



Wave Life Sciences
Fourth Quarter and Full Year
2021 Earnings
March 3, 2022



Forward-looking statements

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Paul Bolno, MD, MBA
President and CEO

Today's agenda

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Opening remarks - Paul Bolno, MD, MBA, President and CEO

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Clinical pipeline - Michael Panzara, MD, MPH, CMO, Head Therapeutics Disc. and Dev.

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AATD AIMer program update – Paloma Giangrande, PhD, VP Biology and Platform Disc.

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4Q 2021 financial results – Kyle Moran, CFO

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2022 outlook - Paul Bolno, MD, MBA, President and CEO

6

Q&A

Wave Life Sciences today

Building a leading genetic medicines company



Innovative RNA therapeutics platform

Stereochemistry, PN chemistry | Silencing, splicing, or editing oligonucleotides | Strong and broad IP¹

Clinical neurology pipeline: ALS/FTD, DMD, HD

- Genetic diseases with high unmet need
- Silencing CNS: ALS, FTD, HD
- Splicing Muscle: DMD

RNA editing with targeted GalNAc delivery: AATD

- Potential first-in-class AIMers for genetic hepatic diseases
- AATD is ideally-suited for an RNA base editing approach

Partnership opportunities to capitalize on assets

- Proprietary PN chemistry modifications improve potency duration, durability of effect of multiple therapeutic modalities
- Internal GMP oligonucleotide manufacturing facility

Wave's PN chemistry highlighted in two Nucleic Acids Research (NAR) manuscripts in 2022 thus far

Nucleic Acids Research, 2022 |
<https://doi.org/10.1093/nar/gkac018>

Control of backbone chemistry and chirality boost oligonucleotide splice switching activity

Pachamuthu Kandasamy^{1,†}, Graham McCloy^{2,†}, Mamoru Shimizu¹, Rowshon Alam¹, Naoki Iwamoto¹, Jayakanthan Kumarasamy¹, Gopa Adam Bezigian¹, Onanong Chivatakarn¹, David C. D. Butler¹, Michae Katarzyna Chwalenka¹, Kay E. Davies¹, Jigar Desai¹, Julli Dilip She Ruth Ellerington¹, Ben Edwards¹, Jack Godfrey¹, Andrew Hoss¹, Far Kenneth Longo^{1,3}, Genliang Lu¹, Subramanian Marappan¹, Jacopo C Erin Purcell Estabrook¹, Chiklu Shivalla¹, Maeve Tieshbeln¹, Tomor Carlo Rinaldi^{1,2,4}, Joana Rajão-Saravila¹, Snehlata Tripathi¹, Hallin Yi Xiansi Zhao¹, Cong Zhou¹, Jason Zhang¹, Luciano Apponi¹, Matthew Chandra Vargese^{1,4*}

¹Wave Life Sciences, Cambridge, MA, USA, ²Department of Paediatrics, University of Oxford OX1 3QX, UK, ³MDUK Oxford Neuromuscular Centre, University of Oxford, Oxford, UK, ⁴Department of Physiology, Anatomy and Genetics, University of Oxford, South Parks Rd

Received September 01, 2021; Revised December 18, 2021; Editorial Decision January 04, 2022; Accepted J.

ABSTRACT

Although recent regulatory approval of splice-switching oligonucleotides (SSOs) for the treatment of neuromuscular disease such as Duchenne muscular dystrophy has been an advance for the splice-switching field, current SSO chemistries have shown limited clinical benefit due to poor pharmacology. To overcome limitations of existing technologies, we engineered chimeric stereopure oligonucleotides with phosphorothioate (PS) and phosphoryl guanidine-containing (PN) backbones. We demonstrate that these chimeric stereopure oligonucleotides have markedly improved pharmacology and efficacy compared with PS-modified oligonucleotides, preventing premature death and improving median survival from 49 days to at least 280 days in a dystrophic mouse model with an aggressive phenotype. These data demonstrate that chemical optimization alone can profoundly impact oligonucleotide pharmacology and highlight the potential for continued innovation around the oligonucleotide backbone. More specifically, we conclude that chimeric stereopure oligonucleotides are a promising splice-switching modality with potential for the treatment of neuro-

muscular and other gene cult to reach tissues such heart.

INTRODUCTION

Exon-skipping oligonucleotide technology, are designed to base pair to pre-mRNA, to bypass exon(s) to restore and protein production is. The recent FDA approval pholino oligomers (PMOs) muscular dystrophy has for the exon-skipping field PMOs, they have demon modifying benefit. Because dystrophin protein and have [/www.fda.gov/news-events/grants-accelerated-approval/rare-duchenne-musculai-dy](https://www.fda.gov/news-events/grants-accelerated-approval/rare-duchenne-musculai-dy) [/www.fda.gov/news-events/approves-targeted-treatment/dystrophy-mutation](https://www.fda.gov/news-events/approves-targeted-treatment/dystrophy-mutation); bit NSPharma_Long-term_1dat continues to be substanti

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The authors wish it to be known that, in their opinion, the first two authors should be regarded as Joint First Authors.

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Nucleic Acids Research, 2022 |
<https://doi.org/10.1093/nar/gkac037>

NAR Breakthrough Article

Impact of guanidine-containing backbone linkages on stereopure antisense oligonucleotides in the CNS

Pachamuthu Kandasamy¹, Yuanjing Liu¹, Vincent Aduda, Sandheep Akare, Rowshon Alam, Amy Andreucci, David Boulay, Keith Bowman, Michael Byrne, Megan Cannon, Onanong Chivatakarn, Julli Dilip Shekhe, Naoki Iwamoto, Tomomi Kawamoto, Jayakanthan Kumarasamy, Sarah Lamore, Muriel Lemaître, Xuena Lin, Kenneth Longo, Richard Looby, Subramanian Marappan, Jake Metterville, Susovan Mohapatra, Bridget Newman, Ik-Hyeon Paik, Saurabh Patil, Erin Purcell-Estabrook, Mamoru Shimizu, Pochi Shum, Stephany Standley, Kris Taborn, Snehlata Tripathi, Hallin Yang, Yuan Yin, Xiansi Zhao, Elena Dale and Chandra Vargese^{1*}

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Received June 30, 2021; Revised December 17, 2021; Editorial Decision January 10, 2022; Accepted January 13, 2022

ABSTRACT

Attaining sufficient tissue exposure at the site of action to achieve the desired pharmacodynamic effect on a target is an important determinant for any drug discovery program, and this can be particularly challenging for oligonucleotides in deep tissues of the CNS. Herein, we report the synthesis and impact of stereopure phosphoryl guanidine-containing backbone linkages (PN linkages) to oligonucleotides using through an RNase H-mediated mechanism, using *Malat1* and *C9orf72* as benchmarks. We found that the incorporation of various types of PN linkages to a stereopure oligonucleotide backbone can increase potency of silencing in cultured neurons under free-uptake conditions 10-fold compared with similarly modified stereopure phosphorothioate (PS) and phosphodiester (PO)-based molecules. One of these backbone types, called PN-1, also yielded profound silencing benefits throughout the mouse brain and spinal cord at low doses, improving both the potency and durability of response, especially in difficult to reach brain tissues. Given these benefits in preclinical models, the incorporation of PN linkages into stereopure oligonucleotides with chimeric backbone modifications has the potential to render regions of the brain beyond the spinal cord more ac-

cessible to oligonucleotides and, consequently, may also expand the scope of neurological indications amenable to oligonucleotide therapeutics.

INTRODUCTION

Antisense oligonucleotides can be designed to promote degradation of a targeted transcript through an RNase H-dependent mechanism. RNase H binds a heteroduplex formed between a DNA-containing oligonucleotide and the targeted RNA and cleaves the RNA (1). Unmodified antisense oligonucleotides have poor pharmacokinetic (PK) properties, so chemical modifications are needed to enable their use in preclinical research and as therapeutics. Phosphorothioate (PS) modification of the phosphodiester (PO) backbone is one of the most common backbone modifications used in research and to improve the properties of oligonucleotide therapeutics (1,2). A consequence of PS modification is the creation of a chiral center, where the molecule can adopt either an Rp or Sp configuration (1,3). We have previously demonstrated that PS-modified stereopure oligonucleotides—with precisely controlled chirality of inter-molecule PS backbone linkages—can outperform stereorandom oligonucleotides—those racemic mixtures derived from traditional chemistries for which the chiral configuration of the backbone is not controlled (4,6). New backbone modifications and methods to control their stereochemistry continue to emerge (3,4,7,9).

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NAR press alert:

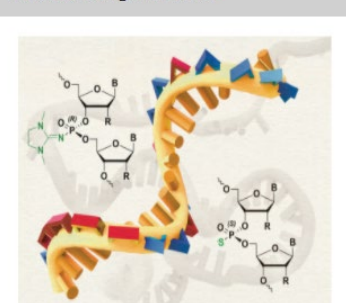
...“these data are the strongest evidence to date for the value of stereopure linkages in oligonucleotides designed to reduce gene expression...”

...Wave has “something unique, the data shows clear advantages of stereopure vs stereorandom P linkages. The duration of gene silencing achieved is impressive...”

...“work offers an encouraging suggestion that novel chemical approaches can push the frontiers of oligonucleotide drug development.”

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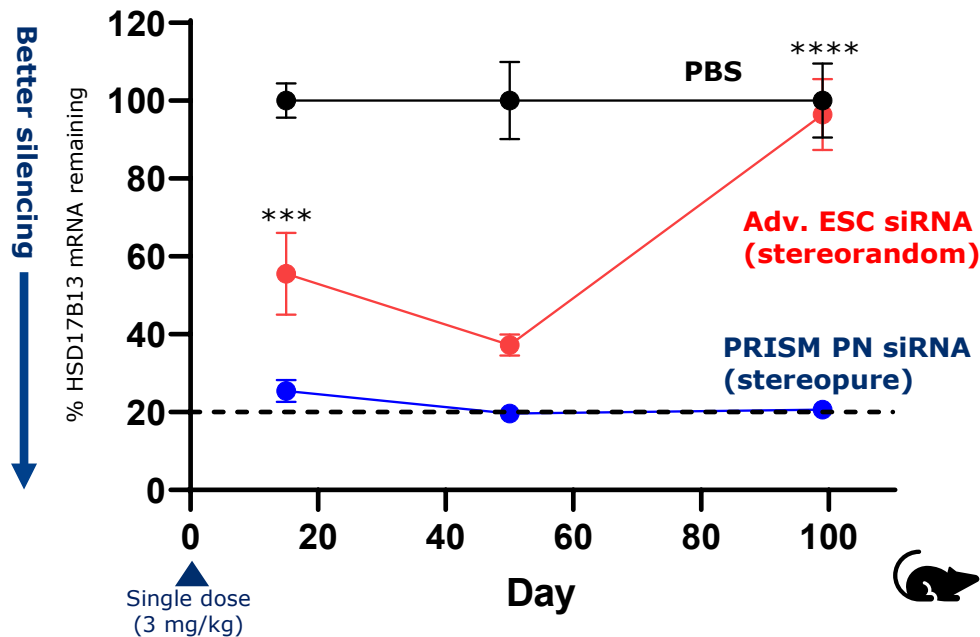
Breakthrough Articles



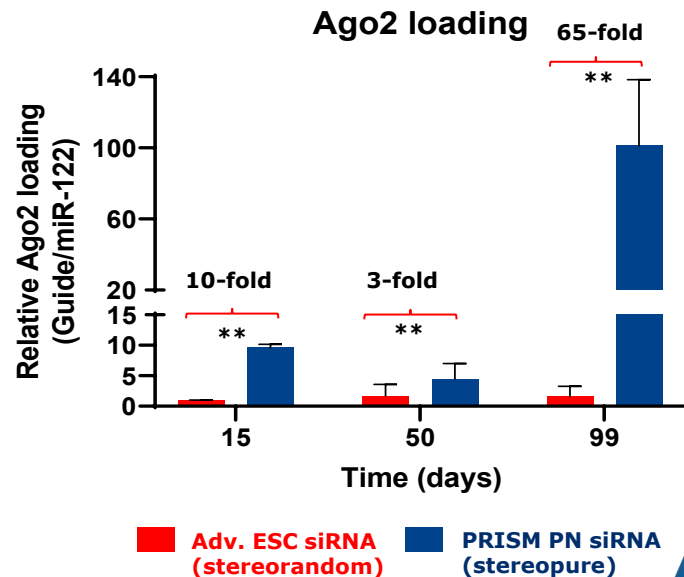
Impact of guanidine-containing backbone linkages on stereopure antisense oligonucleotides in the CNS

PRISM PN siRNA led to unprecedented silencing as compared to state-of-art >3 months after single dose

~80% silencing HSD17B13 mRNA *in vivo* with GalNAc-conjugated PRISM PN siRNA 14 weeks post single dose

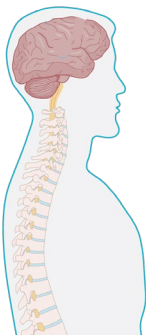


PRISM PN siRNA loaded in RISC is significantly greater than Adv. ESC siRNA

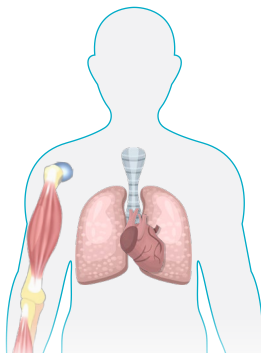


Diversified strategy to rapidly assess the potential of PN candidates across tissue types and modalities

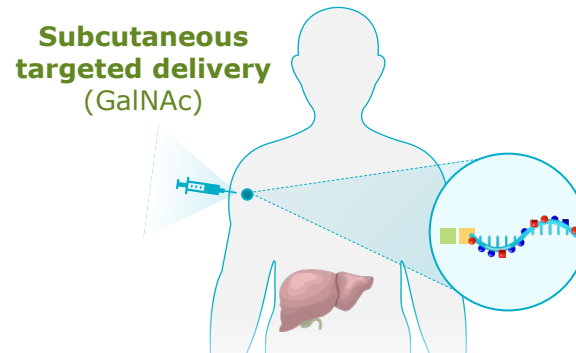
Silencing
CNS



Splicing (exon-skipping)
Muscle



RNA editing (AIMers)
Liver



WVE-004
C9 - ALS/FTD

WVE-003
SNP3 - HD

WVE-N531
Exon 53 - DMD

AIMer
SERPINA1 - AATD

AIMers are short and stable (fully chemically modified) to enable GalNAC-mediated delivery

Clinical data expected in 2022 for WVE-004, WVE-003, WVE-N531 to provide insight into PN chemistry and enable decision making for each program

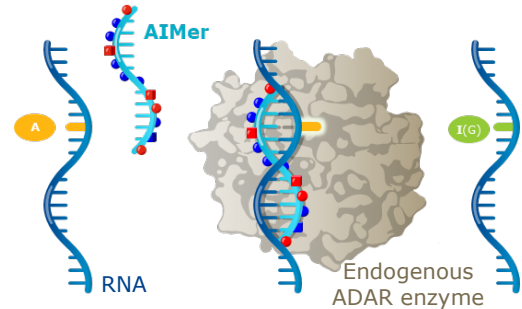
AIMer opportunity: novel RNA editing therapeutics, starting with targeted delivery to hepatocytes

Initial focus on correcting driver mutations of genetic hepatic diseases

Modulate protein interactions with AIMers

Lead program:
Alpha-1
antitrypsin
deficiency
(AATD)

Restore or correct protein function



Upregulate expression

Modify function

Modulate protein-protein interaction

Post-translational modification

Alter folding or processing

Cardiometabolic
Oncology
Immunology
Neurological disorders

- Clinically proven GalNAc-mediated delivery
- Potential for first-in-class therapeutics
- **AATD**: ~200K people in US and EU with mutation in *SERPINA1* Z allele (PI*ZZ)

- Large patient populations
- Human Reference Interactome documents >50K protein-protein interactions involving >8K proteins¹
- >90K Post-translational modifications across ~30K proteins mapped,² thousands associated with disease³

Established internal GMP manufacturing for multiple oligonucleotide modalities

Strong technical knowhow and operating expertise

- Experienced team led by Sridhar Vaddeboina, PhD (SVP Chemistry, Manufacturing, Controls)
- Experts in oligonucleotide synthesis (ASOs, DNAs, RNAs, siRNAs)
- Proven track record scaling complex chemistries; delivered clinical supply for six programs at Wave

Established infrastructure

- State of the art facilities (90,000 sq ft) and expansion space
- Process and analytical development labs
- GMP oligonucleotide (API) manufacturing
- Established Quality and GMP systems (QA, supply chain, logistics, QC testing)



Scalable to support Wave's GMP manufacturing needs, as well as potential new partners



Mike Panzara, MD, MPH
Chief Medical Officer,
Head of Therapeutics
Discovery and Development

PN chemistry enhances potency, tissue distribution and duration of effect of splicing compounds in muscle

Benefits of PN chemistry for exon-skipping

- ✓ Increased exon-skipping activity *in vitro*
- ✓ More stable in cultured myoblasts than PS oligonucleotides
- ✓ Increased muscle exposure in mice and NHPs
- ✓ Increased exon-skipping activity in mice and NHPs
- ✓ Exon-skipping activity well correlated with restored dystrophin expression in mice

Exon 53

WVE-N531

Exon skipping candidate in DMD

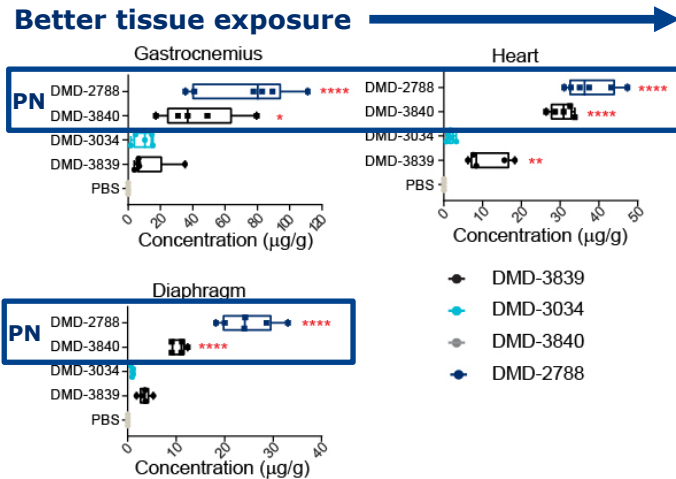
Preclinical mouse models

Non-human primate studies

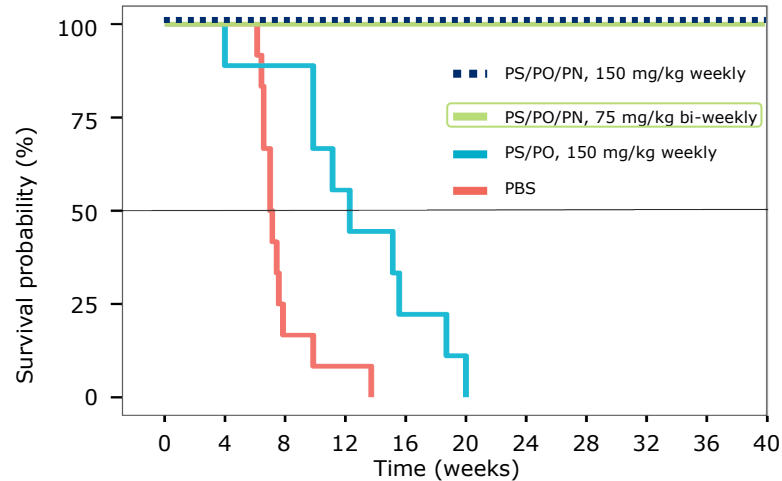
Clinical trial (ongoing)

PN chemistry improved muscle exposure and survival in preclinical mouse models

PN boosted muscle concentrations after single dose, which correlated with exon-skipping activity



Treatment with PN-modified molecules led to 100% survival of dKO mice at time of study termination



Note: Untreated, age-matched mdx mice had 100% survival at study termination [not shown]

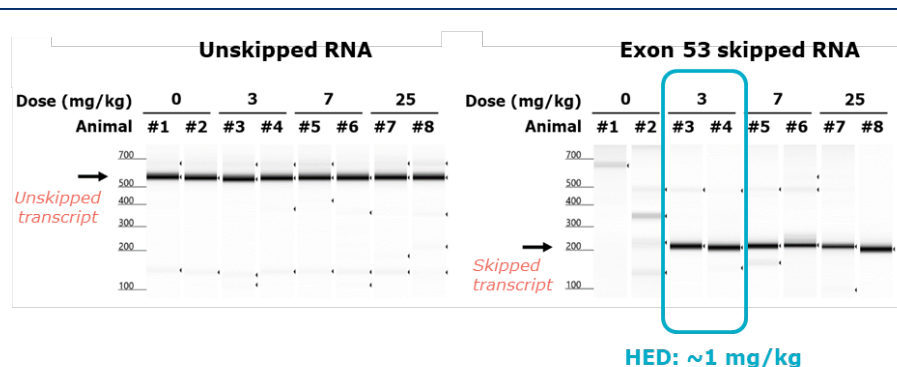
WVE-N531: PN chemistry enhances muscle distribution and exon-skipping in NHPs

Plasma and tissue concentrations of WVE-N531 (PS/PO/PN) significantly higher than suvodirsen (1st-gen PS/PO) in multiple NHP studies

- ✓ Substantially higher muscle concentrations (including heart and diaphragm) as compared to suvodirsen
- ✓ Higher plasma C_{max}, AUC and C_{trough}

WVE-N531 leads to exon-skipping in NHPs at doses significantly lower than suvodirsen

6 weekly doses of 3 mg/kg



Healthy NHPs have normal levels of dystrophin, but target engagement can be assessed by detection of skipped transcript

WVE-N531 plasma concentrations at starting dose significantly improved over suvodirsen

WVE-N531 Phase 1b/2a open-label clinical trial starting dose

Dose escalation is ongoing

	WVE-N531 (PN chemistry) fold increase over suvodirsen at the same dose level	
Plasma:		
C_{max}	~2.5x	↑ Increase in plasma concentrations with single dose
AUC	~4x	
Muscle:	<i>Patient muscle biopsies expected in 2022</i>	

WVE-N531 plasma half-life estimated to be >1 week

(vs. less than 24 hours for suvodirsen)

Dose escalation ongoing in clinical trial of WVE-N531

- Open-label clinical trial of boys with DMD amenable to exon 53 skipping
- Dose level and dosing frequency guided by tolerability and plasma PK

Initial cohort

- Ascending intra-patient doses of WVE-N531
- Up to 4 dose levels (administered ≥ 4 weeks apart) evaluated to select dose level for multidose
- Up to 3 additional doses given every-other-week at selected dose level, followed by muscle biopsy

Cohort expansion to be guided by assessment of muscle biopsies: (drug distribution in muscle and biomarkers)

Possible cohort expansion (up to 15 boys)

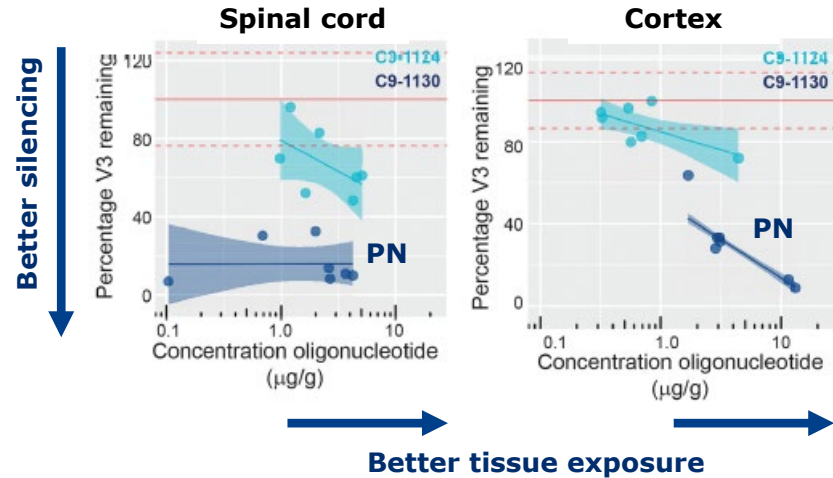
- Additional patients enrolled and dosed every other week at selected dose level
- Up to 7 total doses to be given followed by a minimum 8-week safety monitoring period
- Powered to evaluate change in dystrophin expression

Clinical data, including muscle biopsies, expected in 2022

PN chemistry enhances tissue exposure, potency and duration of silencing in CNS of mice models

Benefits of PN chemistry in CNS

- ✓ Increased potency in neurons
- ✓ Well-tolerated in multiple *in vitro* & *in vivo* assays
- ✓ Increased potency & extended durability of silencing in mice
- ✓ Enhanced tissue exposure in mice



FOCUS-C9 trial recognized as innovative clinical study for patients with C9-FTD, C9-ALS or both



Alzheimer's
Drug Discovery
Foundation



ADDF and AFTD Partner to Support Wave Life Sciences' FTD and ALS Clinical Program

NEWS PROVIDED BY
Alzheimer's Drug Discovery Foundation →
Jan 25, 2022, 11:07 ET

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NEW YORK, Jan. 25, 2022 /PRNewswire/ -- The Alzheimer's Drug Discovery Foundation (ADDF) and The Association for Frontotemporal Degeneration (AFTD) announced today that they have partnered to support Wave Life Sciences' FOCUS-C9 Phase 1b/2a clinical trial investigating WVE-004 as a potential treatment for C9orf72-associated frontotemporal degeneration (C9-FTD), as well as amyotrophic lateral sclerosis (C9-ALS). The partnership provides an investment from ADDF and AFTD that will support the evaluation of fluid biomarkers, functional assessments, and digital biomarkers in FOCUS-C9, potentially leading to clinically meaningful endpoints to inform development of treatments for FTD.

ADDF and AFTD made the decision to support the FOCUS-C9 trial following a review of Wave's clinical research application for the Treat FTD Fund, which supports the development of new medicines to treat FTD. Specifically, members of the Treat FTD Fund Joint Steering Committee, a panel of experts convened by ADDF in collaboration with AFTD, and ADDF's Scientific Review Board reviewed and commented on the Phase 1b/2a study plan, preclinical data supporting the program, and credentials of the study team.

- Alzheimer's Drug Discovery Foundation (ADDF) and The Association for Frontotemporal Degeneration (AFTD) partnered to support FOCUS-C9 clinical trial
 - Evaluation of fluid biomarkers
 - Functional assessments
 - Digital biomarkers

FOCUS  **C9**

Wave's allele-selective approach to Huntington's disease highlighted at CHDI conference

- CHDI's 17th Annual HD Conference (Feb 28th – March 3)
- Interest on the importance of preserving wtHTT in the context of HTT reducing treatments remains high in the HD community
- Wave poster: "A novel quantitative wild-type huntingtin (wtHTT) protein biomarker method for human cerebrospinal fluid"
- Wave presentation: "Innovations that led to SELECT-HD, a Phase 1b/2a clinical trial of an allele-selective therapy for Huntington's Disease"

SELECT HD

WAVE
LIFE SCIENCES

A Novel Quantitative Wild-type Huntingtin (wtHTT) Protein Biomarker Method for Human Cerebrospinal Fluid

Ramakrishna Boyanapalli, Dhanu Xu, Jaya Gopal, Manuul Daddar, Nedra Genovese, Alberto Bresciani, Michael Penczari
Wave Life Sciences, Cambridge, MA, USA; Department of Translational Research 2024 S.p.A.

Summary

- We report the development and validation of a novel method to quantify wild-type huntingtin (wtHTT) protein in human cerebrospinal fluid (CSF) samples.
- Our method comprises three steps:
 - First, total HTT protein is quantified using an optimized 2B7-D7F7 assay to identify and measure the total amount of huntingtin protein.
 - Next, mutant HTT protein is immunodepleted using polyQ Ab magnetic beads to reduce the total amount of mutant HTT protein in the sample.
 - Finally, the remaining wtHTT protein is quantified using the 2B7-D7F7 assay.
- Rigorous analytical validation of our wtHTT method resulted in precise, specific, and reproducible results across a range of wtHTT concentrations in a certified CSF reference material.
- Our method performs reliably when applied to CSF from patients with Huntington's disease.

Application of this method during clinical studies can help to evaluate the importance of preserving wtHTT protein in the context of HTT-reducing therapies. This method is a key component of the SELECT-HD clinical trial (NCT03933248).

Introduction

- A 2016 Huntington's disease expert panel (1) identified that the loss of the wtHTT protein leads to the expression of mutant HTT protein and causes HD in asymptomatic individuals.
- The wtHTT protein confers an neuroprotective (2,3,4) effect that protects against the neurodegeneration of HD.
- Since wtHTT exerts its neuroprotective effect in a dose-dependent manner, it is critical to maintain wtHTT protein levels in the brain (5,6).
- Accurate quantification of wtHTT protein can facilitate proof-of-concept studies and evaluation of neuroprotective allele-selective therapies.
- Our novel method for quantifying wtHTT protein in CSF involves three steps (Figure 1) and is applicable to certified CSF (7) and clinical samples (Table 1, 2).

Figure 1. Strategy for quantifying wtHTT protein in CSF

Assessment of wild-type protein in CSF
Depletion of mutant HTT key to ability to measure wild-type HTT protein

Step 1:

- Quantify tHTT (improved 2B7-D7F7 assay)
- If tHTT ≥ 20fM, go to step 2

Step 2:

- Immunodeplete mHTT with polyQ Ab magnetic beads

Step 3:

- Re-test sample with tHTT assay
- Remaining HTT is quantified as a surrogate of wtHTT protein

Legend: Wild-type HTT (blue square), Mutant HTT (red circle), MW1 (polyQ) Ab magnetic beads (yellow star)

Figure 2. Free MW1 Ab concentration of 1,000 or higher minimizes interference with the 2B7-D7F7 total tHTT assay.

Figure 3. Immunodepletion with MW1 has an acceptable level of specificity when total HTT levels are > 20 fM.

Figure 4. wtHTT quantification method performed reliably within the intended control of use.

Portrait photo: wtHTT, mutant HTT protein; wtHTT, wild-type HTT; polyQ, poly glutamine; CSF, CSF.



Paloma Giangrande, PhD

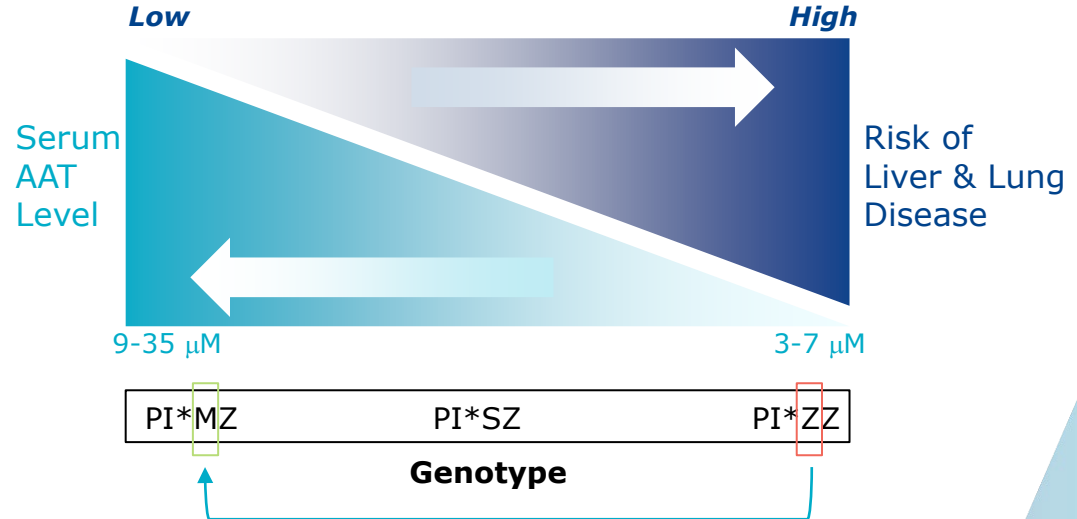
Vice President, Platform
Discovery Sciences, Biology

ADAR editing approach to correct AATD with GalNAc-AIMers addresses lung and liver pathologies

Alpha-1 antitrypsin deficiency (AATD)

- AAT protein produced by hepatocytes
- Point mutation in *SERPINA1* gene (PiZ allele) leads to misfolding and aggregation of mutant Z-AAT protein in hepatocytes in form of globules
- Pathology:
 - Liver damage, fibrosis, cirrhosis, and hepatocellular carcinoma
 - Lung emphysema and bronchiectasis

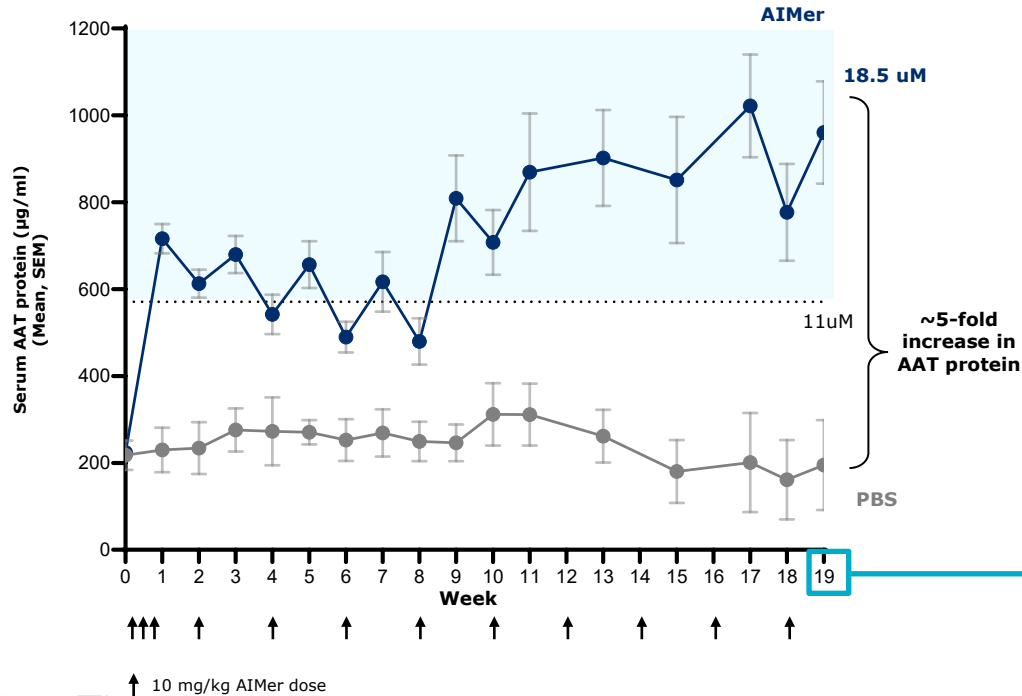
Inverse relationship between circulating AAT levels and disease risk



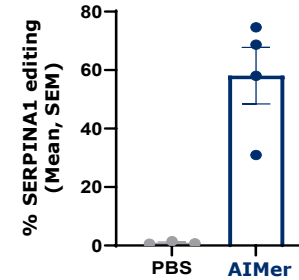
Therapeutic goal: ~50% RNA editing with AIMers to reach MZ phenotype with low risk of lung / liver disease

Preclinical AIMer treatment results in circulating AAT protein levels well above anticipated therapeutic threshold

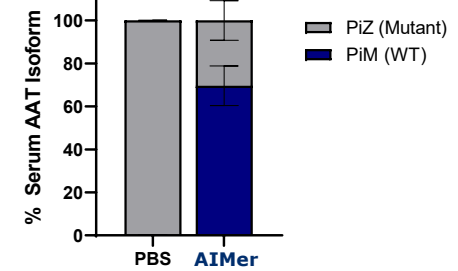
GalNAc-AIMer treatment bi-weekly results in serum AAT protein levels >11 μ M at week 19 in transgenic mice



~60% RNA editing with GalNAc-AIMer treatment

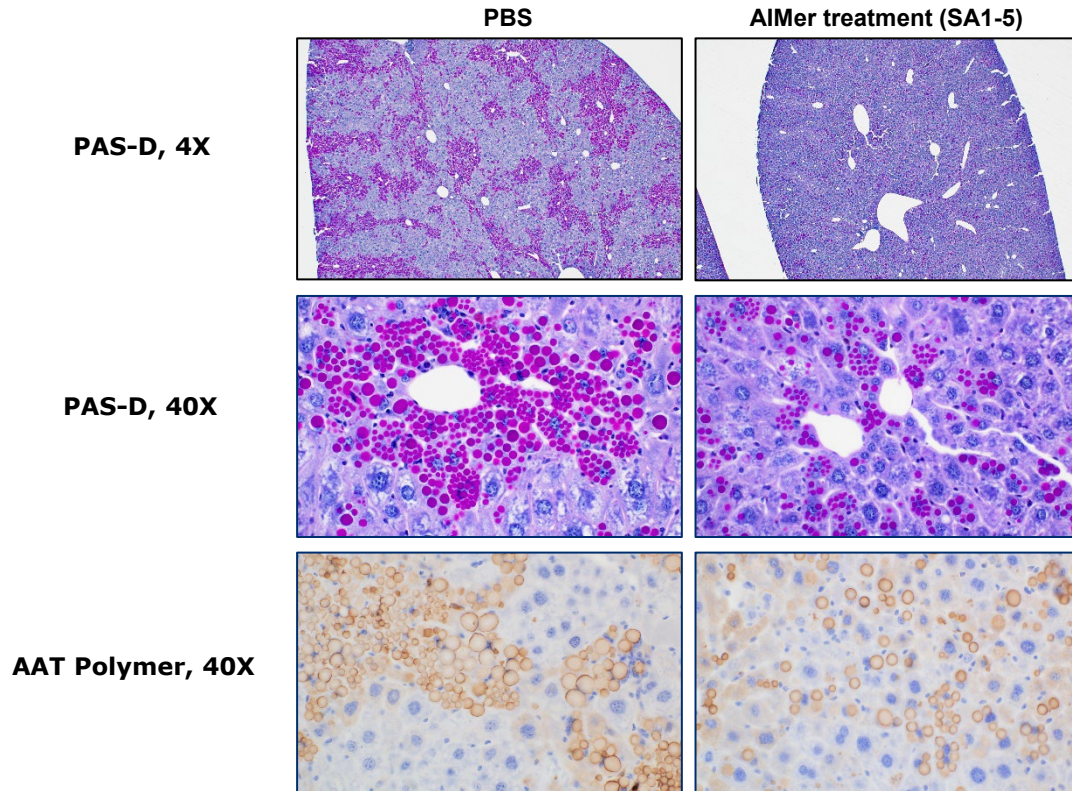


~70% wild-type, M-AAT protein in serum with GalNAc-AIMer treatment

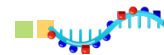


Histological analysis indicates reduction of liver aggregates at 19 weeks with AIMer treatment

Preliminary histological analysis of transgenic mouse liver aggregates (Week 19)



GalNAc-AIMers are uniquely suited to address the key treatment goals for AATD



- ✓ Recruit endogenous ADAR enzyme to edit SERPINA1 Z mRNA with high specificity
- ✓ Restore circulating, functional M-AAT protein above expected therapeutic threshold (11 μ M)
- ✓ Reduce Z-AAT protein aggregation in liver

	AIMers	RNAi	AAT augmentation therapy
Restore circulating functional wild-type AAT	✓		✓
Reduce Z-AAT protein aggregation in liver	✓	✓	
Retain M-AAT physiological regulation	✓		

Expect to select an AATD AIMer development candidate and initiate IND-enabling toxicology studies in 3Q 2022



Kyle Moran
Chief Financial Officer

Fourth quarter 2021 financial results

	Three Months Ended December 31, 2021	Three Months Ended December 31, 2020
<i>Figures are in thousands, except per share amounts</i>		
Revenue	\$1,765	\$9,439
Operating Expenses:		
Research and Development	25,761	30,033
General and Administrative	12,114	9,719
Total Operating Expenses	37,875	39,752
Net Loss from Operations	(36,110)	(30,313)
Total Other Income, Net	1,121	683
Income Tax Benefit, net	204	841
Net Loss	(\$34,785)	(\$28,789)
Net Loss per Share	(\$0.61)	(\$0.59)

As of December 31, 2021

Shares Outstanding: 59.8 million

Cash Balance: \$150.6 million



Paul Bolno, MD, MBA
President and CEO

Data generated in 2022 expected to inform future opportunities and unlock value

WVE-004 C9orf72 ALS & FTD	<ul style="list-style-type: none">Clinical data to enable decision making in 2022	Silencing	CNS <i>(Intrathecal)</i>	
WVE-003 HD SNP3	<ul style="list-style-type: none">Clinical data to enable decision making in 2022		Splicing	Muscle <i>(IV)</i>
WVE-N531 DMD Exon 53	<ul style="list-style-type: none">Clinical data to enable decision making in 2022		ADAR editing	Targeted delivery Liver <i>(Subcutaneous)</i>
AIMer AATD SERPINA1	<ul style="list-style-type: none">Select an AATD AIMer development candidate and initiate IND-enabling toxicology studies in 3Q 2022			

Success with any current program validates platform and unlocks modalities and tissues

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Q&A



Realizing a brighter future for people affected by genetic diseases

For more information:

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