## ARTICLE IN PRESS

HEMATOL TRANSFUS CELL THER. 2023;xxx(xx):1-10



## HEMATOLOGY, TRANSFUSION AND CELL THERAPY



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## Original article

# Applicability and validation of different prognostic scores in allogeneic hematopoietic cell transplant (HCT) in the post-transplant cyclophosphamide era

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#### ARTICLE INFO

#### Article history: Received 20 February 2023 Accepted 21 July 2023 Available online xxx

#### Keywords:

Post-transplant cyclophosphamide Allogeneic hematopoietic cell transplantation Indices Validation

#### ABSTRACT

We investigated the predictive capacity of six prognostic indices [Karnofsky Performance Status (KPS), Hematopoietic Cell Transplant-Specific Comorbidity Index (HCT-CI), Disease Risk Index (DRI), European Bone Marrow Transplantation (EBMT) and Revised Pre-Transplantation Assessment of Mortality (rPAM) Scores and Endothelial Activation and Stress Index (EASIX)] in 205 adults undergoing post-transplant cyclophosphamide (PTCy)-based allo-HCT. KPS, HCT-CI, DRI and EASIX grouped patients into higher and lower risk strata. KPS and EASIX maintained appropriate discrimination for OS prediction across the first 2 years after allo-HCT [receiver operating characteristic curve (area under the curve (AUC) > 55 %)]. The discriminative capacity of DRI and HCT-CI increased during the post-transplant period, with a peak of prediction at 2 years (AUC of 61.1 % and 61.8 %). The maximum rPAM discriminative capacity was at 1 year (1-year AUC of 58.2 %). The predictive capacity of the EBMT score was not demonstrated. This study validates the discrimination capacity for OS prediction of KPS, HCT-CI, DRI and EASIX in PTCy-based allo-HCT.

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https://doi.org/10.1016/j.htct.2023.07.008

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Please cite this article as: M.Q. Salas et al., Applicability and validation of different prognostic scores in allogeneic hematopoietic cell transplant (HCT) in the post-transplant cyclophosphamide era, Hematology, Transfusion and Cell Therapy (2023), https://doi.org/10.1016/j.htct.2023.07.008

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Funding: There is no funding to declare.

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#### Introduction

Prognosis scores predictors of transplant outcomes support clinical decisions in allogeneic hematopoietic cell transplantation (allo-HCT). <sup>1,2</sup> The most widely used prognostic scores in clinical practice include the Karnofsky Performance Status (KPS) and the Hematopoietic Cell Transplantation-Specific Comorbidity Index (HCT-CI). <sup>3,4</sup> The Disease Risk Index (DRI), <sup>5</sup> the European Bone Marrow Transplantation (EBMT) and the Pre-transplantation Assessment of Mortality (PAM) Scores are alternative prognostic tools often used by different transplant institutions. <sup>6,7</sup> More recently, the Endothelial Activation and Stress Index (EASIX) has been proposed as a predictor of transplant outcomes when evaluated before allo-HCT. <sup>8</sup>

Post-transplant cyclophosphamide (PTCy), generally in combination with other immunosuppressant agents, provides an effective prevention of clinically relevant graft versus host disease (GvHD). Secondary to its efficacy, the use of PTCy is expanded to allo-HCT independently of the selected donor source. The PTCY-based allo-HCT has resulted in a novel transplant platform that induces a specific T-cell depletion immunosuppressant effect and transplant-related toxicities, a different infectious risk profile. Consequently, different studies have been conducted exploring whether the use of this novel prophylaxis impacts on relapse risks and transplant success with inconsistent results. 15-18

The prognostic indices listed above were defined and validated in cohorts of patients transplanted using non-PTCy-containing GvHD prophylaxis.<sup>3-8</sup> As the use of PTCy for GvHD prophylaxis is becoming more widespread in the transplant community, the present study investigates whether the predictive capacity/power of these prognostic indices remains adequate in a contemporary cohort of adults undergoing allo-HCT and PTCy for GvHD prevention.

#### **Methods**

#### Indices application

Prognostic scores were calculated for each patient using the definitions and following the methodology described in the original publications.<sup>3-8</sup> The KPS and HCT-CI were evaluated as part of our clinical practice, executed by expert allo-HCT physicians. The DRI, EBMT, rPAM and EASIX scores were calculated retrospectively using data from clinical charts. The PAM score was calculated using the simplified version published in 2015 [revised PAM score (rPAM score)]<sup>7</sup> and the EASIX score was calculated based on the information provided from bloodwork collected during pre-transplant assessments (between days –30 and –7 before allo-HCT).

#### Patient selection

Between January 2014 and February 2021, 205 consecutive adults with hematological malignancies underwent their first peripheral blood allo-HCT combined with PTCy-based GvHD prophylaxis at our Institution. Eligibility criteria for allo-HCT are detailed in the Supplementary Material. Clinical

information was collected retrospectively and updated in January 2022. This study was approved by the Ethics Committee of the Hospital Clinic de Barcelona, Spain and was performed in accordance with standards set by the Declaration of Helsinki.

## Conditioning regimen, graft characteristics and GVHD prophylaxis

Myeloablative conditioning (MAC) regimens generally combined high-dose of busulfan (3.2 mg/kg/day intravenously (IV)  $\times$  four days) or 12 Gy of total body irradiation (TBI) with fludarabine (30 mg/m²/day IV  $\times$  four days). The majority of reduced-intensity conditioning (RIC) regimens contained lower doses of busulfan (3.2 mg/Kg/day IV  $\times$  three days), or 8 Gy of TBI combined with standard doses of fludarabine.

Between January 2014 and May 2015, the Institutional standard GvHD prophylaxis consisted of PTCy 50 mg/kg/day administered on days +3 and +4, followed by mycophenolate mofetil (MMF) from day +5 to day +28, and tacrolimus (TK) started on day +5 at a dose of 0.04/kg/day IV, maintained therapeutically until day +90, and tapered down progressively up to day +120, in the absence of GvHD. Since June 2015, the standard GvHD prophylaxis out of the haplo-HCT setting was modified to PTCY and TK (PTCY-TK). Information related to graft source, infectious prophylaxis and supportive care is reported in the Supplementary Material.

#### Statistical methods

The overall survival (OS) and non-relapse mortality (NRM) were considered the main outcome variables. The main explanatory variables were KPS, HCT-CI, DRI, EBMT score, rPAM and EASIX. Patients were grouped into two risk groups of high- and low-risk strata. This classification was made based on the optimal cutoff value for each score and for the main outcome variable: OS. The cutoff value was calculated using maximally selected rank statistics. In the case of EASIX and following standard practices in the literature, the original value of the index was transformed to a base-2 logarithm value (log2-EASIX) to perform the statistical analysis.

The OS was calculated using the Kaplan-Meier estimator method and the NRM was estimated using the cumulative incidence method, considering relapse as a competing event. The score prediction ability for the OS and NRM was first explored during the entire post-transplant follow-up across the entire cohort and in each of the scores independently, using the concordance index (C-index). This index confirms the predictive power of the evaluated score for values superior to 0.5 with the perfect discrimination of the value of 1.<sup>19</sup> The prognostic indices were compared using the reclassification calibration statistic goodness-of-fit applied to reclassified categories. This method compares the proportions of patients whose estimated risk shifts in the correct and wrong directions based on a  $\chi^2$  goodness-of-fit test within reclassified categories for each score. In this setting, a p-value > 0.05 indicates a good fit.<sup>20</sup> Diagnostic accuracy measures (sensitivity, specificity and predictive values) were calculated at 2 years.

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The impact of the explanatory variables on the OS and NRM and the prediction ability of the five scores were analyzed using the regression analysis and C-index, respectively, and censoring the post-transplant follow-up at the following time points after the allo-HCT: 3 months, 6 months, 1 year and 2 years. The score discrimination was additionally evaluated using the inverse probability of censoring weighting estimation of cumulative time-dependent receiver operating characteristic (ROC) curves and, from them, calculating the area under the curve (AUC) at each time-point. The perfect discrimination corresponds to an AUC of 1.0, meaning that the predicted risk for all individuals who developed the outcome was higher than that for all individuals who did not experience the outcome. An AUC of 0.5 is indicative of an equal probability of risk assigned to individuals who experience the outcome and to individuals who did not, i.e., the prediction is purely random.<sup>21</sup> The p-values were two-sided. For statistical analyses, p < 0.05 was considered to indicate a statistically significant result. Statistical analysis was performed using the R.3.6.1 software (R Foundation for Statistical Computing, Vienna, Austria, http://www.R-project.org).

#### Results

#### Patient characteristics and outcome information

As shown in Table 1, the patients' median age was 52 years (range: 18 - 70), acute myeloid leukemia (38.0 %) was the most prevalent baseline diagnosis, 51.2 % of adults underwent the RIC allo-HCT and 72.2 % of patients received grafts from unrelated donors (UDs), while 19.5 % received from haploidentical donors.

Of the 205 patients included, 202 (98.5 %) were primary engrafted. Three (1.4 %) patients had moderate-severe sinusoidal obstruction syndrome (SOS) and 9 (4.4 %), thrombotic microangiopathy (TMA). The cumulative incidences of grades II to IV and grades III to IV acute GvHD at day +100 were 24.4 % (95 %CI 18.7–30.5) and 4.9 % (95 %CI 2.5–8.4), respectively, while those of moderate/severe chronic GvHD at 2 years was 7.6 (95 % CI 4.2–12.2). With a median follow-up among surviving patients of 35.9 months, 54 (26.3 %) patients relapsed and 66 (32.2 %) died. The main causes of death were relapse (34 patients, 16.6 %) and infection (13 patients, 6.3 %). The estimated 2-year OS, NRM, relapse-free survival and cumulative incidence of relapse were, respectively, 69.9 % (95 %CI 62.8–76.0), 16.1 % (95 %CI 11.4–21.6), 57.5 % (50.1 %–64.2 %) and 26.3 % (95 %CI 20.3–32.8).

#### Clinical scoring systems distribution

Based on the result obtained from the five clinical risk-scoring indices, patients were categorized into two risk groups (high-risk and low-risk) based on the optimal cutoff value for the OS prediction determined using maximally selected rank statistics. As shown in Table 1, according to the KPS, 49 (23.9 %) patients were classified into the high-risk group (KPS values: 70–80 %). According to the HCT-CI, the number of high-risk (HCT-CI: > 3) patients was 45 (22.0 %). Regarding the DRI, 46 (22.4 %) patients were classified into the high-risk group (DRI:

Age       Median (years)       52 (18–70)         Sex       Male       113 (55.1)         Female       92 (44.9)         Baseline Diagnosis       AML       78 (38.0)         MDS       34 (16.6)         ALL       35 (17.1)         MPN       12 (5.9)         CML       5 (2.4)         SLP / CLL       30 (14.6)         PCD       11 (5.4)         Conditioning Regimen (intensity)       MAC         MAC       100 (48.8)         RIC       105 (51.2)         Donor type       10/10 HLA MSD       17 (8.3)         10/10 HLA MUD       75 (36.6)       9/10 MMUD       75 (36.6)         9/10 MMUD       75 (36.6)       9/10 MMUD       73 (35.6)         Haploidentical donor       40 (19.5)       KPS         90-100 %       156 (76.1)       70-80 %       49 (23.9)         HCT-CI       Median (range)       3 (0-7)         0-3       160 (78.0)       > 3         DRI       Low-Intermediate       159 (77.4)       High-Very High       46 (22.4)         EBMT score
Median (years)       52 (18–70)         Sex         Male       113 (55.1)         Female       92 (44.9)         Baseline Diagnosis       34 (16.6)         AML       78 (38.0)         MDS       34 (16.6)         ALL       35 (17.1)         MPN       12 (5.9)         CML       5 (2.4)         SLP / CLL       30 (14.6)         PCD       11 (5.4)         Conditioning Regimen (intensity)       MAC         RIC       100 (48.8)         RIC       107 (51.2)         Donor type       17 (8.3)         10/10 HLA MSD       17 (8.3)         10/10 HLA MUD       75 (36.6)         9/10 MMUD       73 (35.6)         Haploidentical donor       40 (19.5)         KPS       90-100 %       156 (76.1)         70-80 %       49 (23.9)         HCT-CI       Median (range)       3 (0-7)         0-3       160 (78.0)         > 3       45 (22.0) </th
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FROM LINER
Median (range) 4 (1–7)
0–3 137 (66.8)
4–7 68 (33.8)
rPAM score
Median (IQR) 18.0 (14.2–22.1)
` '
$\geq$ 21.0 59 (28.8) <b>EASIX</b>
Median (IQR) 1.289 (0.778–2.500)
Log2-EASIX
Median (IQR) 0.366 (-0.361-1.322)
< 1.900 167 (81.5)
≥ 1.900 38 (18.5)
Additional Information:
Relapse* 54 (26.3)
Dead* 66 (32.2)
Dead without relapse* 32 (15.6)

AML: Acute Myeloid Leukemia; MDS: Myelodysplastic Syndrome; ALL: Acute Lymphoblastic Leukemia; MPN: myeloproliferative neoplasm; CML: Chronic Myeloid Leukemia; NHL: Non-Hodgkin Lymphoma; CLL: Chronic Lymphocytic Leukemia: PCL: Plasma Cell Disorders; MSD: Matched-Sibling Donor: MUD: Matched Unrelated Donor: MMUD: Mismatched Unrelated Donor: KPS: Karnofsky Performance Status; HCT-CI: Hematopoietic Cell Transplantation specific Comorbidity Index; DRI: Disease Risk Index; EBMT: European Society for Blood and Marrow Transplantation; rPAM: Revised Pretransplantation Assessment of Mortality; EASIX: Endothelial Activation and Stress Index. IQR: Interquartile Range.

\* Events during the entire follow-up.

high or very high risk) and, according to the EBMT score, 68 (33.8 %) adults were considered high-risk (EBMT score:  $\geq$  4). The optimal cutoff value of the rPAM score for the prediction of the OS was  $\geq$  21.0 (Supplementary Figure 1) and 59 (28.8 %) adults were classified into the high-risk group. Lastly, 38 (18.5 %) patients with log-2 EASIX  $\geq$  1.900 (pre-logarithmic transformed EASIX equivalent value  $\geq$  3.800) were classified into the high-risk group (Supplementary Figure 2).

# Predictive power and discrimination capacity of the different scoring systems for overall survival

As shown in Table 2 and Figure 1, the classifications of patients in the high- and low-risk strata according to the KPS, HCT-CI, DRI and EASIX indices for the prediction of the OS were statistically significant during the entire post-transplant period. In terms of hazard ratios (HRs) and according to the KPS, high-risk adults were 2.11 (p = 0.004) times more likely to die secondary to any cause than low-risk adults. According to the DRI, patients classified as high-risk were 2.04 (p = 0.005) times more likely to die than low-risk patients. The estimated HR for mortality of high-risk adults according to the HCT-CI index (score > 3) was 1.95 (p = 0.012) and the probability of mortality was 2.15 times superior for patients with an EASIX value  $\geq$  3.800 (log2-EASIX  $\geq$  1.900). According to the C-index value, the ranking of indices in terms of the predictive power of overall mortality, from high to low, was as follows: first the KPS (C-value 60 %), DRI (C-value 58 %), EASIX (C-value 58 %),

HCT-CI (C-value 56 %), rPAM (C-value 55 %) and EBMT score (C-value 54 %).

The discrimination capacity for the OS at 100 days, 6 months, 1 year, and 2 years is summarized in Table 3 and Figure 2. The KPS and EASIX maintained an appropriate discrimination during the four-time points evaluated after the allo-HCT, with values of the AUC-statistic higher than 55 %. The discriminative ability of the DRI and HCT-CI was low at day +100 (AUC values inferior to 50 %) and increased progressively during the post-transplant period, with a peak of discriminative capacity at 2 years (AUC equal to 61.1 % and 61.8 %, respectively). The rPAM showed an acceptable discrimination capacity during the first year after the allo-HCT (AUC increasing from 55.9 % at day +100 to 58.7 % and 58.2 % at 6 months and 1 year), decreasing to as low as 54.6 % at 2 years. These results were consistent with those obtained in the Cox regression analysis as the criterion for statistical significance and the C-value. Lastly, the predictive capacity for the OS of the EBMT score was neither demonstrated during the entire post-transplant period nor at the different time points after the allo-HCT.

#### Sensitivity and specificity for overall survival prediction

Sensitivity and specificity, together with the positive predictive values (PPVs) and negative predictive values (NPVs) were calculated for the prediction of the 2-year OS for each score independently (Table 3). The estimated sensitivity of each score was inferior to 40 %, except for the EBMT score, which

Table 2 – Prediction ability of the selected indices for overall survival and non-relapse mortality during the entire post-transplant follow-up.

	Prediction ability of the five indices for OS											
	Predi	ction		Calibration	Concordance							
	HR (95 % CI) <sup>1</sup>	p-value	$\chi 2^2$	p-value	C-Index <sup>3</sup>	SE (%)						
KPS: 70 - 80 % (vs. 90-100 %)	2.11 (1.27-3.15)	0.004	2.33	0.13	60 %	3.0						
HCT-CI: > 3 (vs. 0-3)	1.95 (1.15-3.28)	0.012	8.35	0.23	56 %	2.8						
DRI: High- Very High (vs. Low – Int)	2.08 (1.23-3.50)	0.005	0.77	0.38	59 %	2.8						
EBMT score: 4-7 (vs. 0-3)	1.42 (0.87-2.36)	0.159	1.38	0.24	54 %	3.0						
rPAM score: $\geq$ 21.0 (vs. < 21.0)	1.37 (0.82-2.29)	0.223	3.36	0.07	55 %	2.8						
Log2-EASIX: ≥1.900 (vs. < 1.900)	2.15 (1.26–3.66)	0.005	3.46	0.06	58 %	3.0						

		Prediction ability of the five indices for NRM											
	HR (95 % CI) <sup>1</sup>	p-value	Statistic <sup>2</sup>	p-value	C-Index <sup>3</sup>	SE							
KPS: 70-80 % (vs. 90-100 %)	3.24 (1.61-6.51)	< 0.001	2.62	0.11	65 %	4.4							
HCT-CI: >3 (vs. 0−3)	1.37 (0.64-2.92)	0.40	0.98	0.32	53 %	3.8							
DRI: High- Very High (vs. Low – Int)	1.90 (0.92-3.89)	0.079	2.18	0.14	56 %	4.0							
EBMT score: 4-7 (vs. 0-3)	1.52 (0.78-3.10)	0.20	2.12	0.15	55 %	4.0							
rPAM score: $\geq$ 21.0 (vs. < 21.0)	1.76 (0.87-3.54)	0.110	0.04	0.85	57 %	4.0							
Log2-EASIX: ≥ 1.900 (vs. < 1.900)	3.32 (1.66-6.64)	< 0.001	0.74	0.39	61 %	4.0							

KPS: Karnofsky Performance Status; HCT-CI: Hematopoietic Cell Transplantation Specific Comorbidity Index; DRI: Disease Risk Index; EBMT: European Society for Blood and Marrow Transplantation; rPAM: Revised Pre-Transplantation Assessment of Mortality; EASIX: Endothelial Activation and Stress Index.

<sup>&</sup>lt;sup>1</sup> Univariate Cox Regression Model.

<sup>&</sup>lt;sup>2</sup> Calibration for OS. Goodness of fit test for survival data (likelihood ratio  $\chi 2$  statistic). Calibration for NRM: Goodness of fit test for proportional sub-distribution hazards model.

Concordance Index for the prediction of OS and NRM and standard error (SE).

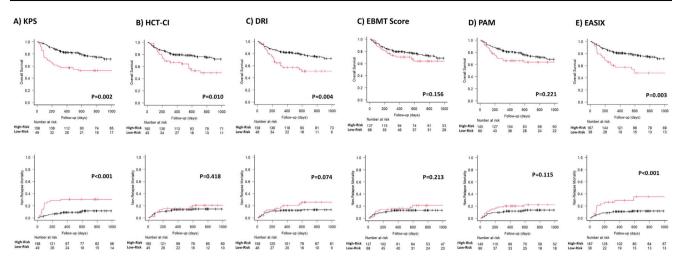


Figure 1-Overall Survival (First Row) and Non-Relapse Mortality (Second Row) Based on the Clinical Scoring System Distribution.

was 44.1 %. The EASIX score, followed by the KPS and HCT-CI, had the highest PPV values (50 %, 46.9 % and 46.6 %, respectively) and the rPAM and the EBMT scores, the lowest (36.2 % and 36 %, respectively). The EASIX, HCT-CI, KPS and DRI were the more specific predictive tools, with proportions superior to 80 % (87 %, 83.6 %, 82.9 % and 82.3 %, respectively) and all these four scales presented an NPV higher than 75 % (76.6 %, 76.8 %, 77.5 % and 76.8 %, respectively) (Table 4).

# Predictive power and discrimination capacity for non-relapse mortality

The predictive ability of the five scores was additionally calculated for the prediction of the NRM across the entire cohort and in each of the scores independently. The study cohort was stratified into the same two groups defined for

the prediction of the main outcome variable, the OS. In the case of the EASIX and rPAM, the cutoff values for the prediction of the NRM were re-calibrated and were found to be close to those defined for the prediction of the OS. For this reason, as well as to increase the clinical application of the conclusions obtained from the present study, the cutoff values for the EASIX and rPAM were maintained at 3.800 (log2-EASIX 1.900) and 21.0, respectively (Supplementary Figures 1 and 2).

As shown in Table 2 and Figure 1, the KPS and EASIX grouped patients into higher and lower risk strata, supporting their use for risk classification for the prediction of the NRM and during the entire post-transplant period. According to the KPS, high-risk adults were 3.24 (p < 0.001) times more likely to die secondary to transplant-related toxicity than low-risk adults. The estimated HR for transplant-related

Table 3 – Prediction ability of the six indices for overall survival predicted at different time-points after allo-HCT.

	Predic	tion at +3	months		Predic	tion at +	6 months		Pre	Prediction at +1 year Prediction at +2 year						
	% OS (95%CI)	HR 95%CI <i>p</i> -value	AUC* 95%CI p-value	C-Index	% OS (95%CI)	HR 95%CI p-value	AUC* 95%CI p-value	C-Index	% OS (95%CI)	HR 95%CI <i>p</i> -value	AUC* 95%CI p-value	C-Index	% OS (95%CI)	HR 95%CI p-value	AUC* 95% CI p-value	C-Index
KPS 90 - 100% 70 - 80%	96.1 (91.6 - 98.2) 73.5 (58.7 - 83.6)	7.58 2.9 - 19.9 < 0.001	73.0 60.7 - 85.2 < 0.001		91.0 (85.2 - 94.6) 67.3 (52.3 - 78.6)	4.26 2.1 - 8.7 < 0.001	67.1 57.7 - 76.5 < 0.001		82.3(78.3 - 87.5) 57.1 (42.2 - 69.6)	3.07 1.73 - 5.44 < 0.001	62.0 54.2-69.7 < 0.001	62%	75.4 (67.3 - 81.8) 52.4 (37.4 - 65.4)	2.63 1.6 - 4.5 0.001	58.8 51.4-66.2 < 0.001	60%
	90.6 (84.9 - 94.2) 91.1 (78.0 - 96.6)	0.91 0.3 - 2.8 0.877	41.7 34.7 - 48.7 0.54		86.8 (80.5 - 91.2) 80.0 (65.1 - 89.1)	1.50 0.7 - 3.3 0.302	54.6 45.8 - 63.4 0.13		79.6 (71.8 - 84.6) 66.6 (50.8 - 78.3)	1.70 0.92 - 3.13 0.088	56.1 48.8-63.4 0.05	55%	75.8 (68.1 - 81.9) 49.3 (33.1 - 68.9)	2.21 1.3 - 3.8 0.004	61.1 53.9-69.3 0.001	58%
	91.2 (85.6 - 94.7) 89.0 (75.5 - 95.3)	1.25 0.5 - 3.7 0.668	48.9 38.3 - 59.5 0.34		87.4 (81.2 - 91.7) 77.9 (62.7 - 87.4)	1.81 0.8 - 3.9 0.125	56.6 47.6 - 65.5 0.06		81.7 (74.7 - 86.9) 57.0 (41.1 - 70.1)	2.59 1.45 - 4.63 0.001	62.1 54.6-69.7 < 0.001	59%	75.3 (67.4 - 81.5) 50.8 (34.7 - 64.9)	2.40 1.4 - 4.1 0.001	61.8 54.9 - 8.7 0.001	58%
	92.0 (86.0 - 95.5) 88.2 (77.8 - 93.9)	1.43 0.6 - 3.6 0.435	50.0 37.6 - 62.5 0.19		86.8 (79.9 - 91.5) 82.4 (71.0 - 89.6)	1.34 0.6-2.8 0.428	53.8 44.4 - 63.3 0.20		79.3 (71.4 - 85.2) 70.4 (58.0 - 79.8)	1.47 0.85 - 2.43 0.170	54.8 46.8-62.8 0.07	54%	73.2 (64.4 - 80.2) 63.6 (50.7 - 74.0)	1.44 0.9 - 2.4 0.170	53.8 45.9-61.7 0.12	54%
	91.7 (85.9 - 95.2) 88.3 (77.0 - 94.2)	1.46 0.6 - 3.7 0.425	55.9 43.1 - 68.8 0.08		88.3 (81.8 - 92.5) 78.4 (65.3 - 86.6)	1.97 0.9 - 4.1 0.065	58.7 49.2 - 68.1 0.03		80.5 (73.0 - 86.1) 65.9 (52.3 - 76.5)	1.94 1.1 - 3.4 0.023	58.2 50.3-66.0 0.02	58%	72.5 (64.0 - 79.4) 63.7 (49.8 - 74.7)	1.6 0.9 - 2.7 0.09	54.6 47.1-62.1 0.09	55%
	93.4 (88.4 - 96.3) 78.9 (62.3 - 88.9)	3.30 1.3 - 8.2 0.010	57.9 45.7 - 70.2 0.003		88.0 (82.1 - 92.1) 73.7 (56.6 - 84.9)	2.37 1.1-5.1 0.026	58.6 49.7 - 67.5 0.01		80.6 (73.7 - 85.9) 57.3 (40.0 - 71.3)	2.55 1.39 - 4.65 0.002	60.0 52.3-67.2 < 0.001	59%	75.1 (67.4-81.2) 47.7 (30.7-62.9)	2.56 1.5 - 4.4 < 0.001	59.4 52.4-66.3 < 0.001	59%
Dead Relapse NRM	19 (9.2%) 7 (3.4%) 12 (5.8%)				30 (14.6%) 7 (3.4%) 23 (11.2%)		. —		48 (23.4%) 12 (5.8%) 36 (17.5%)				58 (28.9%) 13 (6.3%) 45 (21.9%)			

**Legend:** The prediction ability of each score was analyzed in the entire cohort and independently. Each time period accounts for the sum of events documented from day 0 to day +100, day 180, day +365 and day +730. Green color: Good prediction (C-Index  $\geq$  58 %), Yellow color: Acceptable prediction (C-Index > 55 % - < 58 %), Red color: Inaccurate prediction (C-Index  $\leq$  55 %).
\*ROC curve: Null hypothesis: true area = 0.5.

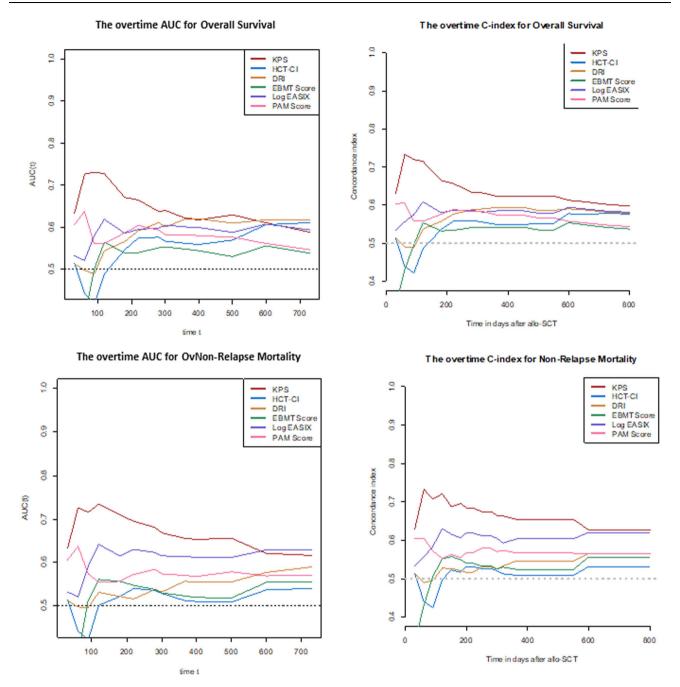


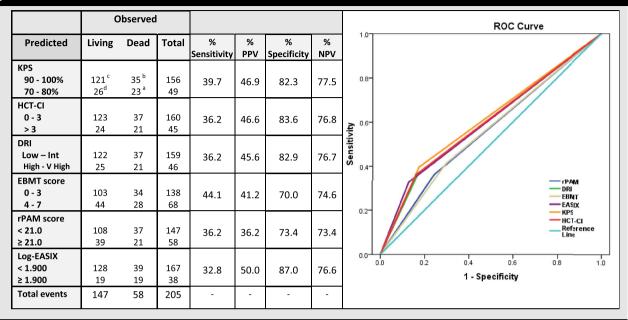
Figure 2 – Overall Survival and Non-Relapse Mortality Predictive Ability of the Six Indices at Different Time-Points after Allo-HCT.

mortality of high-risk adults, according to the EASIX (log2-EASIX > 1.900), was 3.32 (p < 0.001). Additionally, high-risk adults, according to the DRI, were found to present a non-significant trend to a higher NRM, compared to low-risk patients (HR 1.90, p = 0.079). According to the C-index, the ranking of indices in terms of the predictive power of the NRM from high to low was the KPS (65 %), EASIX (61 %), rPAM (57 %), DRI (56 %) EBMT score (55 %) and HCT-CI (53 %).

The KPS and EASIX maintained an appropriate predictive capacity of NRM during the four time points evaluated during the first 2 years after the allo-HCT and with values of the AUC-statistic higher than the cutoff value of 55 % (Table 5,

Figure 2). The KPS was the most accurate prognostic index for the NRM, with an estimated median of AUC ranges > 60 % during the post-transplant period and with a peak of prediction at day +100 and +180 (AUC 71.8 % and 66.2 %, respectively). The rPAM showed an acceptable discrimination capacity at day +180 (AUC of 55.9 %) and at 2 years (56.6 % at 2 years) after the allo-HCT for the prediction of the NRM. In addition, the DRI, HCT-CI and EBMT score were not appropriate predictors for the NRM during the post-transplant period. These results were consistent with those obtained when using the cumulative incidence regression analysis as the criterion for the statistical significance and C-Index.

Table 4 – Sensitivity, positive predictive value, specificity and negative predictive value of each index for the prediction of 2-year overall survival.



Legend: If a predictive tool is less sensitive, risk of mortality can be overestimated (Type I error). If a test less is specific, risk of mortality can be underestimated (Type II error); for these reasons, the present table explores the sensitivity, predictive positive value, specificity and predictive negative value of the selected five prognostic scales for the prediction of mortality at 2 years.

Sensitivity: ability of a screening test to detect a true positive. It refers to the proportion of those who have the condition that received a positive result on this test. Calculation: number of true positives / number of true positives + number of false negatives. Formula:% Sensitivity=  $[a/(a+b)] \times 100$ . Positive Predictive Value (PPV): probability that people with a positive screening test result indeed do have the condition of interest. Calculation: Truly classified as high-risk/Total predicted as high-risk. Formula: % PPV=  $[a/(a+b)] \times 100$ .

Specificity: ability of a screening test to detect a true negative. It refers to the proportion of those who do not have the condition that received a negative result on this test. Calculation: number of true negatives/number of true negatives + number of false positives. Formula:% Specificity= $[c/(c+d)] \times 100$ . Positive Predictive Negative (PPN): probability that people with a negative screening test result indeed do not have the condition of interest. Calculation: Truly classified as high-risk/Total predicted as high-risk. Formula: % PPN= $[c/(c+b)] \times 100$ . Needs to be checked!!!.

Table 5 - Prediction ability of the six indices for non-relapse mortality predicted at different time-points after allo-HCT.

	Predi	ction at +3	months		Predi	ction at +6	months		Pre	diction at	Pred	rediction at +2 years				
	% NRM (95%CI)	HR 95%CI p-value	AUC* 95%CI p-value	C-Index	% NRM (95%CI)	HR 95%CI <i>p</i> -value	AUC* 95%CI p-value	C-Index	% NRM (95%CI)	HR 95%CI <i>p</i> -value	AUC* 95%CI p-value	C-Index	% NRM (95%CI)	HR 95%CI <i>p</i> -value	AUC* 95%CI p-value	C-Index
KPS 90 - 100% 70 - 80%	3.9 <mark>(1L.6</mark> - 7.8) 24.5 (13.5-37.2)	6.94 2.60- 18.4 < 0.001	71.5 58.6 - 84.5 < 0.001	71%	5.8 (2.8 - 10.2) 28.6 (16.7 - 41.7)	5.58 2.43 - 12.8 < 0.001	71.88 61.4 - 82.3 < 0.001	70%	9.1 (5.2 - 14.3) 30.6 (18.3 - 43.8)	3.94 1.91 - 8.43 < 0.001	66.2 56.4 - 75.9 < 0.001	65%	11.6 (7.0 - 17.5) 30.6 (18.3 - 43.8)		62.21 52.5 - 71.9 < 0.001	63%
HCT-CI 0 - 3 > 3	8.8 (5.0 - 13.8) 8.9 (2.8 - 19.5)	0.98 0.33 - 2.92 0.98	42.4 35.0 - 50.0 0.49	42%	10.7 (6.5 - 16.0) 13.3 (5.3 - 25.0)	1.22 0.49 - 3.05 0.66	53.85 44.3 - 63.4 0.31	52%	13.9 (9.0 - 19.8) 15.6 (6.7 - 27.7)	1.12 0.48 - 2.59 0.79	53.23 44.7 - 61.7 0.39	51%	14.7 (9.7 - 20.8) 20.8 (10.1 - 34.1)		56.24 47.3 - 65.1 0.18	53%
DRI Low – Int High - V High	8.2 (4.6 - 13.1) 11.0 (4.0 - 22.1)	1.34 0.48 - 3.73 0.57	50.39 39.2 - 61.6 0.29	50%	10.7 (6.5 - 16.1) 13.3 (5.3 - 24.9)	1.25 0.49 - 3.15 0.72	53.84 44.3 - 63.4 0.33	52%	12.6 (8.0 - 18.3) 20.4 (9.9 - 35.5)	1.62 0.74 - 3.52 0.22	57.59 48.5 - 66.6 0.12	55%	13.4 (8.6 - 19.3) 25.9 (13.7 - 39.9)		61.15 52.0 - 70.3 0.04	57%
EBMT score 0 - 3 4 - 7	7.3 (3.7 - 12.5) 11.8 (5.5 - 20.7)	1.58 0.63 - 3.96 0.33	51.69 38.6 - 64.8 0.15	51%	9.5 (5.3 - 15.2) 14.7 (7.5 - 24.2)	1.54 0.68 - 3.48 0.30	55.98 45.2 - 66.8 0.13	55%	13.3 (8.2 - 19.6) 16.2 (8.6 - 26.0)	1.23 0.58 - 2.60 0.57	52.96 43.2 - 62.7 0.28	52%	13.3 (8.2 - 19.6) 21.2 (12.2 - 31.8)		56.67 46.7 - 66.7 0.08	56%
rPAM score < 21.0 ≥ 21.0	7.5 (4.0 - 12.7) 11.7 (5.1 - 21.4)	1.59 0.6 - 4.1 0.33	57.46 44.1 - 70.9 0.07	57%	9.7 (5.5 - 15.1) 15.1 (7.4 - 25.4)	1.62 0.7 - 3.7 0.25	56.73 46.1 - 67.3 0.09		11.8 (7.2 - 17.7) 20.2 (11.1 - 31.3)	1.80 0.9 - 3.8 0.12	57.46 47.7 - 67.2 0.06	57%	13.5 (8.5 - 19.8) 22.4 (12.6 - 34.0)		57.88 48.2 - 67.6 0.06	56%
Log-EASIX <1.900 ≥ 1.900	6.0 (3.1-10.3) 21.1 (9.8 - 35.2)	3.63 1.6 - 9.1 0.005	58.94 46.0 - 71.8 0.002	59%	8.4 (4.8 - 13.3) 23.7 (11.6 - 38.2)	3.01 1.3 - 6.7 0.009	62.01 51.6 - 72.4 0.004		10.9 (6.7 - 16.1) 29.2 (15.6 - 44.3)	3.0 1.4 - 6.2 0.004	61.36 52.0 - 70.8 0.002	60%	11.6 (7.2-17.1) 35.3 (20.1-50.9)	3.4 1.7 - 6.6 < 0.001	62.46 53.0 - 71.9 < 0.001	62%

Legend: The prediction ability of each score was analyzed in the entire cohort and independently. Each time period accounts for the sum of events documented from day 0 to day +100, day 180, day +365 and day +730. Green color: Good prediction (C-Index  $\geq$  58 %), Yellow color: Acceptable prediction (C-Index > 55 % - < 58 %), Red color: Inaccurate prediction (C-Index  $\leq$  55 %).

#### Discussion

Secondary to its efficacy, the use of PTCy for GvHD prevention is expanding from the haplo-HCT to matched sibling and UD allo-HCT with notable success.<sup>22</sup> This retrospective analysis validates the discrimination capacity for the OS prediction of the KPS, HCT-CI, DRI and EASIX in the PTCy-based allo-HCT and supports their use in clinical practice.

The KPS and HCT-CI maintained an appropriate discriminative ability for the OS prediction, supporting its use as prognostic predictors in the PTCy-containing allo-HCT. When patients were classified into two risk groups, according to the KPS, high-risk patients continued to be those with scores ≤ 80 %. This result correlated with the cutoff used in clinical practice and in different studies performed with cohorts of patients receiving other GvHD prophylaxis.3,23 In the case of the HCT-CI, the most accurate cutoff value for high-risk classification was found to be > 3 and those patients presented a 2-year OS and NRM expected probability of 49.3 % and 20.8 %, respectively. Notably, this cutoff differed in 1 point from the original cutoff proposed by Sorror et al. for the classification of patients into the highest risk group, which was defined as > 2.<sup>24</sup> The prognostic utility of the HCT-CI has been progressively validated by different investigators simultaneously with the expansion of the allo-HCT. 25,26 Nevertheless, the predictive ability of the HCT-CI for the PTCy-containing allo-HCT has not been explored in detail. This analysis suggests that high-risk patients, according to the HCT-CI, would be more appropriately classified if the cutoff used for discrimination be > 3. However, this result needs to be validated in an external cohort of patients transplanted using the PTCy-containing GVHD prophylaxis.

Interestingly, the KPS showed the highest discrimination capacity at day +100 (AUC: 73 %) and subsequently decreased to as low as 58.8 % at 2 years. In contrast, the HCT-CI discrimination ability for the OS increased from low to high during the post-transplant follow-up, with a peak of prediction at 2 years (AUC 61.8 %). Notably, the KPS prediction accuracy was higher during the early post-transplant period, when the patient fitness was probably still impacted by the underlying hematological disorder. However, and based on these results, comorbidities seemed to have more relevance in the patent long-term survivorship, as they persist in the individual during the post-transplant follow-up.

According to the DRI, high-risk patients had a higher risk of mortality, compared to low-risk patients. The DRI, developed only for the primary outcome of the OS evaluated at 2 years after the allo-HCT, is a validated tool to categorize groups of patients undergoing the allo-HCT by disease risk. This prognostic tool unifies a unique variable impact of disease risk and status before the allo-HCT, mainly for research purposes, stratifying patients in broad disease risk categories for retrospective or prospective studies. The prediction ability of the DRI for the OS has been validated by different investigators. Similar to the results reported in the original and other publications, the DRI was not found to be a predictor for the NRM in our analysis and the discriminative ability of the DRI for the OS prediction increased progressively during the post-transplant period with a peak of prediction at

2 years after the HCT. The result of this study supports that disease-related information prior to the allo-HCT remains to be relevant for the OS prediction in patients receiving the PTCy-based GvHD prophylaxis.

The EASIX is a biomarker-based laboratory formula and considered a surrogate parameter of endothelial activation.<sup>29</sup> This index was first designed to predict mortality in patients with acute GvHD and, consecutively, the prognostic utility of this index evaluated before allo-HCT was validated for the prediction of the NRM and OS.8 This study shows that the EASIX was an accurate predictor for the OS and NRM in the allo-HCT using the PTCy-based GVHD prophylaxis. Notably, the predictive accuracy of the EASIX for the OS and NRM estimation persisted stable during the post-transplant period, even though this index was evaluated only once and based on the information provided in the bloodwork performed during the pre-transplant evaluation, as well as taking into consideration the low rates of the SOS, TMA and clinically relevant acute GvHD diagnosed in patients included in the study cohort. This result significantly increases the utility of this new index and enhances the relevancy of the endothelium in the estimation of the probability of presenting posttransplant complications in the allo-HCT using PTCy-containing GvHD prophylaxis. Higher EASIX scores have been associated with an increased risk of mortality. The optimal cutoff valuable for the patient risk classification has not been yet determined. Based on the results provided by this analysis, the best cutoff for the patient risk distribution into two independent groups was 3.90 (log2-EASIX: 1.90). Nevertheless, this value needs to be externally validated in an independent cohort of patients undergoing the allo-HCT and receiving PTCy for GVHD prevention.

Revised PAM and EBMT scores are two composite prognostic indices based on transplant-related factors, patient baseline information and disease-related information.<sup>6,7</sup> The original PAM score was developed in 2006 based on clinical information provided by adults undergoing the allo-HCT between 1990 and 2002 and further refined and recalibrated in 2015 (revised PAM) after exploring its predictive ability at 2 years after the allo-HCT landmark in the cohort of patients transplanted between 2006 and 2009. The revised PAM score was validated for the prediction of the 2-year mortality after the allo-HCT in a cohort of patients transplanted between 2011 and 2015.<sup>25</sup> In our study, the rPAM was shown to exhibit maximal predictive ability for the OS and NRM 1 year after the allo-HCT (1-year AUC: 58 % and 57 %, respectively) and decreased at the 2-year timepoint landmark after the allo-HCT (2-year AUC: 55 % and 56 %, respectively) (Figures 1 and 2), differing from what has been reported in previous analyses.25

The EBMT score was designed in 1998 to predict mortality in patients with chronic myeloid leukemia undergoing the allo-HCT, consecutively, extended for outcome prediction to a broad range of acquired hematological disorders. The prognostic capacity of the EBMT score has not been demonstrated in our study. Based on these results, the impact of the 5 clinical variables comprising the EBMT score was explored individually (Supplementary Table 1 and Supplementary Figure 3) and none of the explanatory variables included in the univariate model were found to be significant predictors for the OS.

These results bolster the findings reported in the present analysis. Nevertheless, the results provided by the present analysis have to be interpreted carefully and need to be validated externally in a larger cohort of patients transplanted using the PTCy for GvHD prevention.

The specificity and sensitivity 30,31 of each index were calculated for the estimation of the 2-year OS. The EASIX, HCT-CI, KPS and DRI specificities and negative predictive values were higher than 80 % and 75 %, respectively, therefore, suggesting that these prognostic tools had a capability to correctly assign a low probability of mortality to patients classified as low-risk patients by the index superior to 80 %. Interestingly, the estimated sensitivity of most of the selected prognostic tools was less than 40 %. This result suggests that only 40 % of patients classified in the high-risk group will die during the first two years after the allo-HCT. These results enhance how sensitivity and specificity of each score should be carefully considered when these indices are used for decision-making, especially for patients classified into the highrisk group, as more than 50 % of these patients would potentially survive after the allo-HCT.

The main limitations of the study were the retrospective nature of the analysis and the relatively limited sample size. Additionally, the study cohort was selected from a single institution, limiting the extrapolation of the conclusions provided by this analysis to other Allo-HCT Programs and needing an external validation of the results to further confirm the findings reported in this analysis.

#### Conclusion

In conclusion, this analysis explores the applicability and the predictive ability of the KPS, HCT-CI, DRI and EASIX for the OS prediction in a cohort of patients undergoing the allo-HCT, using the PTCy-containing GvHD prophylaxis from a single institution. Based on the findings of this analysis, the authors encourage other centers to undertake similar internal analyses to further confirm the results provided in the present analysis and to avoid under or over-risk estimations.

#### Conflicts of interest

The authors declare no relevant conflicts of interest, financial or otherwise.

#### Acknowledgments

We thank our patients and the nursing and support staff in the Hematopoietic Cell Transplant Program at the Hospital Clinic de Barcelona.

#### Supplementary materials

Supplementary material associated with this article can be found in the online version at doi:10.1016/j.htct.2023.07.008.

#### REFERENCES

- Duarte RF, Labopin M, Bader P, Basak GW, Bonini C, Chabannon C, et al. Indications for haematopoietic stem cell transplantation for haematological diseases, solid tumours and immune disorders: current practice in Europe, 2019. Bone Marrow Transplant. 2019;54(10):1525–52.
- 2. Kanate AS, Majhail NS, Savani BN, Bredeson C, Champlin RE, Crawford S, et al. Indications for hematopoietic cell transplantation and immune effector cell therapy: guidelines from the American society for transplantation and cellular therapy. Biol Blood Marrow Transplant. 2020;26(7):1247–56.
- Sorror M, Storer B, Sandmaier BM, Maloney DG, Chauncey TR, Langston A, et al. Hematopoietic cell transplantation-comorbidity index and Karnofsky performance status are independent predictors of morbidity and mortality after allogeneic nonmyeloablative hematopoietic cell transplantation. Cancer. 2008;112(9):1992–2001.
- Sorror ML, Maris MB, Storb R, Baron F, Sandmaier BM, Maloney DG, et al. Hematopoietic cell transplantation (HCT)-specific comorbidity index: a new tool for risk assessment before allogeneic HCT. Blood. 2005;106(8):2912–9.
- Armand P, Gibson CJ, Cutler C, Ho VT, Koreth J, Alyea EP, et al. A disease risk index for patients undergoing allogeneic stem cell transplantation. Blood. 2012;120(4):905–13.
- Gratwohl A. The EBMT risk score. Bone Marrow Transplant. 2012;47(6):749–56.
- Au BK, Gooley TA, Armand P, Fang M, Madtes DK, Sorror ML, et al. Reevaluation of the pretransplant assessment of mortality score after allogeneic hematopoietic transplantation. Biol Blood Marrow Transplant. 2015;21(5):848–54.
- 8. Luft T, Benner A, Terzer T, Jodele S, Dandoy CE, Storb R, et al. EASIX and mortality after allogeneic stem cell transplantation. Bone Marrow Transplant. 2020;55(3):553–61.
- Nunes NS, Kanakry CG. Mechanisms of graft-versus-host disease prevention by post-transplantation cyclophosphamide: an evolving understanding. Front Immunol. 2019;10:2668.
- Williams L, Cirrone F, Cole K, Abdul-Hay M, Luznik L, AS Al-Homsi. Post-transplantation cyclophosphamide: from HLAhaploidentical to matched-related and matched-unrelated donor blood and marrow transplantation. Front Immunol. 2020;11:636.
- Luznik L, O'Donnell PV, Symons HJ, Chen AR, Leffell MS, Zahurak M, et al. HLA-haploidentical bone marrow transplantation for hematologic malignancies using nonmyeloablative conditioning and high-dose, posttransplantation cyclophosphamide. Biol Blood Marrow Transplant. 2008;14(6):641–50.
- 12. Bolanos-Meade J, Reshef R, Fraser R, Fei M, Abhyankar S, Al-Kadhimi Z, et al. Three prophylaxis regimens (tacrolimus, mycophenolate mofetil, and cyclophosphamide; tacrolimus, methotrexate, and bortezomib; or tacrolimus, methotrexate, and maraviroc) versus tacrolimus and methotrexate for prevention of graft-versus-host disease with haemopoietic cell transplantation with reduced-intensity conditioning: a randomised phase 2 trial with a non-randomised contemporaneous control group (BMT CTN 1203). Lancet Haematol. 2019;6(3): e132–e43.
- 13. Khimani F, Ranspach P, Elmariah H, Kim J, Whiting J, Nishihori T, et al. Increased infections and delayed CD4(+) T cell but faster B cell immune reconstitution after post-transplantation cyclophosphamide compared to conventional GVHD prophylaxis in allogeneic transplantation. Transplant Cell Ther. 2021;27(11):940–8.
- 14. Oltolini C, Greco R, Galli L, Clerici D, Lorentino F, Xue E, et al. Infections after allogenic transplant with post-transplant cyclophosphamide: impact of donor HLA matching. Biol Blood Marrow Transplant. 2020;26(6):1179–88.

- 15. Battipaglia G, Labopin M, Hamladji RM, Blaise D, Chevallier P, Brissot E, et al. Post-transplantation cyclophosphamide versus antithymocyte globulin in patients with acute myeloid leukemia undergoing allogeneic stem cell transplantation from HLA-identical sibling donors: a retrospective analysis from the Acute Leukemia Working Party of the European Society for Blood and Marrow Transplantation. Cancer. 2021;127(2):209–18.
- 16. Nagler A, Labopin M, Dholaria B, Wu D, Choi G, Aljurf M, et al. Graft-versus-host disease prophylaxis with post-transplantation cyclophosphamide versus cyclosporine a and methotrexate in matched sibling donor transplantation. Transplant Cell Ther. 2021
- 17. Mehta RS, Saliba RM, Chen J, Rondon G, Hammerstrom AE, Alousi A, et al. Post-transplantation cyclophosphamide versus conventional graft-versus-host disease prophylaxis in mismatched unrelated donor haematopoietic cell transplantation. Br J Haematol. 2016;173(3):444–55.
- 18. Nagler A, Labopin M, Houhou M, Aljurf M, Mousavi A, Hamladji RM, et al. Outcome of haploidentical versus matched sibling donors in hematopoietic stem cell transplantation for adult patients with acute lymphoblastic leukemia: a study from the Acute Leukemia Working Party of the European Society for Blood and Marrow Transplantation. J Hematol Oncol. 2021;14(1):53.
- Longato E, Vettoretti M, Di Camillo B. A practical perspective on the concordance index for the evaluation and selection of prognostic time-to-event models. J Biomed Inform. 2020;108:103496.
- 20. Solari A, le Cessie S, Goeman JJ. Testing goodness of fit in regression: a general approach for specified alternatives. Stat Med. 2012;31(28):3656–66.
- 21. Le Borgne F, Combescure C, Gillaizeau F, Giral M, Chapal M, Giraudeau B, et al. Standardized and weighted time-dependent receiver operating characteristic curves to evaluate the intrinsic prognostic capacities of a marker by taking into account confounding factors. Stat Methods Med Res. 2018;27(11):3397–410.
- 22. Broers AEC, de Jong CN, Bakunina K, Hazenberg MD, van Marwijk Kooy M, de Groot M, et al. Posttransplant cyclophosphamide for prevention of graft-versus-host disease: the prospective randomized HOVON-96 trial. Blood Adv. 2022.

- 23. Guilfoyle R, Demers A, Bredeson C, Richardson E, Rubinger M, Szwajcer D, et al. Performance status, but not the hematopoietic cell transplantation comorbidity index (HCT-CI), predicts mortality at a Canadian transplant center. Bone Marrow Transplant. 2009;43(2):133–9.
- 24. McDonald GB, Sandmaier BM, Mielcarek M, Sorror M, Pergam SA, Cheng GS, et al. Survival, nonrelapse mortality, and relapse-related mortality after allogeneic hematopoietic cell transplantation: comparing 2003-2007 versus 2013-2017 cohorts. Ann Intern Med. 2020;172(4):229–39.
- 25. Shouval R, Fein JA, Shouval A, Danylesko I, Shem-Tov N, Zlotnik M, et al. External validation and comparison of multiple prognostic scores in allogeneic hematopoietic stem cell transplantation. Blood Adv. 2019;3(12):1881–90.
- 26. Barba P, Pinana JL, Martino R, Valcarcel D, Amoros A, Sureda A, et al. Comparison of two pretransplant predictive models and a flexible HCT-CI using different cut off points to determine low-, intermediate-, and high-risk groups: the flexible HCT-CI Is the best predictor of NRM and OS in a population of patients undergoing allo-RIC. Biol Blood Marrow Transplant. 2010;16 (3):413–20.
- 27. Beauverd Y, Roosnek E, Tirefort Y, Nagy-Hulliger M, Bernimoulin M, Tsopra O, et al. Validation of the disease risk index for outcome of patients undergoing allogeneic hematopoietic stem cell transplantation after T cell depletion. Biol Blood Marrow Transplant. 2014;20(9):1322–8.
- Armand P, Kim HT, Logan BR, Wang Z, Alyea EP, Kalaycio ME, et al. Validation and refinement of the Disease Risk Index for allogeneic stem cell transplantation. Blood. 2014;123(23):3664– 71
- Luft T, Benner A, Jodele S, Dandoy CE, Storb R, Gooley T, et al. EASIX in patients with acute graft-versus-host disease: a retrospective cohort analysis. Lancet Haematol. 2017;4(9):e414–e23.
- Simon R. Sensitivity, specificity, PPV, and NPV for predictive biomarkers. J Natl Cancer Inst. 2015;107(8).
- **31.** Trevethan R. Sensitivity, specificity, and predictive values: foundations, pliabilities, and pitfalls in research and practice. Front Public Health. 2017;5:307.