**

Daniel Ajzenberg (Centre hospitalo-universitaire (CHU) de Limoges), Marie-Thérèse Baixench (CHU Cochin, Paris), Marc Bernard (CHU, Rennes), Anne-Lise Bienvenu (CHU, Rouen), Karine Bilger (CHU, Strasbourg), Marie-Elisabeth Bougnoux (CHU Necker-Enfants malades, Paris), Jean-Henri Bourhis (Institut Gustave Roussy, Villejuif), Patrice Ceballos (CHU, Montpellier), Jacques Chandenier (CHU, Tours), Patrice Chevallier (CHU, Nantes), Eric Dannaoui (Hopital Européen Georges Pompidou, Paris), Etienne Daguindau (CHU, Besançon), Gandhi Damaj (CHU, Caen), Marguerite Fines (CHU, Caen), Jean-Pierre Gangneux (CHU, Rennes), Martine Gari-Toussaint (CHU, Nice), Gaelle Guillermin (CHU, Brest),



Françoise Gay Andrieu (CHU, Nantes), Norbert Ifrah (CHU, Angers), Catherine Kauffman (CHU, Poitiers), Claire Lacroix (CHU Lariboisière-Saint-Louis-Fernand Widal, Paris), Bernadette Lebeau (CHU, Grenoble), Valerie Letscher (CHU, Strasbourg), Marie Machouart (CHU, Nancy), Natacha Maillard (CHU, Poitiers), Mauricette Michallet (CHU Edouard Herriot, Lyon), Laurence Million (CHU, Besançon), Noël Milpied (CHU, Bordeaux-Pessac), Frédérique de Monbrison (CHU Edouard Herriot, Lyon), Baptiste Pérard (CHU, Bordeaux-Pessac), Jean-Louis Poirot (CHU Saint-Antoine, Paris), Dorothée Quinio (CHU, Brest), Stéphane Ranque (CHU, Aix-Marseille), Pierre-Simon Rohrllich (CHU L'Archet, Nice), Marie-Thérèse Rubio (CHU, Nancy), Yvon Sterkers (CHU, Montpellier), Gérard Socié (CHU Lariboisière-Saint-Louis-Fernand Widal, Paris), Felipe Suarez (CHU Necker-Enfants malades, Paris), Anne Thiebault-Bertrand (CHU, Grenoble) and Pascal Turlure (CHU, Limoges).

## REFERENCES

1. Atalla A, Garnica M, Maiolino A, Nucci M. Risk factors for invasive mold diseases in allogeneic hematopoietic cell transplant recipients. *Transpl Infect Dis.* 2015;17(1):7-13.
2. Harrison N, Mitterbauer M, Tobudic S, et al. Incidence and characteristics of invasive fungal diseases in allogeneic hematopoietic stem cell transplant recipients: a retrospective cohort study. *BMC Infect Dis.* 2015;15:584-92.
3. Herbrecht R, Bories P, Moulin JC, Ledoux MP, Letscher-Bru V. Risk stratification for invasive aspergillosis in immunocompromised patients. *Ann N Y Acad Sci.* 2012;1272:23-30.
4. Shi JM, Pei XY, Luo Y, et al. Invasive fungal infection in allogeneic hematopoietic stem cell transplant recipients: single center experiences of 12 years. *J Zhejiang Univ Sci B.* 2015;16(9):796-804.
5. Neofytos D, Horn D, Anaissie E, et al. Epidemiology and outcome of invasive fungal infection in adult hematopoietic stem cell transplant recipients: analysis of Multicenter Prospective Antifungal Therapy (PATH) Alliance registry. *Clin Infect Dis.* 2009;48(3):265-73.
6. Baddley JW. Clinical risk factors for invasive aspergillosis. *Med Mycol.* 2011;49 Suppl 1:S7-S12.
7. Kontoyiannis DP, Marr KA, Park BJ, et al. Prospective surveillance for invasive fungal infections in hematopoietic stem cell transplant recipients, 2001-2006: overview of the Transplant-Associated Infection Surveillance Network (TRANSNET) Database. *Clin Infect Dis.* 2010;50(8):1091-100.
8. Lortholary O, Gangneux JP, Sitbon K, et al. Epidemiological trends in invasive aspergillosis in France: the SAIF network (2005-2007). *Clin Microbiol Infect.* 2011;17(12):1882-9.
9. Garcia-Vidal C, Upton A, Kirby KA, Marr KA. Epidemiology of invasive mold infections in allogeneic stem cell transplant recipients: biological risk factors for infection according to time after transplantation. *Clin Infect Dis.* 2008;47(8):1041-50.
10. Marr KA, Carter RA, Boeckh M, Martin P, Corey L. Invasive aspergillosis in allogeneic stem cell transplant recipients: changes in epidemiology and risk factors. *Blood.* 2002;100(13):4358-66.
11. Kojima R, Kami M, Nannya Y, et al. Incidence of invasive aspergillosis after allogeneic hematopoietic stem cell transplantation with a reduced-intensity regimen compared with transplantation with a conventional regimen. *Biol Blood Marrow Transplant.* 2004;10(9):645-52.
12. Kontoyiannis DP, Chamilos G, Lewis RE, et al. Increased bone marrow iron stores is an independent risk factor for invasive aspergillosis in patients with high-risk hematologic malignancies and recipients of allogeneic hematopoietic stem cell transplantation. *Cancer.* 2007;110(6):1303-6.
13. Panackal AA, Li H, Kontoyiannis DP, et al. Geoclimatic influences on invasive aspergillosis after hematopoietic stem cell transplantation. *Clin Infect Dis.* 2010;50(12):1588-97.
14. Fukuda T, Boeckh M, Carter RA, et al. Risks and outcomes of invasive fungal infections in recipients of allogeneic hematopoietic stem cell transplants after nonmyeloablative conditioning. *Blood.* 2003;102(3):827-33.
15. Hol JA, Wolfs TF, Bierings MB, et al. Predictors of invasive fungal infection in pediatric allogeneic hematopoietic SCT recipients. *Bone Marrow Transplant.* 2014;49(1):95-101.
16. Junghanss C, Marr KA, Carter RA, et al. Incidence and outcome of bacterial and fungal infections following nonmyeloablative compared with myeloablative allogeneic hematopoietic stem cell transplantation: a matched control study. *Biol Blood Marrow Transplant.* 2002;8(9):512-20.
17. Labbe AC, Su SH, Laverdiere M, et al. High incidence of invasive aspergillosis associated with intestinal graft-versus-host disease following nonmyeloablative transplantation. *Biol Blood Marrow Transplant.* 2007;13(10):1192-200.
18. Li L, Wang J, Zhang W, et al. Risk factors for invasive mold infections following allogeneic hematopoietic stem cell transplantation: a single center study of 190 recipients. *Scand J Infect Dis.* 2012;44(2):100-7.

19. Mihu CN, King E, Yossepovitch O, et al. Risk factors and attributable mortality of late aspergillosis after T-cell depleted hematopoietic stem cell transplantation. *Transpl Infect Dis.* 2008;10(3):162-7.
20. Thursky K, Byrnes G, Grigg A, Szer J, Slavin M. Risk factors for post-engraftment invasive aspergillosis in allogeneic stem cell transplantation. *Bone Marrow Transplant.* 2004;34(2):115-21.
21. Zhang P, Jiang EL, Yang DL, et al. Risk factors and prognosis of invasive fungal infections in allogeneic stem cell transplantation recipients: a single-institution experience. *Transpl Infect Dis.* 2010;12(4):316-21.
22. Baron F, Ruggeri A, Beohou E, et al. RIC versus MAC UCBT in adults with AML: A report from Eurocord, the ALWP and the CTIWP of the EBMT. *Oncotarget.* 2016;7(28):43027-38.
23. Passweg JR, Baldomero H, Bregni M, et al. Hematopoietic SCT in Europe: data and trends in 2011. *Bone Marrow Transplant.* 2013;48(9):1161-7.
24. Savani BN, Labopin M, Kroger N, et al. Expanding transplant options to patients over 50 years. Improved outcome after reduced intensity conditioning mismatched-unrelated donor transplantation for patients with acute myeloid leukemia: a report from the Acute Leukemia Working Party of the EBMT. *Haematologica.* 2016;101(6):773-80.
25. Ascioglu S, Rex JH, de Pauw B, et al. Defining opportunistic invasive fungal infections in immunocompromised patients with cancer and hematopoietic stem cell transplants: an international consensus. *Clin Infect Dis.* 2002;34(1):7-14.
26. De Pauw B, Walsh TJ, Donnelly JP, et al. Revised definitions of invasive fungal disease from the European Organization for Research and Treatment of Cancer/Invasive Fungal Infections Cooperative Group and the National Institute of Allergy and Infectious Diseases Mycoses Study Group (EORTC/MSG) Consensus Group. *Clin Infect Dis.* 2008;46(12):1813-21.
27. Glucksberg H, Storb R, Fefer A, et al. Clinical manifestations of graft-versus-host disease in human recipients of marrow from HL-A-matched sibling donors. *Transplantation.* 1974;18(4):295-304.
28. Ullmann AJ, Lipton JH, Vesole DH, et al. Posaconazole or fluconazole for prophylaxis in severe graft-versus-host disease. *N Engl J Med.* 2007;356(4):335-47.
29. Benjamini Y HY. Controlling the false discovery rate. a practical and powerful approach to multiple testing. *Statist Soc B (Methodological).* 1995;57:289–300.
30. von Elm E, Altman DG, Egger M, et al. [The Strengthening the Reporting of Observational Studies in Epidemiology (STROBE) statement: guidelines for reporting of observational studies]. *Internist (Berl).* 2008;49(6):688-93.
31. Grow WB, Moreb JS, Roque D, et al. Late onset of invasive aspergillus infection in bone marrow transplant patients at a university hospital. *Bone Marrow Transplant.* 2002;29(1):15-9.
32. Bergeron A, Porcher R, Sulahian A, et al. The strategy for the diagnosis of invasive pulmonary aspergillosis should depend on both the underlying condition and the leukocyte count of patients with hematologic malignancies. *Blood.* 2012;119(8):1831-7.
33. Cordonnier C, Botterel F, Ben Amor R, et al. Correlation between galactomannan antigen levels in serum and neutrophil counts in haematological patients with invasive aspergillosis. *Clin Microbiol Infect.* 2009;15(1):81-6.
34. Garcia-Hermoso D, Alanio A, Cabaret O, et al. High diversity of non-sporulating moulds in respiratory specimens of immunocompromised patients: should all the species be reported when diagnosing invasive aspergillosis? *Mycoses.* 2015;58(9):557-64.
35. Bochud PY, Chien JW, Marr KA, et al. Toll-like receptor 4 polymorphisms and aspergillosis in stem-cell transplantation. *N Engl J Med.* 2008;359(17):1766-77.
36. Cunha C, Aversa F, Lacerda JF, et al. Genetic PTX3 deficiency and aspergillosis in stem-cell transplantation. *N Engl J Med.* 2014;370(5):421-32.
37. White PL, Parr C, Barnes RA. Predicting Invasive Aspergillosis in Hematology Patients by Combining Clinical and Genetic Risk Factors with Early Diagnostic Biomarkers. *J Clin Microbiol.* 2017;56(1):e01122-17.



**Table 1:** Univariate analysis of pre- and post-transplant factors associated with invasive aspergillosis

Pre-transplant factors	Cases (n = 185)	Controls (n = 651)	<i>P</i> <sup>a</sup>	Analyses adjusted for matching variables <sup>b</sup>	
				aOR [95% CI]	<i>P</i>
Age, median (Q1-Q3), years	50.1 (36.7-57.6)	49.3 (38.0-57.6)	0.85		
Female sex	71 (38.4)	257 (39.5)	0.79		
Body mass index (152/542) <sup>c</sup>	24.2 (21.9-26.6)	23.6 (21.2-26.3)	0.11	0.7 [0.2-2.0]	0.47
Karnofsky score = 100 (161/560)	68 (42.2)	248 (44.3)	0.64		
Underlying disease			0.52		
Acute leukemia and myelodysplastic syndrome	122 (66.0)	453 (69.6)			
Lymphoproliferative disorder	41 (22.2)	120 (18.4)			
Myeloproliferative neoplasm	16 (8.7)	48 (7.4)			
Other <sup>d</sup>	6 (3.2)	30 (4.6)			
Disease status (141/483)			0.20		
Complete remission	92 (65.3)	333 (68.9)			
Partial remission	0	9 (1.9)			
Stable disease	2 (1.4)	2 (0.4)			
Failure/relapse	46 (32.6)	134 (27.7)			
Other	1 (0.7)	5 (1.0)			
Stem cell source (185/649)			0.91		
Bone marrow	51 (27.6)	169 (26.0)			

PBSC	111 (60.0)	402 (61.9)			
Cord blood unit	23 (12.4)	76 (11.7)			
Bone marrow + PBSC	0	2 (0.3)			
Unrelated donor (185/649)	106 (57.3)	350 (53.9)	0.42		
Type of conditioning			0.21		
Myeloablative conditioning	10 (5.4)	27 (4.2)			
Fludarabine + 2 Gy TBI	48 (26.0)	135 (20.7)			
Reduced-intensity conditioning	127 (68.7)	489 (75.1)			
D/R CMV serology (183/638)			0.11		
D-/R+	61 (33.3)	156 (24.5)		1.54 [1.1-2.2]	<b>0.02</b>
D-/R-	47 (25.7)	198 (31.0)			
D+/R-	23 (15.6)	89 (14.0)		1	
D+/R+	52 (28.4)	195 (30.6)			
Transplantation during summer <sup>e</sup>	44 (23.8)	124 (19.1)	0.16	1.3 [0.9-2.0]	0.16
ATG-based conditioning	64 (34.6)	277 (45.6)	<b>0.05</b>	0.7 [0.5-1.0]	<b>0.05</b>
TBI (184/646)	110 (94.4)	316 (48.9)	<b>0.01</b>	1.6 [1.1-2.2]	<b>0.01</b>
Clofarabine	6 (3.2)	8 (1.2)	0.10	2.8 [1.0-8.3]	0.06
Invasive fungal infection before transplant (145/524)	12 (8.3)	52 (9.9)	0.55		
Female donor (181/641)	75 (41.4)	272 (42.4)	0.81		
Donor age (149/547), median (Q1-Q3) <sup>f</sup> , years	41.7 (30.2-49.6)	43.6 (32.2-51.4)	0.18	1.3 [0.9-1.9]	0.21

Post-transplant factors	Cases (n=185)	Controls (n=526)	<i>P</i>	aOR [95% CI]	<i>P</i>
-------------------------	---------------	------------------	----------	--------------	----------

Acute graft-versus-host disease (149/516)			<b>&lt;0.001</b>		
Grade 0-1	349 (67.6)	104 (56.2)		1	
Grade 2	108 (20.9)	37 (20.0)		1.2 [0.7-1.8]	0.54
Grade 3-4	59 (11.4)	44 (23.8)		2.6 [1.7-4.1]	<b>&lt;0.001</b>
CMV infection (185/526)	36 (19.5)	113 (21.5)	0.56		
Time of engraftment <sup>g</sup> (185/526), median (Q1-Q3)	19 (16-27)	19 (15-25)	0.76		
Absence of engraftment (182/515)	10 (5.5)	18 (3.5)	0.24		
Relapse after transplant (175/500)	40 (22.9)	7 (1.4)	<b>&lt;0.001</b>	20.7 [9.0-47.5]	<b>&lt;0.001</b>
Secondary neutropenia (138/418)	55 (39.9)	94 (22.5)	<b>&lt;0.001</b>	2.3 [1.5-3.5]	<b>&lt;0.001</b>

Abbreviations: aOR: adjusted odds ratio; CI: confidence interval; PBSC: peripheral blood stem cell; CBU: Cord blood unit; ATG: antithymoglobulin; TBI: Total body irradiation. D: donor; R: recipient. CMV: cytomegalovirus.

<sup>a</sup> *P* values of the chi-squared test Fisher exact test, or nonparametric Kruskal–Wallis test as appropriate

<sup>b</sup> logistic regression analysis adjusted for matching variables, such as HSCT center, age of patient ( $\pm$  5 years), and year of transplant

<sup>c</sup> aOR calculated for body mass index < 18

<sup>d</sup> solid tumor, bone marrow failure, inherited disorder, histiocytic disorder, and hemoglobinopathy

<sup>e</sup> July, August, and September

<sup>f</sup> aOR calculated for age > 60 years

<sup>g</sup> aOR calculated for number of days of neutrophil recovery after transplant > 18 days

**Table 2:** Multivariate analysis of pre- and post-transplant factors associated with invasive aspergillosis

<b>Model 1: Pre-transplant factors (185 cases and 651 controls)<sup>a</sup></b>	<b>aOR [95% CI]</b>	<b>P</b>
CMV D-R+ serology	1.5 [1.0-2.2]	<b>0.03</b>
Clofarabine	4.1 [1.3-13.2]	<b>0.02</b>
TBI	1.6 [1.1-2.4]	<b>0.01</b>
<b>Model 2: Pre- and post-transplant factors (185 cases and 526 controls)<sup>a</sup></b>	<b>aOR [95% CI]</b>	<b>P</b>
Total body irradiation	1.9 [1.1-3.2]	<b>0.02</b>
Acute GvHD		
Grade 2	1.8 [1.0-3.2]	<b>0.05</b>
Grade 3-4	4.3 [2.4-7.7]	<b>&lt;0.001</b>
Relapse	31.0 [10.4-92.4]	<b>&lt;0.001</b>
Secondary neutropenia	1.6 [1.0-2.8]	0.07

Abbreviations: aOR: adjusted odds ratio; CI: confidence interval; CMV: cytomegalovirus; D: donor; R: recipient; TBI: total body irradiation; GvHD: graft-*versus*-host disease;

<sup>a</sup> Multivariate logistic regression adjusted for matching variables, such as HSCT center, age of patient ( $\pm$  5 years), and year of transplant, and variables listed in the table



**Table 3:** Univariate analysis of pre- and post-transplant factors according to the timing of invasive aspergillosis

Pre-transplant factors	Analyses adjusted for matching variables aOR [95% CI] <sup>b</sup>										
	Controls	Early IA	Late IA	Very late IA	<i>P</i> <sup>a</sup>	Early IA		Late IA		Very late IA	
	(n = 651)	(n = 35)	(n = 33)	(n = 117)		OR [95% CI]	<i>P</i> <sup>b</sup>	OR [95% CI]	<i>P</i> <sup>b</sup>	OR [95% CI]	<i>P</i> <sup>b</sup>
Age, median	49.3	47.5	54.8	50.1	0.87						
(Q1-Q3), years	(38.0-57.6)	(35.9-60.8)	(38.8-57.9)	(36.7-57.4)							
Female R	257 (39.5)	15 (42.9)	10 (30.3)	46 (39.3)	0.73						
BMI (542/30/30/92),	23.6	24.1	24.2	24.3	0.24						
median (Q1-Q3)	(21.2-26.3)	(22.5-29.8)	(19.6-27.4)	(21.8-26.5)							
Karnofsky score = 100	312 (55.7)	16 (53.3)	15 (45.5)	62 (63.3)	0.30						
(560/30/33/98)											
Underlying disease:					0.15						
Acute leukemia and											
myelodysplastic	453 (69.6)	22 (62.9)	18 (54.6)	82 (70.1)		0.7 [0.2-3.1]	0.62	0.3 [0.9-1.0]	0.18	5.8 [0.8 -43.1]	<b>0.18</b>
syndrome											
Lymphoproliferative	120 (18.4)	7 (20.0)	7 (21.2)	27 (23.1)		0.8 [0.2-4.2]	0.78	0.5 [0.1-2.1]	0.50	7.3 [0.9-56.7]	0.18
disorder											
Myeloproliferative	48 (7.4)	4 (11.4)	5 (15.2)	7 (6.0)		1.2 [0.2-7.5]	0.96	1.0 [0.2-4.6]	0.96	4.2 [0.5-36.6]	0.57
disorder											
Other <sup>c</sup>	30 (4.6)	2 (5.7)	3 (9.1)	1 (0.9)		1		1		1	
Absence of complete	150 (31.1)	14 (53.9)	6 (25.0)	29 (31.9)	0.09	2.4 [1.1-5.4]	0.09	0.7 [0.3-1.8]	0.65	1.1 [0.7-1.7]	0.82

remission (483/26/24/91)											
Cord blood unit stem cell source (649/35/33/117)	76 (11.7)	10 (28.6)	3 (9.1)	10 (8.6)	<b>0.03</b>	2.9 [1.3-6.4]	<b>0.03</b>	0.7 [0.2-2.3]	0.51	0.8 [0.4-1.6]	0.51
Unrelated donor (649/35/33/117)	350 (53.9)	27 (77.1)	16 (48.5)	63 (53.9)	<b>0.05</b>	2.7 [1.2-6.1]	0.06	0.7 [0.4-1.5]	0.54	1.1 [0.7-1.6]	0.75
Type of conditioning					0.27						
Myeloablative conditioning	27 (4.2)	4 (11.4)	1 (3.0)	5 (4.3)							
Fludarabine + 2 Gy TBI	135 (20.7)	9 (25.7)	10 (30.3)	29 (24.8)							
RIC	489 (75.1)	22 (62.9)	22 (66.7)	83 (70.9)							
D-/R+ CMV serology	156 (24.5)	11 (31.4)	8 (24.2)	42 (36.5)	<b>0.05</b>	1.5 [0.6-3.6]	0.51	1.0 [0.5-2.4]	0.94	1.7 [1.1-2.7]	<b>0.01</b>
Transplantation during summer <sup>d</sup>	124 (19.1)	12 (34.3)	6 (18.2)	26 (22.2)	0.16	2.3 [1.1-4.7]	0.09	1.0 [0.4-2.4]	0.93	1.2 [0.8-2.0]	0.68
ATG-based conditioning	277 (42.3)	20 (57.1)	10 (30.3)	34 (29.1)	<b>0.005</b>	1.5 [0.7-3.2]	0.25	0.4 [0.2-0.9]	0.08	0.6 [0.4-1.0]	0.08
TBI (646/35/33/116)	316 (48.9)	21 (60.0)	18 (54.6)	71 (61.2)	0.07	1.7 [0.9-3.5]	0.20	1.4 [0.7-2.9]	0.33	1.6 [1.1-2.4]	0.09
Clofarabine	8 (1.2)	3 (8.6)	2 (6.1)	1 (0.9)	<b>0.008</b>	6.3 [1.5-25.6]	<b>0.03</b>	3.9 [0.8-19.6]	0.15	0.9 [0.1-7.0]	0.88
IFI before transplant (524/30/31/84)	52 (9.9)	2 (6.7)	3 (9.7)	7 (8.3)	0.96						
Female donor (641/34/33/114)	272 (42.4)	15 (44.1)	9 (27.3)	51 (44.7)	0.34						
Donor age (547/23/29/97), median	41.7 (30.2-49.6)	40.5 (26.8-48.1)	45.4 (36.5-53.3)	43.6 (33.8-50.8)	0.26						

(Q1-Q3), years

Post-transplant factors	Controls (n = 526)	Early IA (n = 35)	Late IA (n = 33)	Very late IA (n = 117)	P	Early IA (n = 35)	P	Late IA (n = 33)	P	Very late IA (n = 117)	P
Acute GvHD (516/35/33/117)					<b>&lt;0.001</b>						
Grade 0-1	349 (67.6)	33 (94.3)	14 (42.4)	57 (48.7)		1		1			
Grade 2	108 (20.9)	1 (2.9)	4 (12.1)	32 (27.4)		0.1 [0.0-0.8]	<b>0.05</b>	1.0 [0.3-3.1]	1.00	1.7 [1.1-2.9]	<b>0.05</b>
Grade 3-4	59 (11.4)	1 (2.9)	15 (45.5)	28 (23.9)		0.2 [0.0-1.3]	0.09	6.2 [2.8-13.7]	<b>&lt;0.001</b>	3.2 [1.9-5.5]	<b>&lt;0.001</b>
CMV infection	113 (21.5)	1 (2.9)	5 (15.2)	30 (25.6)	<b>0.03</b>	0.1 [0.0-0.8]	0.09	0.7 [0.3-1.8]	0.45	1.3 [0.8-2.1]	0.41
Time of engraftment <sup>e</sup> (117/35/33/117), median (Q1-Q3)	19 (15-25)	25 (17-40)	17 (15-34)	19 (15-23)	0.13	1.6 [0.8-3.3]	0.32	0.6 [0.3-1.2]	0.32	0.9 [0.6-1.4]	0.67
Absence of engraftment (515/34/32/116)	18 (3.5)	8 (23.5)	2 (6.3)	0	<b>&lt;0.001</b>	8.7 [3.4-22.0]	<b>&lt;0.001</b>	1.8 [0.4-8.4]	0.65	2.6 10 <sup>-7</sup> [0-]	0.99
Relapse after transplant (500/32/33/110)	7 (1.4)	3 (9.4)	2 (6.1)	35 (31.8)	<b>&lt;0.001</b>	8.0 [1.9-32.6]	<b>0.004</b>	5.0 [1.0-25.3]	<b>0.05</b>	31.1 [13.2-73.1]	<b>&lt;0.001</b>
Secondary neutropenia (418/23/25/90)	94 (22.5)	5 (21.7)	8 (32.0)	42 (46.7)	<b>&lt;0.001</b>	0.9 [0.3-2.6]	0.91	1.5 [0.6-3.6]	0.57	3.2 [2.0-5.2]	<b>&lt;0.001</b>

Abbreviations: IA: invasive aspergillosis; aOR: adjusted odds ratio; CI: confidence interval; R/ recipient; BMI: body mass index; TBI: total body irradiation; RIC: reduced-intensity conditioning; ATG: antithymocyte globulin; GvHD: graft-versus-host disease; CMV: cytomegalovirus; IFI: invasive fungal infection.

<sup>a</sup> P values of the chi-squared test, Fisher exact test, or nonparametric Kruskal–Wallis test as appropriate comparing the four groups overall

<sup>b</sup> multinomial logistic regression adjusted for matching variables, such as center, age of patient ( $\pm 5$  years), and year of transplant, comparing each group to the control population (reference category)

<sup>c</sup> solid tumor, bone marrow failure, inherited disorder, histiocytic disorder, and hemoglobinopathy

<sup>d</sup> July, August, and September

<sup>e</sup> aOR calculated for number of days of neutrophil recovery after transplant  $> 18$  days.

**Table 4:** Multivariate analysis of pre- and post-transplant factors for early, late, and very late invasive aspergillosis

	Early IA		Late IA		Very late IA	
	aOR [95% CI]	<i>P</i> <sup>a</sup>	aOR [95% CI]	<i>P</i> <sup>a</sup>	aOR [95% CI]	<i>P</i> <sup>a</sup>
<b><u>Pre-transplant factors</u></b>						
Absence of complete remission	2.4 [0.9-6.3]	0.07				
Unrelated donor	2.4 [0.9-6.4]	0.10				
D-/R+ CMV serology					1.6 [1.0-2.5]	<b>0.04</b>
ATG-based conditioning			0.4 [0.2-1.0]	0.06		
<b><u>Pre- and post-transplant factors</u></b>						
Absence of complete remission	2.6 [0.9-7.2]	0.06				
Acute GvHD						
Grade 2			1.2 [0.3-4.4]	0.76	2.6 [1.3-5.3]	<b>0.006</b>
Grade 3-4			4.7 [1.8-12.0]	<b>0.001</b>	6.9 [3.1-15.3]	<b>&lt;0.001</b>
Absence of engraftment	16.4 [2.9-92.6]	<b>0.002</b>				
Relapse after transplant			11.4 [0.9-142.5]	0.06	24.1 [7.3-79.6]	<b>&lt;0.001</b>
Secondary neutropenia					2.2 [1.1-4.1]	<b>0.02</b>

Abbreviations: aOR: adjusted odds ratio; CI: confidence interval; IA: invasive aspergillosis; GvHD: graft-versus-host disease; D: donor; R: recipient; CMV: cytomegalovirus; ATG: antithymoglobulin

<sup>a</sup> *P* value from multinomial logistic regression adjusted for matching variables, such as HSCT center, age of patient ( $\pm$  5 years), and year of transplant, and for the variables listed in the table comparing each group to the control population (reference category)

**Figure 1:** Flow chart

