Stoke Therapeutics

September 2019



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Stoke is pioneering a new way to treat the underlying causes of severe genetic diseases by precisely upregulating protein expression

We Believe Stoke's Approach is Highly Differentiated and Positioned for Success

Novel technology

Antisense oligonucleotides (ASOs) target pre-mRNA splicing to <u>restore target protein</u> to near normal levels Diseasemodifying therapies

ASOs are designed to <u>address the underlying</u> <u>cause</u> of severe genetic diseases, including autosomal dominant haploinsufficiencies



Efficient

syndrome leverages <u>validated</u> <u>ASO chemistry</u>, a <u>well-defined</u> <u>patient population</u>, and <u>learnings from recently</u> <u>approved drugs</u>

Phase 1/2 trial expected to begin in 1H 2020; preliminary <u>efficacy data</u>, expected in 2021 Potential broad applications

Emerging pipeline spans severe genetic diseases of the <u>CNS, eye, liver and</u> <u>kidney</u>

Plan to <u>nominate second</u> <u>candidate</u> to treat an additional genetic disease for preclinical development by 1H 2020



- Rapidly advance our lead program, STK-001, to clinical proof-of-concept, approval and commercialization
- **Prioritize genetic epilepsies** for near-term development efforts
- Expand our pipeline into other disease areas to fully exploit the potential of our proprietary platform
- Continue to strengthen and expand our IP portfolio
- Maintain broad commercial rights to our product candidates
- **Opportunistically evaluate potential collaboration arrangements** with a pharmaceutical or biotechnology company



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Huw Nash, Ph.D. – Chief Operating Officer and Chief Business Officer	AILERON	neogenesis	
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Autosomal Recessive	Autosomal Dominant Gain-of-function / Dominant Negative	Autosomal Dominant Haploinsufficiency
 Disease examples: Phenylketonuria Lysosomal storage disorders Beta-thalassemia Cystic fibrosis 	 Disease examples: Huntington's disease Parkinson's disease Spinocerebellar ataxia Autosomal dominant hypocalcemia 	 Disease examples: Dravet syndrome Optic atrophy Polycystic kidney disease Tuberous sclerosis
Gene therapy • Small molecules Gene editing Modified mRNA Protein-based drugs	 Current / emerging approaches: Gene therapy Oligonucleotides Gene editing Small molecules Protein-based drugs 	Emerging approach: • Stoke's TANGO technology STERAPEUTICS

Existing precision medicine platforms are poorly suited to address haploinsufficiency diseases. Consequently, there has been little focus on drug development for these diseases despite a significant unmet medical need

Source: globalgenes.org; Mathura et al., Science, 2019; Lykken et al., Journal of Neurodevelopmental Disorders, 2018; Company websites

We Believe Stoke's TANGO Technology Offers Key Advantages Based on Preclinical Studies



Ability to address underlying genetic cause of disease



Applicability to most loss-offunction mutations



Utility across small and large gene targets

 No observed unwanted offtarget effects



Ability to control dose level and duration



Utility across a wide array of diseases and tissue types



Simple and scalable manufacturing

TANGO exploits unique, patented mechanisms for antisense-mediated modulation of splicing to precisely upregulate protein expression, thereby addressing the underlying genetic cause of the disease rather than merely alleviating the symptoms of the disease

ASOs upregulate expression of the wild-type allele, meaning the TANGO mechanism does not rely on targeting a specific mutation

ASOs upregulate protein expression regardless of gene size and are not constrained to smaller gene targets

TANGO-mediated upregulation of protein expression only occurs where the gene is being naturally transcribed, limiting the likelihood of expression in non-native tissues

ASOs provide the ability for dose titration, thereby allowing for dose-dependent and reversible control of level and duration of protein expression. The ability to titrate dosage will enable us to deliver the right dose, at the right location, for each indication

ASO delivery to the CNS, eye, kidney and liver is well-established, enabling Stoke to address a broad range of genetic diseases. FDA-approved ASO (SPINRAZA) demonstrates ASO delivery to the CNS, and there are other ASOs in clinical development

ASOs are synthesized by highly scalable, solid-phase chemical synthesis and leverage a well-established, global manufacturing base

Source: Stoke data based on preclinical studies to date. Our product candidate has not been approved by the FDA.

Stoke's TANGO Technology Targets Retained Introns to Upregulate Protein Expression

TANGO mechanism for increasing protein synthesis: Retained Intron



Retained introns are found in ~60% of gene transcripts and are part of the wild-type sequence of the gene. Non-productive mRNA remains in the nucleus and is not translated into protein



Source: Stoke data based on preclinical studies to date

Stoke's TANGO Technology Targets Retained Introns to Upregulate Protein Expression

TANGO mechanism for increasing protein synthesis: Retained Intron



Stoke's ASOs bind to the pre-mRNA and redirect the splicing machinery to remove the retained intron. This splice-switching decreases non-productive mRNA and increase productive mRNA, which is translated into increased protein expression from the wild-type allele



Stoke's TANGO Technology Targets NMD Exons to Upregulate Protein Expression

TANGO mechanism for increasing protein synthesis: Nonsense mediated decay (NMD) exon



NMD exons are found in over 25% of gene transcripts and are part of the wild-type sequence of the gene. Non-productive mRNA is degraded in the cytoplasm by NMD and is not translated into protein

Note: NMD denotes nonsense-mediated mRNA decay Source: Stoke data based on preclinical studies to date



Stoke's TANGO Technology Targets NMD Exons to Upregulate Protein Expression

TANGO mechanism for increasing protein synthesis: NMD exon



Stoke's ASOs bind to the pre-mRNA and redirect the splicing machinery to prevent inclusion of the NMD exon. This splice-switching decreases non-productive mRNA and increase productive mRNA, which is translated into increased protein expression from the wild-type allele



Target identification process



- Next generation RNA sequencing yields proprietary database of approximately 85,000 non-productive events in the human transcriptome
- Cross-referencing with genetic disease databases identifies approximately 2,900 monogenic diseases amenable to TANGO
- Approach is highly predictive and enables rapid and systematic identification of clinically relevant targets



Source: Stoke data

- Patents for TANGO technology exclusively licensed from University of Southampton and Cold Spring Harbor Laboratory
- Multi-national allowed and pending claims for the TANGO mechanisms
- Multi-national pending claims for more than 140 genetic diseases amenable to TANGO
- Leveraging previously-validated ASO chemistries



50 million people globally affected by epilepsy

>30%

of patients are refractory to medical treatment, especially those with a genetic epilepsy

of patients with epilepsy have significant cognitive problems



of epilepsies have an identified genetic cause and many of these are haploinsufficiencies

Diagnostic work-up of epilepsy routinely includes genetic testing for more than

80 disease associated genes

While genetic mechanisms are often well understood ...

genetically-targeted therapies for epilepsies are available

Source: WHO 2018 fact sheet; Sirven, Cold Spring Harbor Perspectives in Medicine 2015; Pal et al., *Nature Reviews Neurology* 2010; Chen et al., *JAMA Neurology* 2018; Lagae et al., *Developmental Medicine & Child Neurology* 2017; Vlaskamp et al., *Neurology* 2019; Reddy SD et al., J Pharmacol Exp Ther 2018; NIH Genetics Home Reference; Company websites



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Up to

We Believe Many Genetic Epilepsies are Amenable to Stoke's TANGO Technology





Lead Program

Source: Modified from McTague et al., *Lancet* 2016; Stoke data



RARE & CATASTROPHICEPILEPSY IN 16,000 REGRESSION L OPMENT DEL AYS SPEECH ISSUES SWALLOWING AUTISIM UCED LIFE EXPECTANCY ATAXIA BALANCE ISSUES CULTIES WILL NOT OUTGROW

- Autosomal dominant condition caused by more than 1,250 *de novo* mutations in *SCN1A*, resulting in 50% Na_v1.1 protein expression
- Caused by pathogenic mutation or deletion of the SCN1A gene in ~85% of patients
- Existing antiepileptic drugs only address the occurrence of seizures, and more than 90% of Dravet syndrome patients still report suffering from incomplete seizure control
- No disease-modifying therapies in clinical development
- ~35,000 patients across U.S., Canada, Japan, Germany, France and the UK



Source: 2018 Health Advances Report; Djémié et al., Molecular Genetics & Genomic Medicine 2016; Lagae et al., Developmental Medicine & Child Neurology 2017; Nabbout et al., Orphanet Journal of Rare Diseases 2013

Transformative Potential of TANGO Technology in Dravet Syndrome



TANGO mechanism for increasing protein synthesis in a prospective patient with Dravet syndrome

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Source: Escayg and Goldin, Epilepsia 2010; Stoke data based on preclinical studies to date

Transformative Potential of TANGO Technology in Dravet Syndrome



TANGO mechanism for increasing protein synthesis in a prospective patient with Dravet syndrome

STOKE THERAPEUTICS

Source: Escayg and Goldin, Epilepsia 2010; Stoke data based on preclinical studies to date

STK-001 Increases *Scn1a* mRNA and Na_v1.1 Protein in Wild-type Mice





Source: Stoke data

STK-001 Selectively Upregulates *Scn1a* Gene in Wild-type Mice



Scn gene family members

STK-001 is very specific for *Scn1a* among the highly homologous family of sodium channel genes, limiting the likelihood of off-target activities



STK-001 Restores Na_v1.1 to Near Normal Levels for >3 Months in Dravet Syndrome Mice



In preclinical studies, STK-001 exhibits long-lasting exposure, suggesting the potential for a favorable dosing regimen of as few as two to three administrations per year in humans

Note: Na_v1.1 protein quantification based on standard curve obtained from untreated wild-type mouse brain as a reference control Source: Stoke data; University of Michigan (in-life study)



STK-001 Significantly Reduces Premature Mortality in Dravet Syndrome Mice



	Placebo wild-type	Placebo Dravet syndrome	STK-001 wild-type	STK-001 Dravet syndrome
Total n through 90 days	49	62	27	34
Number of deaths	0	48	0	1

Note: Neonate Dravet syndrome and wild-type mice were administered a single injection dose of either placebo (consisting of a phosphate-buffered solution) or 20 ug of STK-001 by intracerebroventricular injection (placebo: n=49 wild-type mice, n=62 Dravet syndrome mice; STK-001: n=27 wild-type mice, n=34 Dravet syndrome mice) Source: Stoke data, University of Michigan



STK-001 Therapeutic Dosing also Significantly Reduces Premature Mortality



	Placebo wild-type	Placebo Dravet syndrome	STK-001 wild-type	STK-001 Dravet syndrome
Total n through 35 days	68	45	41	46
Number of deaths	0	13	0	1

Note: Neonate Dravet syndrome and wild-type mice were administered a single injection dose of either placebo (consisting of a phosphate-buffered solution) or 60 ug of STK-001 by intracerebroventricular injection (placebo: n=68 wild-type mice, n= 45 Dravet syndrome mice; STK-001: n= 41 wild-type mice, n=46 Dravet syndrome mice). Data is preliminary Source: Stoke data, University of Michigan

Key safety measures

No complement activation	\checkmark
No decrease in platelet counts	\checkmark
No change in hepatic function	\checkmark
No clinical signs or symptoms over 28 day period after administration	\checkmark
Normal histopathology in key organs	\checkmark





- Increase in *Scn1a* mRNA expression and Na_v1.1. protein levels in wild-type mice
- Selective upregulation of *Scn1a*, and not closely related ion channels in wild-type mice
- Restoration of Na_v1.1 protein to near normal levels in Dravet syndrome mice
- Effects persisting for at least 14 weeks in Dravet syndrome mice
- Dramatic reduction in mortality in Dravet syndrome mice
- Well-tolerated at a pharmacologically-active dose level in non-human primates



Efficient Clinical Program for STK-001 in Dravet Syndrome



Phase 1/2 trial

- Analogous trial design and endpoints to recently approved antiepileptic drugs for Dravet syndrome
- Change in seizure frequency over 12-week treatment period, cognitive function, and quality of life will be included as secondary endpoints
- Preliminary clinical data for primary and secondary endpoints of Phase 1/2 SAD study expected in 2021



Note: Timelines are estimates and subject to change; ¹ 'SAD' denotes Single Ascending Dose; 'MAD' denotes Multiple Ascending Dose

Strong Momentum for the Year Ahead

- Nominated lead target
- Demonstrated *in vivo* proof of concept in mouse model for Dravet syndrome
- Completed pre-IND meeting for Dravet syndrome
- Completed Series B financing

2018



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2019 – 2020

- Completed Initial Public Offering
- Received FDA orphan drug designation for STK-001
- Enrolled first patient in BUTTERFLY, observational study for Dravet syndrome
- Complete GLP toxicology studies to support STK-001 IND application for Dravet syndrome in early 2020
- □ Initiate Phase 1/2 clinical trial for Dravet syndrome in 1H 2020
- Nominate second candidate to treat an additional genetic disease for preclinical development by 1H 2020
- Opportunistically secure first pharma partnership as early as 2019

Stoke is Building a Pipeline of Precision Medicines





Cash Expected to be Sufficient to Fund Operations into 2023

Cash, Cash Equivalents and Restricted Cash as of 06/30/2019	\$242.9 million
Shares Outstanding	32,718,134
as of 08/14/2019	(basic shares)

Raised \$151.9 million in net proceeds in June 2019 initial public offering





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