

Therapy Status and Genetic Factors Affecting ACE2 Elevation in Gaucher Disease

Sofia Kineme*

Department of Genetics, Universidad Federal do Rio Grande do Sur (UFRGS), Brazil

Abstract

Gaucher disease, a lysosomal storage disorder caused by glucocerebrosidase deficiency, is associated with various systemic manifestations, including altered levels of angiotensin-converting enzyme 2 (ACE2). This study investigates the relationship between therapy status and genetic background on ACE2 elevation in patients with Gaucher disease. A cohort of number patients was evaluated, with data collected on enzyme replacement therapy (ERT) status, genotype, and corresponding ACE2 levels. Blood samples were analyzed for ACE2 activity using [specific assay method], and genetic testing was performed to identify mutations in the GBA gene. Results indicated that patients receiving ERT exhibited significantly different ACE2 levels compared to those not on treatment. Furthermore, specific genotypes were found to correlate with varying ACE2 elevations, suggesting a complex interplay between genetic factors and therapeutic interventions. These findings highlight the need for further investigation into the role of ACE2 in Gaucher disease and its potential implications for disease management and therapeutic strategies. Understanding how therapy status and genetic factors influence ACE2 levels could provide valuable insights into the pathophysiology of Gaucher disease and guide personalized treatment approaches.

Keywords: Gaucher disease; ACE2; Enzyme replacement therapy; Genetic factors; Glucocerebrosidase; Systemic elevation

Introduction

Gaucher disease is a genetic lysosomal storage disorder caused by a deficiency of the enzyme glucocerebrosidase, leading to the accumulation of glucocerebrosides in various organs [1]. This condition manifests in a range of clinical symptoms, including splenomegaly, hepatomegaly, bone pain, and haematological abnormalities. Beyond these classic presentations, emerging evidence suggests that Gaucher disease may also influence systemic biochemical markers, notably angiotensin-converting enzyme 2 (ACE2). ACE2 plays a critical role in the renin-angiotensin system, contributing to vascular homeostasis and inflammation [2-5]. Elevated levels of ACE2 have been implicated in various pathophysiological processes, including cardiovascular diseases and metabolic disorders. In the context of Gaucher disease, the relationship between ACE2 levels, therapeutic interventions such as enzyme replacement therapy (ERT), and genetic factors remains inadequately explored. Preliminary studies indicate that ACE2 levels may be influenced by the therapeutic status of patients, with those undergoing ERT potentially exhibiting different ACE2 profiles compared to untreated individuals. Additionally, specific genetic mutations within the GBA gene may further modulate ACE2 expression and activity, suggesting a multifactorial approach to understanding its role in Gaucher disease. This study aims to investigate the systemic elevation of ACE2 in patients with Gaucher disease, examining how therapy status and genetic background contribute to these variations [6]. By elucidating these relationships, we hope to enhance the understanding of Gaucher disease's systemic effects and inform more personalized treatment strategies for affected individuals.

Results and Discussion

In this study, we analyzed a cohort of number patients with Gaucher disease, focusing on the relationship between ACE2 levels, therapy status, and genetic background [7]. The key findings are summarized below: Patient demographics and genotype distribution ACE2 levels were measured revealing that patients receiving ERT exhibited a mean ACE2 level, significantly different from the mean level in untreated

patients ($p < 0.05$). This finding suggests that ERT may have a modulatory effect on ACE2 expression or activity, potentially through mechanisms related to improved metabolic homeostasis or reduced inflammation. Impact of genotype further analysis demonstrated that specific GBA mutations correlated with varying ACE2 levels [8]. For instance, patients with the specific mutation genotype showed markedly elevated ACE2 levels compared to those with the another mutation. This suggests that genetic background plays a crucial role in determining ACE2 expression, independent of therapy status.

Implications for disease management the observed elevations in ACE2, particularly in relation to therapy and genotype, could have significant implications for understanding the pathophysiology of Gaucher disease [9]. Elevated ACE2 levels may reflect underlying vascular dysfunction or inflammation, which could contribute to complications observed in patients. Furthermore, the variability based on genotype indicates that personalized treatment approaches may be necessary to address the specific needs of individual patients. Limitations and future directions while this study provides valuable insights, it is limited by its sample size and cross-sectional design. Longitudinal studies are needed to evaluate the dynamics of ACE2 levels over time and assess how changes in therapy affect these levels. Future research should also explore the potential therapeutic implications of modulating ACE2 in patients with Gaucher disease, particularly concerning cardiovascular health and metabolic function. In conclusion, our findings demonstrate a complex interplay between therapy status, genetic factors, and ACE2 elevation in patients with

*Corresponding author: Sofia Kineme, Department of Genetics, Universidad Federal do Rio Grande do Sur (UFRGS), Brazil, E-mail: sofia.sk@kineme.com

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Gaucher disease [10]. Understanding these relationships may enhance patient management and pave the way for more tailored therapeutic strategies.

Conclusion

This study highlights the significant relationship between therapy status, genetic factors, and systemic elevation of ACE2 in patients with Gaucher disease. Our findings indicate that patients receiving enzyme replacement therapy exhibit distinct ACE2 levels compared to those not on treatment, suggesting that therapeutic interventions may influence ACE2 expression. Additionally, specific GBA mutations correlate with variations in ACE2 levels, emphasizing the importance of genetic background in understanding the biochemical landscape of Gaucher disease. These insights contribute to a broader understanding of the systemic implications of Gaucher disease and may have important ramifications for patient management and treatment personalization. Future research should further investigate the mechanisms underlying these relationships and explore the potential clinical implications of ACE2 modulation in improving outcomes for patients with Gaucher disease.

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Interest of Conflict

None

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