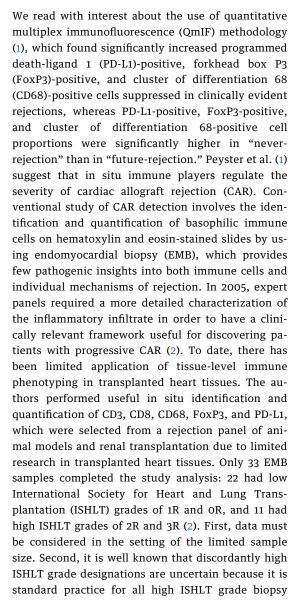
Letters

TO THE EDITOR

In Situ Immune Profiling Identifies Immune Players Involved in Allograft Rejection

A Call for Precision Medicine





events to receive some form of altered immunosuppression regardless of the presence of altered clinical data. Third, the most interesting findings support the existence of different "immunobiologies" in comparison to concordantly high ISHLT grade cases. This concept is not entirely new because the presence of distinct phenotypes with distinct fates was proposed in 2003 (3). On the other hand, the relatively high rate of a technical failure of QmIF analysis (26% of EMB) is a real concern in common clinical practice, but as the authors recognized, "it may reflect our dependence on residual material following routine clinical processing and the 6- to 12-year interval between EMB sampling and QmIF analysis" (1). Because of intrinsic limitations of the study and future reorganization of costs for the health care system due to the emergent coronavirus 2019 (COVID-19) pandemic, Peyster et al. (1) performed a valuable study of pathogenesis of clinical interest in the surveillance of CAR. Additionally, it would be interesting to analyze their data also in correlation to human leukocyte antigen-DR isotype (HLA-DR) matching at the time of transplantation, because this index may influence outcomes (4). Nevertheless, cost rationing is an inevitable occurrence where the potential demand for effective high-cost techniques will exceed supply. Despite this rather harmful consideration, the future need is to investigate prospectively whether integration of EMB with tissue (1) and liquid biopsy (5) could act synergistically to form a novel medicine paradigm (5) leading to the optimized management of patients undergoing heart transplantation.

*Claudio Napoli, MD, PhD Francesco Donatelli, MD Ciro Maiello, MD

*Clinical Department of Internal Medicine and Specialistic Units, and University Department of Advanced Medical and Surgical Sciences

University of Campania "Luigi Vanvitelli"

Piazza Miraglia 1 Naples 80138

Italy

E-mail: claudio.napoli@unicampania.it

https://doi.org/10.1016/j.jacbts.2020.05.012

© 2020 The Authors. Published by Elsevier on behalf of the American College of Cardiology Foundation. This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0/)

Please note: The authors have reported that they have no relationships relevant to the contents of this paper to disclose.

The authors attest they are in compliance with human studies committees and animal welfare regulations of the authors' institutions and Food and Drug Administration guidelines, including patient consent where appropriate. For more information, visit the *JACC: Basic to Translational Science* author instructions page.

REFERENCES

- **1.** Peyster EG, Wang C, Ishola F, et al. In Situ immune profiling of heart transplant biopsies improves diagnostic accuracy and rejection risk stratification. J Am Coll Cardiol Basic Trans Science 2020;5:328–40.
- **2.** Stewart S, Winters GL, Fishbein MC. Revision of the 1990 working formulation for the standardization of nomenclature in the diagnosis of heart rejection. J Heart Lung Transplant 2005;24:1710-20.
- 3. Klingenberg R, Koch A, Schnabel PA. Allograft rejection of ISHLT grade ≥3A occurring late after heart transplantation—a distinct entity? J Heart Lung Transplant 2003;22:1005-13.
- **4.** Cacciatore F, Amarelli C, Sabia C, et al. Effect on long-term mortality of HLA-DR matching in heart transplantation. J Card Fail 2019;25:409-11.
- **5.** Benincasa G, Mansueto G, Napoli C. Fluid-based assays and precision medicine of cardiovascular diseases: the "hope" for Pandora's box? J Clin Pathol 2019;72:785-99.

REPLY: In Situ Immune Profiling Identifies Immune Players Involved in Allograft Rejection



A Call for Precision Medicine

We appreciate the interest shown by Dr. Napoli and colleagues in our recent publication in *JACC: Basic to Translational Science*. We also appreciate their clear recognition of the need for improved biological characterization of transplanted endomyocardial biopsy tissues, as well as their appreciation of the in situ methodology we used in our work. Their letter raises several important points which we address in this response.

In our publication, approximately one-fourth of cases failed technical quality control. There were several reasons for this, some of which are mentioned in our paper (e.g., sample age), but other more technical factors are likely to improve with further experience and optimization. For example, the automated quantitative multiplex immunofluorescence (QmIF) staining workflow had not previously been applied to samples of heart tissue, and the default temperature of a heating step used for most tissues caused coverslips to become loose in heart tissues and created artifacts. Issues like this are relatively easily addressed once identified, and we expect a higher quality control pass rate moving forward.

The second point by Dr. Napoli and colleagues refers to the difficulty in determining if a high-grade asymptomatic case is truly "discordant" or the development of overt graft dysfunction is avoided by early treatment. It is true that the widespread convention of

treating high-grade endomyocardial biopsy tissue with augmented immunosuppression based on histology alone makes it difficult to control for this potential confounder in retrospective investigations. Nevertheless, the clear differences in in situ immune profiles between high-grade endomyocardial biopsies with and without evidence of graft dysfunction and much higher expression of the anti-inflammatory mediators PD-L1 and FoxP3 in the clinically silent cases suggests that there are real biological differences between these groups. Although this cannot be proven conclusively until a prospective investigation is performed, it is compelling circumstantial evidence.

Finally, Dr. Napoli and colleagues discuss the issue of cost containment in the context of advanced diagnostic approaches such as QmIF. Although cost considerations may influence adoption of new technologies, if QmIF substantially improves the accuracy of rejection diagnosis and aids in risk stratification, then initial assay costs may be offset by reduced complications and improved patient outcomes. In low-risk populations, minimizing low-yield procedures saves money, especially if associated with reductions in excess immunosuppression which can predispose patients to iatrogenic injury. In high-risk populations, more aggressive surveillance, prevention, and treatment strategies can reduce hospitalizations and major complications and potentially reduce cost. Whether this potential will be realized will, of course, require further investigation.

Eliot G. Peyster, MD, MSc Michael D. Feldman, MD, PhD *Kenneth B. Margulies, MD

*Perelman School of Medicine University of Pennsylvania Translational Research Center, Room 11-101 3400 Civic Center Boulevard, Building 421 Philadelphia, Pennsylvania 19104

 $\label{lem:eq:constraint} \textbf{E-mail: kenneth.margulies@pennmedicine.upenn.edu} \\ \text{https://doi.org/10.1016/j.jacbts.2020.05.008}$

© 2020 The Authors. Published by Elsevier on behalf of the American College of Cardiology Foundation. This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0/)

Please note: Supported by National Center for Advancing Translational Sciences of the National Institutes of Health award TL1TR001880 and Akoya Biosciences. Dr. Feldman is an equity holder and has technology licensed to both Elucid Bioimaging and Inspirata; and is a consultant for Inspirata, Phillips Healthcare, XFIN, and Virbio; and is a member of the advisory board of Inspirata. Dr. Feldman is consultant for Phillips Healthcare, XFIN, and Virbio. Dr. Margulies has received research grants from Sanofi-Aventis USA and GlaxoSmithKline; and is a consultant for and an advisory board member for Pfizer and MyoKardia.

The authors attest they are in compliance with human studies committees and animal welfare regulations of the authors' institutions and Food and Drug Administration guidelines, including patient consent where appropriate. For more information, visit the *JACC: Basic to Translational Science* author instructions page.