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# Myeloablative vs Reduced-Intensity Conditioning Allogeneic Hematopoietic Cell Transplantation for Chronic Myeloid Leukemia

Saurabh Chhabra

*Medical College of Wisconsin*

Kwang Woo Ahn

*Medical College of Wisconsin*

Zhen-Huan Hu

*Medical College of Wisconsin*

Sandeep Jain

*Medical University of South Carolina*

Amer Assal

*Columbia University*

*See next page for additional authors*

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**Authors**

Saurabh Chhabra, Kwang Woo Ahn, Zhen-Huan Hu, Sandeep Jain, Amer Assal, Jan Cerny, Edward A. Copelan, Andrew Daly, Zachariah DeFilipp, Shahinaz M Gadalla, Robert Peter Gale, Siddhartha Ganguly, Betty K. Hamilton, Gerhard C. Hildebrandt, Jack W. Hsu, Yoshihiro Inamoto, Abraham S. Kanate, H. Jean Khoury, Hillard M. Lazarus, Mark R. Litzow, Sunita Nathan, Richard F. Olsson, Attaphol Pawarode, Olle Ringden, Jacob M. Rowe, Ayman Saad, Bipin N. Savani, Harry C. Schouten, Sachiko Seo, and Nirav N. Shah

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# Myeloablative vs reduced-intensity conditioning allogeneic hematopoietic cell transplantation for chronic myeloid leukemia

Saurabh Chhabra,<sup>1</sup> Kwang Woo Ahn,<sup>2,3</sup> Zhen-Huan Hu,<sup>3</sup> Sandeep Jain,<sup>4</sup> Amer Assal,<sup>5</sup> Jan Cerny,<sup>6</sup> Edward A. Copelan,<sup>7</sup> Andrew Daly,<sup>8</sup> Zachariah DeFilipp,<sup>9</sup> Shahinaz M. Gadalla,<sup>10</sup> Robert Peter Gale,<sup>11</sup> Siddhartha Ganguly,<sup>12</sup> Betty K. Hamilton,<sup>13</sup> Gerhard Carl Hildebrandt,<sup>14</sup> Jack W. Hsu,<sup>15</sup> Yoshihiro Inamoto,<sup>16</sup> Abraham S. Kanate,<sup>17</sup> H. Jean Khoury,<sup>18</sup> Hillard M. Lazarus,<sup>19</sup> Mark R. Litzow,<sup>20</sup> Sunita Nathan,<sup>21</sup> Richard F. Olsson,<sup>22,23</sup> Attaphol Pawarode,<sup>24</sup> Olle Ringden,<sup>22</sup> Jacob M. Rowe,<sup>25</sup> Ayman Saad,<sup>26</sup> Bipin N. Savani,<sup>27</sup> Harry C. Schouten,<sup>28</sup> Sachiko Seo,<sup>29</sup> Nirav N. Shah,<sup>1</sup> Melhem Solh,<sup>30</sup> Robert K. Stuart,<sup>4</sup> Celalettin Ustun,<sup>31</sup> Ann E. Woolfrey,<sup>32</sup> Jean A. Yared,<sup>33</sup> Edwin P. Alyea,<sup>34</sup> Matt E. Kalaycio,<sup>13</sup> Uday Popat,<sup>35</sup> Ronald M. Sobecks,<sup>13</sup> and Wael Saber<sup>3</sup>

<sup>1</sup>Division of Hematology/Oncology, Department of Medicine, <sup>2</sup>Division of Biostatistics, Institute for Health and Society, and <sup>3</sup>Center for International Blood and Marrow Transplant Research, Department of Medicine, Medical College of Wisconsin, Milwaukee, WI; <sup>4</sup>Division of Hematology/Oncology, Department of Medicine, Medical University of South Carolina, Charleston, SC; <sup>5</sup>Columbia University Medical Center, New York, NY; <sup>6</sup>UMass Memorial Medical Center, Worcester, MA; <sup>7</sup>Department of Hematologic Oncology and Blood Disorders, Levine Cancer Institute, Carolinas HealthCare System, Charlotte, NC; <sup>8</sup>Tom Baker Cancer Centre, Calgary, AB, Canada; <sup>9</sup>Blood and Marrow Transplant Program, Massachusetts General Hospital, Boston, MA; <sup>10</sup>Division of Cancer Epidemiology & Genetics, National Cancer Institute Clinical Genetics Branch, National Institutes of Health, Rockville, MD; <sup>11</sup>Hematology Research Centre, Division of Experimental Medicine, Department of Medicine, Imperial College London, London, United Kingdom; <sup>12</sup>Blood and Marrow Transplantation, Division of Hematology and Oncology, University of Kansas Medical Center, Kansas City, KS; <sup>13</sup>Department of Hematology and Medical Oncology, Cleveland Clinic Foundation, Cleveland, OH; <sup>14</sup>Markey Cancer Center, University of Kentucky, Lexington, KY; <sup>15</sup>Division of Hematology & Oncology, Department of Medicine, Shands HealthCare & University of Florida, Gainesville, FL; <sup>16</sup>Division of Hematopoietic Stem Cell Transplantation, National Cancer Centers Hospital, Tokyo, Japan; <sup>17</sup>Osborn Hematopoietic Malignancy and Transplantation Program, West Virginia University, Morgantown, WV; <sup>18</sup>Emory University Hospital, Atlanta, GA; <sup>19</sup>Seidman Cancer Center, University Hospitals Cleveland Medical Center, Case Western Reserve University, Cleveland, OH; <sup>20</sup>Division of Hematology and Transplant Center, Mayo Clinic Rochester, Rochester, MN; <sup>21</sup>Rush University Medical Center, Chicago, IL; <sup>22</sup>Division of Therapeutic Immunology, Department of Laboratory Medicine, Karolinska Institutet, Stockholm, Sweden; <sup>23</sup>Centre for Clinical Research Sormland, Uppsala University, Uppsala, Sweden; <sup>24</sup>Blood and Marrow Transplantation Program, Division of Hematology/Oncology, Department of Internal Medicine, The University of Michigan Medical School, Ann Arbor, MI; <sup>25</sup>Department of Hematology, Shaare Zedek Medical Center, Jerusalem, Israel; <sup>26</sup>Division of Hematology/Oncology, Department of Medicine, University of Alabama at Birmingham, Birmingham, AL; <sup>27</sup>Division of Hematology/Oncology, Department of Medicine, Vanderbilt University Medical Center, Nashville, TN; <sup>28</sup>Department of Hematology, Academische Ziekenhuis, Maastricht, The Netherlands; <sup>29</sup>Department of Hematology & Oncology, National Cancer Research Center East, Chiba, Japan; <sup>30</sup>The Blood and Marrow Transplant Group of Georgia, Northside Hospital, Atlanta, GA; <sup>31</sup>Division of Hematology, Oncology and Transplantation, Department of Medicine, University of Minnesota Medical Center, Minneapolis, MN; <sup>32</sup>Fred Hutchinson Cancer Research Center, Seattle, WA; <sup>33</sup>Blood & Marrow Transplantation Program, Division of Hematology/Oncology, Department of Medicine, Greenebaum Cancer Center, University of Maryland, Baltimore, MD; <sup>34</sup>Center of Hematologic Oncology, Dana-Farber Cancer Institute, Boston, MA; and <sup>35</sup>MD Anderson Cancer Center, Houston, TX

## Key Points

- RIC is a reasonable alternative to MAC for CML patients in the TKI era.
- In CML patients, RIC results in similar survival as MAC, albeit at the expense of increased early posttransplant relapse.

Allogeneic hematopoietic cell transplantation (allo-HCT) is a potentially curative treatment of chronic myeloid leukemia (CML). Optimal conditioning intensity for allo-HCT for CML in the era of tyrosine kinase inhibitors (TKIs) is unknown. Using the Center for International Blood and Marrow Transplant Research database, we sought to determine whether reduced-intensity/nonmyeloablative conditioning (RIC) allo-HCT and myeloablative conditioning (MAC) result in similar outcomes in CML patients. We evaluated 1395 CML allo-HCT recipients between the ages of 18 and 60 years. The disease status at transplant was divided into the following categories: chronic phase 1, chronic phase 2 or greater, and accelerated phase. Patients in blast phase at transplant and alternative donor transplants were excluded. The primary outcome was overall survival (OS) after allo-HCT. MAC (n = 1204) and RIC allo-HCT recipients (n = 191) from 2007 to 2014 were included. Patient, disease, and transplantation characteristics were similar, with a few exceptions. Multivariable analysis showed no significant difference in OS between MAC and RIC groups. In addition, leukemia-free survival and nonrelapse mortality did not differ significantly between the 2 groups. Compared with MAC, the RIC

group had a higher risk of early relapse after allo-HCT (hazard ratio [HR], 1.85;  $P = .001$ ). The cumulative incidence of chronic graft-versus-host disease (cGVHD) was lower with RIC than with MAC (HR, 0.77;  $P = .02$ ). RIC provides similar survival and lower cGVHD compared with MAC and therefore may be a reasonable alternative to MAC for CML patients in the TKI era.

## Introduction

With the remarkable success of tyrosine kinase inhibitors (TKIs) for the treatment of patients with chronic myeloid leukemia (CML), the use of allogeneic hematopoietic cell transplantation (allo-HCT) since the turn of the century has dramatically decreased.<sup>1-4</sup> Nonetheless, allo-HCT is a useful and potentially curative treatment option for a subset of CML patients who are refractory to or intolerant of TKIs and those who present in accelerated phase (AP) or blast phase (BP).<sup>5-8</sup> Traditionally, myeloablative conditioning (MAC) is the standard intensity for CML patients in need of allo-HCT.<sup>8-10</sup> MAC is, however, characterized by a high risk of toxicity and nonrelapse mortality (NRM), especially among patients with comorbid conditions and advanced age. This prompted exploration of reduced-intensity/nonmyeloablative conditioning (RIC) regimens.<sup>11,12</sup>

Retrospective studies comparing MAC with RIC in patients with acute myeloid leukemia (AML) or myelodysplastic syndromes suggested that RIC was associated with increased relapse but reduced NRM, resulting in similar overall survival (OS), even though patients receiving RIC were older and/or less fit.<sup>13-21</sup> In contrast, a randomized phase 3 study (BMT CTN protocol 0901) demonstrated that in fit (hematopoietic cell transplant-comorbidity index [HCT-CI]  $\leq 4$ ) patients with AML or myelodysplastic syndromes in remission between the ages of 18 and 65 years receiving allo-HCT from HLA-identical sibling or unrelated donors, RIC resulted in lower NRM but a significant disadvantage in leukemia-free survival (LFS) compared with MAC.<sup>13</sup> It is remarkable that in the era of TKIs, there is a dearth of evidence pertaining to the role of conditioning intensity on outcomes after allo-HCT for CML that may guide practice patterns. To date, no prospective or large observational study has evaluated outcomes after MAC and RIC allo-HCT for CML. We conducted a registry analysis from the observational database of the Center for International Blood and Marrow Transplant Research (CIBMTR) comparing outcomes after RIC and MAC for allo-HCT in the era of TKIs. We hypothesized that RIC allo-HCT is as efficacious as MAC allo-HCT in CML patients for survival outcomes, considering the evidence for the graft-versus-leukemia effect of allo-HCT.<sup>22</sup>

## Patients and methods

### Data sources

The CIBMTR is a combined research program of the Medical College of Wisconsin and the National Marrow Donor Program, which consists of a voluntary network of more than 450 transplantation centers worldwide that contribute detailed data on consecutive allogeneic and autologous transplantations to a centralized statistical center. Observational studies conducted by the CIBMTR are performed in compliance with all applicable federal regulations pertaining to the protection of human research participants. Protected health information issued in the performance

of such research is collected and maintained in the CIBMTR's capacity as a Public Health Authority under the Health Insurance Portability and Accountability Act Privacy Rule.

### Patients

Patients with CML between 18 and 60 years of age who underwent allo-HCT using a sibling or unrelated donor<sup>23</sup> between 2007 and 2014 were included in the study. Donors were matched to the recipients at the allele level at HLA-A, HLA-B, HLA-C, and HLA-DRB1 loci or mismatched at a single HLA locus. An upper age limit of 60 years was introduced as an inclusion criterion to restrict the patient population to a cohort where by age criteria both MAC and RIC would be feasible. Patients in the chronic phase (CP) or AP<sup>24</sup> were included. Those in BP at allo-HCT were excluded to reduce bias, because inferior survival outcomes would be expected with an RIC (vs MAC) regimen in BP patients, as suggested by the results of the prospective CTN 0901 study.<sup>13</sup> Patients with haploidentical or cord blood transplant were excluded not only to limit the heterogeneity of the study population but also because of the small number of patients. Conditioning intensity was determined per Consensus CIBMTR criteria: MAC regimens were defined by total body irradiation (TBI)  $\geq 5$  Gy single dose or  $\geq 8$  Gy fractionated or busulfan (Bu) dose  $> 8$  mg/kg oral or  $> 6.4$  mg/kg IV, whereas RIC regimens were defined by Bu dose  $\leq 8$  mg/kg oral or IV equivalent, melphalan (Mel)  $\leq 150$  mg/m<sup>2</sup> or TBI  $\leq 2$  Gy.<sup>25</sup>

### Study end points

The primary end point of the study was OS. OS was defined as the time from transplantation to death from any cause or last follow-up. Death from any cause was considered an event. Surviving patients were censored at last follow-up. Secondary end points included LFS, NRM, and cumulative incidence of relapse and chronic graft-versus-host disease (cGVHD). LFS was defined as time from transplantation to either relapse or death from any cause, and alive patients were censored at the time of relapse or last follow-up. NRM was defined as death from any cause in continuous remission and was summarized by cumulative incidence estimate with relapse as competing risk. Relapse was summarized by cumulative incidence estimate with NRM as the competing risk. For relapse and NRM, patients in continuous complete remission were censored at last follow-up. cGVHD was graded per Consensus criteria.<sup>26</sup> For cGVHD, death without the event was considered a competing event.

### Statistical analysis

This is a retrospective comparative cohort study comparing outcomes after RIC vs MAC allo-HCT using related and unrelated donors for patients with CML in CP or AP. Eligible patients were stratified according to RIC vs MAC. The objective of this analysis was to compare these 2 types of conditioning regimens and their

effect on outcomes after allo-HCT. The outcomes studied were OS, LFS, relapse rate, and incidence of cGVHD and NRM. Probabilities of NRM, relapse, and cGVHD were calculated by cumulative incidence function accounting for competing risks. Comparisons of cumulative incidence across time cohorts were performed via Gray's test. Multivariable analyses were performed to evaluate associations among patient-related (age, Karnofsky Performance Score [KPS], HCT-CI, and Sorror comorbidity index<sup>27</sup>), disease-related (time from diagnosis to transplant and disease status at transplant), donor-related (donor type, cytomegalovirus [CMV] match, and sex match) and transplantation-related (year of transplant, GVHD prophylaxis, graft source, and in vivo T-cell depletion) variables and outcomes of interest (OS, LFS, NRM, GVHD, and relapse) using a Cox proportional hazards regression model. Forward stepwise selection was used to identify significant covariates that influenced outcomes. Disease status at transplant was categorized into chronic phase 1 (CP1), CP2 and beyond (CP2+), and AP. Covariates with  $P < .05$  were considered significant. The proportional hazards assumption for Cox regression was tested by adding a time-dependent covariate for each risk factor and each outcome. Covariates violating the proportional hazards assumption were added as time-dependent covariates in the Cox regression model. Adjusted probabilities for OS, LFS, NRM, cGVHD, and relapse were calculated based on the final Cox models.<sup>28,29</sup> The findings of multivariate analysis of cGVHD, NRM, and relapse by the Cox regression model were confirmed using the Fine-Gray model of competing risks. Interactions between the main effect (conditioning regimen) and significant covariates were examined. Power analysis was conducted; it was assumed that OS at 5 years for RIC was 60%. The median censoring time was assumed to be 7 years. Assuming OS of 45% at 5 years for MAC (relative risk, 1.56), the study had 92% statistical power. All analyses were performed using SAS 9.4.

## Results

### Patient, disease, and transplantation characteristics

Using the eligibility criteria, a total of 1395 patients were included in the study population, of which 1204 received MAC and 191 received RIC. Patient, disease, and transplantation characteristics were similar, with a few exceptions (Table 1). The median ages in MAC and RIC groups were 43 (range, 18-60) and 51 (range, 19-60) years, respectively. There were more males in the MAC group (60%) than the RIC group (47%). A greater proportion of MAC patients (70%) had KPS  $\geq 90$  compared with RIC (58%), and 20% and 30% of MAC and RIC recipients, respectively, had HCT-CI  $\geq 3$ . As might be expected, the RIC cohort was enriched for older patients with lower KPS and higher HCT-CI. Median time from diagnosis to transplant was 23 months in the MAC cohort and 27 months in the RIC cohort. In both the cohorts, three-fourths of the patients received a peripheral blood graft. CP1 was the most common status prior to allo-HCT in both the MAC (42%) and RIC (48%) cohorts ( $P = .34$ ). TKI was used for treatment before allo-HCT in 93% of MAC patients and 94% of RIC patients. In the MAC cohort, Bu/Cy was the most common regimen employed (40%), followed by Flu/Bu4 (24%) and Cy/TBI (24%). In the RIC cohort, Flu/Bu2 was the most common regimen (51%) followed by Flu/Mel (27%). In vivo T-cell depletion using ATG or alemtuzumab was used less commonly in the MAC cohort (22%)

than the RIC cohort (47%). The majority of patients in both cohorts (88% in MAC and 90% in RIC) were from the North America or South America. Approximately 5% of patients were from Europe. Eastern Mediterranean, Southeastern Asian, and Western Pacific countries made up  $\sim 5\%$  of both cohorts. The median follow-up of survivors was 52 and 60 months in the MAC and RIC cohorts, respectively.

Of the 596 patients categorized as CP1 at the time of allo-HCT (504 in MAC and 92 in RIC cohort) (Table 1), the indication for allo-HCT in 78% ( $n = 467$ ) was failure to achieve deeper (cytogenetic and/or molecular) remission with nontransplant therapies. The indication was not available for the other 22% and is presumed to be intolerance to TKIs. A total of 465 patients were in AP at the time of allo-HCT (405 in MAC [34%] and 60 in RIC [31%]) (Table 1). A total of 230 patients (57%) in the MAC cohort and 31 patients (52%) in the RIC cohort had presented with AP or BP and achieved at least a hematologic response prior to allo-HCT. The baseline cytogenetic information (pre-HCT) was not available in the 2 cohorts given the type of data forms used for this study (Transplant Essential Data).

### OS

On multivariable analysis, no significant difference in OS was seen between the MAC and RIC cohorts (HR, 0.99,  $P = .95$ ) (Table 2; Figure 1D). The adjusted probabilities of 5-year OS were 53% (95% confidence interval [CI], 50%-56%) and 53% (95% CI, 45%-60%) for the MAC and RIC cohorts, respectively ( $P = .98$ ) (Table 3). Variables associated with higher mortality included unrelated donor (matched or partially matched), CMV (donor seronegative/recipient seropositive and both donor and recipient seropositive), peripheral blood graft, and KPS  $< 90\%$  (Table 2). Additionally, compared with those in CP1, patients in AP (HR, 1.59;  $P = .0006$ ) and CP2+ (HR, 1.18;  $P = .0008$ ) had an increased risk of mortality after allo-HCT.

### LFS

On multivariable analysis, no significant difference was observed when comparing the MAC and RIC cohorts (HR, 1.13;  $P = .29$ ) (Table 2; Figure 1C). The adjusted 5-year LFS rates for MAC and RIC were 44% (95% CI, 41%-47%) and 43% (95% CI 36%-51%), respectively ( $P = .81$ ; Table 3). Variables associated with higher risk of treatment failure (inverse of LFS) included disease states of AP (HR, 1.47;  $P < .0001$ ) and CP2+ (HR, 1.43;  $P = .0002$ ) at allo-HCT (compared with CP1) and KPS  $< 90\%$  (HR, 1.49;  $P < .0001$ ).

### NRM and relapse

On multivariable analysis, there was no significant difference in NRM between the MAC and RIC groups (Cox regression model: HR, 1.01;  $P = .97$ ; Fine-Gray model: HR, 0.92;  $P = .57$ ) (Table 2; Figure 1A; supplemental Table 2). The adjusted 5-year NRM rates were 32% (95% CI, 29%-35%) and 29% (95% CI, 22%-36%) in the MAC and RIC cohorts, respectively ( $P = .53$ ; Table 3). Independent of the conditioning intensity, recipient age (40-49 years: HR, 1.47;  $P = .02$ ; 50-59 years: HR, 1.66;  $P = .003$  [compared with 18-29 years age group]), donor type (matched unrelated and partially matched unrelated compared with matched sibling), CMV (donor and recipient seropositivity), and

**Table 1. Characteristics of patients aged 18 to 60 years undergoing allo-HCT using MAC vs RIC for CML, 2007-2014**

Variable	MAC (n = 1204)	RIC (n = 191)	P
Follow-up of survivors, median (range), mo	52 (4-102)	60 (7-101)	
Number of centers	163	76	
<b>Patient related</b>			
Age at transplant, median (range), y	43 (18-60)	51 (19-60)	<.001
Age at transplant, y			<.001
18-29	212 (18)	17 (9)	
30-39	276 (23)	21 (11)	
40-49	404 (34)	47 (25)	
50-59	312 (26)	106 (55)	
Sex			<.001
Male	722 (60)	90 (47)	
Female	482 (40)	101 (53)	
KPS, %			<.001
90-100	845 (70)	110 (58)	
<90	267 (22)	72 (38)	
Missing	92 (8)	9 (5)	
WHO region			.60
Africa	1 (<1)	0	
Americas	1054 (88)	172 (90)	
Eastern Mediterranean	22 (2)	0	
Europe	78 (6)	10 (5)	
Southeastern Asian	21 (2)	4 (2)	
Western Pacific	27 (2)	5 (3)	
Missing	1 (<1)	0	
Sorrer comorbidity index (HCT-CI)			.003
0	525 (44)	64 (34)	
1	132 (11)	27 (14)	
2	125 (10)	25 (13)	
3+	246 (20)	58 (30)	
Missing	176 (15)	17 (9)	
<b>Disease related</b>			
Disease status prior to transplant			.20
CP1	504 (42)	92 (48)	
AP	405 (34)	60 (31)	
CP2+	278 (23)	38 (20)	
CP, NOS	17 (1)	1 (<1)	
Prior treatment			.85
No	9 (<1)	1 (<1)	
Yes	1185 (98)	189 (99)	
Missing	10 (<1)	1 (<1)	
Use of TKI prior to HCT			.89
No	73 (6)	11 (6)	
Yes	1121 (93)	179 (94)	
Missing	10 (<1)	1 (<1)	
TKI used pre-HCT			.11
No TKI used	73 (6)	11 (6)	

Data are presented as n (%) of patients unless otherwise indicated.

ATG, anti-thymocyte globulin; BM, bone marrow; CP, chronic phase; CSA, cyclosporine; Cy, cyclophosphamide; F, female; Flu, fludarabine; M, male; NOS, not otherwise specified; PB, peripheral blood; TAC, tacrolimus; URD, unrelated donor; WHO, World Health Organization.

**Table 1. (continued)**

Variable	MAC (n = 1204)	RIC (n = 191)	P
Imatinib + dasatinib + nilotinib	250 (21)	52 (27)	
Imatinib + dasatinib	295 (25)	55 (29)	
Imatinib + nilotinib	73 (6)	13 (7)	
Dasatinib + nilotinib	35 (3)	8 (4)	
Imatinib	395 (33)	43 (23)	
Dasatinib	52 (4)	7 (4)	
Nilotinib	21 (2)	1 (<1)	
Missing	10 (<1)	1 (<1)	
Time to HCT from diagnosis, mo	23 (<1-608)	27 (4-281)	.03
Time to HCT from diagnosis, mo			.12
0-12	316 (26)	41 (21)	
12-36	496 (41)	75 (39)	
≥36	391 (32)	74 (39)	
Missing	1 (<1)	1 (<1)	
<b>Donor-related</b>			
Donor type			.29
HLA-identical sibling	539 (45)	73 (38)	
Well-matched unrelated	447 (37)	85 (45)	
Partially matched unrelated	125 (10)	21 (11)	
Mismatched unrelated	9 (<1)	2 (1)	
Unrelated (matching indeterminable)	84 (7)	10 (5)	
URD-recipient HLA-matching			.10
3	0	1 (<1)	
5	1 (<1)	0	
6	5 (<1)	0	
7	101 (15)	19 (16)	
8	407 (61)	79 (67)	
Missing	151 (23)	19 (16)	
Donor-recipient sex match			.003
M-M	456 (38)	63 (33)	
M-F	271 (23)	66 (35)	
F-M	264 (22)	27 (14)	
F-F	209 (17)	34 (18)	
Missing	4 (<1)	1 (<1)	
Donor-recipient CMV status			.36
+/+	377 (31)	67 (35)	
+/-	99 (8)	18 (9)	
-/+	272 (23)	47 (25)	
-/-	287 (24)	41 (21)	
URD age, median (range), y	30 (18-61)	29 (20-59)	.22
URD age, y			.15
18-29	271 (41)	59 (50)	
30-39	137 (21)	27 (23)	
40-49	103 (15)	17 (14)	
50-59	37 (6)	2 (2)	

Data are presented as n (%) of patients unless otherwise indicated.

ATG, anti-thymocyte globulin; BM, bone marrow; CP, chronic phase; CSA, cyclosporine; Cy, cyclophosphamide; F, female; Flu, fludarabine; M, male; NOS, not otherwise specified; PB, peripheral blood; TAC, tacrolimus; URD, unrelated donor; WHO, World Health Organization.

**Table 1. (continued)**

Variable	MAC (n = 1204)	RIC (n = 191)	P
≥60	2 (<1)	0	
Missing	115 (17)	13 (11)	
<b>Transplant related</b>			
Year of transplant			.08
2007-2008	261 (22)	30 (16)	
2009-2010	311 (26)	60 (31)	
2011-2012	302 (25)	56 (29)	
2013-2014	330 (27)	45 (24)	
Graft type			.58
BM	293 (24)	43 (23)	
PB	911 (76)	148 (77)	
Conditioning regimen			
	TBI ± Cy ± others (376; 31%)	Bu2 + Flu ± others (97; 51%)	
	Bu + Cy ± others (492; 41%)	Flu + Mel ± other (53; 28%)	
	Bu4 + Flu ± others (302; 25%)	Cy ± Flu ± TBI ± others (18; 9%)	
	Others (34; 3%)	TBI ± Flu ± others (11; 6%)	
		Others (12; 6%)	
ATG/alemtuzumab			<.001
ATG alone	258 (21)	72 (38)	
Alemtuzumab alone	15 (1)	18 (9)	
No ATG or alemtuzumab	929 (77)	101 (53)	
Missing	2 (<1)	0	
GVHD prophylaxis			.26
Ex vivo T-cell depletion/CD34 selection	20 (2)	1 (<1)	
TAC based	712 (59)	124 (65)	
CSA based	414 (34)	55 (29)	
Other	53 (4)	11 (6)	
Missing	5 (<1)	0	

Data are presented as n (%) of patients unless otherwise indicated.

ATG, anti-thymocyte globulin; BM, bone marrow; CP, chronic phase; CSA, cyclosporine; Cy, cyclophosphamide; F, female; Flu, fludarabine; M, male; NOS, not otherwise specified; PB, peripheral blood; TAC, tacrolimus; URD; unrelated donor; WHO, World Health Organization.

peripheral blood graft were associated with a significantly higher risk of NRM.

Multivariable analysis confirmed that the risk of relapse between the 2 cohorts was time dependent; RIC had a significantly increased risk of relapse early after allo-HCT (in the first 5 months) (HR, 1.85;  $P = .001$ ), but no statistically significant difference was observed during the late (>5 months) allo-HCT course (Table 2; Figure 1B). Similar results were obtained with the Fine-Gray model ( $\leq 5$  months: HR, 1.86;  $P = .001$ ; >5 months: HR, 0.60;  $P = .06$ ; supplemental Table 2). The adjusted cumulative incidence rates of relapse at 5 years were 26% (95% CI, 23%-28%) and 25% (95% CI, 19%-31%) for MAC and RIC, respectively ( $P = .96$ ; Table 3). In addition to the conditioning intensity, disease status, KPS, and donor type affected the relapse risk. AP (HR 1.87,  $P < .0001$ ) and CP2+ (HR, 1.79;  $P < .0001$ ) carried an increased risk of relapse relative to CP1. Well-matched and partially matched unrelated donors and KPS  $\geq 90\%$  were associated with a lower relapse risk. An evaluation of the causes

of death in the 2 cohorts (supplemental Table 1) did not reveal any significant differences between the cohorts; disease relapse (~30% in both) and GVHD (29% in both) were the dominant causes of death in both cohorts.

### cGVHD

Multivariable analysis showed a significantly reduced risk of cGVHD (Cox model: HR, 0.77; 95% CI, 0.61-0.97;  $P = .02$ ; Fine-Gray model: HR, 0.78;  $P = .03$ ) with RIC relative to MAC (Table 2; supplemental Table 2). The adjusted cumulative incidence rates of cGVHD at 1 and 5 years were 50% (95% CI, 47%-53%) and 59% (95% CI, 56%-62%) after MAC compared with 41% (95% CI, 33%-48%) and 51% (95% CI, 44%-59%) with RIC ( $P = .02$  and  $.07$ , respectively) (Table 3). In patients without in vivo T-cell depletion, the incidence of cGVHD was not significantly different relative to those receiving ATG (HR, 1.13;  $P = .18$ ) but was significantly higher compared with those receiving alemtuzumab (HR, 2.76;  $P = .002$ ). Compared with ATG, alemtuzumab



**Table 2. Multivariable analysis for patients aged 18 to 60 years undergoing allo-HCT using MAC vs RIC for CML, 2007-2014**

	No. of patients (evaluable)	HR	95% CI	P
<b>OS</b>				
MAC	1204	1.0		
RIC	191	0.99	0.79-1.25	.95
<b>Disease status</b>				
CP1	596	1.0		
AP	465	1.59	1.13-2.24	.0006
CP2+	316	1.18	0.85-1.63	.0008
CP, NOS	11	1.96	0.71-5.36	.28
Missing	7	0.87	0.63-1.20	.17
<b>KPS</b>				
≥90%	955	1.0		
<90%	339	1.41	1.18-1.69	.0002
Missing	101	1.08	0.76-1.54	.66
<b>Graft type</b>				
BM	336	1.0		
PB	1059	1.36	1.10-1.66	.004
<b>Donor type</b>				
HLA-identical sibling	612	1.0		
Well-matched unrelated	532	1.32	1.09-1.60	.004
Partially matched unrelated	146	1.45	1.10-1.89	.007
Mismatched unrelated	11	1.65	0.73-3.74	.23
URD matching TBD	94	1.29	0.09-1.83	.15
<b>CMV match</b>				
+/+	444	1.0		
+/-	117	0.93	0.68-1.26	.63
-/+	319	1.11	0.90-1.37	.33
-/-	328	0.76	0.60-0.95	.02
Missing	187	1.04	0.76-1.42	.79
<b>LFS</b>				
MAC	1190	1.0		
RIC	191	1.13	0.92-1.39	.29
<b>Disease status</b>				
CP1	593	1.0		
AP	459	1.47	1.23-1.74	<.0001
CP2+	311	1.43	1.19-1.73	.0002
CP, NOS	11	1.74	0.86-3.53	.12
Missing	7	1.44	0.53-3.86	.47
<b>KPS</b>				
≥90%	948	1.0		
<90%	337	1.49	1.26-1.75	<.0001
Missing	96	0.98	0.73-1.32	.90
<b>Relapse ≤5 mo</b>				
MAC	1183	1.0		
RIC	191	1.85	1.27-2.70	.001
<b>Relapse &gt;5 mo</b>				
MAC	922	1.0		
RIC	137	0.64	0.37-1.08	.097

TBD, to be determined.

**Table 2. (continued)**

	No. of patients (evaluable)	HR	95% CI	P
<b>Disease status</b>				<.0001
CP1	593	1.0		
AP	459	1.87	1.44-2.41	<.0001
CP2+	311	1.79	1.34-2.39	<.0001
CP, NOS	11	1.58	0.50-5.02	.43
<b>KPS</b>				<.0001
≥90%	944	1.0		
<90%	334	1.81	1.42-2.29	<.0001
Missing	96	1.06	0.69-1.62	.80
<b>Donor type</b>				.005
HLA-identical sibling	604	1.0		
Well-matched unrelated	528	0.64	0.50-0.82	.0004
Partially matched unrelated	143	0.64	0.43-0.95	.03
Mismatched unrelated	10	0.82	0.20-3.33	.78
URD matching TBD	89	0.75	0.45-1.23	.25
<b>NRM</b>				
MAC	1190	1.0		
RIC	191	1.01	0.76-1.34	.97
<b>Age, y</b>				.02
18-29	226	1.0		
30-39	295	1.27	0.88-1.82	.20
40-49	445	1.47	1.05-2.05	.02
50-59	415	1.66	1.18-2.33	.003
<b>Graft type</b>				.02
BM	333	1.0		
PB	1048	1.34	1.04-1.72	.02
<b>Donor type</b>				.0002
HLA-identical sibling	608	1.0		
Well-matched unrelated	531	1.56	1.23-1.99	.0003
Partially matched unrelated	143	1.89	1.28-2.53	.0001
Mismatched unrelated	10	2.73	0.81-5.99	.03
URD matching TBD	89	1.46	1.08-2.29	.09
<b>CMV match</b>				.02
+/+	440	1.0		
+/-	117	1.01	0.69-1.48	.95
-/+	316	1.22	0.94-1.58	.13
-/-	326	0.74	0.55-0.97	.04
Missing	182	0.97	0.67-1.39	.85
<b>cGVHD</b>				
MAC	1177	1.0		
RIC	187	0.77	0.61-0.97	.02
<b>Graft type</b>				<.0001
BM	327	1.0		
PB	1037	1.75	1.46-2.10	<.0001
<b>GVHD prophylaxis</b>				.0005
TAC	821	1.0		
T-cell depletion/CD34	19	0.26	0.11-0.58	.001

TBD, to be determined.

**Table 2. (continued)**

	No. of patients (evaluable)	HR	95% CI	P
CSA	457	0.82	0.70-0.96	.01
Other	64	0.67	0.46-0.99	.04
<b>ATG/alemtuzumab</b>				<b>.004</b>
ATG alone	319	1.0		
Alemtuzumab alone	33	0.41	0.21-0.78	.007
No ATG/alemtuzumab	1012	1.13	0.94-1.36	.18
<b>Sex match</b>				<b>.007</b>
M-M	508	1.0		
M-F	331	0.91	0.75-1.10	.33
F-M	284	1.30	1.08-1.58	.006
F-F	241	1.06	0.87-1.30	.57

TBD, to be determined.

significantly decreased the risk of cGVHD (HR, 0.41;  $P = .007$ ). In addition, female donor/male recipient had a 30% higher cGVHD risk than male donor and recipient ( $P = .006$ ).

### Subgroup analysis

**CP2+ and AP.** Multivariable analysis of a prespecified subgroup of CP2+ and AP at the time of allo-HCT was done to see if there were differences in outcomes in this group of patients with higher-risk disease (Table 4). The analysis showed that in the RIC cohort, OS (HR, 1.18;  $P = .26$ ) and LFS (HR, 1.27;  $P = .08$ ) were not significantly different when compared with MAC. NRM was also not affected by the conditioning intensity in this high-risk subgroup (HR, 1.32;  $P = .17$ ). The risk of relapse, however, varied by the time from allo-HCT: RIC patients with CP2+/AP carried a higher risk of early relapse (HR, 2.06;  $P = .002$ ) but had a relapse risk similar to MAC later (>5 months) in the course (HR, 0.46;  $P = .06$ ). The cumulative incidence of cGVHD did not differ significantly between MAC and RIC (HR, 0.83;  $P = .26$ ).

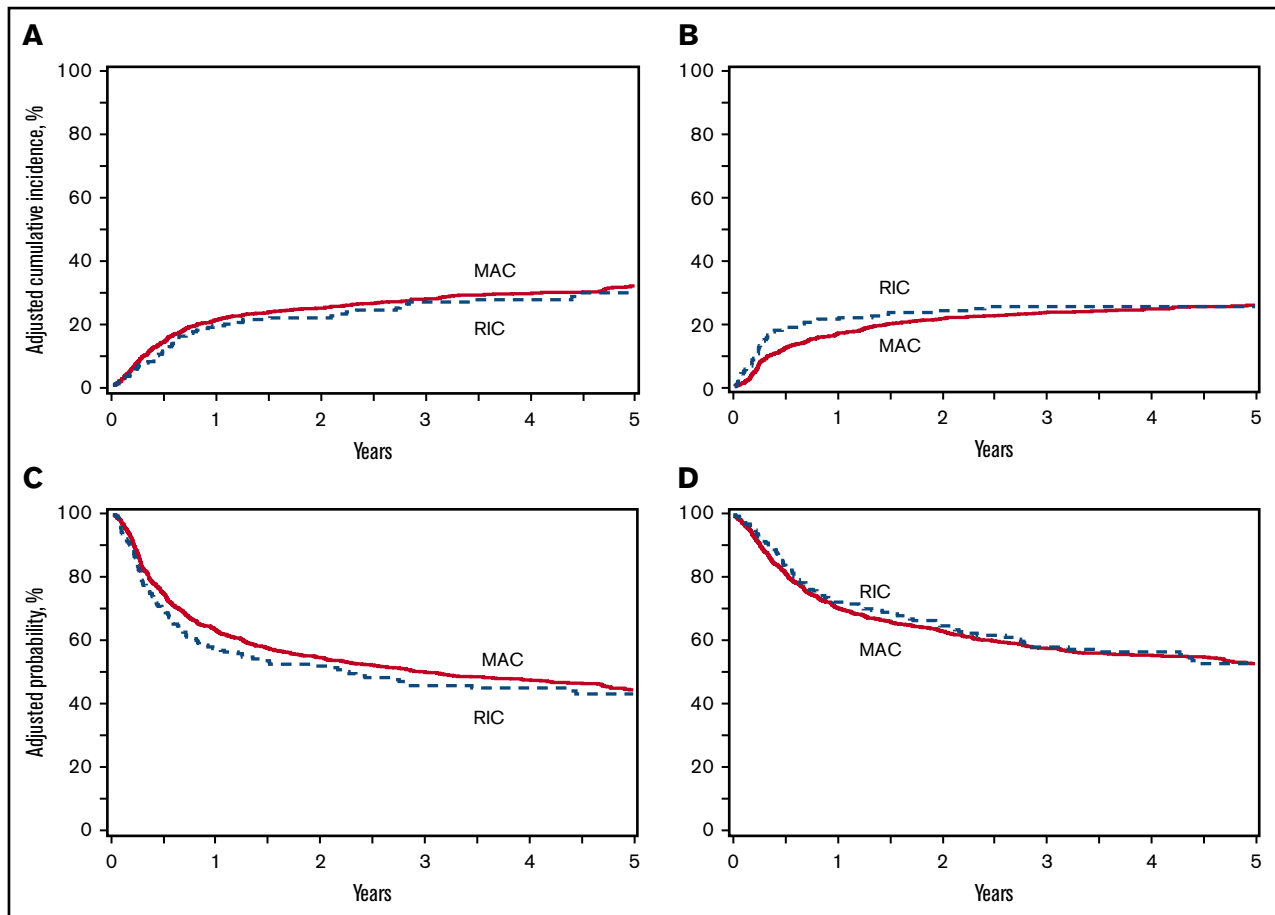
### Discussion

Although allogeneic transplantation is no longer a first-line treatment of CML in CP1 in the United States, many patients who are resistant to or intolerant of TKIs continue to require it.<sup>2,30</sup> The European Leukemia Network, in addition, recommends allo-HCT for all patients with AP or BP at diagnosis.<sup>2</sup> Since the approval of the TKI imatinib mesylate in 2001 based on the phase 3 IRIS study,<sup>31,32</sup> the great majority of CML patients undergo allo-HCT later in their course. Nonetheless, the question about the optimal allogeneic transplant conditioning regimen intensity for CML is relevant and has been addressed for the first time in this observational study. Our data illustrate that in the era of TKIs, RIC is an appropriate alternative to MAC for CML patients given the equivalent survival.

Studies have shown that RIC allo-HCT for CML patients can prolong survival with acceptable NRM. A retrospective study by Crawley et al reported the outcomes after RIC allo-HCT and demonstrated its feasibility in 186 CML patients.<sup>11</sup> The median age of patients was 50 years; 64% patients were in CP1, 13% in

CP2, 17% in AP, and 6% in BP. OS and progression-free survival at 3 years were 58% and 37%, respectively. Warlick et al described outcomes in 306 CML patients of age  $\geq 40$  years undergoing RIC allo-HCT and reported to CIBMTR.<sup>12</sup> Of these, 38% patients were aged 40 to 49 years, 39% were 50 to 59 years, and 23% were  $\geq 60$  years. The 3-year OS (54%, 52%, and 41%) and 1-year NRM (18%, 20%, and 13%) were similar across age groups. LFS and relapse were similar across age groups in CP1 patients.

Our study makes several important observations. Patients with AP and CP2+ at allo-HCT had higher mortality compared with those in CP1, which could be explained by the advanced disease outstripping the ensuing graft-versus-leukemia effect early in the course. In addition, LFS did not differ significantly between the 2 conditioning intensities. The analysis also failed to show any difference in NRM between the 2 cohorts. RIC was associated with 85% greater risk of early after allo-HCT relapse but did not have a statistically significant difference beyond 5 months after allo-HCT. The higher early relapse risk with RIC allo-HCT could be explained by the modest cytoreduction engendered by the conditioning, enabling early relapse. However, the increased early relapse risk in the RIC cohort did not translate into worse OS due to potentially successful salvage treatment with TKIs and/or donor lymphocyte infusion (DLI) in those relapsing. Furthermore, in multivariable analysis, adjusting for many factors, including in vivo T-cell depletion, the cumulative incidence of cGVHD was notably lower in the RIC cohort than the MAC cohort in the first year after allo-HCT but lost statistical significance by 3 years. We can speculate that the use of nonmyeloablative (18%) and reduced-intensity conditioning (82%) may be associated with lower incidence of cGVHD, through potentially less tissue injury and cytokine release compared with MAC, as was also reported in BMT CTN 0901 study,<sup>13</sup> although the strength of evidence in support of this argument would be considered weak based on other published studies.<sup>33-37</sup> It is also possible that a higher frequency of TKI use after RIC allo-HCT for treatment or prevention of CML relapse may have curbed the cGVHD risk.<sup>38-40</sup> In multivariable analysis, no significant interaction was found between conditioning intensity and disease status at the time of allo-HCT. The



**Figure 1. Outcomes after MAC versus RIC alloHCT for CML.** (A) Adjusted curves for NRM after allo-HCT for CML using MAC vs RIC. (B) Adjusted curves for disease relapse after allo-HCT for CML using MAC vs RIC. (C) Adjusted curves for LFS after allo-HCT for CML using MAC vs RIC. (D) Adjusted curves for OS after allo-HCT for CML using MAC vs RIC.

survival outcomes in patients who were in AP or CP2+ at allo-HCT were not significantly different between the conditioning cohorts. Furthermore, no significant interaction was detected between conditioning intensity and other covariates such as age, KPS, HCT-CI, donor type, donor-recipient CMV status, graft source, in vivo T-cell depletion, year of HCT, and time from diagnosis to HCT; this has been demonstrated in a forest plot (Figure 2).

This observational study has limitations. The major limitation of this analysis is that the information on CML therapies (including TKIs and DLI) used after HCT was not available. The use of TKIs as maintenance therapy or for treatment of relapse/progression of CML and of DLI preemptively or for salvage, thereby preventing relapse and/or improving OS in relapsed CML patients in the RIC cohort, is certainly within the realm of possibility. Having 5 Food and Drug Administration–approved TKIs in the United States by 2012 suggests that the great majority of study patients had a decent probability of access to effective TKIs after HCT, and not having that data is a serious limitation. Nonetheless, the study aimed at evaluating the impact of conditioning intensity on allo-HCT outcomes and did not intend to examine the role of prophylactic or maintenance after allo-HCT TKIs. Even though centers have been employing maintenance TKIs,<sup>41-44</sup> a recent

CIBMTR study demonstrated no significant impact of maintenance TKI therapy on risk of cGVHD, NRM, relapse, LFS, and OS following allo-HCT in CML patients.<sup>45</sup> Patients who underwent MAC and RIC alloHCT between 2007 and 2014, and received TKI pre-HCT were included in that study. A landmark analysis was performed at day 100 after HCT. A total of 390 patients were included in the analysis (TKI maintenance, n = 89; no TKI maintenance, n = 301). A major proportion of both cohorts underwent MAC allo-HCT (85% in TKI cohort and 83% in no TKI cohort). With so few RIC patients, the study findings cannot be interpreted as lack of evidence in favor of maintenance TKIs after RIC allo-HCT as a relapse risk-reduction strategy.

An additional caveat of the study is that information on the depth of response prior to allo-HCT (ie, cytogenetic and/or molecular response to therapy) was not available. Having this useful data may further enable demonstration of the impact of the depth of response on outcomes and interaction with conditioning intensity and therefore, may theoretically play a role in the conditioning selection process. In addition, a potential demerit of the study is that the conditioning intensity (MAC vs RIC) selection criteria were not known; we can speculate that the treating physician made the decision to use a specific conditioning regimen based on certain clinical variables, the prevalent institutional guidelines,

**Table 3. Adjusted probabilities of outcomes in CML patients aged 18 to 60 years undergoing allo-HCT using MAC vs RIC, 2007-2014**

Outcomes	MAC (n = 1204)		RIC (n = 191)		P
	No. of patients (evaluable)	Probability (95% CI), %	No. of patients (evaluable)	Probability (95% CI), %	
<b>OS, y</b>	1204		191		
1	831	71 (68-73)	133	72 (66-78)	.61
3	501	58 (55-60)	86	58 (51-65)	.91
5	272	53 (50-56)	49	53 (45-60)	.98
<b>LFS, y</b>	1190		191		
1	731	63 (60-66)	106	58 (51-65)	.15
3	423	50 (47-53)	70	46 (39-53)	.31
5	222	44 (41-47)	40	43 (36-51)	.81
<b>Relapse, y</b>	1190		191		
1	731	16 (14-18)	106	21 (16-27)	.10
3	423	23 (21-26)	70	25 (19-31)	.49
5	222	26 (23-28)	40	25 (19-31)	.96
<b>NRM, y</b>	1190		191		
1	731	21 (18-23)	106	19 (14-24)	.50
3	423	28 (25-30)	70	26 (20-33)	.71
5	222	32 (29-35)	40	29 (22-36)	.53
<b>cGVHD, y</b>	1182		188		
1	316	50 (47-53)	64	41 (33-48)	.02
3	122	58 (55-61)	27	51 (43-58)	.08
5	57	59 (56-62)	16	51 (44-59)	.07

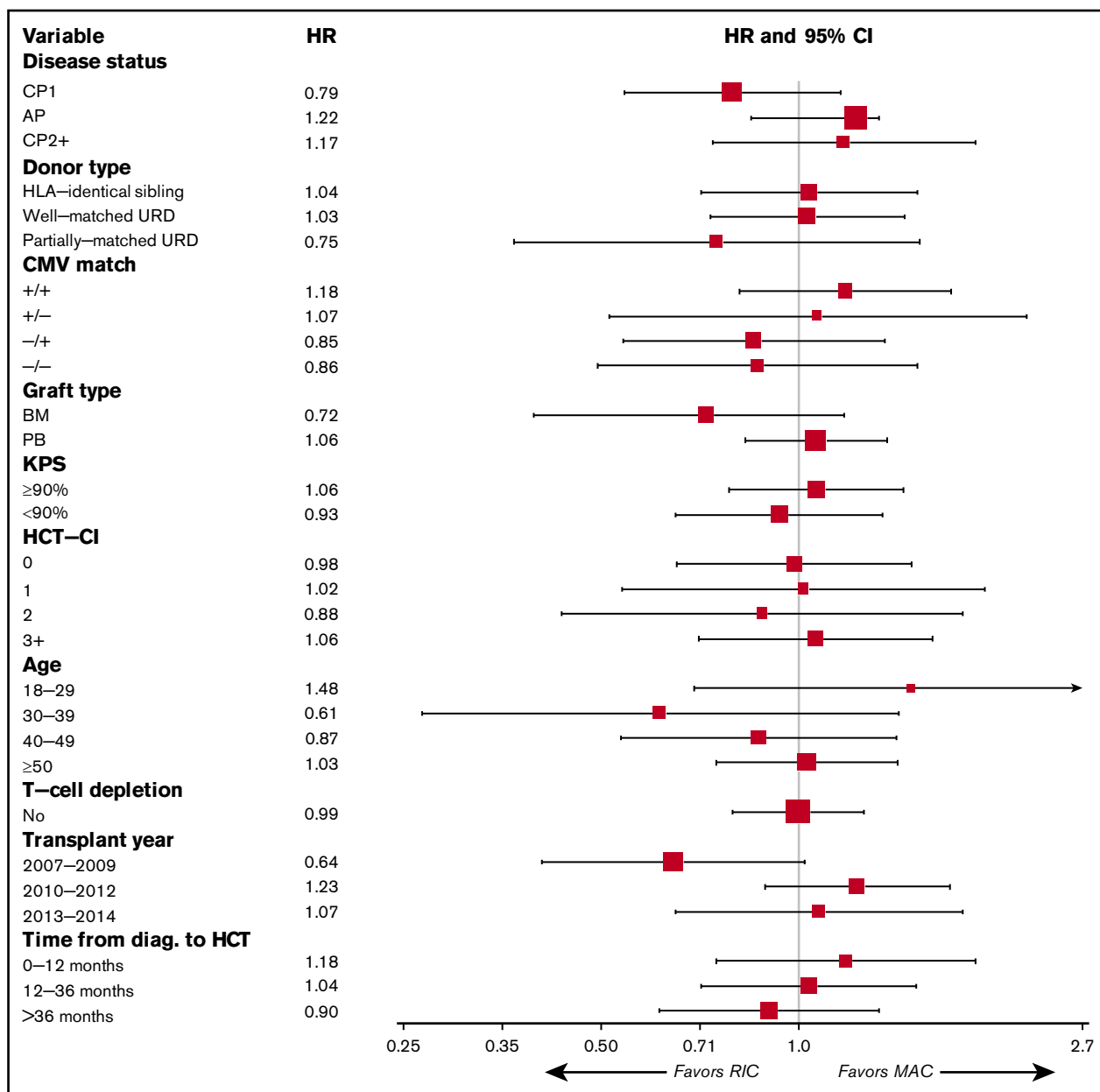
and the available clinical trials. The only way to prevent the selection bias would be to conduct a prospective randomized study. We tried to adjust for confounding caused by the different variables by taking them into account in the multivariable analysis, but the possibility of residual confounding remains. Finally, we analyzed the data in patients who underwent the allo-HCT but could not ascertain the justification for it in all patients. We can safely assume that the decision to proceed to allo-HCT was appropriate in those who were diagnosed with CML in AP or BP, but the reason for allo-HCT in others is not apparent in the data forms.

In summary, the current study represents the comparison of CML patients undergoing MAC and RIC allo-HCT in the era of TKIs and supports the notion that RIC allo-HCT is an appropriate alternative to MAC for CML. However, the lack of difference in OS between the MAC and RIC allo-HCT in the analysis is not to be construed as an argument in favor of selecting RIC over MAC for CML patients. On the contrary, the analysis revealed that compared with MAC, RIC allo-HCT afforded similar survival, albeit at the expense of increased early post-allo-HCT relapse risk. It is paramount to recognize that the intent of the study was not to identify a superior conditioning intensity between MAC and RIC but to demonstrate that OS and LFS in CML patients who are unable to proceed to MAC allo-HCT on clinical grounds are similar, should they receive RIC allo-HCT. Examining the interactions between conditioning intensity and several clinical variables did not suggest any advantage for RIC over MAC or vice versa to help make a recommendation on selection of conditioning intensity in CML patient subgroups based on these clinical variables. To conclude,

this study is significant in light of the fact that a confirmation of the findings through a prospective randomized controlled trial is not likely to be attempted in the current times, where patients have

**Table 4. Multivariable analysis for patients aged 18 to 60 years in AP/CP2+ undergoing allo-HCT using MAC vs RIC for CML, 2007-2014**

	n	HR	95% CI	P
<b>cGVHD</b>				
MAC	668	1		
RIC	97	0.83	0.60-1.15	.26
<b>Relapse ≤5 mo</b>				
MAC	672	1		
RIC	98	2.06	1.31-3.23	.002
<b>Relapse &gt;5 mo</b>				
MAC	503	1		
RIC	99	0.46	0.20-1.05	.06
<b>NRM</b>				
MAC	672	1		
RIC	98	1.32	0.89-1.94	.17
<b>LFS</b>				
MAC	672	1		
RIC	98	1.27	0.97-1.66	.08
<b>OS</b>				
MAC	683	1		
RIC	98	1.18	0.88-1.57	.26



**Figure 2.** Prespecified subgroup analysis according to baseline characteristics showing HRs and 95% CIs for OS after allo-HCT for CML patients using MAC vs RIC, 2007-2014.

access to an increasing number of potent TKIs and there is a possibility of long-term remission with treatment-free intervals and, consequently, fewer patients in need of allo-HCT.

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## Authorship

Contribution: S.C., K.W.A., Z.-H.H., S.J., and W.S. designed research; K.W.A. and Z.-H.H. collected data and performed statistical analysis; S.C., K.W.A., Z.H.H., S.J., and W.S. interpreted data; S.C.

drafted the manuscript; and K.W.A., Z.-H.H., S.J., A.A., J.C., E.A.C., A.D., Z.D., S.M.G., R.P.G., S.G., B.K.H., G.C.H., J.W.H., Y.I., A.S.K., H.J.K., H.M.L., M.R.L., S.N., R.F.O., A.P., O.R., J.M.R., A.S., B.N.S., H.C.S., S.S., N.N.S., M.S., R.K.S., C.U., A.E.W., J.A.Y., E.P.A., M.E.K., U.P., R.M.S., and W.S. critically reviewed and revised the manuscript.

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ORCID profiles: S.C., 0000-0001-9117-8696; J.C., 0000-0002-6602-5505; S.M.G., 0000-0002-3255-8143; S.N., 0000-0001-7920-4265; A.S., 0000-0003-0003-0130.

Correspondence: Saurabh Chhabra, Division of Hematology/Oncology, Department of Medicine, Medical College of Wisconsin, 9200 W Wisconsin Ave, Milwaukee, WI 53226; e-mail: schhabra@mcw.edu.

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