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Glucocerebrosidase Enzyme Activity in *GBA* Mutation Parkinson Disease

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Abstract

Mutations in the glucocerebrosidase (*GBA1*) gene, the most common genetic contributor to Parkinson's disease (PD), are associated with an increased risk of PD in heterozygous and homozygous carriers. While glucocerebrosidase enzyme (GCase) activity is consistently low in Gaucher disease, there is a range of leukocyte GCase activity in healthy heterozygous *GBA1* mutation carriers. To determine whether GCase activity may be a marker for PD with heterozygous *GBA1* mutations (*GBA1* mutation PD, *GBA* PD), *GBA* PD patients (n=15) were compared to PD patients without heterozygous *GBA1* mutations (idiopathic PD; n=8), heterozygous *GBA1* carriers without PD (asymptomatic carriers; n=4), and biallelic mutation carriers with PD (Gaucher disease with PD, GD1 PD; n=3) in a pilot study. GCase activity (nmol/mg protein/hour) in GD1 PD (median [interquartile range]; minimum–maximum: 6.4 [5.7]; 5.3–11) was lower than that of *GBA* PD (p=0.01), while GCase activity in *GBA* PD (16.0 [7.0]; 11–40) was lower than idiopathic PD (28.5 [15.0]; 16–56) (p=0.01) and asymptomatic carriers (25.5 [2.5]; 23–27) (p=0.04). Therefore, GCase activity appears to be a possible marker of

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Conflicts of Interest/Disclosures

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heterozygous *GBA1* mutation PD, and larger studies are warranted. Prospective studies are also necessary to determine whether lower GCCase activity precedes development of PD.

Keywords

Biomarker; Gaucher; GBA; GBA1; GBA enzyme activity; Glucocerebrosidase; Parkinson's disease

1. Introduction

Heterozygous glucocerebrosidase enzyme (GCCase) mutations (from the glucocerebrosidase gene [*GBA1*]) encode lysosomal GCCase and increase the risk of Parkinson's disease (PD) and dementia with Lewy bodies [1,2]. However the penetrance of *GBA1* mutations is low [1,2]. While GCCase activity is consistently decreased in type 1 Gaucher disease (GD1), a range of GCCase activity in heterozygous *GBA1* mutation carriers exists [3,4]. To address the relationship between GCCase activity and PD, this pilot study assesses GCCase activity in *GBA1* mutation carriers, with and without PD, and non-carriers with PD (idiopathic PD).

2. Methods

Participants were recruited from a genetics study of PD in Ashkenazi Jews [6]. GCCase activity from fresh blood was assayed using 4-methylumbelliferyl- β -D-glucopyranoside leukocyte assay [4] for 15 PD patients with heterozygous mutations (*GBA* PD) (12 N370S, one of each 84GG, R496H, and E326), eight idiopathic PD patients, four asymptomatic *GBA1* carriers (N370S, 84GG, R496H, and E326K) and three GD1 patients with PD (GD1 PD) (two previously reported; N370S/N370S, N370S/R496H, and N370S/A456P) [6]. Groups were compared using Mann–Whitney and Fisher's exact tests with Bonferroni correction. The internal review board approved this study.

3. Results

GCCase activity (nmol/mg protein/hour) in *GBA* PD patients (median, interquartile range: 16.0, 7.0) was lower than idiopathic PD (28.5, 15.0) ($p=0.01$) and asymptomatic carriers (25.5, 2.5) ($p=0.04$), although the latter difference was not significant using a correction of $p<0.017$ (Fig. 1, Table 1). GCCase activity in GD1 PD patients (6.4, 5.7) was lower than *GBA* PD patients ($p=0.01$).

4. Discussion

Lower peripheral GCCase activity is associated with *GBA* PD, suggesting that reduced peripheral GCCase activity has potential as a marker of *GBA* PD. While the causal mechanism for GCCase to produce synucleinopathies has not been fully explicated, a bi-directional loop has been reported, where deficiency of GCCase promotes α -synuclein via excess substrate, resulting in stabilization of α -synuclein into oligomers, and greater GCCase dysfunction [5]. This is supported by our finding of decreased GCCase activity in *GBA* PD. Gain of function has also been proposed, as GCCase co-localizes with α -synuclein in Lewy

bodies in *GBA* PD [7]. Alternatively, gain and loss of function mechanisms may co-exist. Additional study including a larger sample, a greater distribution of mutation types and asymptomatic carriers, and assessment of sphingolipid substrate and products may lend insight into leukocyte GCCase activity as a marker and its mechanism.

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Highlights

1. As glucocerebrosidase mutations increase the risk of Parkinson Disease (PD), glucocerebrosidase enzyme activity in different groups was evaluated.
2. The primary focus of this pilot study was on PD patients with and without *GBA1* mutations.
3. Glucocerebrosidase enzyme activity was lower in *GBA1* heterozygous mutation carriers with PD when compared to non-*GBA1* mutation PD.
4. Larger prospective studies are warranted determine whether enzyme activity is associated with the development of PD.

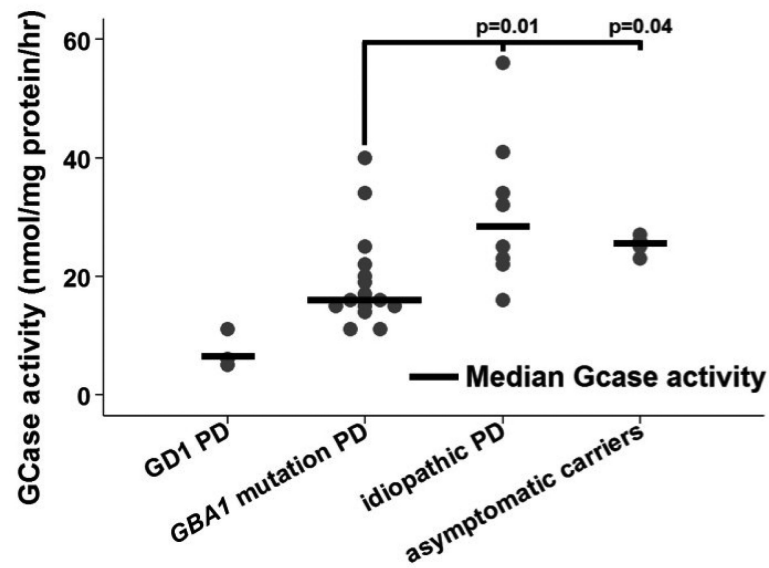


Fig. 1. Glucocerebrosidase enzyme (GCase) activity

Asymptomatic carriers = *GBA1* mutation carriers without Parkinson's disease, *GBA1* mutation PD = monoallelic *GBA1* mutation carriers with Parkinson's disease, GD1 PD = biallelic *GBA1* mutation carriers with Parkinson's disease, idiopathic PD = Parkinson's disease participants without *GBA1* mutations.

** GCase activity in GD1 PD (median 6.4, interquartile range 5.7) was lower than *GBA1* mutation PD ($p=0.01$). Each point represents the GCase activity measured for an individual.

Table 1

Summary of clinical features and glucocerebrosidase enzyme activity in Parkinson's disease and GBA1 mutation groups

	GD1 PD (n=3)	GBA1 PD (n=15)	IPD (n=8)	Non-PD carriers (n=4)	p value		
					GBA1 PD vs. IPD	GBA1 PD vs. non-PD carriers	IPD vs. non-PD carriers
Women, % (n)	66.7 (2)	40.0 (6)	25.0 (2)	50.0 (2)	0.66	0.57	0.55
Age, years	58.0 (7.0)	67.0 (10.0)	63.5 (11.5)	64.0 (4.5)	0.92	0.76	0.67
Onset age, years	53.0 (10.0)	59.0 (17.0)	57.5 (5.5)	---	0.90	---	---
Duration, years	5.0 (3.0)	10.5 (8.0)	5.0 (8.0)	---	0.67	---	---
H&Y stage	2.0 (1.5)	2.0 (1.0)	2.0 (0.0)	---	0.17	---	---
Activity^a	6.4 (5.7)	16.0 (7.0)	28.5 (15.0)	25.5 (2.5)	0.01	0.04	0.73

Data are presented as median (interquartile range) unless otherwise indicated. *GBA1* PD = monoallelic GBA1 mutation carriers with Parkinson's disease, GD1 PD = biallelic GBA1 mutation carriers with Parkinson's disease, H&Y = Hoehn and Yahr, IPD = idiopathic Parkinson's disease participants without GBA1 mutations, Non-PD carriers: GBA1 mutation carriers without Parkinson's disease, *vs.* = *versus*.

^a nmol/mg protein/hour.