



Assessment of Acute Rejection by Global Longitudinal Strain and Cardiac Biomarkers in Heart-Transplanted Patients

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OPEN ACCESS

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Specialty section:

This article was submitted to
Alloimmunity and Transplantation,
a section of the journal
Frontiers in Immunology

Received: 22 December 2021

Accepted: 24 February 2022

Published: 25 March 2022

Citation:

Clemmensen TS, Firooznia N,
Olawi FM, Løgstrup BB, Poulsen SH
and Eiskjær H (2022) Assessment of
Acute Rejection by Global Longitudinal
Strain and Cardiac Biomarkers in
Heart-Transplanted Patients.
Front. Immunol. 13:841849.
doi: 10.3389/fimmu.2022.841849

Aims: The aim of this study was to evaluate left ventricular global longitudinal strain (LVGLS), N-terminal pro brain natriuretic peptide (Nt-ProBNP), and Troponin T as non-invasive markers for acute cellular rejection (ACR) diagnosis and severity assessment after heart transplantation (HTx).

Methods: We retrospectively included all HTx patients transplanted from 2013 to 2019. At each visit, the patients were subjected to endomyocardial biopsy (EMB), measurement of Nt-ProBNP and Troponin T, and protocolized echocardiography with assessment of LVGLS. Sudden drop in graft function (SDGF) was defined as a drop in LVGLS \geq -2% in combination with either an increase in Troponin T \geq 20% or Nt-ProBNP \geq 30% compared with levels at the latest visit.

Results: We included 1,436 EMBs from 83 HTx patients. The biopsies were grouped as 0R (n = 857), 1R (n = 538), and \geq 2R (n = 41). LVGLS was lower and Troponin T and Nt-ProBNP higher in the 2R group than in the 0R and 1R groups (LVGLS: $-12.9 \pm 3.8\%$ versus $-16.9 \pm 3.1\%$ and $-16.1 \pm 3.3\%$; Troponin T: 79 [33;230] ng/l versus 27 [13;77] ng/l and 27 [14;68] ng/l; Nt-ProBNP: 4,174 [1,095;9,510] ng/l versus 734 [309;2,210] ng/l and 725 [305;2,082], all p < 0.01). A SDGF was seen at 45 visits of which 19 had \geq 2R ACR. EMBs showed ACR in 20 cases without SDGF. Finally, neither was SDGF seen nor did the EMB show rejection in 1,136 cases. Thus, the sensitivity of SDGF for \geq 2R ACR detection was 49% (32–65) and specificity 98% (97–99). The positive predictive value (PPV) was 42% (31–55) and the negative predictive value (NPV) 98% (98–99). The diagnostic value improved in a sub-analysis excluding EMBs within 3 months after HTx, clinically interpreted false positive \geq 2R ACR cases, and cases with \geq 2R ACR who recently (<2 weeks) were treated with intravenous methylprednisolone due to \geq 2R ACR (sensitivity 75% (48–93), specificity 97% (96–98), NPV 99% (99–100), and PPV 39% (27–52).

Conclusions: Patients with $\geq 2R$ ACR have lower LVGLS and higher Troponin T and Nt-ProBNP than patients without 2R rejection. A non-invasive model combining changes in LVGLS and Troponin T or Nt-ProBNP showed excellent negative predictive value and moderate sensitivity and may be used as a gatekeeper to invasive biopsies after HTx.

Keywords: heart transplantation, acute rejection, echocardiography, speckle tracking imaging, cardiac biomarker, donor-specific antibodies

INTRODUCTION

Acute rejection remains a significant complication after heart transplantation (HTx) (1). Rejections occur predominantly during the first 6 months but can be seen at any time after HTx (2–4). Symptoms are uncommon in the early course, but eventually symptoms of heart failure or sudden cardiac death may occur. Endomyocardial biopsies are still the golden standard for detection of acute rejection (5, 6). However, the endomyocardial biopsy procedure is invasive and carries a well-known risk of complications such as pneumothorax, cardiac tamponade, lung embolism, and lesions on the tricuspid valve. Furthermore, rejections are often patchy and biopsies may not detect or underestimate the rejection severity. Moreover, the histological interpretation is difficult with significant interobserver variation. Therefore, non-invasive methods for the evaluation of acute rejection are highly desirable to improve the diagnostic setup and the severity assessment. Echocardiographic assessment of left ventricular global longitudinal strain (LVGLS) has been demonstrated to be reduced during rejection episodes and to improve during resolving of the rejection (7). However, the overall reported sensitivity (56%–87%) and specificity (75%–81%) for rejection detection by LVGLS are at most moderate (7–10). Elevated N-terminal pro brain natriuretic peptide (Nt-ProBNP), a non-invasive plasma marker of elevated filling pressure, has previously been associated with acute rejection (11, 12). Furthermore, the degree of rejection-induced myocyte necrosis may be reflected in plasma levels of cardiac Troponin T (13). However, the roles of combined assessment of LVGLS and cardiac biomarkers are unknown.

Thus, the present study aimed at evaluating LVGLS, Nt-ProBNP, and high-sensitivity Troponin T as non-invasive markers for diagnosis and severity assessment of acute rejection.

METHODS

Patient Population

In this retrospective study, we included all patients receiving heart transplantation at Aarhus University Hospital, Denmark, from 2013 to 2019. A total of 86 patients were transplanted during this period. However, three patients died prior to first visit with LVGLS assessment and were not included. Thus, we included 83 patients subjected to a total of 1,476 endomyocardial biopsies during a follow-up of 24 months [15;25]. A total of 40 biopsies were not included in the analysis as these were taken without assessment of

LVGLS or cardiac serological markers. None of these biopsies showed treatment demanding rejection.

All patients received induction therapy with anti-thymocyte globulin for the first 3 postoperative days. The standard immunosuppressive regime comprised a calcineurin inhibitor, tacrolimus, in combination with mycophenolate mofetil. All patients received high-dose methylprednisolone for the first 2 postoperative days followed by prednisolone (0.3 mg/kg/day), which was gradually reduced during the first postoperative year. Steroid treatment was stopped in most patients during the second year after HTx.

The post-transplant routine rejection surveillance included endomyocardial biopsies and comprehensive echocardiography and blood samples including the cardiac biomarkers high-sensitivity Troponin T and Nt-ProBNP.

Demographic and clinical data were collected from electronic patient files. Biopsy results were collected from the histological reports.

Endomyocardial Biopsy

Biopsies were taken by standard local hospital protocol using the right internal jugular vein or a femoral vein. Patients underwent routine biopsies at the first 2 postoperative years. Biopsies were scheduled weekly during the first 6 weeks, every 2 weeks until 3 months, every month until 6 months, and every 2 months for the rest of the first postoperative year. Biopsies were taken every 3 months beyond the first year until year 2. Afterward, biopsies were only taken, when rejection was clinically suspected. Biopsies were routinely examined for histological signs of antibody-mediated rejection (AMR) with examination for C4D and CD68 deposits in case of histological signs of AMR, clinical signs or symptoms, or *de-novo* donor-specific antibodies.

Experienced cardiac pathologists analyzed all biopsies. Acute cellular rejection (ACR) was histopathologically graded as no (0R), mild (1R), moderate (2R), and severe rejection (3R), according to guidelines of ISHLT (14). Acute cellular rejections $\geq 2R$ were treated with intravenous methylprednisolone 1 g daily for 3 days. In contrast, we did not routinely treat 1R rejection episodes with changes in immunosuppression.

Donor-Specific Antibodies

Patients were routinely screened for donor-specific antibodies (DSA) between the first and second years after HTx using LABScreen® Mixed and LABScreen® Single Antigen (One Lambda Inc., CA, USA). HLA antibody specificities above 1,000 MFI were considered positive.

Echocardiography

We used a commercially available ultrasound system (Vivid E9 or E95, GE Healthcare Horten, Norway) with a 3.5-MHz phased array transducer (M5S). Echocardiography was performed by trained sonographers within 24 h of the biopsy and always before medical treatment of rejection. The sonographers were instructed to optimize image quality for assessment of myocardial deformation by LVGLS. LVGLS was obtained by frame-by-frame tracking of speckle patterns throughout the left-sided myocardium in standard 2D cine loops. The region of interest was manually adjusted for optimal tracking results. We excluded segments with unacceptably low tracking quality due to poor image acquisition or artifacts. LVGLS was calculated as the average of peak longitudinal systolic strain in a 17-segment myocardial model (15). LVGLS was assessed when the tracking quality was adequate in at least 5 of 6 segments in each view. The higher negative value of strain equals a higher magnitude of strain.

Statistics

Normally distributed data are presented as mean \pm standard deviation (SD); non-normally distributed data are presented as median and interquartile range [IQR]. Categorical data are presented as absolute values or percentages. Histograms and Q-Q plots were used to check continuous values for normality of the data distribution.

Our analysis plan involved the following:

- A. Comparison of myocardial function and cardiac markers between three different biopsy score groups; 0R, 1R, and $\geq 2R$. Between-group differences were assessed using a mixed model for continuous variables due to the unequal number of observations between patients. We used receiver operating characteristic curves to calculate the diagnostic value of continuing variables for acute rejection prediction. The optimal cutoff points were defined as the intersection points of sensitivity and specificity in the receiver operating characteristic curves.
- B. Serial characterization of LVGLS and cardiac biomarkers before, during, and after $\geq 2R$ rejection. We used repeated ANOVA analysis to compare between visit changes.
- C. Comparison of differences in myocardial function and cardiac biomarkers between patients with sudden drop in graft function with and without biopsy-proven rejection $\geq 2R$ and patients without sudden drop in graft function but biopsy-proven rejection $\geq 2R$.

Sudden drop in graft function was predefined as a drop in LVGLS magnitude $\geq -2\%$ in combination with either an increase in Troponin T $\geq 20\%$ or Nt-ProBNP $\geq 30\%$ compared with levels at the patient's last visit. Changes within the normal range (LVGLS $\geq -18\%$, Troponin T < 14 ng/l or Nt-ProBNP < 300 ng/l) were not considered as sudden drop in graft function.

p values < 0.05 were considered statistically significant. Analyses were performed using STATA (StataCorp LP, College Station, TX, USA).

RESULTS

Demographics

Baseline characteristics are shown in **Table 1**. During follow-up, 28 patients (34%) had at least one $\geq 2R$ rejection episode. A total of 1,436 biopsies were included. The biopsies were grouped as 0R (n = 856), 1R (n = 537), and $\geq 2R$ (n = 43). In the 2R group, 41 patients suffered acute cellular rejection and two patients suffered antibody-mediated rejection.

Rejection Groups and Prediction of Acute Rejection

Table 2 shows hemodynamics, myocardial function, and serological markers in the three rejection groups. As depicted, systolic and diastolic blood pressure was slightly lower in the $\geq 2R$ rejection group than in the 0R and 1R groups and the transmitral Doppler flows revealed a more restrictive LV-filling pattern in the $\geq 2R$ group than in the 0R and 1R groups. LVGLS magnitude was significantly lower and Troponin T and Nt-ProBNP significantly higher in the 2R group than in the 0R and 1R groups. Notably, LVGLS magnitude was significantly lower in the 1R than 0R rejection groups (**Figure 1**).

The abilities of LVGLS, Troponin T, and Nt-ProBNP to predict $\geq 2R$ acute rejection are depicted in **Figure 2**. The normal range cutoff points for rejection prediction by LVGLS ($< -18\%$), Troponin T (> 14 ng/l), and Nt-ProBNP (> 300 ng/l) provided high sensitivity (88.4%, 92.3%, 95.4%, respectively) but low specificity (35.3%, 26.3%, 24.2%, respectively). The optimal cutoff points for rejection prediction were LVGLS: -14.9% (sensitivity 72% and specificity 73%), Troponin T: 43 ng/l (sensitivity and specificity both 64%), and Nt-ProBNP 1,553 ng/l (sensitivity and specificity both 67%). By combining LVGLS, Troponin T, and Nt-ProBNP in a logistic model, the diagnostic accuracy increased providing AUC of 0.80 (0.72–0.88).

TABLE 1 | Baseline characteristics according to rejection groups.

	Baseline (n = 83)	12-month follow-up (n = 67)	p-value
Male sex, n (%)	64 (77)		
Age at HTx, years	54 [40–63]		
Reason for HTx			
Cardiomyopathy, n (%)	48 (58)		
IHD, n (%)	20 (24)		
Congenital heart disease, n (%)	9 (11)		
Other, n (%)	6 (7)		
Weight, kg	81 \pm 17	84 \pm 19	<0.01
Body mass index, kg/m ²	25.7 \pm 0.5	26.7 \pm 0.6	<0.01
Hypertension, n (%)	33 (47)	53 (75)	<0.01
Sensitized prior HTx, n (%)	25 (30)		
Medication			
Prednisolone, n (%)	83 (100)	66 (99)	0.28
Ciclosporine, n (%)	0 (0)	4 (6)	<0.05
Tacrolimus, n (%)	82 (99)	64 (96)	0.24
Mycophenolate, n (%)	83 (100)	65 (97)	0.12
mTOR inhibitor, n (%)	5 (6)	17 (25)	<0.01

Data are presented as mean \pm SD, percentages, or median and [IQR]. HTx, heart transplantation; IHD, ischemic heart disease.

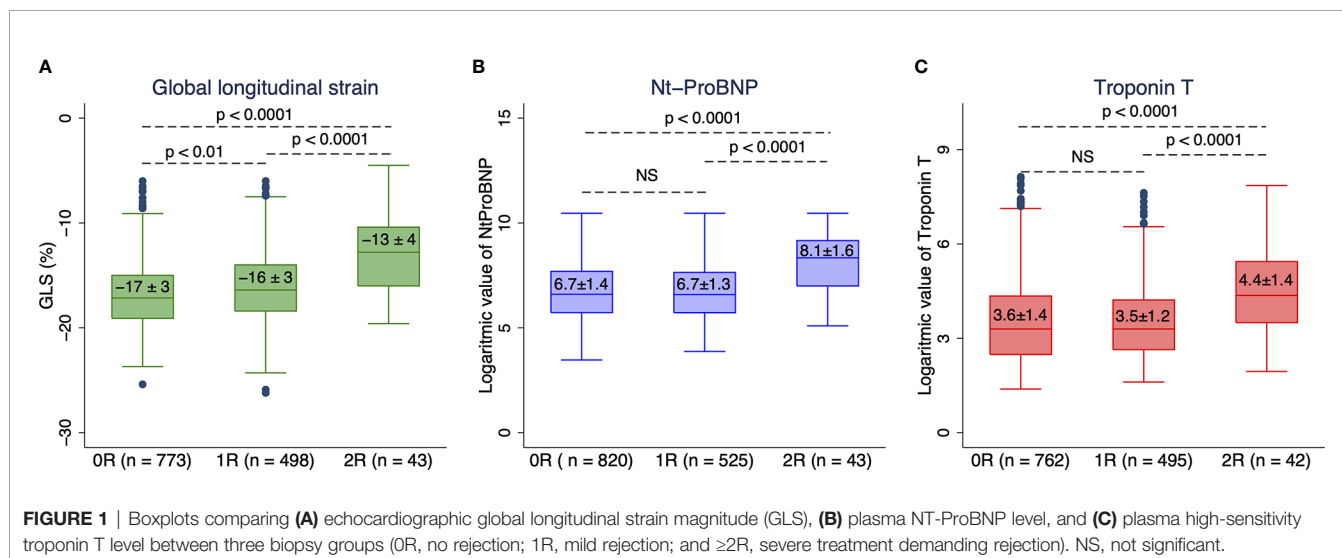
TABLE 2 | Hemodynamics, myocardial function, and serological markers according to rejection group.

	OR (n = 856)	1R (n = 537)	2R (n = 43)	Mixed-model p-value
Hemodynamics				
Heart rate, bpm	86 ± 12	87 ± 12	89 ± 17	0.26
Systolic blood pressure, mmHg	138 ± 15	138 ± 16	131 ± 18	0.01
Diastolic blood pressure, mmHg	85 ± 11	86 ± 11	81 ± 14	0.01
Systolic function				
LV-EF, %	60 ± 5	60 ± 5	56 ± 11	<0.0001
LV-GLS, %	16.9 ± 3.1	-16.1 ± 3.3	-12.7 ± 3.7	<0.0001*
TAPSE, mm	14.6 ± 3.5	14.3 ± 3.4	12.9 ± 4.0	0.03
Diastolic function				
E/A ratio	2.0 ± 0.9	1.9 ± 0.6	2.3 ± 0.9	0.02
Deceleration time, ms	162 ± 32	163 ± 33	145 ± 37	0.02
IVRT, ms	80 ± 17	78 ± 16	81 ± 23	0.49
Biomarkers				
Creatinine, mmol/L	102 [81;130]	94 [77;117]	93 [72;122]	0.002*
Hemoglobin, mmol/L	7.3 [6.5;8.2]	7.5 [6.9;8.3]	6.6 [5.8;8.4]	0.09
Troponin T, ng/L	27 [12;76]	27 [14;68]	79 [32;232]	0.001
NT-ProBNP, ng/L	734 [307;2190]	713 [305;2082]	4174 [910;14448]	<0.0001

Mixed-model p-values using the mixed model to adjust for unequal number of observations per individual. Data are presented as mean ± SD, percentages, or median and [IQR].

*p < 0.05 comparing OR versus 1R.

BPM, beats per minute; LV, left ventricular; EF, ejection fraction; GLS, global longitudinal strain; TAPSE, tricuspid annular plane systolic excursion; IVRT, isovolumetric relaxation time.



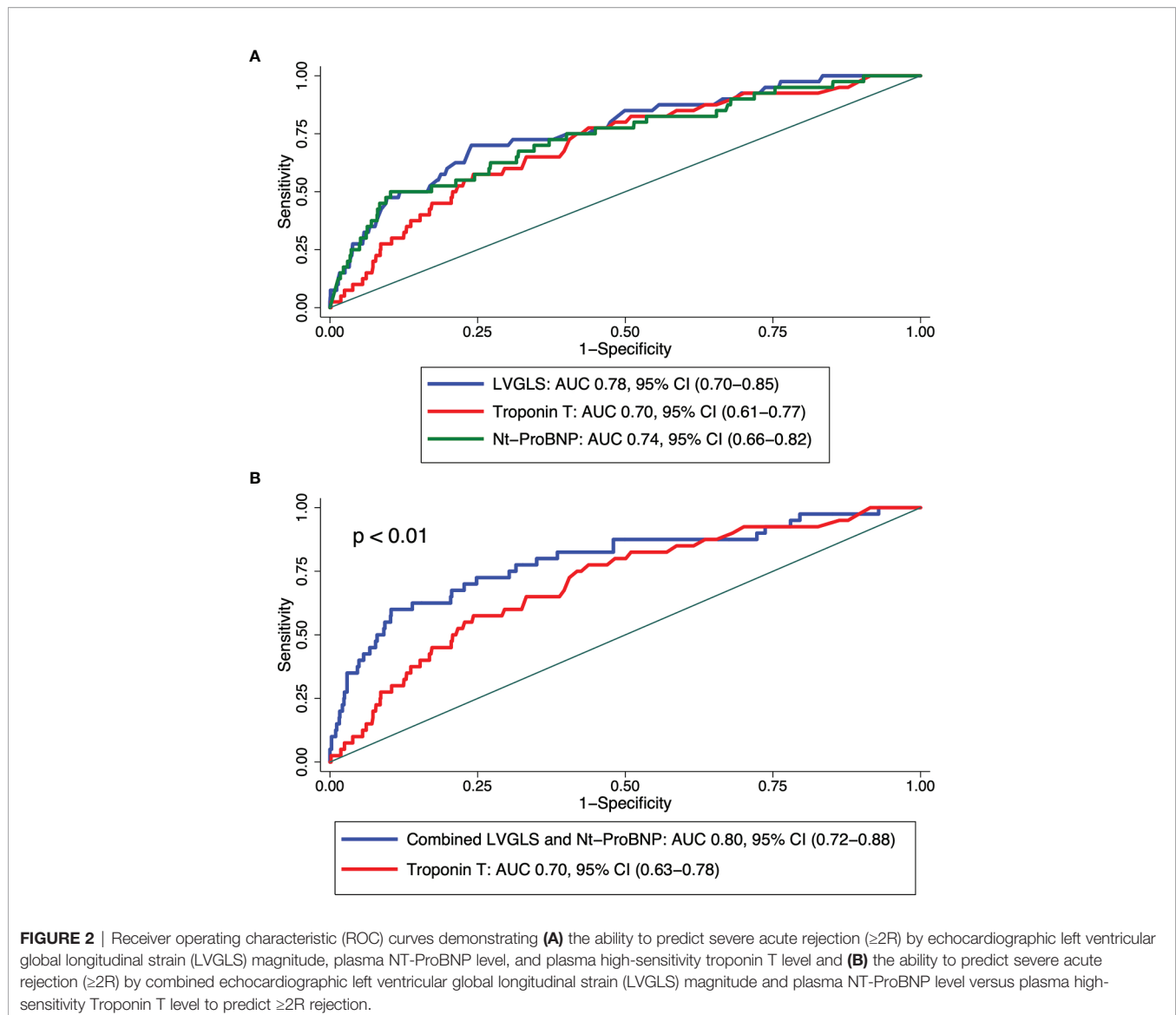
In a logistic model, LVGLS [odds ratio (OR) 1.4 (1.2–1.5)], the logistic value of Troponin T [OR 1.5 (1.2–1.8)], and the logistic value of Nt-ProBNP [OR 1.9 (1.6–2.4)] all predicted $\geq 2R$ acute rejection (all $p < 0.0001$). However, in a multivariate model, only LVGLS [OR 1.3 (1.2–1.4)] and the logistic value of Nt-ProBNP [OR 1.9 (1.3–2.7)] were independently associated with $\geq 2R$ acute rejection.

Serial Changes in LVGLS and Cardiac Markers During and After Acute Rejection

Figure 3 shows serial changes in LVGLS and the logarithmic value of Troponin T and Nt-ProBNP from the last biopsy before $\geq 2R$ rejection, the time of rejection, and the first two biopsies in the resolving phase afterward. As depicted, LVGLS magnitude significantly decreased and Troponin T and Nt-ProBNP significantly increased.

The median time from HTx to the patients last biopsy was 24 months [15;25]. At this time point, patients who had suffered at least one $\geq 2R$ rejection episode had a lower magnitude of LVGLS (-16.9% [-13.1;-18.8] versus -18.5 [-16.1;-19.6], $p = 0.03$) and higher Troponin T (19.0 ng/l [7.0;35.5] versus 12.0 ng/l [7;18], $p = 0.02$) and Nt-ProBNP (398 ng/l [226;1,278] versus 250 ng/l [111;583], $p = 0.02$) than patients who remained free from treatment demanding rejection.

DSA was detected in 12 patients between the first and second years post HTx of which 10 patients had HLA class II DSA and 2 patients had HLA class I DSA. The median MFI in patients with DSA was 1,800 [1,450;2,400]. Only 5 patients (18%) with acute rejection developed DSA whereas 23 patients (82%) did not develop DSA. Furthermore, we observed no difference in LVGLS ($p = 0.90$), Troponin T ($p = 0.62$), or Nt-ProBNP ($p = 0.15$) between patients with and without DSA development.



Sudden Drop in Graft Function With and Without Acute Rejection

Based on predefined changes in LVGLS, Troponin T, and Nt-ProBNP, the biopsy visits were grouped as the following:

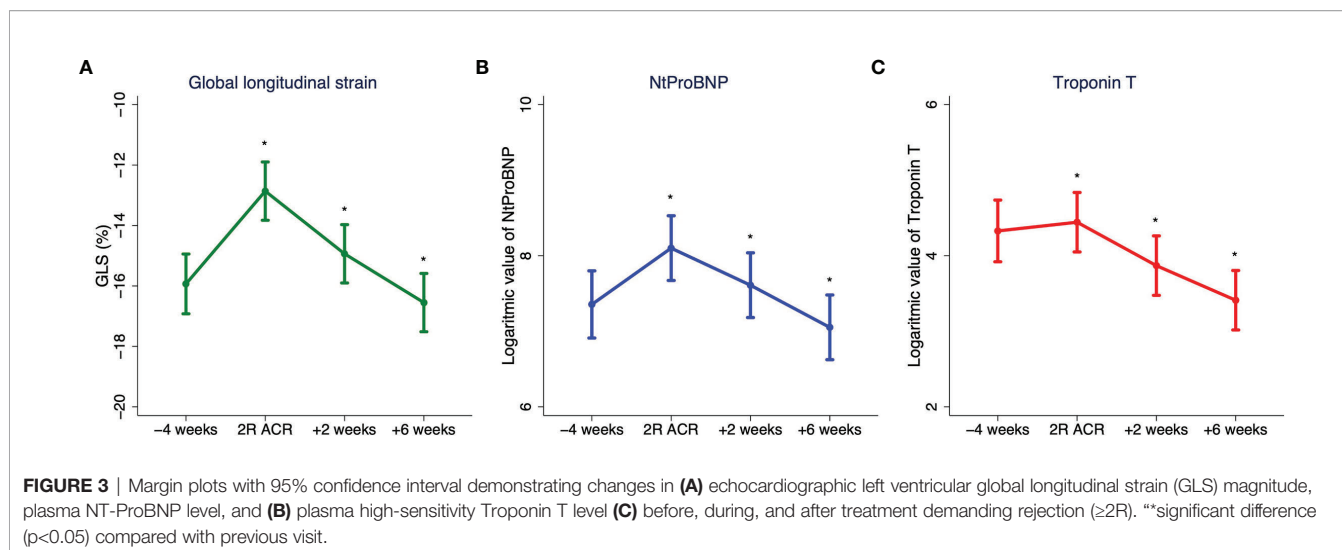
- Group A) Sudden drop in graft function and biopsy-detected acute rejection ($\geq 2R$), $n = 21$
- Group B) Sudden drop in graft function but no biopsy-detected acute rejection ($\geq 2R$), $n = 24$
- Group C) No sudden drop in graft function but biopsy-detected acute rejection ($\geq 2R$), $n = 20$
- Group D) No sudden drop in graft function and no biopsy-detected acute rejection ($\geq 2R$), $N = 1,136$

A sudden drop in graft function for detection of acute rejection ($\geq 2R$) showed a sensitivity of 51% (35–67), a specificity of 98% (97–99), a positive predictive value of 47% (35–59), and a negative predictive value of 98% (98–99). The overall accuracy of the model

was 96% (95–97). The diagnostic performance increased in a subanalysis excluding biopsies taken within the first 3 months after HTx (number: group A = 14; group B = 1,714; group C = 8; group D = 671). Thus, we found sensitivity of 64% (41–83) and specificity 98% (96–99). The positive predictive value was 45% (32–59) and the negative predictive value 99% (98–99). Overall accuracy was 96% (95–98%).

The clinical characteristics of group B are presented in **Supplementary Table 1**. As demonstrated, the vast majority of cases with a sudden drop of graft function without biopsy-proven rejection either were clinically interpreted as acute cellular or humoral rejection, had significant cardiac allograft vasculopathy, had decline in renal function, had acute arrhythmia, or were later diagnosed with acute rejection or suffered cardiac death during follow-up.

The clinical characteristics of group C are presented in **Supplementary Table 2**. Only 5/20 cases in group C had



reduced (LVGLS $< -18\%$). The biopsy result was clinically interpreted as false positive in three cases in which the patients did not receive intravenous methyl prednisolone. Notably, time since transplantation was significantly lower in group C (2.0 months [0.9;5.1]) than group A or B (4.9 months [1.3;23.3] and 6.6 months [2.3;18.6], respectively) ($p < 0.0001$).

In order to evaluate the changes in LVGLS, Troponin T, and Nt-ProBNP, we calculated the product of the absolute change and the percent increase or decrease. In a logistic model, the product of LVGLS change [odds ratio (OR) 2.4 (1.6–3.4)], Troponin T change [OR 3.5 (1.6–7.7)], and Nt-ProBNP change [OR 1.1 (1.1–1.2)] all predicted $\geq 2R$ acute rejection (all $p < 0.001$). However, in a multivariate model, only Nt-ProBNP change [OR 1.1 (1.1–1.2)] was independently associated with $\geq 2R$ acute rejection.

DISCUSSION

In this comprehensive study evaluating the relation between acute rejection, myocardial deformation, and cardiac biomarkers, several important findings were revealed: 1) acute rejections ($\geq 2R$) were rare and only observed in only 3% of biopsies; 2) patients with severe treatment demanding rejection ($\geq 2R$) have reduced LVGLS magnitude and increased Troponin T and Nt-ProBNP levels; 3) in the resolving phase after acute rejection, LVGLS improves and Troponin T and Nt-ProBNP decrease but remain elevated as compared to patients without acute rejection; 4) absolute isolated values of LVGLS, Troponin T, and Nt-ProBNP provide moderate sensitivity and specificity for acute rejection detection; and 5) changes in LVGLS and Troponin T and Nt-ProBNP provide excellent specificity and negative predictive value but at most moderate sensitivity for rejection detection.

Acute cellular rejection is histologically characterized by lymphocyte infiltration leading to myocardial necrosis and

local edema (4, 16). We have previously demonstrated that the rejection severity is reflected by the LVGLS magnitude (7). The mechanism and link between acute rejection and LVGLS are not fully clarified but may be explained by the longitudinal orientation of the sub-endomyocardial fibers that is believed to be particularly sensitive to ischemia, edema, and fibrosis. Nt-ProBNP is a well-known heart failure marker reflecting LV filling pressures and wall stress (17). During rejection, the Nt-ProBNP level may increase due to a direct myocyte damage and/or a secondary increase following the development of myocardial dysfunction and elevated filling pressures. Thus, Nt-ProBNP has been demonstrated a valuable tool for rejection evaluation (11, 12). It is well known that severe rejection episodes lead to myocyte necrosis. Therefore, it is reasonable to expect that high-sensitivity cardiac Troponins would be elevated as seen in other myocardial inflammatory diseases with myocyte necrosis such as myocarditis (18) and cardiac sarcoidosis (19). By combining LVGLS, Troponin T, and Nt-ProBNP, we obtain a non-invasive model that could potentially have a central role in graft surveillance in relation to acute rejection by reflecting the degree of myocardial edema, myocyte necrosis, and inflammation. Potentially, such a model could be used as gatekeeper to invasive biopsies by ruling in and out patients with respect to rejection surveillance. The results of the present study supported our hypothesis. Thus, LVGLS, Troponin T, and Nt-ProBNP were all significantly correlated with rejection degree. However, the sensitivity and specificity for each variable with respect to rejection were at most moderate using fixed cutoff points. This is expectable, as LVGLS, Troponin T, and Nt-ProBNP are all affected by other factors than rejection such as transport damage, the surgical trauma, renal dysfunction, elevated filling pressure, endomyocardial fibrosis, and development of cardiac allograft vasculopathy. These confounders are important and result in a wide range of individual values for LVGLS, Troponin T, and Nt-ProBNP, especially in the early phase after HTx. We previously demonstrated that LVGLS magnitude increases significantly during the first 3 months after HTx (20). Therefore,

caution should be taken using specific cutoff points for rejection suspicion in this period.

In the serial interpretation of data, we noted a significant decline in LVGLS magnitude and increases in Troponin T and Nt-ProBNP during $\geq 2R$ rejection. However, the sensitivity was at most moderate using these parameters for rejection assessment. This may be explained by the natural changes in LVGLS, Troponin T, and Nt-ProBNP in the first months after HTx. Thus, in the early phase after HTx, the rejection-induced LVGLS, Troponin T, and Nt-ProBNP affection must be substantial in order for the model to raise suspicion for acute rejection. Therefore, the majority of cases with biopsy-proven rejection ($\geq 2R$) but no sudden drop in graft function were in the first months after HTx. Furthermore, 25% of cases with $\geq 2R$ rejection but no sudden drop in graft function were taken within 2 weeks from a treated rejection episode at which point the graft function was already affected by acute rejection. Importantly, not all asymptomatic 2R rejection episodes need intensive intravenous methylprednisolone treatment (21). We speculate that cases with biopsy-proven 2R rejection but without affected LVGLS, Troponin T, and Nt-ProBNP do not need intravenous methylprednisolone treatment. In our study, we only omitted intravenous methylprednisolone treatment in three cases with biopsy-proven rejection but no drop in graft function. All had a favorable prognosis during follow-up, but larger studies are needed to evaluate the safety of omitted intravenous methylprednisolone treatment in cases without sudden drop in graft function.

In our study, we noted a strong negative predictive value of the non-invasive model combining LVGLS, Troponin T, and Nt-ProBNP assessment. Furthermore, the model identified other important hazards such as cardiac allograft vasculopathy and potential false negative biopsies. However, the proposed non-invasive model missed several $\geq 2R$ rejections, which could be reduced by use of the model only after the first 3 months and patients with a recent treatment demanding rejection. Importantly, our non-invasive rejection model was based on predefined cutoff values. In clinical practice, evaluation of trends may be useful as a gradual reduction in LVGLS magnitude or increase in Troponin T and Nt-ProBNP over time should raise suspicion of rejection. Thus, we noted a significant relation between the magnitude of LVGLS, Troponin T, and Nt-ProBNP change and the risk of treatment demanding rejection. Potentially, the non-invasive model evaluating a sudden drop in graft function may significantly reduce the number of biopsies and the associated risk of pneumothorax, cardiac tamponade, lung embolism, tricuspid valve damage, and procedure-related death. Furthermore, the model could be cost effective, but there is a need of a randomized trial evaluating the outcome and safety of non-invasive rejection monitoring LVGLS, Troponin T, and Nt-ProBNP versus traditional biopsy surveillance.

Limitations

This study is a single-center experience in a small cohort of patients. However, we used repeated measurements with a predefined standard protocol for rejection monitoring by speckle tracking imaging and serological cardiac markers.

We have previously reported low interobserver variation for LVGLS (7). However, the interobserver variation for the histological rejection graduation is unknown. Thus, it is possible that some episodes with a sudden drop in graft function but no 2R rejection were false negative and that some episodes with 2R rejection but no sudden drop in graft function were false positive.

CONCLUSIONS

Treatment demanding rejection was observed in only 3% of biopsies. Cases with $\geq 2R$ rejection had a lower LVGLS magnitude and higher Troponin T and Nt-ProBNP levels than cases without 2R rejection. In addition, significant changes in these parameters were observed during the course of grade 2R rejection compared to pre-rejection levels. A non-invasive model combining changes in LVGLS, Troponin T, or Nt-ProBNP showed excellent negative predictive value and may be used as a cost-effective non-invasive gatekeeper to the invasive biopsies after HTx.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT

Ethical review and approval was not required for the study in accordance with the local legislation and institutional requirements. The regional data protection agency approved the study. Written informed consent for participation was not required for this study in accordance with the national legislation and the institutional requirements.

AUTHOR CONTRIBUTIONS

Designed the study: all authors. Data collection: TC, NF, FO, HE. Data management: TC. Writing of the paper: all authors. All authors contributed to the article and approved the submitted version.

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fimmu.2022.841849/full#supplementary-material>

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