

Synthesis and application of stereopure guanidine-containing backbone to multiple oligonucleotide modalities in preclinical studies

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PRISM platform enables rational drug design



PRISM backbone modifications



Advances in stereopure monomer synthesis & manufacturing



Efficient scalable process for synthesis







DEA wash

Rp



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- Std CNF amidite for PO
- Chiral amidite for both PS and PN
- High diastereoselectivity High coupling efficiency

Standard C&D process

- On column DEA wash **
- Ammonium hydroxide *

Scalable process

- 96-well plate to GMP * manufacturing
- Comparable yield to std oligo *

Characterization of stereopure dimers



Potency is enhanced with addition of PN modifications across modalities







Left: Experiment was performed in iPSC-derived neurons *in vitro*; target mRNA levels were monitored using qPCR against a control gene (HPRT1) using a linear model equivalent of the $\Delta\Delta$ Ct method; Middle: DMD patient-derived myoblasts treated with PS/PO or PS/PO/PN stereopure oligonucleotide under free-uptake conditions. Exon-skipping efficiency evaluated by qPCR. Right: Data from independent experiments

WVE-004 treatment resulted in durable reduction of poly-GP biomarker in mouse spinal cord & cortex^{1,2}



*: $p \le 0.05$, **: $p \le 0.01$, ***: $p \le 0.001$. DPR, dipeptide repeat protein. ¹Liu et al., 2022 *Mol Ther Nucleic Acids*. ²Liu et al. Poster i182 presented at ENCALS Meeting 2022. Poly-GP assay: **Wilson et al., 2022** *J Neurol Neurosurg Psychiatry*. Silencing

PN modification improves silencing for GalNAcsiRNAs by increasing Ago2 loading in mice



Proprietary human HSD17B13 transgenic mouse model dosed as indicated on day 1 ($n \ge 3$) (Left) Percentage HSD17B13 mRNA expression normalized to mouse Hprt. Stats: Linear Mixed Effects Model with Random subject effects *** p<0.001, **** p<0.0001; (Right) Ago2 loading. Stats: ** p<0.01, 2-way ANOVA

Silencing

PN modification increases exon skipping, restores dystrophin expression and prolongs survival in dKO mice



Increased dystrophin protein expression in dKO mice







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WVE-006 directs RNA base editing to increase serum AAT protein above potential therapeutic threshold in mice



LIFE SCIENCES WVE-006: GalNAc-conjugated AIMer; NSG-PiZ mice dosed with WVE-006 (n=5). (Left) SERPINA1 editing quantified from 13-week liver biopsies by Sanger sequencing; (Right) Total serum AAT protein quantified by ELISA

Impact of PN chemistry highlighted in three highimpact publications this year



• Thanks to all colleagues and contributors from Wave Life Sciences

