# **Tolerability of HTT-Lowering: Lessons Learned from Nonhuman Primates**

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#### Abstract

Background: Huntington's disease (HD) is a neurodegenerative disease caused by CAG expansion in the huntingtin (HTT) gene, resulting in expression of mutant HTT (mHTT) protein. HTT-lowering is a potential approach for the treatment of HD actively being explored. A long-standing question is the extent to which HTT-lowering can be tolerated in adults. **Aims:** We sought to explore the tolerability of deep and sustained HTT-lowering in wild-type (wt) non-human primates (NHPs) over 6 months after single and repeated intrathecal administration of ALN-HTT02, a C16-conjugated small interfering RNA (siRNA) targeting HTT. Methods: Tolerability of HTT-lowering was assessed in NHPs through in-life clinical observations, neurological exams, detailed histopathology assessments, and multiple CSF parameters. HTT protein levels were assessed in CNS tissues at 3- and 6-months to confirm extent of lowering. Results: Treatment of NHPs with ALN-HTT02 achieved robust reduction in HTT protein expression (> 90% lowering in the cortex) and appeared to be well tolerated throughout the study period. There were no treatment-related clinical observations, abnormalities in neurological exams or histopathological changes, and no changes in any measured CSF parameters, including NfL, total protein, and immune cell counts. Conclusions: Our results suggest that deep, sustained lowering of HTT protein expression in the CNS of non-human primates by a C16-siRNA conjugate can be well tolerated for at least 6 months, supporting future clinical evaluation of ALN-HTT02, a C16-conjugated siRNA, as a novel platform approach to HTT-lowering in the CNS.

## Figure 1: Targeting *HTT exon1* with ALN-HTT02



## Figure 4: Sustained HTT-Lowering with Multiple Doses of ALN-HTT02 in NHP; Encouraging Safety and Tolerability Profile (Study 4)

**HTT Protein Lowering in Frontal Cortex** after Multiple IT Doses (Q3m x2)



**6m Multidose study** 

No in-life neurological abnormalities

Targeting exon 1 reduces expression of all pathogenic forms of mHTT in addition to wtHTT. ALN-HTT02 targets a conserved mRNA sequence in the first exon of HTT and is designed to reduce expression of all HTT mRNA species, resulting in subsequent HTT protein reduction. Adapted from Bates et al. (2015)<sup>1</sup>

#### Figure 2: Evidence of mHTT Exon 1 Isoform Target Engagement in **HD Mouse Model**

Control

Het

nge

100

50

Α. **Q175 Striatal Tissue Full-Length Htt mRNA** (Exons 50-53) \*\*\*\* 150 \*\*\*\* Het Control 50 ~



(5' region of Intron 1)

\*\*\*\*



- No adverse CSF parameter changes
- No adverse microscopic findings

No adverse findings, even after deep HTT-lowering (>90%)

HTT protein levels relative to aCSF control in prefrontal cortex at 6 months following multiple IT doses of ALN-HTT02 in cynomolgus monkeys.

Figure 5: Transient CSF NfL Elevation Returned to Baseline Following a Single IT Dose of ALN-HTT02 (Study 2)





Q175 HET; siHTT-FL (n=10) Q175 HET; Control (n=5) Q175 HET; siHTT-Ex1 (n=10)

Htt mRNA levels in the striatum of Q175 mice 3 months after ICV administration of vehicle or 300ug of a siRNA targeting HTT in exon 1 (Ex1) or downstream in the full-length (FL) region as detected by a QuantiGene multiplex panel.<sup>2</sup> A. Reduction of Full-length *Htt* mRNA by both siHTT-FL and siHTT-Ex1 as detected by probe set spanning exons 50-53 of the spliced transcript. B. Reduction of the Exon 1 fragment Htt mRNA transcript by siHTT-Ex1 as detected by a probe set designed against the intron 1 region 5' of previously identified cryptic polyA site.<sup>3</sup> Statistics performed on Normalized Mean Fluorescence Intensity (MFI) Values.

## Figure 3: Widespread, Dose-Dependent & Durable HTT-Lowering in NHP CNS After a Single IT Dose of ALN-HTT02 (Study 3)



**Timepoint (days)** 

NfL displayed as percent change from individual animal predose mean values. Early, transient increase was observed between Days 1 and 7 following IT dosing, with recovery to baseline by 1 month. Abbreviations: aCSF=artificial cerebrospinal fluid; CSF=cerebrospinal fluid; Nfl=neurofilament light chain. Note: Data is presented as the mean normalized to baseline with individual data points displayed; n=3-8/group; Dotted line denotes baseline

#### Table 1: Summary of HTT Lowering in 2-4 Year Old NHP by ALN-HTT02

Study	Peak Cortical Knockdown	Duration	Tolerability Assessments	Key Result
1	90%	59d		ALN-HTT02 was well tolerated
2	96%	85d, 172d	In-life Clinical and Neurological Evaluations	Well tolerated at all dose levels with no adverse treatment associated findings
3	84%	85d, 169d	Longitudinal CSF Analysis Post-Mortem Histopathology	Well tolerated at all dose levels with no adverse treatment associated findings
4	94%	86d, 170d		Well tolerated at all dose levels with no adverse treatment associated findings

#### Summary

HD is caused by CAG repeat expansion in the first exon of the HTT gene resulting in two distinct mHTT isoforms: full-length and a shorter Exon 1 fragment

HTT protein levels normalized to tissue from aCSF (control) animals with individual datapoints displayed as measured by ELISA A. Day 85 after single Low or Mid-level IT administration of ALN-HTT02. B. Day 85 or 169 after single Mid-level IT dose. Data are displayed as mean (±SD). aCSF, artificial CSF; CSF, cerebrospinal fluid; DLPFC, dorsolateral prefrontal cortex; HTT, huntingtin; IT, intrathecal(ly); OFC, orbitofrontal cortex; Rel., relative; SD, standard deviation.

- ALN-HTT02 leverages Alnylam's clinically validated C16-siRNA delivery platform<sup>4</sup> and aims to lower both mHTT isoforms by targeting a conserved sequence within Exon 1
- ALN-HTT02 has been evaluated in four independent NHP studies with no adverse findings to date
  - Widespread distribution across CNS regions
  - Durable HTT-lowering, supporting infrequent dosing in clinic
  - Encouraging safety profile through at least 6 months
  - No adverse findings, even after deep (>90%) HTT-lowering
- Encouraging profile supports further development of ALN-HTT02

#### References

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