

Inter- and intra-day variability in β -glucocerebrosidase activity and pathway biomarkers in healthy volunteers and patients with Parkinson's disease with a *GBA1* mutation

Jurrian P. van der Valk¹, Daniël B. Dumas¹, Eva Thijssen¹, Diana R. Pereira¹, Wieke Grievink¹, Yalçın Yavuz¹, Cedric Pesch¹, Daniel Ysselstein², Mackenzie Hagey², Omer Siddiqui², Jesse M. Cedarbaum², Kevin W. Hunt², Philip H.C. Kremer¹

1. Centre for Human Drug Research (CHDR) and Leiden University Medical Centre (LUMC), Leiden, the Netherlands
2. Vanqua Bio, Chicago, IL, U.S.A.

See all CHDR posters:



Background

Heterozygous mutations in the *GBA1* gene, which result in reduced β -glucocerebrosidase (GCCase) activity, are a major risk factor for Parkinson's disease (PD). Decreased GCCase activity is associated with impaired lysosomal function and alpha synuclein aggregation. A challenge for therapies targeting GCCase is the measurement of target and pathway engagement. In this study we used a novel approach to measure lysosomal GCCase assay to determine the suitability of this approach for target engagement with a GCCase targeted therapeutic and assessed plasma glucosyl β -sphingosine (GluSphing) as a potential pathway engagement marker.

Objectives

1. Evaluate a novel approach to assess lysosomal GCCase activity in whole blood samples from healthy volunteers (HVs) and PD patients with GBA mutation (GBA-PD), including inter- and intra-day variability
2. Determine plasma GluSphing levels in HVs and GBA-PD patients, including inter- and intra-day variability

Methods

- Whole blood was obtained from 8 HVs and 12 GBA-PD patients at 3 time points (Day 1, 0 and 4h, and Day 8, time matched to 0h).
- Lysosomal GCCase activity was assessed using the GCCase substrate PFB-FDGLu in whole blood, followed by flow cytometry analysis to assess activity in monocytes (Figure 1).
- A mass spectrometry assay was developed and qualified to enable quantification of plasma Glucosyl- and Galactosyl-Sphingosine.

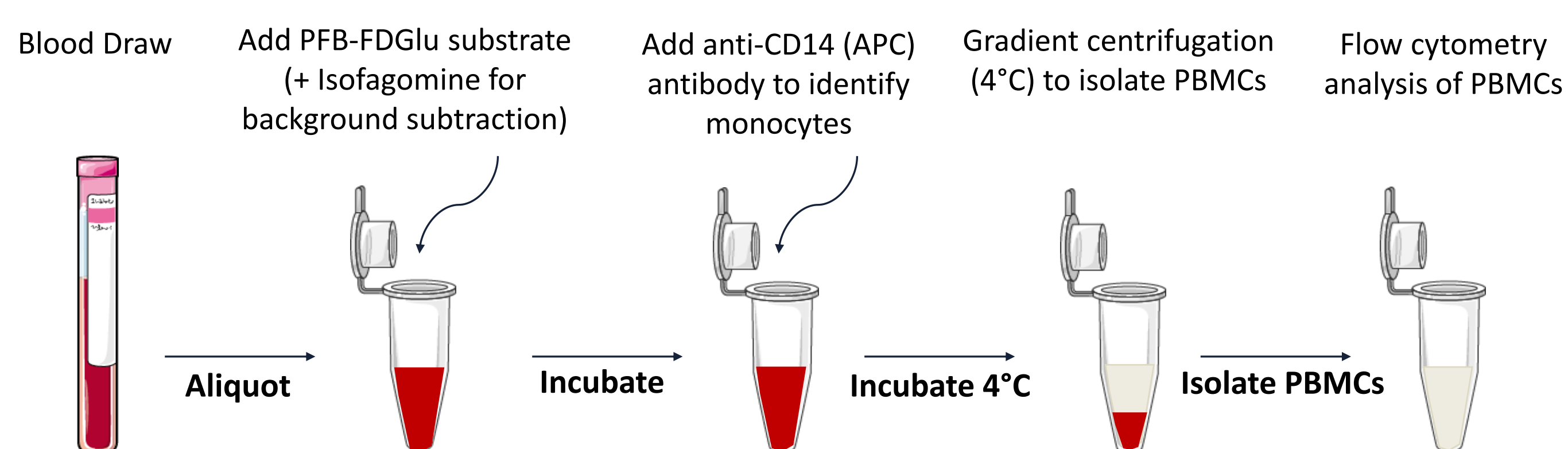


Figure 1. Overview of the approach to directly assess lysosomal GCCase activity in monocytes from whole blood samples

Results

Participant Information

	HVs	GBA-PD patients
Number	8	12
Age (y)	71.1 (2.6)	67.3 (6.6)
BMI (kg/m ²)	25.9 (2.1)	24.9 (3.5)
Gender, male (%)	88%	42%
Hoehn and Yahr	-	2.2 (0.9)

Table 1. Demographics

Mutation	Count (%)
E326K	3 (25%)
E326K/D140H	3 (25%)
T369M	3 (25%)
L444P	1 (8.3%)
R120W	1 (8.3%)
H490R	1 (8.3%)

Table 2. *GBA1* mutations

Lysosomal GCCase activity is reduced in GBA-PD relative to HVs

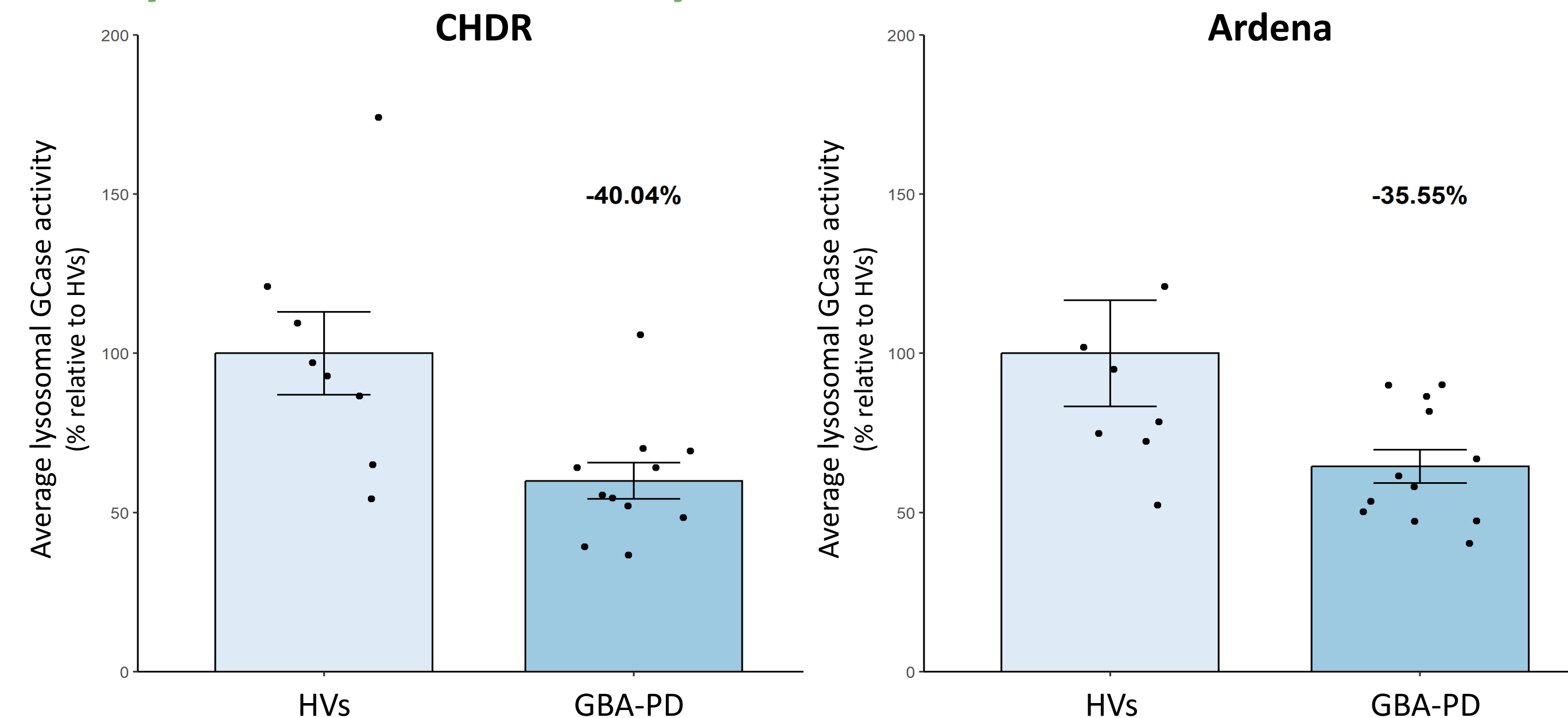


Figure 2. Lysosomal GCCase activity in GBA-PD patients is significantly reduced in matched blood samples analysed at two independent labs. Data is presented as the percent GCCase activity relative to HV average, \pm SEM, $p < 0.05$.

GCCase activity is consistently reduced in GBA-PD over time

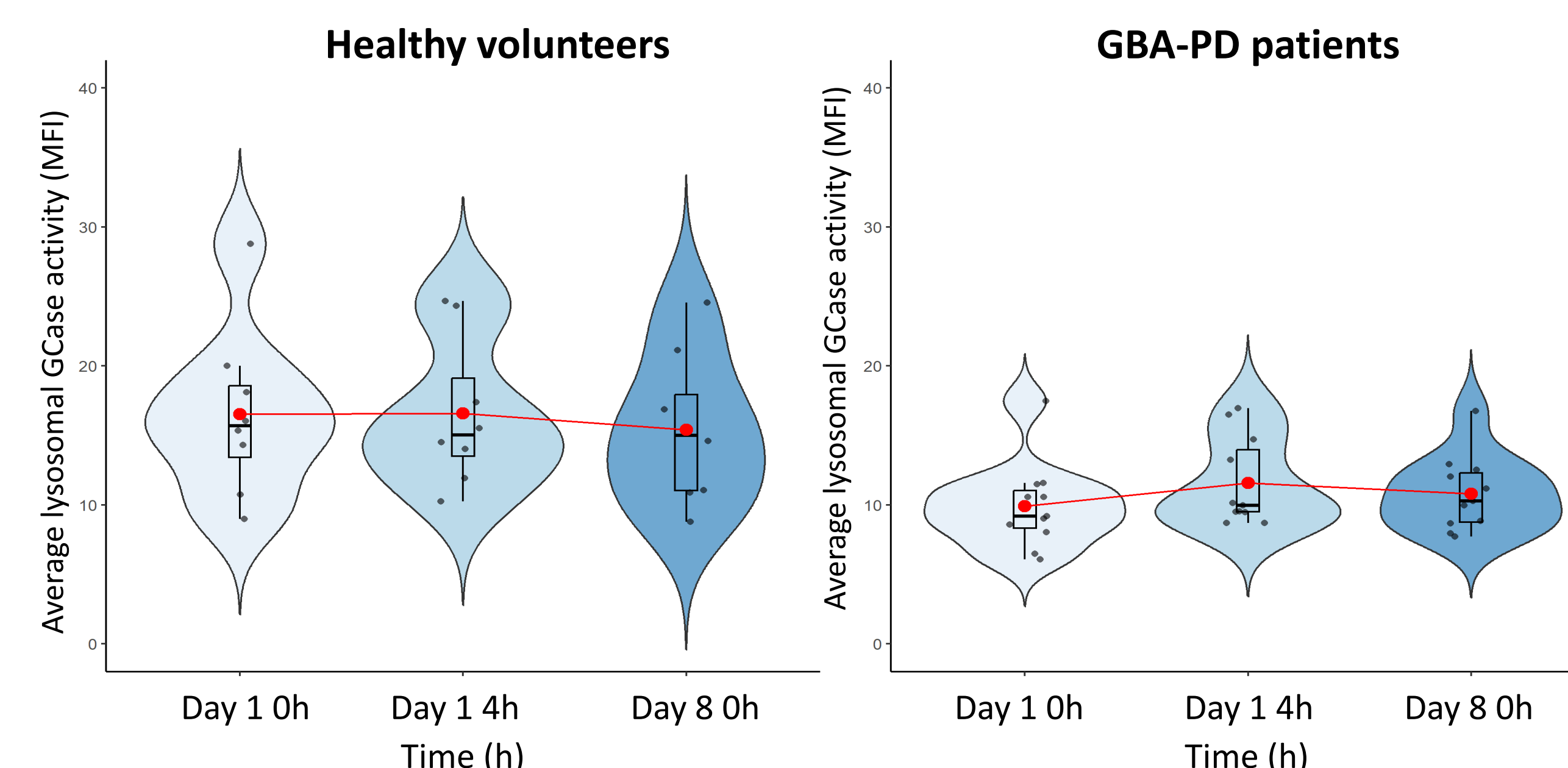


Figure 3. Average lysosomal GCCase activity measured in blood samples on Day 1 (0 and 4h) and Day 8 (0 h) was 33% lower across all time points in GBA-PD compared to HVs with minimal inter- and intra-day variability ($p = 0.004$).

Plasma Glucosylsphingosine is elevated in GBA-PD patients

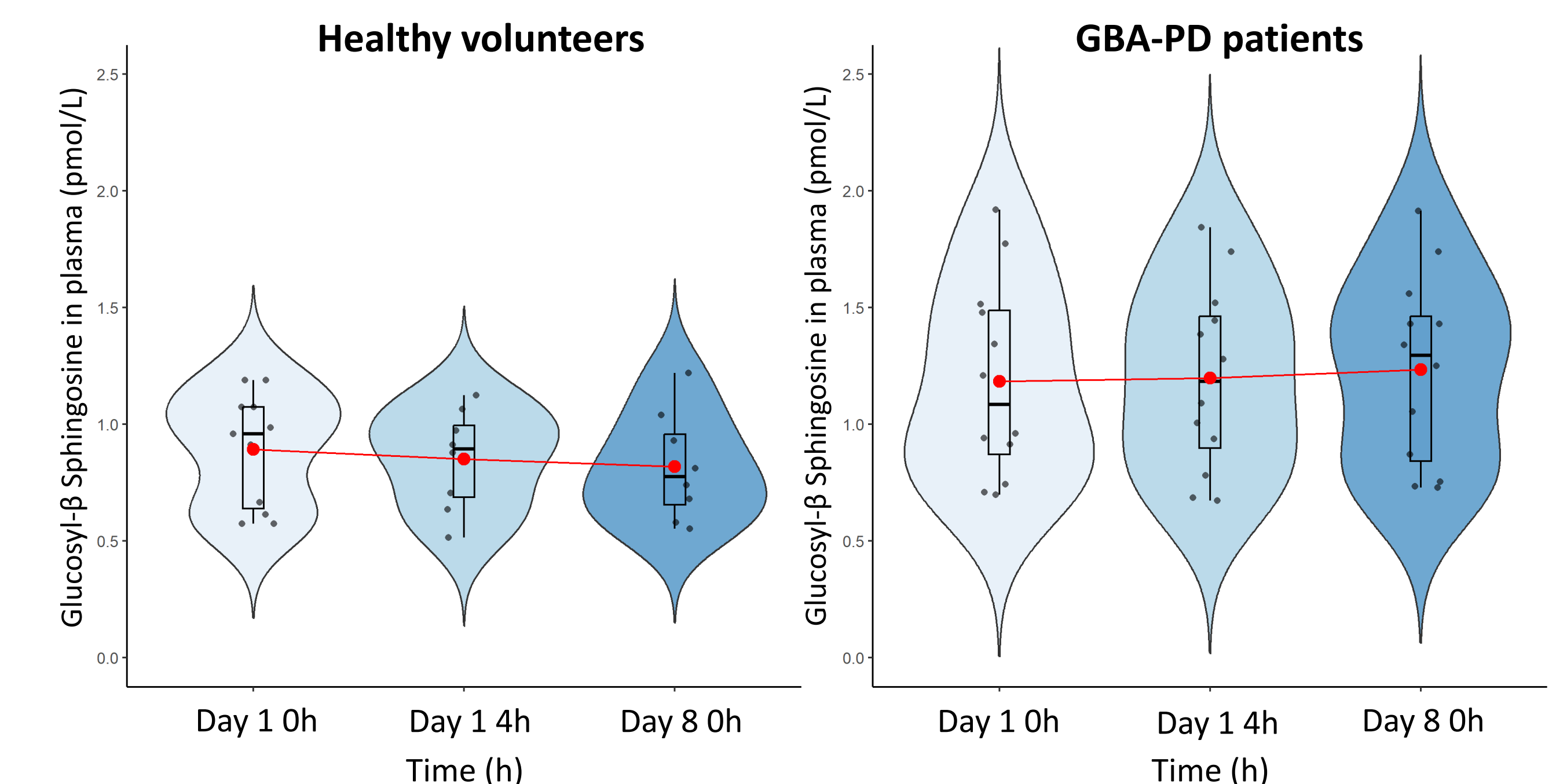


Figure 4. Plasma GluSphing concentrations in HVs and GBA-PD patients on Day 1, (0 and 4h), and Day 8, (time matched to 0h). GluSphing is 42% higher in GBA-PD but not statistically significant relative to HVs (0.31 pmol/mL (95% CI: -0.014, 0.64), $p = 0.06$).

Group	GCCase activity		Glucosyl- β Sphingosine	
	HV	GBA-PD	HV	GBA-PD
Inter-day (%CV)	24.0	37.1	2.4	1.7
Intra-day (%CV)	16.8	25.9	0.0	0.0

Table 3. Variability (%CV) in GCCase activity and Glucosyl- β sphingosine in HVs and GBA-PD

Conclusion

- Using a novel lysosomal GCCase activity assay, we observe biochemically relevant reductions in GCCase activity in whole blood samples from GBA-PD patients relative to HVs with minimal inter- and intra-day variability.
- Comparable GCCase activity data was generated by two independent labs demonstrating that this approach is transferrable and reproducible.
- Plasma glucosylsphingosine was increased in GBA-PD but was not statistically significant in this small cohort.
- These results support the use of this assay to assess target engagement of GCCase therapeutics in early clinical development.
- A further optimized and validated version of this assay is currently being used to assess target engagement of an allosteric GCCase activator in Phase 1 clinical studies.

Acknowledgements:

- We would like to acknowledge the bioanalytical lab Ardena for sample analysis.