

Targeting glucocerebrosidase with structurally targeted allosteric regulators corrects abnormal phenotypes in models of Parkinson's disease

A.M. García-Collazo¹, N. Pérez¹, E. Cubero¹, M. Montpeyó², A. Navarro-Romero², J. Riera², B. Guzman¹, M. Pons-Vizcarra¹, S. Morales¹, A. Ruano¹, A. Delgado¹, N. Callizot³, A. Henriques³, M. Martinez-Vicente², R. Maj¹, X. Barril¹

¹Gain Therapeutics, Lugano (Switzerland) and Barcelona (Spain).

²Vall d'Hebron Research Institute (VHIR), Neurodegenerative Diseases, Barcelona (Spain).

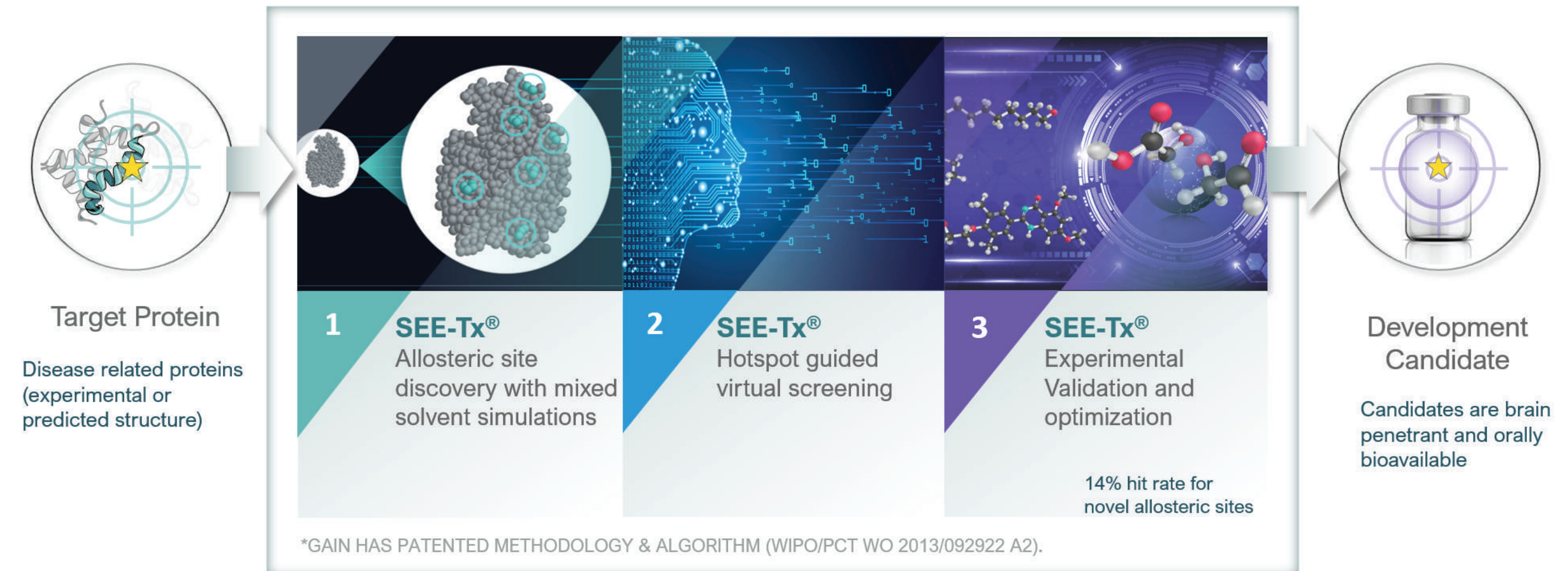
³Neuro-Sys SAS, Gardanne (France)

amgarcia@gaintherapeutics.com

Introduction

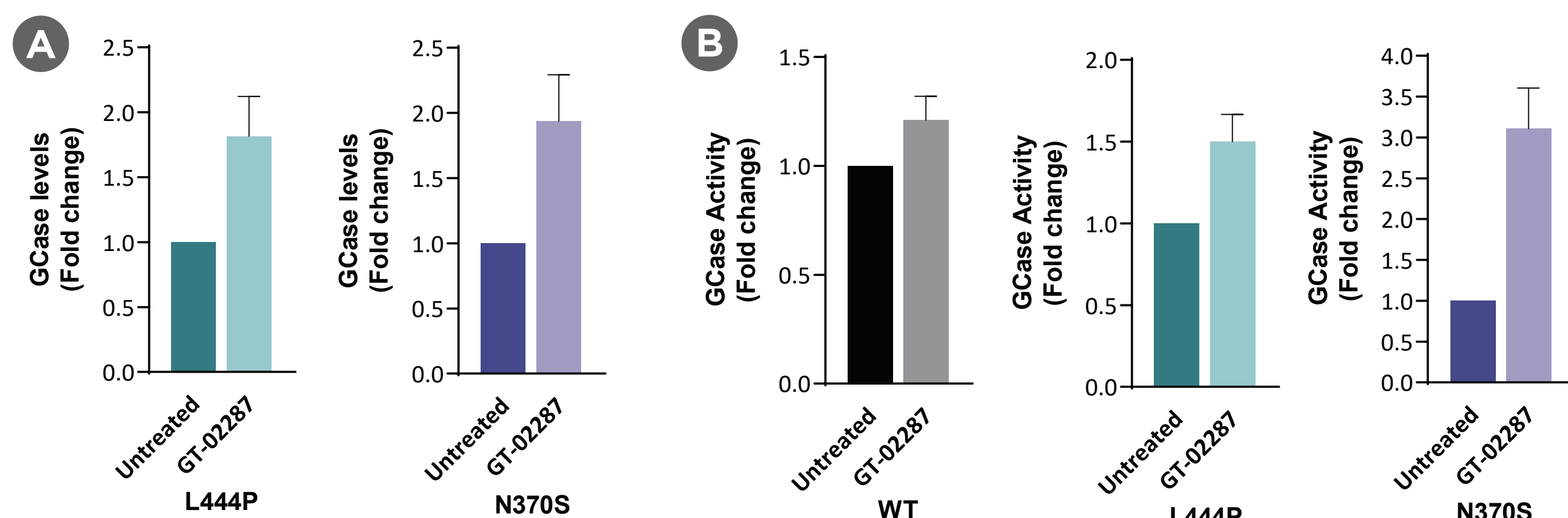
Mutations in the *GBA1* gene encoding the acid β -glucocerebrosidase (GCase) represent the most common genetic risk factor for Parkinson's disease (PD). The hallmark of PD is the presence of alpha-synuclein (α -syn) accumulation in specific areas of the brain. Interestingly, there appears to be an inverse relationship between GCase and α -syn levels: reduced GCase function is associated with increased α -syn accumulation as well as a change from its soluble form to its aggregated form, and it has been postulated that α -syn accumulation may reduce overall GCase activity. For these reasons, decreased GCase activity and levels may contribute to PD pathogenesis and restoring dysfunctional GCase may therefore represent a potential therapeutic strategy.

SEE-Tx™ Technology



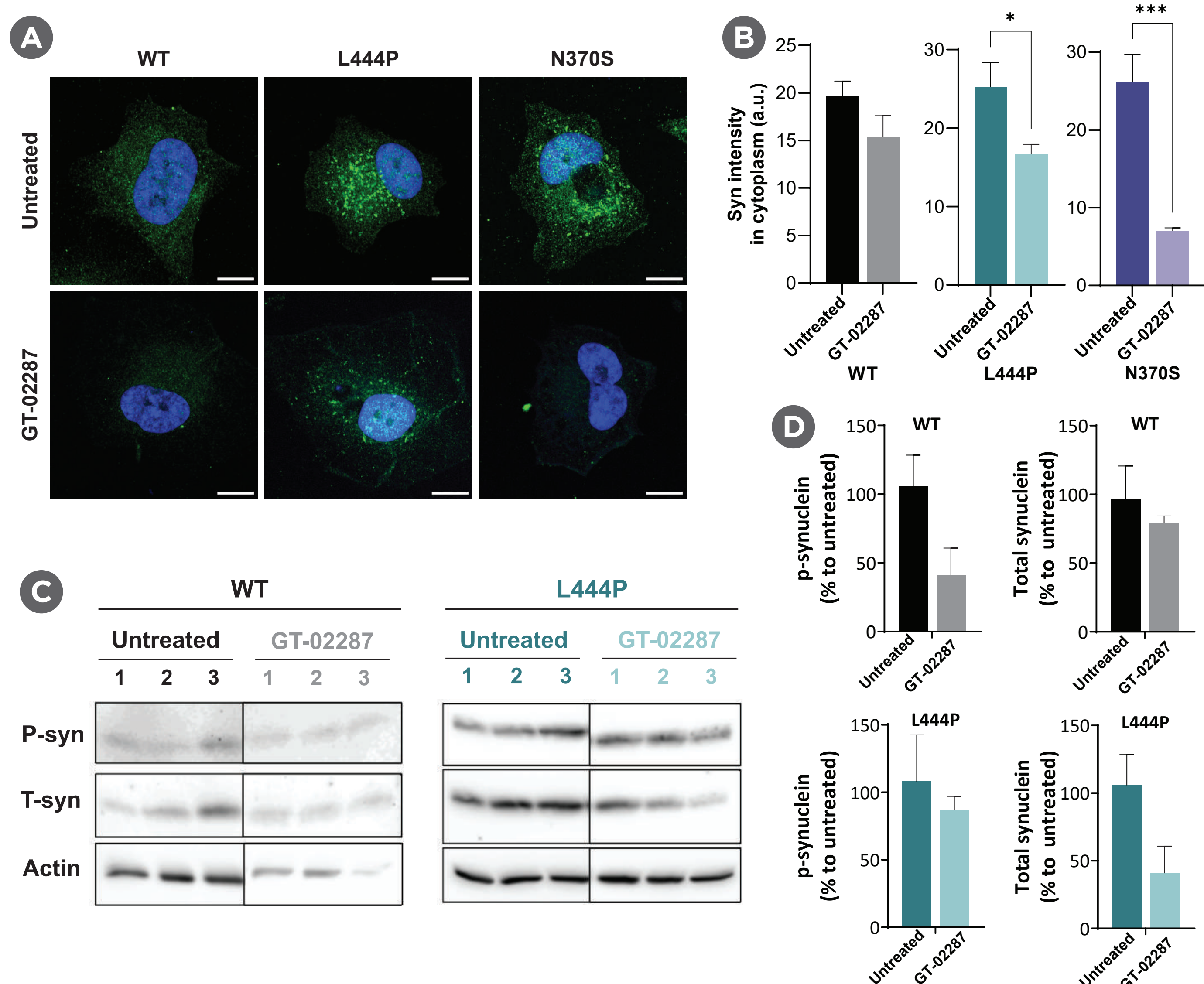
Aim: Restore GCase function using allosteric regulators to slow or stop PD progression

1 GT-02287 increases GCase levels and activity in a dopaminergic neuronal model



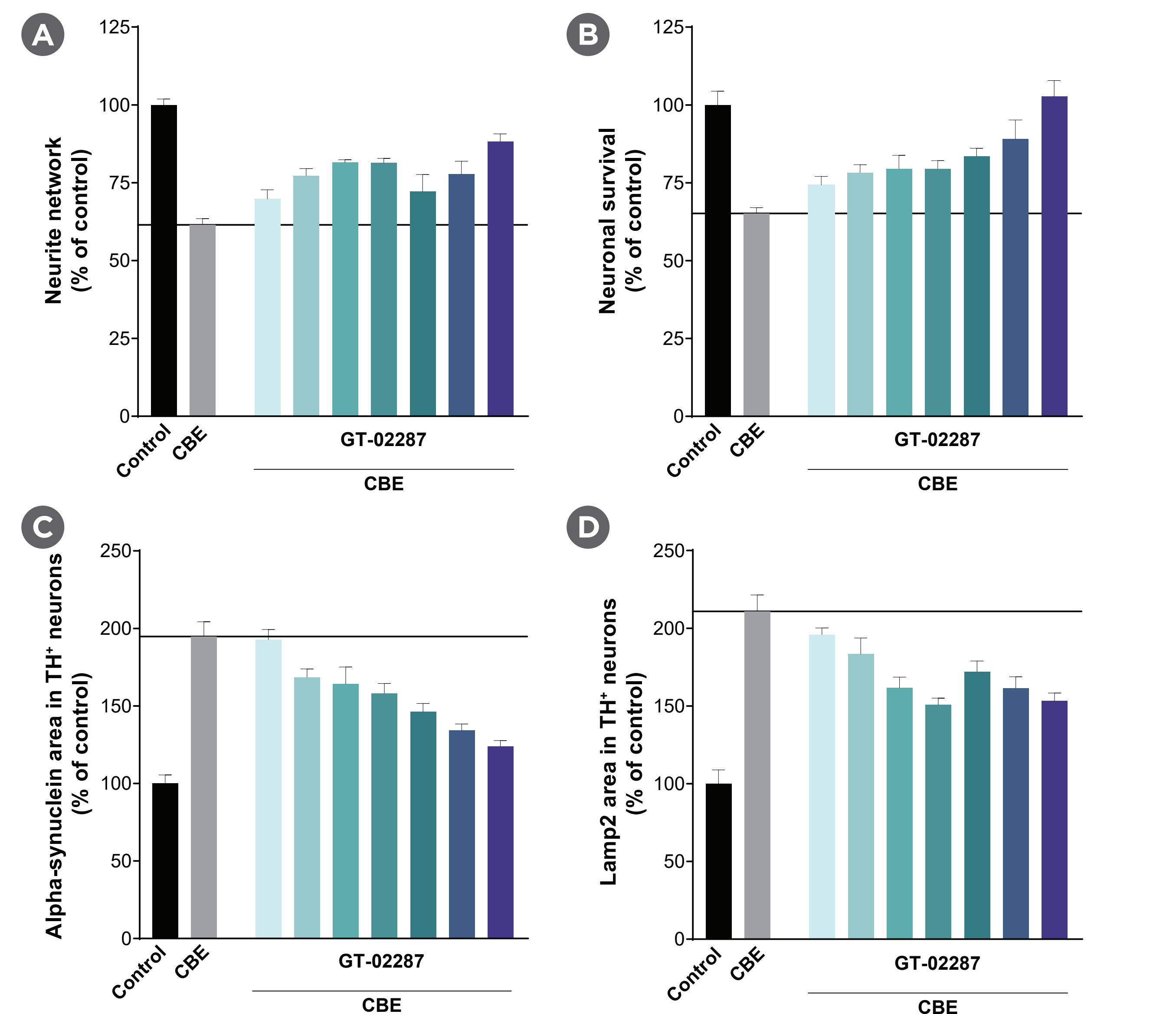
Dopaminergic neurons BE(2)-M17, WT or carrying either L444P or N370S *GBA1* mutations, were treated for 4 days with 25 μ M GT-02287. (A) GCase levels were evaluated in homogenates by western blot and normalized by Ponceau. (B) GCase activity was measured using 4-methylumbelliferyl- β -D-glucopyranoside and normalized to untreated. Results are presented as mean+SEM from 3 independent experiments.

2 GT-02287 reduces alpha-synuclein levels in WT, L444P and N370S BE(2)-M17 cells



Dopaminergic neurons BE(2)-M17, WT or carrying either N370S or L444P *GBA1* mutations, were treated for 10 days with 25 μ M GT-02287. (A) Cells were stained with anti- α -synuclein oligomer specific (Syn33) antibody (green) and nuclei were counterstained with DAPI (blue). Representative overlay images are shown. Scale bars: 10 μ m. (B) Syn intensity inside the cytoplasm was measured. (C) Typical WB of phosphorylated α -syn (p-syn) and total synuclein (T-syn). (D) Quantification of α -syn and phosphorylated α -syn (Ser 129 Ab). Results are presented as mean+SEM. One-way ANOVA (Welch correction) were used comparing each column with its corresponding untreated. Significance is denoted: * p <0.5, *** p <0.001

3 GT-02287 is neuroprotective and improves lysosomal health as well as synuclein pathology



Primary cultures of rat mesencephalic neurons were established. On day 6, GT-02287 was applied and after 24 hours, CBE (400 μ M) was added to the culture medium for 48 hours. On day 8, the culture was fixed and stained for tyrosine hydroxylase (TH), a marker for dopaminergic neurons. (A) Neurite network, (B) neuronal survival, (C) aggregated synuclein and (D) lysosomal area were evaluated.

Legend for Figure 3:
 ■ GT-02287 (10 nM)
 ■ GT-02287 (25 nM)
 ■ GT-02287 (50 nM)
 ■ GT-02287 (100 nM)
 ■ GT-02287 (500 nM)
 ■ GT-02287 (1 μ M)
 ■ GT-02287 (12.5 μ M)

Conclusions

SEE-Tx™ is a fast and cost-effective solution that has allowed us to identify structurally targeted allosteric regulators (STARs) of the GCase enzyme that are orally bioavailable and brain-penetrant.

GT-02287:

- Enhances GCase levels and activity in dopaminergic cells
- Effectively reduces alpha-syn in a neuronal cell model
- Increases neuronal viability and reduces lysosomal area and pathogenic synuclein in dopaminergic neurons

GT-02287 restores GCase-related key biological activities found to be impaired in many forms of PD, thus warranting further development towards the clinic.