

Systemic AAV gene therapy with CNS-targeted engineered capsids achieves significant GCase activity increases in the primate brain to support potential treatment of PD-GBA

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Introduction

Biallelic mutations in the GBA1 gene cause autosomal recessive Gaucher's Disease, the most common lysosomal storage disorder. More recently, epidemiological studies have revealed a role for mutations in this gene in the development of Parkinson's disease (PD), and current evidence suggests that heterozygous mutations within the GBAI gene are present in up to 15% of PD patients, making it the most significant genetic risk factor¹.

The GBA1 gene encodes a lysosomal hydrolase enzyme, glucocerebrosidase (GCase), which metabolizes glycolipids within the lysosomal compartment. Mutations in this gene result in decreased GCase activity, leading to accumulation of glycolipid species which are thought to contribute either directly or indirectly to α -synuclein pathology, the pathological hallmark of PD^{2, 3}. While certain aspects of the pathogenesis underlying PD-GBA remain unclear, substantial preclinical evidence suggests that interventions targeted to restore GCase activity in the brain could slow or stop the progression of PD-GBA^{4, 5, 6}.

Capsida is developing a gene supplementation therapy candidate (CAP-003) to be administered as a single intravenous (IV) infusion to PD-GBA patients. CAP-003 consists of an engineered AAV capsid that is designed to deliver the functional human GBA1 gene broadly across the CNS while de-targeting the liver and DRGs, which have been shown to be associated with safety issues in previous gene therapy programs. Using a loss-of-function mouse model we provide proof of concept pharmacology demonstrating that administration of the therapeutic cargo using a surrogate capsid results in dose-dependent increases in GCase activity that coincide with dosedependent decreases in glycolipid accumulation. In non-human primates using the clinical candidate, we show that administration of CAP-003 at low to moderate doses results in robust expression across the entire CNS along with increases in levels of GCase protein and activity which show strong correlations with similar measurements in biofluids. Given that a ~30% decrease in brain GCase activity is expected in the patient population⁷, the observed increases are expected to normalize activity levels, reduce glycolipids, and potentially slow or stop the progression of PD-GBA.

Methods and Materials

enabling assessment of target engagement engagement biomarkers. in CNS tissue achieved by a capsid that crosses the BBB in mice.

GBA1 LOF mouse model: GBA D409V KI (Jax **NHPs:** The CAP-003 development candidate was Strain# 019106) and WT mice (Jax Strain# administered IV to WT male and female 005304) received a retroorbital IV injection at cynomolgus macaques at ~33 months of age. 8-weeks of age. After 4-weeks cargo DNA and After 6-weeks, DNA biodistribution and mRNA RNA were assessed using PCR. GCase protein, expression were assessed using ddPCR. GCase activity, and glycosphingolipid content were protein, GCase activity, and glycosphingolipid assessed in CNS tissue using mass content were measured via mass spectrometry in spectrometry. A surrogate capsid (CAP.B10) both tissue as well as plasma and CSF to allow for was used to deliver the therapeutic cargo early assessment of planned clinical target



Figure 1. NHP-Driven Capsid Engineering Platform utilized to identify Capsid for CAP-003 for PD-GBA Capsida's high-throughput screening process in NHPs identifies capsids that target desired tissues and cell types while de-targeting undesired tissues such as liver and dorsal root ganglion which have been associated with adverse events

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Results





Figure 2. IV administration of CAP-003 in primates results in robust DNA biodistribution and mRNA expression across key areas of interest in the CNS while de-targeting peripheral organs including the liver and DRGs reducing safety risks associated with WT AAV9. AAV9 was IV delivered at a similar low dose. Representative histology in the substantia nigra was derived from a study that administered an epitope tagged version of CAP-003 to enable histological quantification of transduced cells.





Figure 3. AAV treatment results in dose-dependent increase in GBA1 protein and GCase activity. These increases coincide with significantly reduced GluSph levels in the CNS and validate target engagement of the hGBA1 cargo. GluSph 18:1 = Glucosylsphingosine, GalSph 18:1 = Galactosylsphingosine



• Capsida's engineering platform in NHPs has identified novel capsids with significant improvements in CNS penetrance and peripheral de-targeting relative to AAV9; CAP-003 is well tolerated and shows a favorable safety profile

- plans for IND filing in 1H 2025.

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Results

Increases in GCase protein and activity in the NHP brain significantly surpass expected therapeutic threshold and correlate with key biomarkers

Conclusions

• Proof-of-concept pharmacology in a GBA1 LOF mouse demonstrates dose-dependent increases in GCase protein levels and activity following administration of the hGBA1 cargo. These increases coincide with dose-dependent decreases in GluSph and indicate successful target engagement • CAP-003 achieves broad CNS distribution in non-human primates, including high expression in regions impacted by PD-GBA pathology and increases in GCase activity expected to normalize levels in the PD-GBA patient population. Importantly, exploratory biomarkers show strong relationships to GCase activity in the brain which sets expectations for clinical outcomes in similar measures • Capsida has initiated IND-enabling studies with the PD-GBA development candidate (CAP-003) with

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