

SELECT-HD: an adaptive first-in-human clinical trial to evaluate WVE-003, an investigational allele selective mHTT-lowering oligonucleotide, in early manifest Huntington's disease

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Forward-looking statements

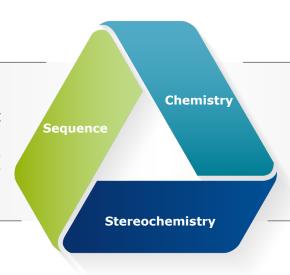
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PRISM Unlocking the body's own ability to treat genetic disease

DESIGN

Unique ability to construct stereopure molecules and control three structural features of oligonucleotides to efficiently engage biological machinery



OPTIMIZE

Provides the resolution to observe this structural interplay and understand how it impacts key pharmacological properties

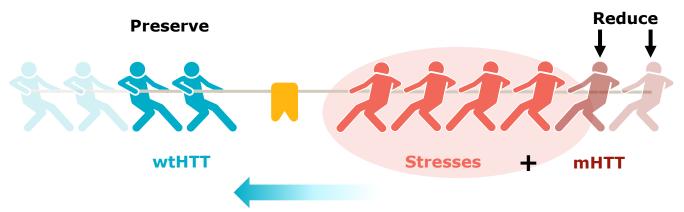
Built-for-Purpose Candidates to Optimally Address Disease Biology

Silencing | Splicing | RNA Editing



Allele-selective approach to treating HD

Preserve neuroprotective effects of wildtype HTT and reduce toxic mutant HTT





wtHTT supports healthy brain function, especially in the context of stress



Promotes neuronal survival



Essential role in synaptic protein transport



Supplies BDNF to striatum to regulate synaptic plasticity



Critical role in cilia function underlying CSF circulation needed to clear catabolites & maintain homeostasis

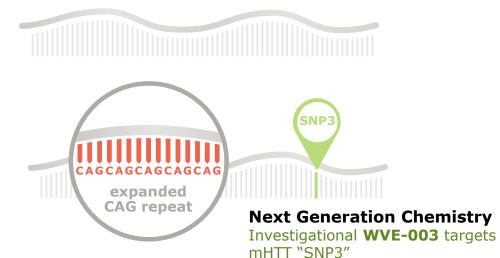


Allele-selectivity achieved by targeting downstream SNPs

Target mHTT transcript to selectively reduce mHTT protein with antisense oligonucleotides

Wildtype huntingtin RNA

Mutant huntingtin RNA







Introducing WVE-003, an investigational allele-selective oligonucleotide

Contains new chemistry with potential to address limitations of first-generation molecules

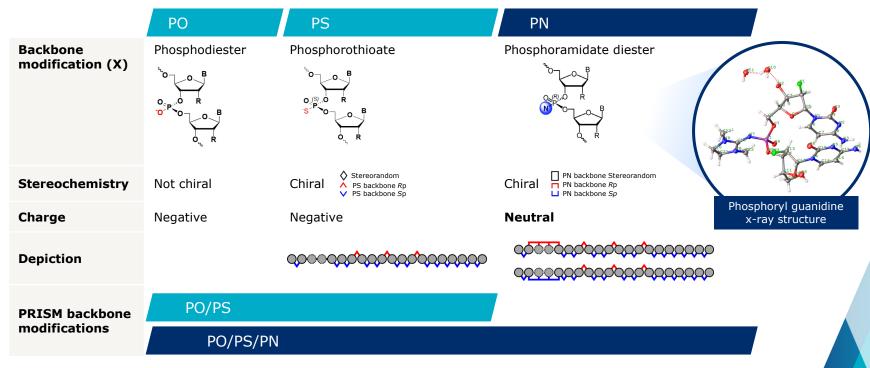
- ✓ Allele-selective approach designed to lower mutant HTT & preserve wildtype HTT
- New PN chemistry has demonstrated enhanced potency, tissue exposure & durability in CNS
- Preclinical pharmacological profile available in multiple species to inform clinical starting dose



WVE-003 contains backbone modifications with novel PN backbone chemistry



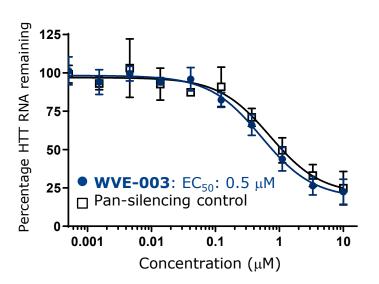
Backbone linkages



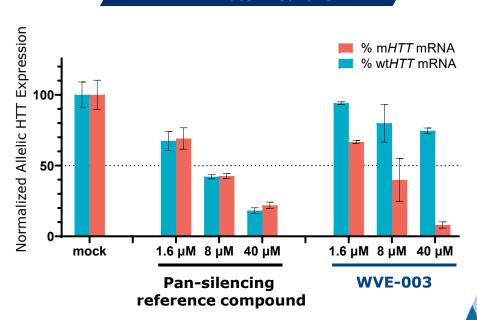


WVE-003 is potent and allele selective in vitro





Selectively reduces mHTT in HD motor neurons





Left: dose-response for HTT remaining in iPSC-derived motor neurons homozygous for SNP3, mean ± SD, n=4. Right: mHTT and wtHTT RNA expression in iPSC-derived motor neurons heterozygous for SNP3, total HTT knockdown quantified by qPCR and normalized to HPRT1 and mock treated sample. mean ± sem, n=4. iPSCs (induced pluripotent stem cells) generated from HD patient cells mHTT, mutant HTT; wtHTT, wild-type HTT

HD mouse models

Evaluation of potency



BACHD Mouse¹

HTT	Key characteristics
mHTT (human)	97 CAA-CAG repeatsMultiple copies
	❖ Subset of copies contain SNP3
wtHTT (mouse)	❖ Mouse genomic Hdh
	Lacks SNP3

Evaluation of selectivity



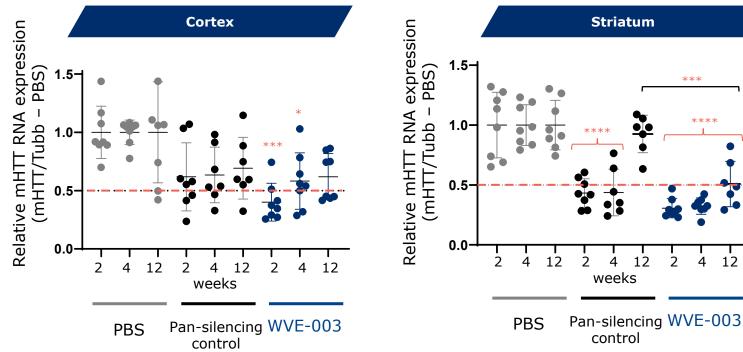
Hu97/18 Mouse²

HTT	Key characteristics
mHTT (human)	❖ 97 CAA-CAG repeats
	Multiple copies
	Subset of copies contain SNP3
wtHTT (human)	❖ 18 CAG repeats
	Lacks SNP3
	❖ Lacks mouse Hdh



WVE-003 has potent and durable effects in cortex and striatum

Maximum knockdown of 75% with ~50% knockdown persisting for at least 3 months



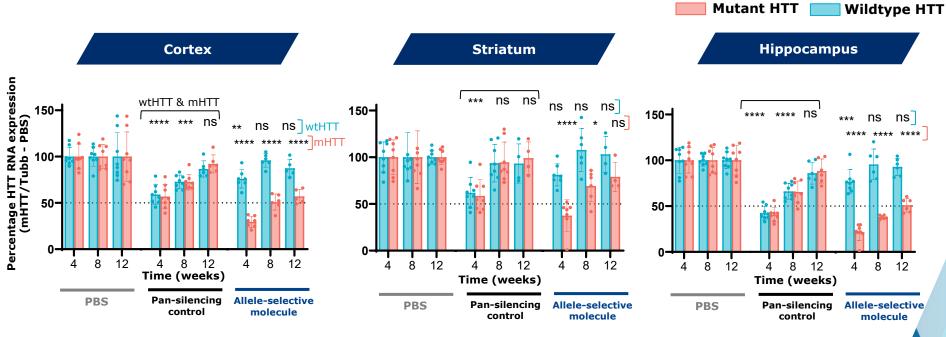


BACHD mice administered 3x100 µg intracerebroventricular doses PBS or oligonucleotide. (Left) Relative mHTT RNA in cortex at 2, 4 and 12-weeks post-dosing. (Right): Relative mHTT in striatum at same time points as cortex. BACHD contains SNP3 only in some mHTT transgenes. Data are mean ± SD, n=8. *P<0.0332, ***P<0.0002, ****P<0.0001 versus PBS unless otherwise noted). P values were calculated via 1-way analysis of variance, mHTT, mutant HTT; Tubb, tubulin

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Allele-selective activity in CNS of Hu97/18 mice

Allele-selective molecule **decreases mutant HTT, spares wildtype HTT**; whereas pan-acting molecule uniformly decreases wildtype HTT and mutant HTT





Preclinical pharmacological modeling available to inform clinical starting dose

BACHD

NHP

Human (cortex, striatum)







Ascending dose studies

- PK & mHTT knockdown data
- IC₅₀ determination

Concentrations in cortex and **striatum** sufficient for target engagement

Anticipated mHTT knockdown in cortex and striatum



Advancing investigational WVE-003 into a Phase 1b/2a clinical trial

Leverage learnings from PRECISION-HD clinical trials

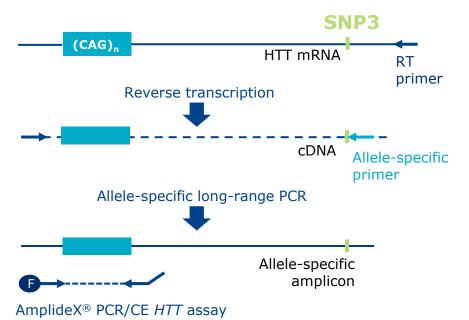


- √ Validated genotyping assay in CLIA setting to improve efficiency of patient identification
- ✓ Qualified biomarker assay to measure wildtype HTT protein in CSF
- ✓ Clinical experience of sites from PRECISION-HD



Rapid patient identification

Investigational assay enables SNP genotyping and phasing with CAG-repeat expansion



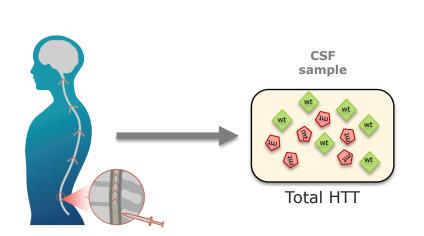
- CAG length, SNP zygosity and phasing information from a single assay
- PCR-based assay using in vitro diagnostic device ready capillary electrophoresis (CE) platform
- CLIA validated
- 1-2 weeks turn around time

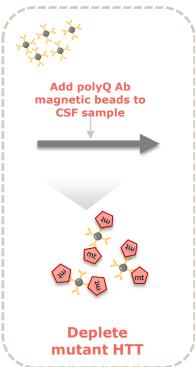




Assessment of wild-type HTT protein in CSF

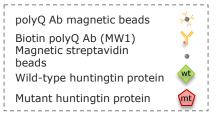
Immunodepletion of mHTT to measure wtHTT protein







Wild-type HTT

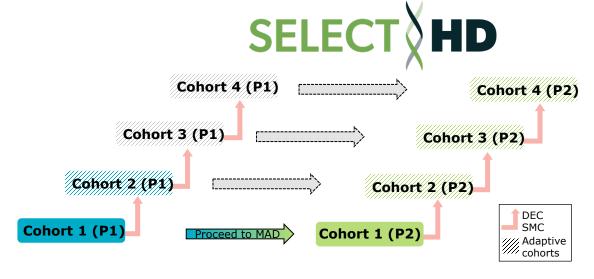


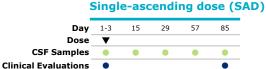


SELECT-HD: Adaptive first-in-human study for WVE-003

Ph 1b/2a global, multicenter, randomized, double-blind placebo-controlled trial

Eligible PRECISION-HD participants can transition to this study after wash out







Patients

- Targeting 36 patients
- ≥25 and ≤60 years of age
- Confirmed early manifest HD diagnosis with SNP3 variant

Primary Objectives

Safety and Tolerability

Secondary Objectives

- Plasma PK profile
- CSF exposure

Exploratory Objectives

Biomarkers

Clinical Endpoints

- mHTT NfL
- UHDRS
- wtHTT MRI



Adaptive cohorts: dose escalation and dosing interval guided by independent safety monitoring committee.

wtHTT, wild-type HTT; mHTT, mutant HTT; Nfl, neurofilament light chain; SNP, single nucleotide polymorphism; UHDRS, United Huntington's Disease Rating Scale; MRI, Magnetic
Resonance Imaging; PK, pharmacokinetic; PD pharmacodynamic; CSF, cerebrospinal fluid; SNP, single nucleotide polymorphism; DEC, Dose Escalation Committee; SMC, Safety

Monitoring Committee

Wave remains committed to an allele-selective approach in HD

SELECT HD

- Advancing WVE-003 into SELECT-HD, a Phase 1b/2a clinical trial
 - New chemistry has potential to address limitations of first-generation molecules in PRECISION-HD
 - Preclinical pharmacological profile available in multiple species in vivo
 - Initiate at pharmacologically active dose
 - Incorporates learnings from PRECISION-HD trials
 - Sites being activated, first patient has been dosed (Sept 2021)



Acknowledgements

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- EVOTEC
- IRBM
- Wave's scientists and Study Team

For additional information on SELECT-HD contact <u>clinicaltrials@wavelifesci.com</u> or visit clinicaltrials.gov (NCT05032196)

