F→KF THERAPEUTICS

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BACKGROUND

- DS is a severe and progressive genetic epilepsy characterized by frequent, prolonged, and refractory seizures, typically beginning within the first year (y) of life
- Available therapies do not adequately control seizures in 90% of patients with DS, and they do not address other comorbidities, including intellectual disability, ataxia/motor abnormalities, behavioral problems, speech impairment, sleep disturbances, and a high risk for sudden unexpected death
- Disease complications often contribute to a poor quality of life for patients and their caregivers
- In approximately 85% of cases, DS is caused by spontaneous, heterozygous loss of function mutations in the SCN1A gene, which encodes the voltage-gated sodium channel type 1 α subunit $(Na_v 1.1)$ protein
- Upregulating Na_v1.1 protein may restore functioning neurons and prevent seizures and reduce nonseizure related comorbidities in DS

MECHANISM OF ACTION

- The proprietary TANGO (Targeted Augmentation of Nuclear Gene Output) platform aims to increase protein production from the healthy gene
- In DS, patients have one functional gene (wild type) copy and one mutated copy, resulting in half as much protein as needed to maintain health
- STK-001 is an investigational proprietary ASO designed to upregulate Na_V1.1 protein expression by leveraging the non-mutant (wild type) copy of SCN1A to restore physiological Na_v1.1 protein levels
- SCN1A is transcribed into pre-messenger RNA (premRNA) that is spliced to generate productive mRNA (which is translated into Na_V1.1 protein) and nonproductive mRNA due to the inclusion of an exon that leads to nonsense-mediated mRNA decay (NMD). TANGO ASOs bind to specific stretches of SCN1A pre-mRNA to prevent the inclusion of the non-productive exon thereby increasing productive mRNA level
- Increased level of productive mRNA from the functional gene copy increases Na_v1.1 protein production restoring it to near normal levels
- STK-001 may be the first disease-modifying therapy to address the genetic cause of DS by upregulating $Na_{v}1.1$ protein levels

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SWALLOWTAIL: An Open-Label Extension (OLE) Study for Children and Adolescents with Dravet Syndrome (DS) who Previously Participated in a Study of Antisense Oligonucleotide (ASO) STK-001

STUDY DESIGN

OLE study conducted at 18 sites in the US (NCT04740476) following participation in Single (SAD) or Multiple Ascending Dose (MAD) study, MONARCH (NCT04442295) Patients receive intrathecal (IT) doses of STK-001, once every 4 months and patients currently receive

30 mg per dose

PATIENT DEMOGRAPHICS

Demographics (N=44)									
Age at Screening, y									
Mean (SD)	11.2 (5.03)								
ian (min, max)	13.0 (2, 19)								
Age Group, N (%)									
2–12y	21 (47.7)								
3y and older	23 (52.3)								
Sex									
emale, n (%)	23 (52.3)								
Male, n (%)	21 (47.7)								
Race, N (%)*									
Asian	4 (9.09)								
ick or African American	4 (9.09)								
White	37 (84.1)								
refer not to answer	3 (6.82)								
Ethnicity, N (%)									
Not spanic/Latino	36 (81.8)								

Able to choose >1 option

PATIENT EXPOSURE

tal STK-001 doses administered (N=44)							
10 mg	10						
20 mg	14						
30 mg	82						
45 mg	36						



Analysis of VABS-III data also indicated an improvement in Gross Motor function with mean change in GSV of +12.2 [3.09, 21.4] at Month 12 (not shown) Both Clinicians and Caregivers rated patient's overall condition improved on the Global Impression of Change (CGIC and CaGIC) scales with mean scores of 2.5 [1.92, 3.08] and 1.53 [0.893, 2.17] (not shown)

*Clinical assessment analyses included patients who completed treatment in MONARCH and received only doses of 30 mg or 45 mg in SWALLOWTAIL. Vineland Adaptive Behavior Scales-III (VABS-III) and Behavior Rating Inventory of Executive Function Preschool (BRIEF-P) data analyzed in mixed model repeated measures with AR(1) covariance structure. Baseline covariates in BUTTERFLY matched to SWALLOWTAIL. BUTTERFLY sample size: n=36 at screening and n=27 for VABS-III and n=30 for BRIEF-P at Month 12; SWALLOWTAIL sample size: n=24 for VABS-III and n=25 for BRIEF-P at screening; n=9 at Week 48 and n=5 at Week 64. GSV=Growth Scale Value; Brackets indicate 95% CI; Rating on CGIC and CaGIC ranges from 1=very much improved to 7=very much worse. A rating of 4=No change

PATIENT DISPOSITION		SAFETY							
Assigned Cohort (MONARCH)Enrolled (% total patients whoAssigned Onge	Enrolled (%		Number (N (%)) of Patients with	2-12y	13-18y	Total	TEAEs Reported in >10% Patients		
	Ongoing		N=21	N=23	N=44	Preferred Term	N (%)		
	ongoing	TEAEs	16 (76.2)	22 (95.7)	38 (86.4)	CSF protein increased	14 (31.8)		
	MONARCH)		TEAEs related to study drug (all	6 (28.6)	11 (47.8)	17 (38.6)	COVID-19	13 (29.5)	
10 mg SAD*	5 (100)	3	mild or moderate) [*]	- ()			Post lumbar puncture		
20 mg SAD	4 (100)	3	drug administration	6 (28.6)	13 (56.5)	19 (43.2)	syndrome	7 (15.9)	
30 mg SAD	7 (100)	6	≥Grade 3 TEAEs	1 (4 76)	0	1 (2 27)	Pyrexia	7 (15.9)	
45 mg SAD	5 (100)	5	Scrade 3 TEAEs related to study	. (. ()	Seizure	7 (15.9)	
20 mg MAD	4 (80.0)	3	drug	0	0	0	Upper respiratory tract		
20 mg MAD	14 (100)	13	Serious TEAEs	3 (14.3)	2 (8.70)	5 (11.4)	infection	6 (13.6)	
	T4 (100)	-	Serious TEAE related to study	0		0	Headache	5 (11.4)	
45 mg MAD	5 (83.3)	5	drug	0	0	0	Nasal congestion	5 (11.4)	
*1 patient in 10 mg SAD cohort received an incorrect dose in MONARCH but received 3 correct doses in SWALLOWTAIL *Only Treatment-Related TEAE in >1 patient was CSF protein increased: 14/44 (31.8%) patients					Puncture site pain	5 (11.4)			

CLINICAL ASSESSMENTS*



OVERALL SUMMARY

44 of 46 (95.7%) patients who completed MONARCH enrolled in SWALLOWTAIL, and 38 of 44 (86.4%) were ongoing in the study Patients received up to 7 doses of STK-001 ranging from 10 mg-45 mg/dose in SWALLOWTAIL • Multiple doses of STK-001 up to 45 mg/dose given every 4 months IT were generally well tolerated • Data indicated durable reductions in convulsive seizure frequency and substantial improvements in cognition and behavior over 12m







Improving