



Initial Safety, Tolerability, Pharmacokinetics, and Pharmacodynamics Data from an Ongoing Open-Label Study Investigating FTX-6058 in Adults Living with Sickle Cell Disease

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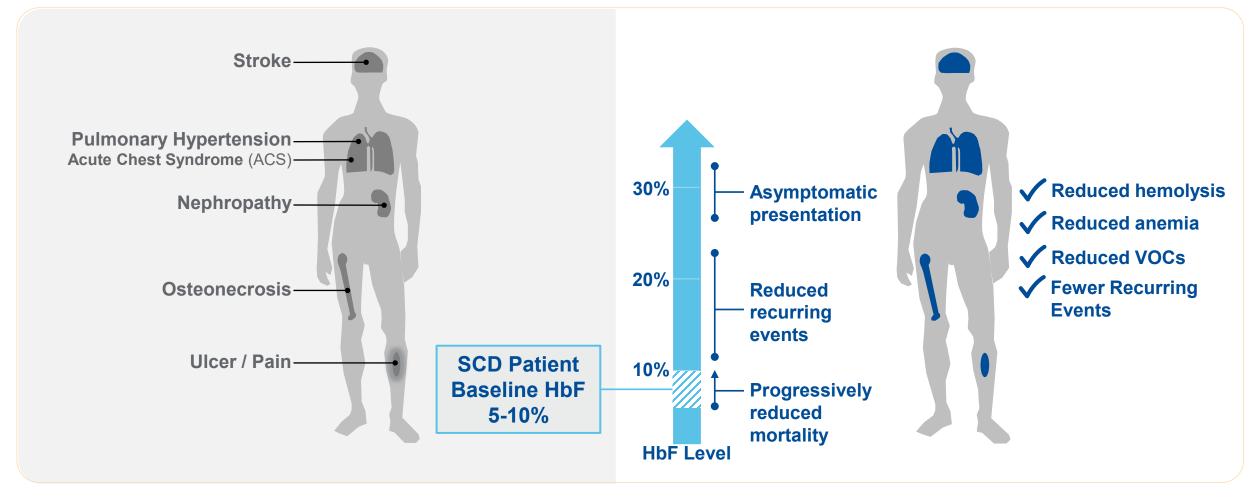
Background

FTX-6058 is a potent and selective oral small-molecule inhibitor of Embryonic Ectoderm Development (EED) that has demonstrated robust HbF (fetal hemoglobin) protein induction in human primary cell and murine models of sickle cell disease (SCD). SCD is a genetic disorder of the red blood cells caused by a mutation in the hemoglobin beta (HBB) gene, which results in red blood cell sickling, hemolysis, vaso-occlusive crises (VOCs), and other complications. HbF ($\alpha_2 \gamma_2$) prevents the pathologic polymerization of sickle hemoglobin (HbS) in deoxygenated environments. Individuals with SCD who also have hereditary persistence of HbF (HPFH) have attenuated pathology. HPFH provides genetic and clinical evidence that increasing HbF has the potential to prevent or reduce disease-related pathophysiology, including hemolysis, pain, and VOCs.

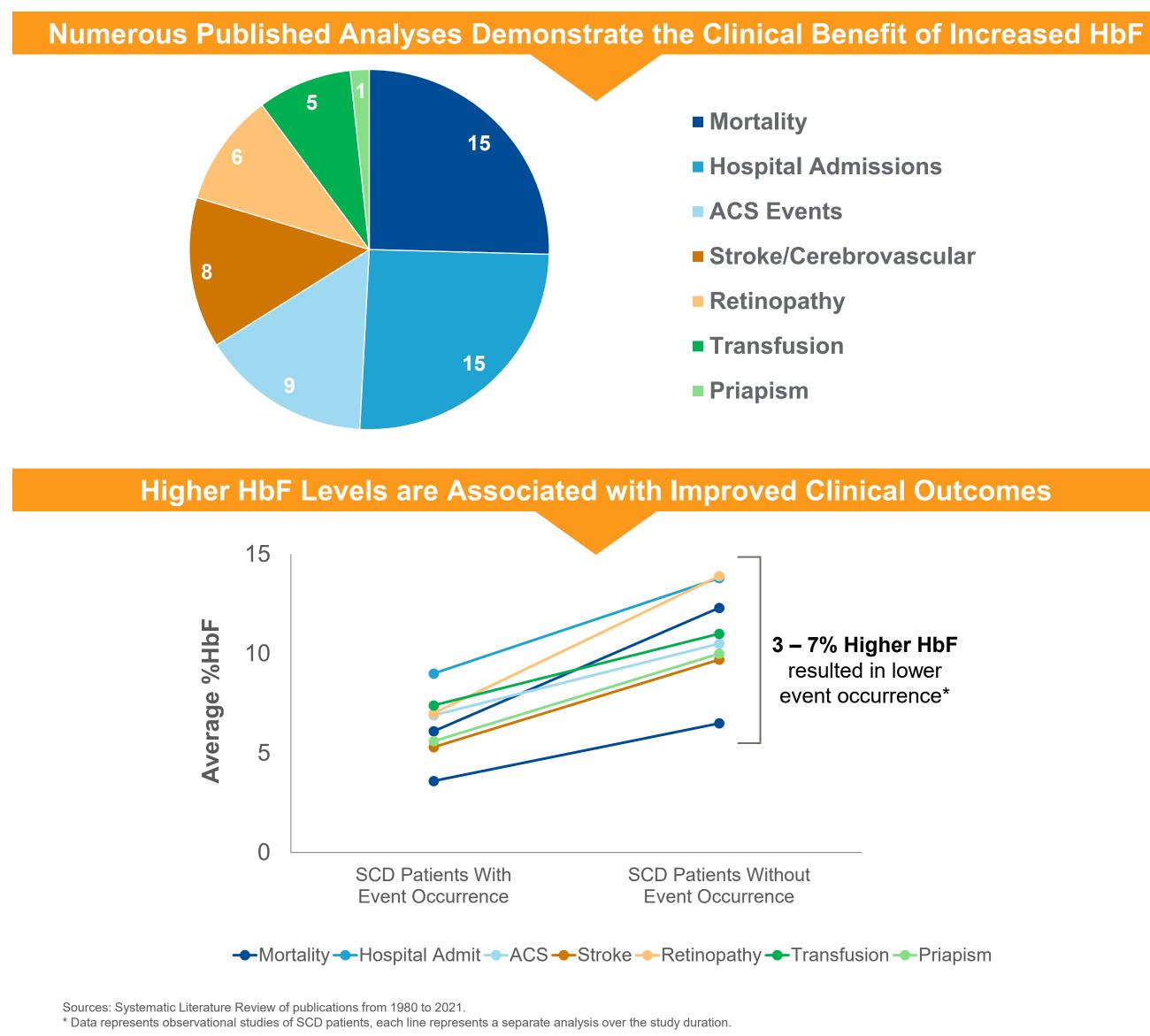
FTX-6058 has demonstrated consistent 2-3 fold HbF induction with strong correlation between HBG mRNA and HbF protein in pre-clinical studies, which has the potential to positively impact important clinical outcomes in SCD. Clinically, multiple ascending dose (MAD) data from a Phase 1, randomized, double-blind, placebo-controlled study of FTX-6058 in healthy volunteers demonstrated robust target engagement (H3K27me3 reduction) and potent induction of HBG (hemoglobin subunit γ) mRNA, translation of which is required for HbF expression.

HbF Increases 5-10% Above Baseline (10-20% Total HbF) Have **Been Shown to Broadly Improve Outcomes in SCD**

Typical SCD Patient



Increases in HbF Protein can Provide Broad Clinical Benefit



SCD Patient with HPFH

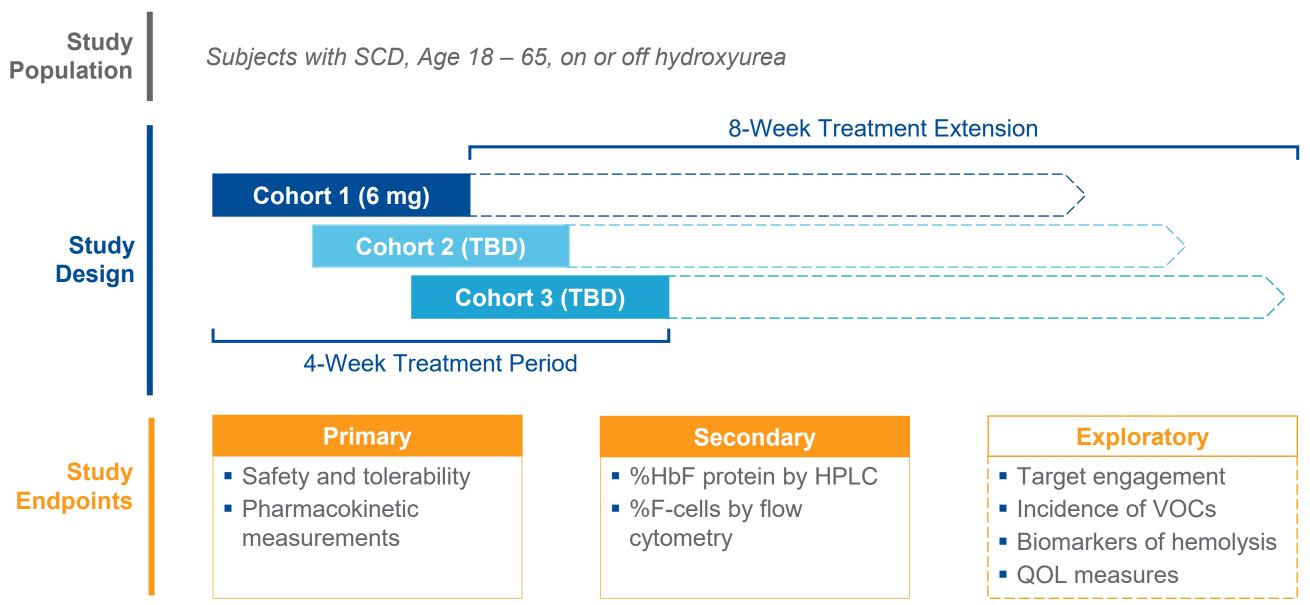
VOCs: Vaso-occlusive crisis; HPFH: Hereditary Persistence of Fetal Hemoglobin

3 – 7% Higher HbF resulted in lower event occurrence*

Methodology

The ongoing Phase 1b open-label dose ranging study investigating FTX-6058 is a multiple cohort trial that is being conducted in adults with SCD on and off hydroxyurea. In the first cohort, up to 10 subjects will receive FTX-6058 6 mg once daily orally for 4 weeks. Eligible subjects may choose to continue dosing for an additional 8 weeks at the same dose via an extension study. Up to two additional cohorts with doses selected between 2 – 20 mg inclusive may be added to the study. The primary endpoints of the study are the safety and tolerability of FTX-6058 as measured by the frequency of adverse events and the pharmacokinetic profile of FTX-6058 in subjects with SCD. Secondary and exploratory endpoints include the effect of FTX-6058 on HbF induction in peripheral blood and other relevant SCD biomarkers and clinical endpoints.

Ongoing Phase 1b Clinical Trial in SCD Subjects



Demographics

SCD Phase 1b Demographics

	Phase 1b Study (6 mg Cohort)
Number of subjects enrolled, n	6
Average age, years (range)	31 (25, 48)
Gender, Male (%)	1 (16%)
Mean baseline HbF (range)	5.7% (3.7, 9.2)
Genotype, n (%)	
HbSS	6 (100%)
HbSβ ⁰	0 (0%)
HbSβ+	0 (0%)
Hydroxyurea Utilization, n (%)	0 (0%)

Data analysis cutoff as of May 25th, 2022; One subject was omitted from PD analysis due to protocol deviation

- Mean baseline HbF of 5.7% is consistent with recent SCD clinical studies and published data
- No hydroxyurea utilization among enrolled subjects
- All subjects enrolled to date have the HbSS genotype

Results

Day 1 Pharmacokinetic Parameters for FTX-6058 6 mg Dose

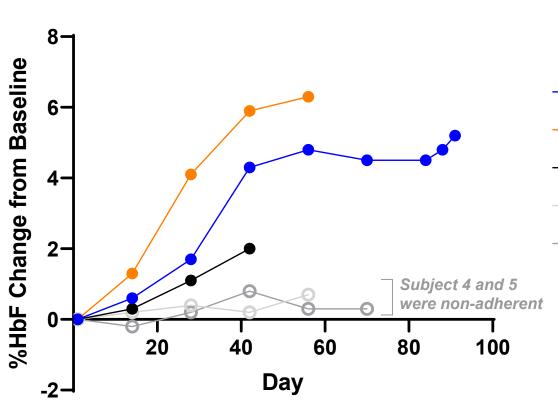
	Phase 1 HV Values (SD)	Phase 1b SCD Values (SD)				
Cmax (ng/mL)	24.9 (9.3)	16.9 (3.5)				
Tmax (hr)	4.08 (0.80)	2.50 (0.84)				
T1/2 (hr)	5.60 (0.36)	5.47 (0.72)				
AUC 0-last (hr*ng/mL)	192.5 (57.02)	134 (45.4)				
 Potentially lower exposure (25 – 30%) observed in SCD subjects versus healthy volunteers (HVs) Preliminary observations suggest differences in absorption (vs metabolism) 						

In SCD subjects, exposure (AUC 0-last) from 6 mg dose is similar to ~4 mg dose in HVs

ndary		Exploratory	
in by HPLC flow		 Target engagement Incidence of VOCs 	
		Biomarkers of hemolysisQOL measures	

Results (continued)

FTX-6058 Achieved Up To 6.3% Absolute Increase in HbF

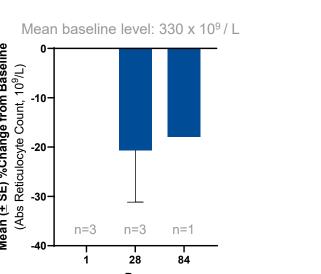


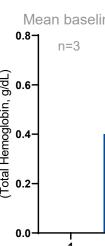
- benefit to people living with SCD
- Observed measurable increases in HbF protein as early as 14 days after treatment initiation
- Exposure appears to correlate with efficacy
- clinical data

Data analysis cutoff as of May 25th, 2022; Subject 1 was not dispensed drug after Day 56; Subject 3 withdrew from the study; *Per SAP, Subjects 4 and 5 were determined to be non-adherent based on vestigator and patient reporting, and/or pill count; Subject 6 was discontinued due to protocol deviation

FTX-6058 Decreases Hemolysis

Absolute Reticulocyte Count





Reductions in reticulocytes and increases in total hemoglobin indicate less anemia and hemolysis

FTX-6058 has been Generally Well Tolerated Summary of All Treatment Emergent Adverse Events (TEAEs)

TEAE	Severity
Swelling of Legs and Feet	Mild (Grade 1)
Light Headache	Mild (Grade 1)
Lower Back Pain	Mild (Grade 1)
Sore Throat	Mild (Grade 1)
Abdominal Pain	Moderate (Grade 2)
UTI	Moderate (Grade 2)
Tonsilitis	Moderate (Grade 2)
VOC (L Lower Leg Pain Crisis)	Moderate (Grade 2)
	Swelling of Legs and Feet Light Headache Lower Back Pain Sore Throat Abdominal Pain UTI Tonsilitis

- All TEAEs are non-serious, resolved, and were deemed to be unrelated to study drug
- No treatment emergent SAEs reported, and no discontinuation reported due to TEAEs
- Unlike hydroxyurea, no myelosuppression observed
- VOC observed in non-adherent subject (i.e., Subject 4)

Conclusions

- Initial FTX-6058 data in SCD subjects demonstrates compelling proof-of-concept • FTX-6058 rapidly and robustly induces HbF
- Initial subjects from first dose cohort achieved up to 6.3% HbF induction over baseline
- HbF levels increasing at last measured timepoint
- clinical data FTX-6058 improved biomarkers of hemolysis
- Data supports the potential for FTX-6058 to provide broad benefit to SCD patients

References

Adeodu, et al. Mediterr J Hematol Infect Dis. 2017 Akinsheve, I. Blood 2011 Curtis, et al. PLoS One 2016 Duan, et al. Am J Ophthalmol 2019 Kato, G.J., et al., Sickle cell disease. Nat Rev Dis Primers, 2018. 4: p. 18010, Reference 4 Piel, F.B., M.H. Steinberg, and D.C. Rees, Sickle cell disease. N Engl J Med, 2017. 376(16): p. 1561-1573 Platt, OS. NEJM 1994 Powars, DR, Blood 1984 Odenheimer, et al. Am J Hum Genet. 1987 Shurafa, et al. Am J Hematol. 1982



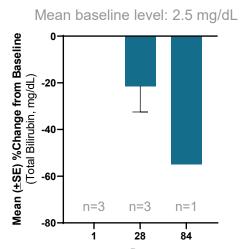
 Subject 1 Subject 2 	Subject	Baseline %HbF	%HbF at Data Cutoff	∆%HbF
 Subject 3 	Subject 1	9.2	14.4	5.2
- Subject 4*	Subject 2	3.7	10	6.3
-O- Subject 5*	Subject 3	6.2	8.3	2.1
	Subject 4*	4.6	5.3	0.7
	Subject 5*	7.0	7.3	0.3

• Achieved absolute HbF increases within the 5 – 10% range which has been shown to provide transformational

• HBG mRNA and HbF protein fold induction correlation (approximately 1:1) is consistent across preclinical and

Total Hemoglobin

Mean baseline level: 7.5 g/dl



Total Bilirubin

Bilirubin decreases indicate less hemolysis

Data analysis cutoff as of May 25th, 2022; Data presented are from Subjects 1 - 3

• HBG mRNA and HbF protein fold induction correlation (approximately 1:1) is consistent across preclinical and

• FTX-6058 was generally well-tolerated in SCD subjects with up to 3 months exposure