

COMMENTARY

Glucosamine regulates macrophage function in heart failure

Marcin Wysoczynski¹ | Jonah Stephan¹ | Shizuka Uchida² ¹Diabetes and Obesity Center, University of Louisville School of Medicine, Louisville, Kentucky, USA²Center for RNA Medicine, Department of Clinical Medicine, Aalborg University, Copenhagen SV, Denmark**Correspondence**

Marcin Wysoczynski, Diabetes and Obesity Center, Delia Baxter Building, Room 204B, University of Louisville, 580 South Preston Street, Louisville, KY 40202, USA.

Email: marcin.wysoczynski@louisville.edu

Shizuka Uchida, Center for RNA Medicine, Department of Clinical Medicine, Aalborg University, Frederikskaj 10B, 2 (building C), DK-2450 Copenhagen SV, Denmark.

Email: heart.lncrna@gmail.com or suc@dcm.aau.dk**Funding information**

Novo Nordisk Fonden, Grant/Award Number: NNF18OC0033438; National Institutes of Health, Grant/Award Numbers: R01 HL141191, P01 HL078825

KEYWORDS

glucosamine, heart failure, macrophage, signaling

1 | BACKGROUND

Healing myocardium after ischemic injury due to infarction requires sequential recruitment of immune cells to facilitate necrotic tissue clearance and subsequent activation of mesenchymal compartment to initiate deposition of extracellular matrix resulting in formation of a collagen-based scar tissue. Although post-myocardial infarction (MI) leukocytosis is associated with adverse outcomes, the last two decades of investigation revealed that immune cells actively participate in tissue repair and homeostasis after MI.¹ Macrophages are the central component of the healing myocardium. Tissue-resident macrophages deposited during embryonic development actively maintain tissue homeostasis, but also orchestrate immune cells infiltration after ischemic injury. In addition to tissue-resident macrophages, circulating monocytes of bone marrow and splenic origin are rapidly recruited to the heart after MI.² Following extravasation, monocytes differentiate to macrophages and facilitate necrotic tissue clearance via efferocytosis. Macrophage plasticity allows

them to acquire various secretory specialised phenotype. In general, in response to the local tissue environment, monocyte-derived macrophages can differentiate into classical pro-inflammatory or alternative reparative and pro-resolving phenotype.² The general macrophage contributions to infarct healing have been established; however, mechanistic aspects involved in regulation of macrophage number and function remain cryptic.

Among numerous post-translational protein modifications, there is emerging evidence that O-linked β -N-acetylglucosamine protein modification has immunoregulatory functions.³ O-linked β -N-acetylglucosaminylation (O-GlcNAcylation) is a subtype of glycosylation that involves addition of O-GlcNAc onto serine and threonine residues of nuclear or cytoplasmic proteins by O-GlcNAc transferase (OGT). O-GlcNAcylation overlaps the sites of protein phosphorylation.³ This suggests that GlcNAcylation may antagonise protein actions mediated by phosphorylation. However, there is no consensus regarding the functional outcome of protein GlcNAcylation in macrophages. Several studies demonstrated

This is an open access article under the terms of the [Creative Commons Attribution](https://creativecommons.org/licenses/by/4.0/) License, which permits use, distribution and reproduction in any medium, provided the original work is properly cited.

© 2022 The Authors. *Clinical and Translational Medicine* published by John Wiley & Sons Australia, Ltd on behalf of Shanghai Institute of Clinical Bioinformatics

that macrophage protein GlcNAcylation regulates various aspects of macrophage biology. To name a few, GlcNAcylation of STAT3 on threonine 717 (T717) antagonises phosphorylation and reduces IL-10 expression in lipopolysaccharide (LPS)-stimulated macrophages. By this means, it suppresses anti-inflammatory actions of macrophages.⁴ In contrast, other studies show that hyper-O-GlcNAcylation correlates with anti-inflammatory actions of macrophages in sepsis. Glucosamine (GlcN) stimulation induces hyper-O-GlcNAcylation of proteins and reduces transcriptional activity of NF κ B signaling in macrophages upon LPS stimulation.⁵ Similarly, GlcN treatment has been associated with M2 macrophage polarization in an LPS-induced septic lung injury via O-GlcNAcylation of nucleocytoplasmic proteins.⁶ Together, these data strongly indicate that protein GlcNAcylation has context-dependent effects on macrophage effector function in various systemic inflammatory diseases. However, the role of macrophage protein GlcNAcylation in pathophysiology of cardiovascular disease remains understudied.

2 | COMMENTARY

In the recent issue of “*Clinical and Translational Medicine*”, Zhou⁷ and colleagues demonstrate that GlcN administration preserves ventricular structure and function in the mouse model of MI-induced heart failure. Two treatment protocols were tested: early (starting 1 day before MI) and late (starting 3 days after MI). Each protocol consisted of six daily intraperitoneal administrations of GlcN (300 mg/kg/day). Ventricular function measured with echocardiography revealed that GlcN administration improved left ventricular ejection fraction (EF) and fractional shortening (FS) at 7, 14 and 28 days after MI in both early and late treatment groups. Histological analysis of the heart sections at 28 days after MI exhibited reduction of the scar function in the treatment groups compared to the vehicle control group. Immunoprofiling with flow cytometry shows that the GlcN treatment increased Ly6C^{Low} and decreased Ly6C^{High} macrophage contents. Additionally, histological analysis demonstrated accumulation of CD206^{Pos} macrophages but no change in total macrophage content in the hearts of mice treated with GlcN compared with vehicle controls. These data suggest that the GlcN treatment impacts skewing of macrophage towards reparative and pro-resolving phenotype, and as a consequence facilitate myocardial repair after MI. Mechanistic studies revealed that the GlcN treatment increased protein O-GlcNAcylation in myeloid cells. More detailed analysis shows an increase in STAT1 O-GlcNAcylation, which results in increased expression of Cx3cr1. Thus, the authors concluded that GlcN administration facilitates

myocardial recovery after ischemic injury via increase in recruitment of CX3CR1^{Pos} macrophages skewed towards a reparative and pro-resolving phenotype.⁷

This study adds new knowledge to the existing literature regarding immunomodulatory effects of GlcN supplementation. In addition to the effects of GlcN administration on myocardial repair, the novel observation suggests that the classical proinflammatory actions of STAT1 in macrophages are antagonised by O-GlcNAcylation.^{7–9} That further support the hypothesis that protein O-GlcNAcylation may have antagonistic effect on protein phosphorylation. However, multitude of question remains. Although the authors focus on the role of GlcN supplementation on STAT1 signaling, there are numerous of other pathways that are affected by GlcN and protein O-GlcNAcylation.¹⁰ Thus, it remains inconclusive whether enhanced myocardial repair after GlcN treatment in fact is solely due to STAT1 O-GlcNAcylation or via other mechanisms. More mechanistic studies would be warranted to explore this mechanism. Furthermore, GlcN shows clear benefit when injected acutely during ongoing inflammatory phase and scar formation. Because accumulation of macrophages in the chronic phase after MI contributes to progressive scarring and exacerbates ventricular dysfunction, it would be imperative to test the impact of GlcN administration on chronic non-resolving inflammation and remodelling in chronic models of heart failure.² Nevertheless, the current study warrants large preclinical studies in large animals (e.g., pigs and nonhuman primates) and perhaps clinical studies in MI patients to evaluate the efficacy of GlcN supplementation on the outcomes of post-MI heart failure.

ACKNOWLEDGEMENTS

This work was supported by the National Institutes of Health [grant numbers R01 HL141191, P01 HL078825] and the Novo Nordisk Foundation (NNF18OC0033438).

ORCID

Shizuka Uchida  <https://orcid.org/0000-0003-4787-8067>

REFERENCES

1. Steffens S, Nahrendorf M, Madonna R. Immune cells in cardiac homeostasis and disease: Emerging insights from novel technologies. *Eur Heart J*. 2021. <https://pubmed.ncbi.nlm.nih.gov/34897403/>
2. Swirski FK, Nahrendorf M. Cardioimmunology: The immune system in cardiac homeostasis and disease. *Nat Rev Immunol*. 2018;18:733–744.
3. Dong H, Liu Z, Wen H. Protein O-GlcNAcylation regulates innate immune cell function. *Front Immunol*. 2022;13:805018.
4. Li X, Zhang Z, Li L, et al. Myeloid-derived cullin 3 promotes STAT3 phosphorylation by inhibiting OGT expression and

- protects against intestinal inflammation. *J Exp Med*. 2017;214:1093–1109.
5. Hwang SY, Hwang JS, Kim SY, Han IO. O-GlcNAcylation and p50/p105 binding of c-Rel are dynamically regulated by LPS and glucosamine in BV2 microglia cells. *Br J Pharmacol*. 2013;169:1551–1560.
 6. Hwang JS, Kim KH, Park J, et al. Glucosamine improves survival in a mouse model of sepsis and attenuates sepsis-induced lung injury and inflammation. *J Biol Chem*. 2019;294:608–622.
 7. Zhou W, Jiang X, Tang Q, et al. Glucosamine facilitates cardiac ischemic recovery via recruiting Ly6C(low) monocytes in a STAT1 and O-GlcNAcylation-dependent fashion. *Clin Transl Med*. 2022;12:e762.
 8. Kim HS, Kim DC, Kim HM, et al. STAT1 deficiency redirects IFN signalling toward suppression of TLR response through a feedback activation of STAT3. *Sci Rep*. 2015;5:13414.
 9. Tugal D, Liao X, Jain MK. Transcriptional control of macrophage polarization. *Arterioscler Thromb Vasc Biol*. 2013;33:1135–1144.
 10. Qiang A, Slawson C, Fields PE. The role of O-GlcNAcylation in immune cell activation. *Front Endocrinol*. 2021;12:596617.

How to cite this article: Wysoczynski M, Stephan J, Uchida S. Glucosamine regulates macrophage function in heart failure. *Clin Transl Med*. 2022;12:e819. <https://doi.org/10.1002/ctm2.819>