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# Prospective Validation of the Predictive Power of the Hematopoietic Cell Transplantation Comorbidity Index: A CIBMTR<sup>®</sup> Study

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# Abstract

Prospective validation of the hematopoietic cell transplantation-comorbidity index (HCT-CI) using contemporary patients treated with HCT across the Unites States is necessary to confirm its widespread applicability. We performed a prospective observational study including all patients (8115 recipients of allogeneic and 11,652 recipients of autologous HCT) who underwent first HCT that was reported to the CIBMTR between 2007 and 2009. In proportional hazards models, increased HCT-CI scores were independently associated with increases in hazard ratios for NRM (p<0.0001) and overall mortality (p<0.0001) among recipients of allogeneic HCT. HCT-CI Scores of 3 were uniformly associated with higher risks for outcomes in both allogeneic and autologous HCT, and all subgroups regardless of diagnoses, age, and conditioning intensity. Recipients of allogeneic HCT with scores of 1–2 who were aged <18 or were treated with lower intensity conditioning regimens had similar outcomes compared to those with score 0. Higher risks for

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overall mortality, but not for NRM, were observed among recipients of autologous HCT with scores of 1–2 versus 0. Our results confirm the validity the HCT-CI in both allogeneic and autologous HCT. The index should be used as a valid standard-of-care health measure in counseling patients for HCT, in clinical trial design, and in adjusting outcome analyses.

#### Keywords

Bone marrow transplantation; hematopoietic cell transplantation; autologous; allogeneic; HCT-CI; comorbidities; validity; inter-rater reliability

#### Introduction

Organ dysfunctions (comorbidities) impact both treatment decisions and treatment outcomes in oncology, and is particular salient in hematopoietic cell transplantation (HCT), where the morbidity associated with the procedure is high.[1] Until 2004, age alone had been widely used as the primary measure of a patient's ability to tolerate the conditioning regimens for allogeneic HCT.[2] Recently, the HCT-CI was designed as a health measure suited for capturing the burden and complexity of organ dysfunctions among recipients of allogeneic HCT. The index was modeled to predict non-relapse mortality (NRM) and initial analysis validated its ability to discriminate risks for NRM as well as overall mortality in an independent randomly selected cohort from the same institution.[3] Subsequently, comorbidity evaluation integrated in transplant-related analyses have demonstrated the importance of risk assessment prior to HCT[4–7] or even conventional therapies[8–11], and to better select patients for different regimen intensities.[12,13] Additional studies suggested that comorbidities may have more important role than calendar age in determining HCT eligibility.[14]

While some investigators have confirmed the prognostic significance of the HCT-CI in their respective patients, [15–18] others did not confirm the importance of the HCT-CI.[19,20] Therefore, it became important to study the prognostic significance of the HCT-CI in a prospective, well designed, multi-center setting to confirm its utility as a prognostic health-status measure of HCT outcomes. Further, there has been only limited number of studies that assessed the performance of the HCT-CI in the autologous HCT settings.[21,22] If the utility of the HCT-CI is confirmed in adequately designed large validation studies, this index would allow for consistent integration of comorbidities into the design of randomized clinical trials in HCT, adjustment of clinical trial results across transplant institutions, and better understanding of the biological causes of post-HCT morbidities.

We hereby summarize the results from a large multi-institutional prospective study gathering information from all United States transplant centers that report to the Center of International Blood and Marrow Transplantation Research (CIBMTR). The study was aimed to determine the discriminative capacity of the HCT-CI among recipients of allogeneic and autologous HCT, and the effectiveness of the HCT-CI in stratifying outcomes among HCT patients with different diagnoses, age groups, and conditioning intensities.

# **Patient and Methods**

#### Data Source

The CIBMTR<sup>®</sup> (Center for International Blood and Marrow Transplant Research<sup>®</sup>) is a research affiliate of the International Bone Marrow Transplant Registry, Autologous Blood and Marrow Transplant Registry, and the National Marrow Donor Program (NMDP) established in 2004. It comprises a voluntary working group of more than 450 transplantation centers worldwide that contribute data on consecutive allogeneic and autologous HCT procedures to a statistical center at the Medical College of Wisconsin in Milwaukee and the NMDP Coordinating Center in Minneapolis, Minnesota. Participating centers report longitudinal data on all transplants and compliance is monitored by on-site audits. 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 used in the performance of such research is collected and maintained in CIBMTR's capacity as a Public Health Authority under the Health Insurance Portability Act Privacy Rule.

# Study Design and Patients

In 2007, a new prospective multi-institutional observational study was initiated at the CIBMTR to collect comorbidities from all transplant centers by their respective evaluators and to validate the predictive power of the HCT-CI in a large sample of patients. The HCT-CI was adapted into the Pre-Transplant Essential Data (pre-TED) collection form #2400. Data managers from all institutions attended an education session on comorbidity coding per the HCT-CI at the 2007 Tandem BMT Meeting in Keystone, Colorado. This session was then made public to all data managers at the CIBMTR website "www.cibmtr.org/Meetings/Materials/CRPDMC/Pages/feb2007sorror.aspx".

The study was designed to score comorbidities prospectively for all patients meeting the following criteria: 1) diagnoses of hematological malignant diseases, 2) treatment with autologous or allogeneic HCT between December 1<sup>st</sup> 2007 and December 31<sup>st</sup> 2009, 3) receiving conditioning regimens of any intensity or composition, 4) receiving grafts from HLA-matched related or unrelated donors, and 5) given marrow or granulocyte-colony stimulating factor-mobilized peripheral blood mononuclear cells (G-PBMC) grafts. No upper limit was stated for the number of patients to be enrolled into the study. Figure 1 is an organization chart depicting patient eligibility and enrollment into the study. Among 23,876 recipients (Figure 1) of first HCT in the United States between 2007 and 2009 who were reported to CIBMTR, final samples of 8,115 recipients of allogeneic HCT and 11,652 recipients of autologous HCT contributed to this analysis.

## Study end points and definitions

The primary outcomes studied were non-relapse (NRM) and overall mortality. NRM was defined as post-transplantation death that was not preceded by disease progression or relapse. Progression was defined as >50% increase in the burden of primary disease compared to pre-transplant disease status and/or development of disease at new sites.

Relapse was defined as reappearance of primary disease following achievement of post-HCT complete remission. For survival, patients were considered to have an event at time of death from any cause; survivors were censored at last contact. Conditioning regimens were classified into high-dose, reduced-intensity (RIC), or nonmyeloablative (NMA) intensity based on the previously published criteria.[23] Comorbidities were evaluated by respective staff at each site, while total scores were assigned by the CIBMTR statistical team following previously published guidelines.[3] The HCT CI score was derived directly from the presence of a comorbidity per the HCT-CI as collected in the TED forms. Additional comorbidities that were not part of the HCT-CI, but collected in free text fields under the "other" category, were not considered for the validation of this score. We analyzed a subset of these "other comorbidity" fields to assess discrepancies between what was documented in the free text field and the HCT CI components. We found that the content in this free text field could potentially change the overall HCT CI score in less than 5% of cases. Consequently, the "other comorbidity" field was not used to modify the score reported in the HCT-CI specific fields. To further rule out the contribution of these write-in entries, patients with a HCT CI score of 0 but with any "other comorbidity" reported in the free text field were analyzed as a separate risk group in the statistical models.

## Statistical methods

Cumulative incidence and Kaplan Meier estimates were calculated to evaluate the unadjusted associations between the HCT-CI scores and NRM and survival, respectively. Relapse or progression of the primary disease was treated as a competing risk for NRM and vice versa. Because this study investigates the impact of the HCT-CI on outcomes following the first transplant, all outcomes were censored at the second transplant.

Proportional hazards models were used to estimate the hazard ratio (HR) for NRM and survival associated with HCT-CI scores among the whole patient population as well as among adults versus children, high-dose versus RIC/NMA regimens, and among patients with different diagnoses. The models were adjusted for patient-related risk factors: age, Karnofsky performance status (KPS) score, race, and cytomegalovirus (CMV) serology results; disease-related risk factors: diagnosis category, sensitivity to last chemotherapy among patients with lymphoma, and disease status among patients with acute leukemia, and interval between diagnosis and HCT; and transplant-related risk factors: donor type/HLA typing, stem cell source, conditioning regimen, and GVHD prophylaxis regimen. Multivariate p-values for a variable were based on adjustment for all other variables in the model. All p-values were derived from likelihood ratio statistics and were two-sided. In these multivariate analyses, the HCT-CI was primarily modeled as a categorical variable with group stratifications of 0, 1-2, and 3 similar to the initially recommended model to allow for almost uniform distribution of patient samples per risk group. A subset analysis using categorization of 0, 1, 2, 3, 4, and 5 was also performed with nested comparisons of both stratification models.

Three samples of 58, 80, and 77 patients were selected randomly from patients reported from Dana Farber Cancer Institute, Roswell Park Cancer Institute, and Fred Hutchinson Cancer Research Center to assess the magnitude of inter-rater reliability. These samples of

patients were re-evaluated for comorbidity coding and score assignment by the study coinvestigators V.T.H., P.L.M, and M.L.S, respectively. Score assignments were compared between two raters from each institution; one that collected original data reported to the CIBMTR and another that assigned scores independently.

Kappa statistic is a measure used to analyze inter-rater agreement,[24,25] and it adjusts for the degree of agreement that would be expected to occur by chance, and is therefore more appropriate than Pearson's product moment, Spearman's correlation, or percent agreement. [26] It is reported from 0.0 to 1.0. Weighted Kappa statistic (Kw),[27] which assigns less weight to agreement as risk categories are further apart, was computed with Fleiss-Cohen weights[28] to analyze the magnitude of inter-rater agreement between two raters on assignment of patients to the HCT-CI risk-categories of 0–1, 2, 3, and 4. Standard errors (S.E.) for kappa and Kw statistics were calculated as previously described.[29] Weighted Kw has been used to compare agreement between two raters.[27] The Landis scale was used for interpretation of the magnitude of Kw statistics where values <0 indicate no agreement; 0.0–0.20, slight; 0.21–0.40, fair; 0.40–0.60, moderate; 0.61–0.80, substantial; and 0.81–1.00, almost perfect agreement, respectively.[30]

## Results

#### **Patient Characteristics**

Patient-, disease-, and transplant-related characteristics are described in Table 1 for recipients of high-dose allogeneic (n=5,460), RIC/NMA allogeneic (n=2,655), and autologous HCT (n=11,652). Median (range) ages were 45 (<1-74), 59 (1-78), and 56 (<1-83) years, respectively. Median (range) intervals between diagnoses and HCT were 7(<1-377), 13 (1–347), and 11 (<1–389) months, respectively. Cyclophosphamide (Cy) combined with high-dose total body irradiation (TBI, 33%) and busulfan (Bu) combined with Cy (26%) or fludarabine (20%) were the most frequently used regimens among recipients of high dose conditioning, while fludarabine (Flu) combined with Bu (31%), melphalan (20%), or low-dose TBI (23%) were the most frequent regimens among recipients of RIC/NMA conditioning. Methotrexate combined with tacrolimus was the most frequently used graftversus-host disease (GVHD) prophylaxis regimen among recipients of both high-dose (53%) and RIC/NMA regimens (41%). Other frequent GVHD prophylaxis regimens included MTX/cyclosporine (CSP, 15%) among high dose regimens, and mycophenolate mofetil combined with CSP (16%) or tacrolimus (15%) among the RIC/NMA regimens. Among recipients of autologous HCT, the most common conditioning regimens were melphalanbased (50%) and carmustine, etoposide, cytarabine, and melphalan combination (BEAM, 25%).

#### Prevalence of Comorbidities and Distribution of Comorbidity Scores

Overall, comorbidities were common among recipients of RIC/NMA allogeneic HCT when compared with those of high-dose allogeneic and autologous HCT. Patients with scores 0 were found in 41% compared to 52% and 51%, respectively, while those with scores of 1–2 and 3 were found in 28% and 31% compared to 25% and 23%, and 27% and 22%, respectively. Overall, pulmonary comorbidities were the most prevalent among the three

groups of patients (26% compared to 22% and 21%, respectively) followed by psychiatric (14% compared to 13% and 11%, respectively) and combined cardiac comorbidities (18% compared to 9% and 12%, respectively) that included arrhythmia, heart failure, low ejection fraction, and heart valve disease (Figure 2). Overall, 11% samples within each of the three groups were reported as having other comorbidities that did not acquire a score per the HCT-CI.

#### Validation of the HCT-CI among recipients of allogeneic HCT

In Cox regression models adjusted for other co-variables, HCT-CI scores of 1–2 and 3 were statistically significantly associated with increased risks for NRM and overall mortality compared to score 0 (Table 2 and Figure 3A). Patients with scores of 0, 1–2, and 3 had 1-year probabilities of NRM of 17%, 21%, and 26% (p<0.001) and overall survival of 69%, 62%, and 56%, respectively, (p<0.001). The corresponding figures for 3-year NRM were 24%, 28%, and 35% (p<0.001) and for 3-year overall survival were 54%, 47%, and 38%, respectively (p<0.001). When the HCT-CI was categorized into 0, 1, 2, 3, 4, 5, each increased score was statistically significantly associated with higher risks for NRM and overall mortality with the exception of the associations between scores 1 and 2 with NRM that did not reach statistical significance (Table 2 and Figure 3B and C). Figure 4A and B demonstrated increasing probabilities of NRM and decreasing overall mortality, respectively, with each increment in HCT-CI scores. A nested comparison between the two categorization models suggested that the later had statistically significant better stratification power for both NRM (p=0.005) and overall mortality (p<0.001).

Overall, patients who were assigned score 0 with versus without "other comorbidities that are not included within the HCT-CI" reported similar risks for NRM and overall mortality.

#### Performance of the HCT-CI among Subgroups of Recipients of Allogeneic HCT

In Cox regression models adjusted for other co-variables, recipients of high-dose conditioning and allogeneic HCT with HCT-CI scores of 1–2 and 3 had statistically significantly higher risks for NRM and overall mortality compared to those with score 0 (Figure 5A). Likewise, patients with scores of 1–2 and 3 experienced respectively higher probabilities of NRM and overall survival when conditioned with high-dose regimens (Figures 5C and D). By contrast, patients with scores of 1–2 tolerated RIC/NMA regimens equally well compared to those with score 0, but those with scores of 3 experienced higher risks and probabilities for NRM and overall mortality compared to score 0 (Figures 5B, E, and F).

All recipients of allogeneic HCT were categorized into children and adults. Among adults, scores of 1–2 and 3 were associated with statistically significantly increased risks for NRM and overall mortality compared to score 0 (Appendix 2A). Children with HCT-CI scores of 3 were associated with statistically significant higher risks for NRM and overall mortality compared to score 0, while the higher hazard ratios (HRs) associated with scores of 1–2 did

not reach statistical significance (Appendix 2B).

Increasing HCT-CI scores were also associated with increased risks for NRM and overall mortality among patients diagnosed with myeloid or lymphoid malignancies (Appendix 3).

#### Validation of the HCT-CI among recipients of autologous HCT

Recipients of autologous HCT with scores of 1–2 had statistically non-significant higher HRs for NRM but statistically significant higher HRs for overall mortality compared to those with score 0 (Table 2). Patients with scores of 3 experienced statistically significantly higher HRs for both outcomes compared to those with score of 0. Nested comparison between this stratification system and a more detailed one (0, 1, 2, 3, 4, and 5) did not show additional benefit in regards to prediction of NRM (p=0.297) or overall mortality (p=0.433).

Probabilities of NRM at 1-year were 3%, 3%, and 5% (p<0.001) for those with scores of 0, 1–2, and 3; while the figures for 3-year probabilities were 5%, 6%, and 9%, respectively (p<0.001). One-year and 3-year rates of survival were 91%, 88%, and 86% (p<0.001), respectively and 79%, 73%, and 70% (p<0.001), respectively (Figures 4C and D).

# Performance of the HCT-CI among recipients of autologous HCT Diagnosed with Lymphoma versus Multiple Myeloma

Similar to the general population of recipients of autologous HCT, the associations between scores 1–2 among those with lymphoma or multiple myeloma only reached statistical significance for overall mortality but not for NRM (Appendix 1). Alternatively, scores of 3 were uniformly associated with higher HRs for both outcomes in both groups of patients. Day 100 NRM for the 3 HCT-CI score groups were 1%, 2%, and 3% (p<0.001), respectively among patients with multiple myeloma and 3%, 4%, and 6% (p<0.001), respectively, among those with lymphoma. At 3-years, NRM probabilities were 4%, 6%, and 7% (p=0.007), respectively, for myeloma patients, while survival rates were 84%, 76%, and 74%, respectively (p<0.001). Among lymphoma patients, the 3-year NRM probabilities were 5%, 7%, and 10% (p<0.001), respectively, while survival rates were 77%, 72%, and 67% (p<0.001), respectively.

### Reliability of the HCT-CI

Weighted kappa statistic estimates were 0.54 in DFCI, 0.81 in RPCI, and 0.47 in FHCRC data samples, suggesting a fair-moderate agreement rate among evaluators.

# Discussion

This prospective, multi-center study has generated several key findings about the performance of the HCT-CI as a prognostic comorbidity model for HCT outcomes. First, The HCT-CI stood the test of time as it was shown here to predict outcomes among a group of patients treated about almost a decade after those who originally contributed to its design. Second, it was valid in predicting both NRM and survival among recipients of both allogeneic and autologous HCT. The ability of the HCT-CI to predict NRM after autologous HCT is an important finding, especially considering that the idnex was originally designed based on data from a cohort of allogeneic HCT recipients, and considering the substantial

differences between both transplant strategies. Other comorbidity indices have also been shown to be useful in settings beyond those from which they were developed.[2,31,32] Third, we found that patients with score 0 plus additional comorbidities coded under "other", in aggregate, did not influence outcomes compared to patients with scores of 0 alone. These results confirm the original design of the index that dropped most of these comorbidities from consideration due to lack of statistical association. Fourth, the index in this prospective contemporary patient cohort demonstrated sensitive stratification of outcomes that varied based on criteria of age or conditioning intensity. Specifically, patients undergoing RIC/NMA allogeneic HCT or children tolerated allogeneic HCT equally well when they had scores of 1–2 versus 0. This confirms the benefit of the style that was used to build the HCT-CI, where its association with increased risks of NRM was meant to be a range of values that would vary based on other variables such as conditioning intensity, age, or disease-risk. Finally, the index performed well among subgroups of diagnoses, age categories, and conditioning intensities. The findings of this large study, in conglomerate, have affirmed the adaptability and integrity of the HCT-CI in the real world HCT setting.

Since its development, the HCT-CI has been tested in a number of retrospective analyses with conflicting results. Many of these retrospective analyses suffered from limited sample size, lack of complete comorbidity data, and apparent inaccurate coding of comorbidities. [19,20,33–35] There has been an unmet need for prospective evaluation of the comorbidity impact in a well-designed and appropriately powered study. A recent prospective study from Italy confirmed validity of the index.[36] Our study is the first prospective study to evaluate the performance of the HCT-CI among a large number of patients treated at transplant centers across the US. The proven value of the HCT-CI in the current study should encourage investigators and community physicians to incorporate comorbidity assessment in their daily practice.

The prospective nature of this study, together with the inclusion of large groups of patients from various transplant centers, would ensure generalizability of the study findings. In 3 randomly selected samples, we have found that the rate of agreement between evaluators ranged from fair to moderate. Variable IRR is a common problem among comorbidity indices[37–39] and it was recently underscored for the HCT-CI promoting the production of comprehensive guidelines for comorbidity coding.[40] The guidelines were summarized in a web-based application (www.hctci.org ), and has been validated to improve the IRR among novice evaluators to an excellent magnitude.[40] It is interesting that the fair-moderate degree of IRR in the current study did not obscure the validity of the index in predicting outcomes. Still, it would be important to use the new guidelines consistently in order to standardize comorbidity coding across institutions, which is critical when using the index to adjust comparisons of center performances or to test new associations with outcomes. Nested comparison analyses showed that each digit increase in the score of the HCT-CI between 0- 5 was associated with statistically significant increases in risks for NRM and survival among overall recipients of allogeneic HCT. Nevertheless, the stratification of 0, 1-2, 3 is preferable for patient counseling and assessment of outcomes at relatively smaller studies given the ease of use and given the limitations in sample size outside of registry studies.

The prospective validation of the HCT-CI as achieved in this study should promote consistent use of this index in the stratification of patients in future randomized HCT studies. The index could also be incorporated as a variable to adjust comparisons of outcomes and performances across transplant centers. This strategy is already being used by the Center of International Blood and Marrow Transplantation Research (CIBMTR) in the determination of the center specific outcomes comparisons, and should prove valuable to patients, insurers, and investigators. Finally, the HCT-CI would be helpful to clinicians in counseling potential HCT recipients about their risks of NRM, and choosing the appropriate conditioning regimen.

Patients with score 1–2 has similar NRM risks to those with score 0 when patients with children or when they received either autologous HCT or allogeneic HCT after reduced-intensity/nonmyeloablative regimens. These results validate findings from previous studies about the performance of the HCT-CI in these specific cohorts of patients.[41,42] These results also highlight the sensitivity of the index in differentiation between adults versus children and recipients of high-intensity versus lower intensity regimens in regards to tolerability of HCT. This differentiation is important in the clinic to help transplant physicians to decide between the variable options of transplant strategies. Patients with HCT-CI scores of 3 consistently have higher risks for NRM and overall mortality, compared to those with scores of 0, regardless of transplant strategy, conditioning intensity, diagnoses, or age groups.

In the future, we could achieve finer discrimination of outcomes by combining the HCT-CI with other relevant metrics. This could be specifically important for vulnerable patients such as those of age 60 years or older. For example, one could potentially further stratify risks for mortality by combining the HCT-CI scores with biomarkers,[6,33,43] performance status, [44] and/or some components of a geriatric assessment (GA) for older patients.[45] The recent Bone Marrow Transplant-Clinical Trial Network (BMT-CTN)-State of the Science Symposium (SOSS) has suggested a prospective study to develop a novel risk assessment tool comprising of the HCT-CI, performance status model, biomarkers and GA to improve risk-assessment prior to HCT.[46] In summary, our study confirms that the HCT-CI is a powerful tool for predicting NRM and survival after HCT. Co-morbidity assessments should be applied in future research and clinical care of HCT recipients. This is of particular importance considering the increasing age and vulnerability of the population eligible for HCT in the United States.

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# Here are the Highlights for this manuscript as requested

- 1. High HCT CI is linked to mortality after allogeneic and autologous transplants
- 2. HCT CI as a prognostic tool is validated for all transplants for malignancies
- 3. Only HCT CI 3 were prognostic for mortality in children and in RIC HCT

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# Figure 1. Organization chart of patient eligibility and enrollment into the prospective observational study

Among a total sample of 23,876 patients who received hematopoietic cell transplantation in United States between 12/2007 and 12/2009, 8,115 recipients of allogeneic and 11,652 recipients of autologous HCT contributed to the study analyses.



Figure 2. Prevalence of comorbidities as captured by the HCT-CI among recipients of high-dose allogeneic, reduced-intensity/nonmyeloablative allogeneic, and autologous HCT Pulmonary, psychiatric and heart comorbidities were the most prevalent among all three groups of patients

Figure 3A







Figure 3C







**Figure 3. Illustration of the independent associations between HCT-CI score groups and risks of NRM and overall mortality among recipients of HCT using Cox regression models** Among recipients of allogeneic HCT, (Figure 3A) HCT-CI scores of 1–2 and 3 were statistically significantly associated with increased risks for NRM and reduced overall mortality compared to score 0. In When the HCT-CI was categorized into 0, 1, 2, 3, 4, 5, for prediction of (Figure 3B) NRM and (Figure 3C) overall mortality among recipient of allogeneic HCT, each increased score was statistically significantly associated with higher risks for these outcomes with the exception of the associations between scores 1 and 2 with NRM that did not reach statistical significance. Among recipients of autologous HCT, HCT-

CI score stratification of 1–2 and 3 was used for prediction of risks for NRM and overall mortality in (Figure 3D).

Figure 4A



Figure 4B



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Figure 4. Stratification of probabilities of outcomes by the HCT-CI scores among recipients of allogeneic or autologous HCT

Among recipients of allogeneic HCT (Figures 4A and 4B), HCT-CI scores of 0, 1–2, 3, 4, and 5 stratified well probabilities of (Figure 4A) NRM and (Figure 4B) survival. Among recipients of autologous HCT, HCT-CI scores of 0, 1–2, 3 stratified well probabilities of (Figure 4C) NRM and (Figure 4D) survival.





Figure 5B





Figure 5D









Figure 5. Subgroup validation of the predictive power of the HCT-CI among recipients of allogeneic HCT following high-dose versus reduced-intensity/nonmyeloablative conditioning regimens

**Figures 5A and 5B demonstrate results from** Cox regression models focusing on NRM and overall mortality. HCT-CI scores of 1–2 and 3 were statistically significantly associated with increased risks for NRM and overall mortality among recipients of (Figure 5A) high-dose conditioning regimens, while only scores of 3 were associated with the same outcomes among recipients of (Figure 5B) reduced-intensity/nonmyeloablative conditioning regimens compared to those with score of 0.

Figures 5C, 5D, 5E, and 5F demonstrate probabilities of TRM and survival. HCT-CI scores of 1–2 and 3 stratified well probabilities of (Figure 5C) NRM and (Figure 5D) overall survival among recipients of high-dose conditioning.

Only patients with HCT-CI scores of 3 experienced (Figure 5E) increased probabilities of NRM and (Figure 5F) decreased probabilities of survival among recipients of reduced-intensity/nonmyeloablative conditioning regimens compared to those with score of 0.





B: Allo Malignant Multivariate Analysis: Children, TRM and OS (HCT – CI=0, 1-2, 3+)



#### Appendix 2.

Illustration of the independent associations between HCT-CI scores of 0, 1–2, and 3 and risks of NRM and overall mortality among recipients of allogeneic HCT, who were A) adults or B) children using Cox regression models.

A: Allo Malignant Multivariate Analysis: Myeloid, TRM and OS (HCT – CI=0, 1-2, 3+)



B: Allo Malignant Multivariate Analysis: Lymphoid, TRM and OS (HCT – CI=0, 1-2, 3+)



#### Appendix 3.

Illustration of the independent associations between HCT-CI scores of 0, 1–2, and 3 and risks of NRM and overall mortality among recipients of allogeneic HCT, who were diagnosed with A) myeloid or B) lymphoid malignancies using Cox regression models.

## Table 1

Characteristics of US patients who received an allogeneic or autologous HCT for malignant diseases between 2007 and 2009, registered with the CIBMTR

Characteristics of patients	High-dose n=5460 (%)	RIC/NMA n=2655 (%)	Autologous n=11,652 (%)
Age of patients			
>0-19	894 (16)	50 (2)	811 (7)
20-39	1315 (24)	191 (8)	1423 (13)
40-49	1179 (22)	327 (12)	1695 (15)
50–59	1438 (26)	853 (32)	3420 (29)
60	634 (12)	1224 (46)	4303 (37)
Race of patients			
Caucasian	4233 (78)	2274 (86)	9022 (77)
African-American	324 (6)	120 (5)	1301 (11)
Asian/Pacific Islander	181 (3)	76 (3)	289 (2)
Hispanic	636 (12)	148 (6)	920 (8)
Others	86 (2)	37 (1)	120 (1)
Karnofsky score, %			
< 90	1615 (30)	937 (35)	3752 (32)
90	3566 (65)	1578 (59)	7201 (62)
Missing	279 (5)	140 (5)	699 (6)
Disease			
Acute myelogenous leukemia	2391 (44)	926 (35)	268 (2)
Acute lymphoblastic leukemia	1228 (22)	134 (5)	21 (<1)
Other leukemia	148 (3)	364 (14)	8 (<1)
Chronic myelogenous leukemia	298 (5)	59 (2)	0
Myelodysplastic	600 (11)	344 (13)	0
Myeloprolifterative disorders	161 (3)	131 (5)	0
Lymphomas	481 (9)	652 (25)	4763 (41)
Myelomas	73 (1)	21 (1)	5717 (49)
Other Malignancies	80 (1)	24 (1)	875 (8)
AML/ALL disease status at transplant			
Never treated	29 (<1)	15 (1)	0
Primary Induction Failure	406 (11)	101 (10)	292 (99)
Complete Remission	2756 (76)	857 (81)	0
Relapse	426 (12)	85 (8)	3 (1)
Missing	2 (<1)	2 (<1)	0
Lymphoma disease status prior to HCT			
Sensitive	352 (73)	513 (79)	4335 (91)
Resistant	120 (25)	129 (20)	377 (8)

Characteristics of patients	High-dose n=5460 (%)	RIC/NMA n=2655 (%)	Autologous n=11,652 (%)
Unknown/untreated	9 (2)	10 (2)	51 (1)
Donor/recipient CMV status			
_/_	1548 (28)	675 (25)	-
+/+	1637 (30)	823 (31)	-
+/	596 (11)	327 (12)	-
_/+	1568 (29)	746 (28)	-
unknown	111 (2)	84 (3)	-
Graft type			
Marrow	1343 (25)	271 (10)	72 (<1)
G-PBMC	4117 (75)	2384 (90)	11580 (99)
Donor type			
HLA-identical sibling	2266 (42)	1012 (38)	-
Other related	257 (5)	225 (8)	-
Unrelated donor	2882 (53)	1399 (53)	-
Twins	51 (1)	19 (<1)	-
URD HLA match status			
8/8	1853 (64)	976 (70)	-
7/8	483 (17)	198 (14)	-
6/8	66 (2)	15 (1)	-
5/8	10 (<1)	1 (<1)	-
Missing	470 (16)	209 (15)	-
HCT-CI score			
0	2825 (52)	1088 (41)	5851 (50)
1	767 (14)	436 (16)	1714 (15)
2	610 (11)	320 (12)	1375 (12)
3	611 (11)	345 (13)	1306 (11)
4	341 (6)	217 (8)	673 (6)
5	296 (5)	246 (9)	667 (6)
Missing	10 (<1)	3 (<1)	66 (<1)

RIC indicates reduced-intensity conditioning; NMA, nonmyeloablative; AML, acute myeloid leukemia; ALL, acute lymphocytic leukemia; HCT, hematopoietic cell transplantation; CMV, cytomegalo-virus; G-PBMC, granulocyte colony stimulating factor-mobilized peripheral blood mononuclear cells; HLA, human leukocyte antigen; URD, unrelated donor; CI, comorbidity index.

# Table 2

autologous HCT. Three group stratification of HCT-CI scores predicted outcomes well among both cohorts of patients but the six group stratification Cox regression models for associations of HCT-CI scores with risks of non-relapse (NRM) and overall mortality among recipients of allogeneic and model performed better for the former patients

			NRM			Overall mortality	<b>v</b>
	HCT-CI scores	u	HR* (95% CI)	d	u	HR* (95% CI)	d
Recipients of allogeneic HCT	* overall			<0.0001			<0.0001
	0	2887	1.00		3026	1.00	
	0, but other comorbidity reported	826	0.93 (0.79–1.10)	0.385	866	0.96 (0.85–1.08)	0.474
	1–2	2036	1.12 (1.00–1.26)	0.053	2126	1.12 (1.03–1.22)	800.0
	3+	1936	1.47 (1.31–1.65)	<0.0001	2051	1.36 (1.25–1.48)	<0.0001
	* overall			<0.0001			<0.0001
	0	2887	1.00		3026	1.00	
	0, but other comorbidity reported	826	0.93 (0.79–1.10)	0.396	866	0.96 (0.85–1.08)	0.483
	1	1149	1.12 (0.97–1.28)	0.122	1199	1.13 (1.02–1.24)	0.017
	2	887	1.13 (0.98–1.32)	0.097	927	1.12 (1.00–1.24)	0.048
	3	903	1.31 (1.13–1.51)	<0.0001	953	1.22 (1.10–1.36)	<0.0001
	4	527	1.52 (1.28–1.80)	<0.0001	558	1.39 (1.23–1.57)	<0.0001
	5+	506	1.77 (1.50–2.10)	<0.0001	540	1.62 (1.43–1.83)	<0.0001
Recipients of autologous HCT	* overall			0.001			<0.0001
	0	4090	1.00		4621	1.00	
	0, but other comorbidity reported	1086	0.93 (0.67–1.28)	0.641	1229	0.85 (0.73-1.00)	0.047
	1–2	2811	1.16 (0.93–1.44)	0.200	3089	1.23 (1.11–1.37)	<0.0001
	3+	2385	1.49 (1.20–1.85)	0.000	2645	1.37 (1.23–1.52)	<0.0001

The Cox regression models were adjusted for diagnosis category, disease status for acute leukemia, chemo-sensitivity for lymphoma, donor type/HLA matching, stem cell source, KPS percentage, CMV serology status, conditioning regimen, GVHD prophylaxis regimen, recipient age, recipient race, and interval between diagnosis and HCT. Cox regression models for associations of HCT-CI scores with risks of non-relapse (NRM) and overall mortality among recipients of autologous HCT and within subgroups of lymphoma versus multiple myeloma.

HCT-CT scores         n         HR (95% CI           Lymphoma subgroup $*$ overall         1         1           Lymphoma subgroup $*$ overall         1         1           0         0         1243         1.00           1         0         0         10         1.00           1         0         0         1243         1.00           1         0         0         1243         1.00           1         1         0         0         1.24         0.85-1.7           1         1         1         0         977         1.24         0.86-1.7           1         1         1         2         1.24         0.88-1.7         1.24         0.88-1.7           1         1         1         1         2         1.24         0.88-1.7           Myeloma subgroup $*$ overall         1         1.24         0.88-1.7         1.26         1.25         1.16           Myeloma subgroup $*$ overall $*$ overall         1.56         1.26         1.20         1.20           Myeloma subgroup $*$ overall $*$ overall         1.65         1.00         1.00	INKIN		<b>Overall mortality</b>	y
Lymphoma subgroup $^{\circ oerall}$ $^{\circ oerall}$ $^{\circ oerall}$ $^{\circ oerall}$ $^{\circ 0}$	HR (95% CI) 1	u d	HR (95% CI)	Ρ
0         0         1243         1.00 $0, but other comorbidity reported         311 0.96 (0.57-1.6) 1-2 1-2 977 1.24 (0.88-1.7) 1-2 3+ 977 1.24 (0.88-1.7) 3+ 3 832 1.56 (1.12-2.1)           Myeloma subgroup         *_{overall} 832 1.56 (1.12-2.1)           Myeloma subgroup         *_{overall} 832 1.56 (1.12-2.1) 0, but other comorbidity reported         832 1.56 (1.12-2.1) 0, but other comorbidity reported         586 0.95 (0.59-1.5) 1-2 1-2 1.32 (0.93-1.8) $	)	.043		0.001
0, but other comorbidity reported         311 $0.96 (0.57-1.6$ $1-2$ $1-2$ $977$ $1.24 (0.88-1.7)$ $1-2$ $3+$ $832$ $1.56 (1.12-2.1)$ Myeloma subgroup         *overall $832$ $1.56 (1.12-2.1)$ Of $972$ $1.56 (1.12-2.1)$ $1.00$ Myeloma subgroup         *overall $1.57 (0.93-1.5)$ $1.00$ O $0.104$ or	243 1.00	1264	1.00	
	11 0.96 (0.57–1.61) (	318).863 318	0.99 (0.77–1.28)	0.959
3+         3-         1.56 (1.12-2.1           Myeloma subgroup         * overall         832         1.56 (1.12-2.1           Myeloma subgroup         * overall         1.00         1.00           0         0. but other comorbidity reported         586         0.55 (0.59-1.5)           1-2         1-2         1.32 (0.93-1.8)         1.32 (0.93-1.8)	77 1.24 (0.88–1.73) 0	988 988	1.23 (1.05–1.46)	0.013
Myeloma subgroup         * overall         1           0         1657         1.00           0, but other comorbidity reported         586         0.95 (0.59-1.5)           1-2         1.32         1.32 (0.93-1.8)	32 1.56 (1.12–2.17) (	.009 838	1.37 (1.15–1.62)	<0.0001
0         1.00           0, but other comorbidity reported         586         0.95 (0.59–1.5)           1-2         1.32         0.93–1.8	)	.048		<0.0001
0, but other comorbidity reported         586         0.95 (0.59–1.5)           1-2         1.32         1.32 (0.93–1.8)	557 1.00	1949	1.00	
1–2 1382 1.32 (0.93–1.8	36 0.95 (0.59–1.55) (	).844 682	0.76 (0.59–0.98)	0.032
	382 1.32 (0.93–1.87) (	0.119 1620	1.39 (1.18–1.64)	<0.0001
3+ 1168 1.55 (1.09–2.2	168 1.55 (1.09–2.20) (	0.015 1386	1.43 (1.21–1.69)	<0.0001

The Cox regression models were adjusted for diagnosis category, chemo-sensitivity for lymphoma, stem cell source, KPS percentage, conditioning regimen, recipient age, recipient race, and interval between diagnosis and HCT.