

FTX-6058, A Novel HbF Inducer: Phase 1 Healthy Volunteer Trial Update

December 6th, 2021



Fulcrum
Therapeutics



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Agenda

- **Introduction**

Bryan Stuart

- **Overview of Fetal Hemoglobin Repression**

Gerd Blobel, MD, PhD

- **FTX-6058 Preclinical Pharmacology and Mechanistic Insights**

Judith Dunn, PhD

- **FTX-6058 Phase 1 Healthy Volunteer Update**

Christopher Morabito, MD

- **Q&A Session**

Introduction

Bryan Stuart

President and Chief Executive Officer



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Overview of Fetal Hemoglobin Repression

Gerd Blobel, MD, PhD

Frank E. Weise III Endowed Chair in Pediatric Hematology at Children's Hospital of Philadelphia



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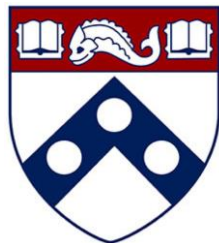


Pursuing HbF Elevation as a Therapeutic Strategy

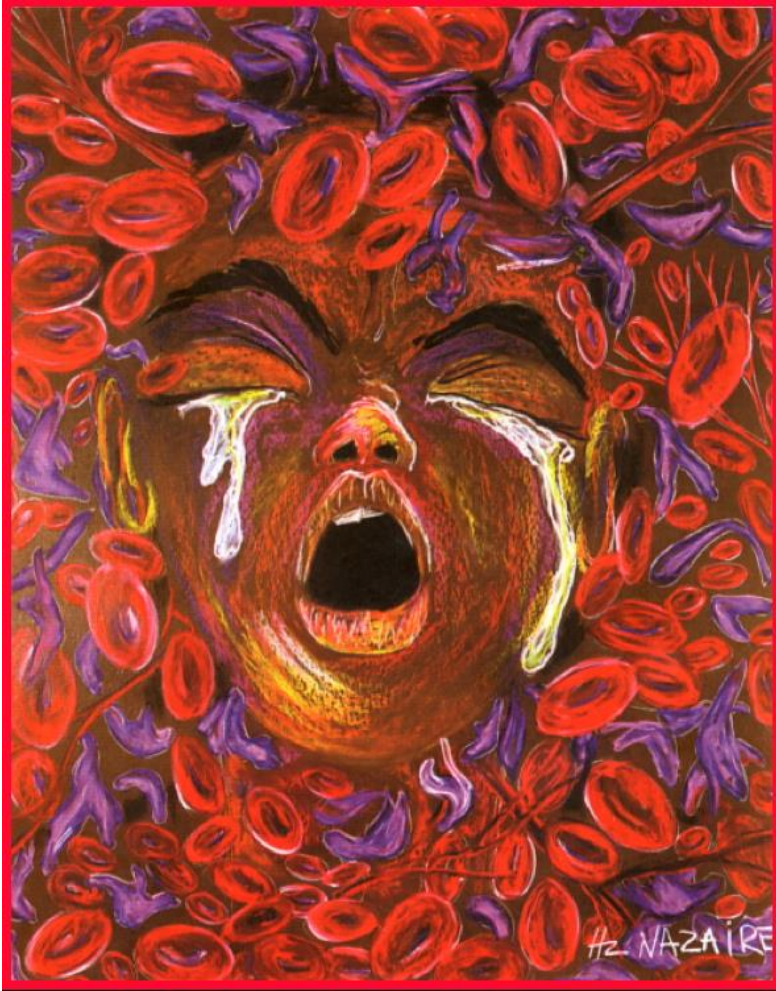


Gerd Blobel, M.D., Ph.D.

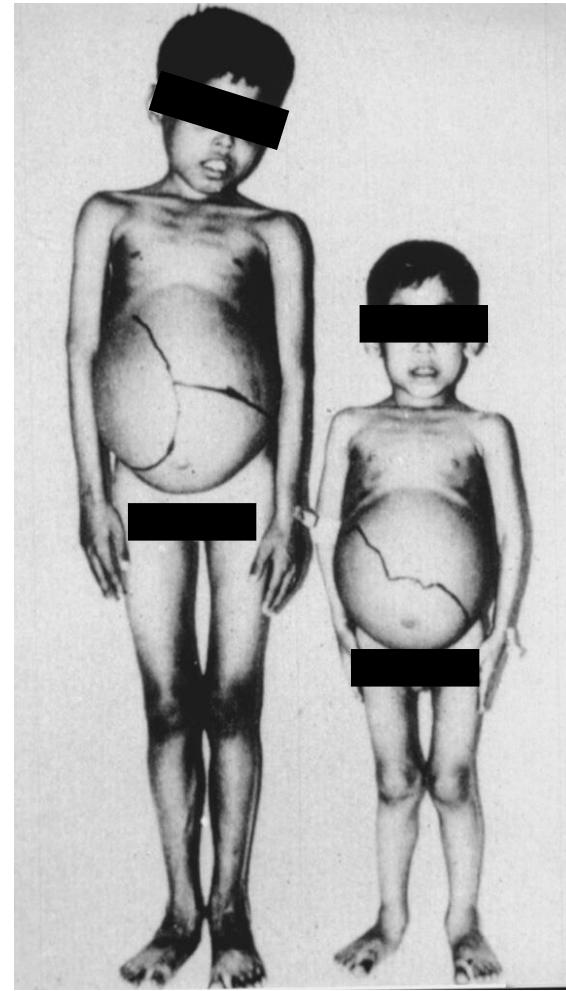
Frank E. Weise III Professor of Pediatrics
The Children's Hospital of Philadelphia
Co-director U-Penn Epigenetics Institute
Perelman School of Medicine
University of Pennsylvania



Hemoglobinopathies: Diseases of multi-system complications



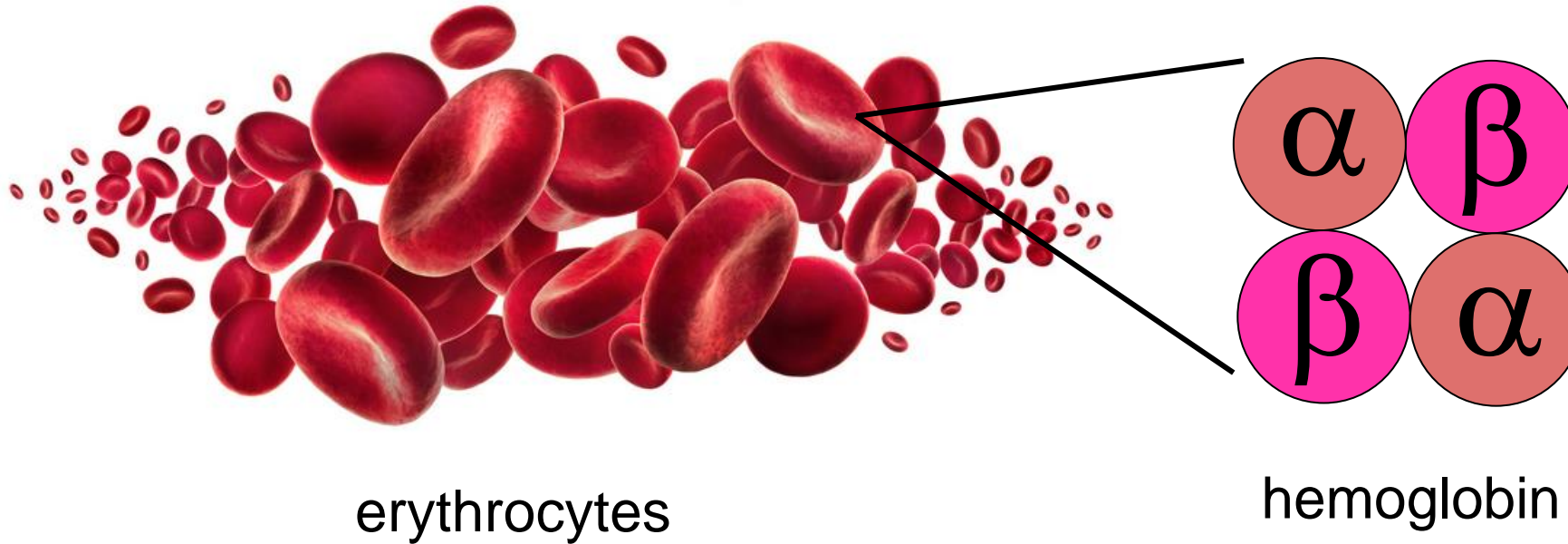
Sickle cell disease



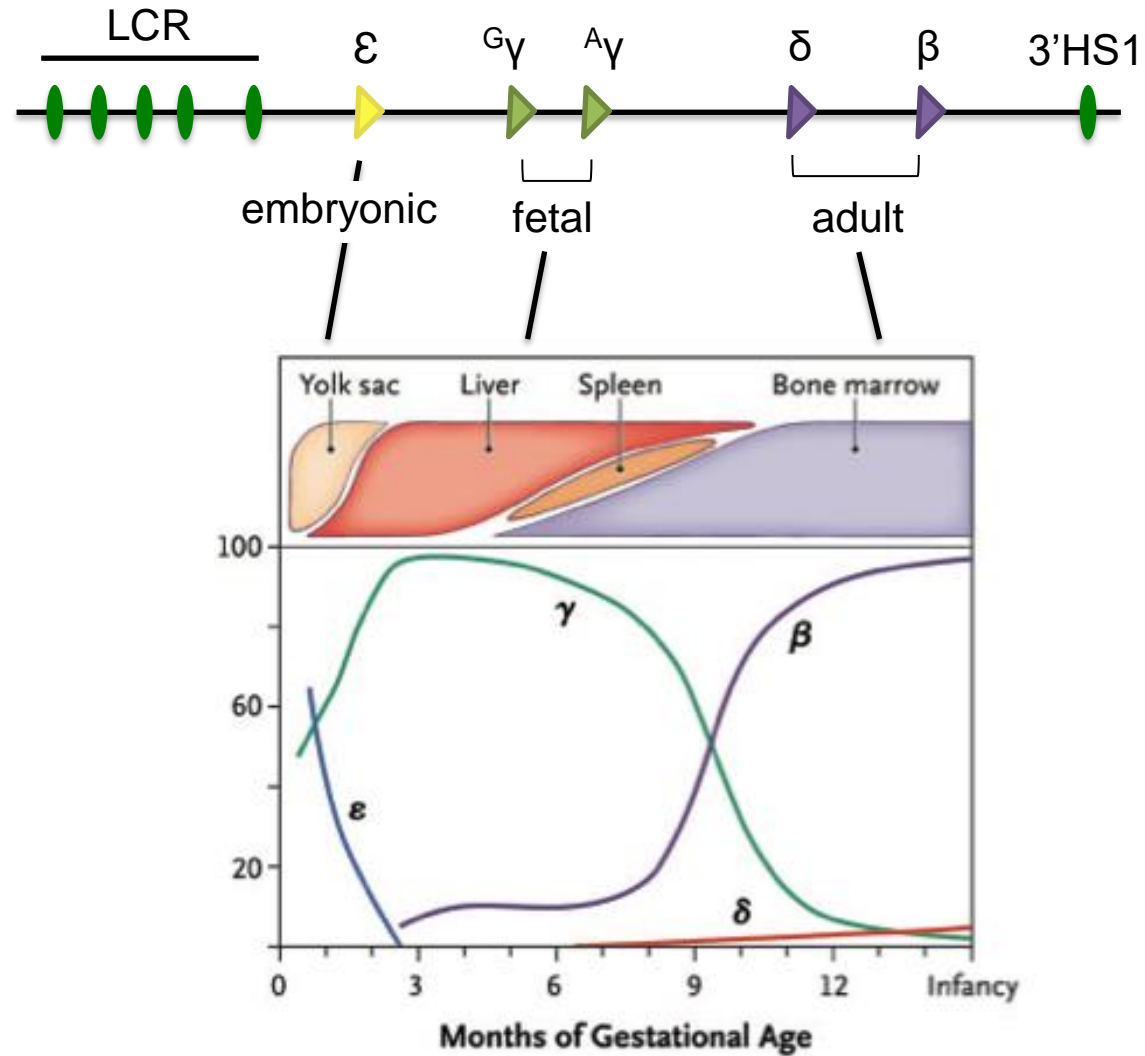
Thalassemia intermedia



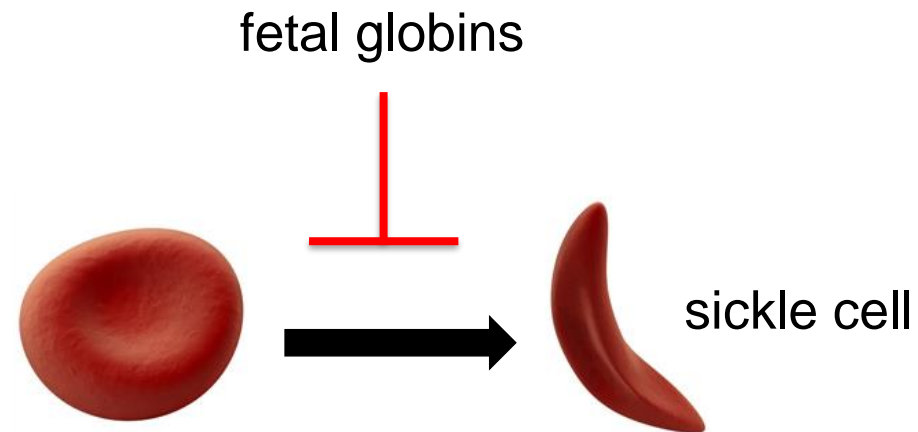
Red blood cells contain hemoglobin



Human hemoglobin switching



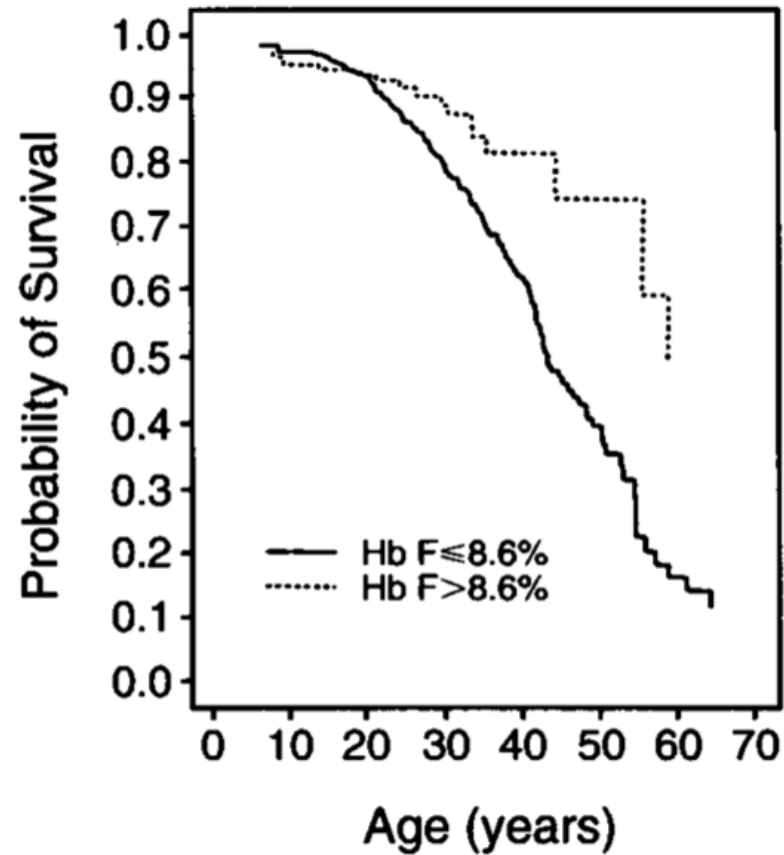
Elevated levels of fetal hemoglobin (HbF) improve the clinical course of patients with sickle cell disease



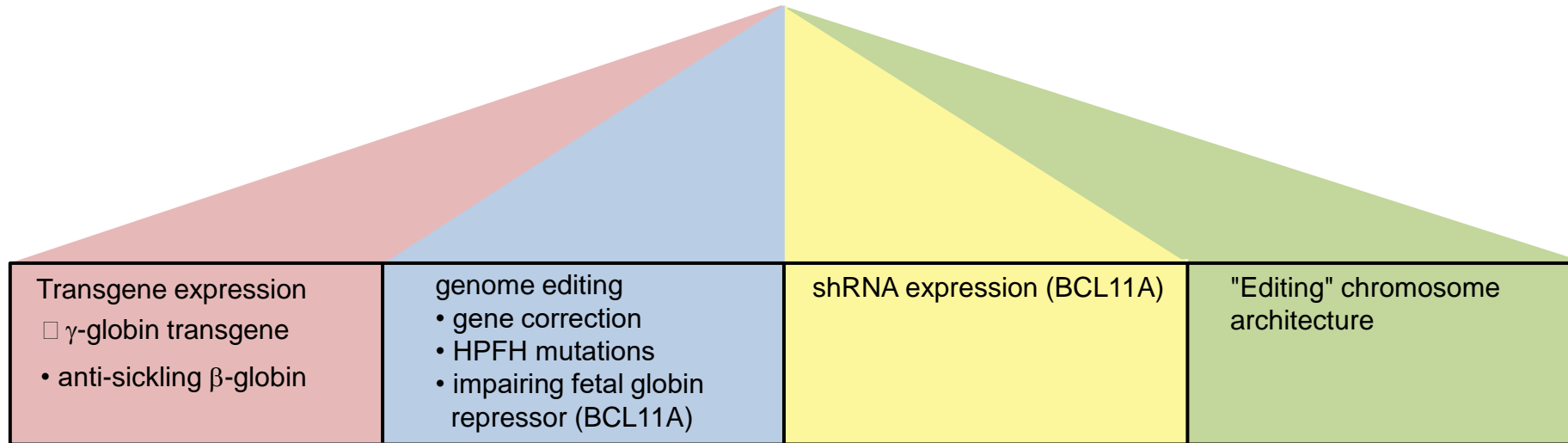
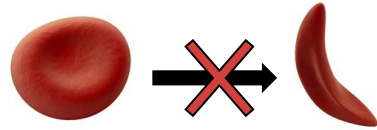
hereditary persistence of fetal hemoglobin (HPFH)



Elevated levels of fetal hemoglobin (HbF) improve the clinical course of patients with sickle cell disease



Therapeutic approaches for sickle cell disease (a brief scenic tour)



All of these approaches require autologous BM transplants



Goal: Identify HbF regulators that could be targeted with small molecules



What controls the fetal-to-adult switch?

Insights from GWAS

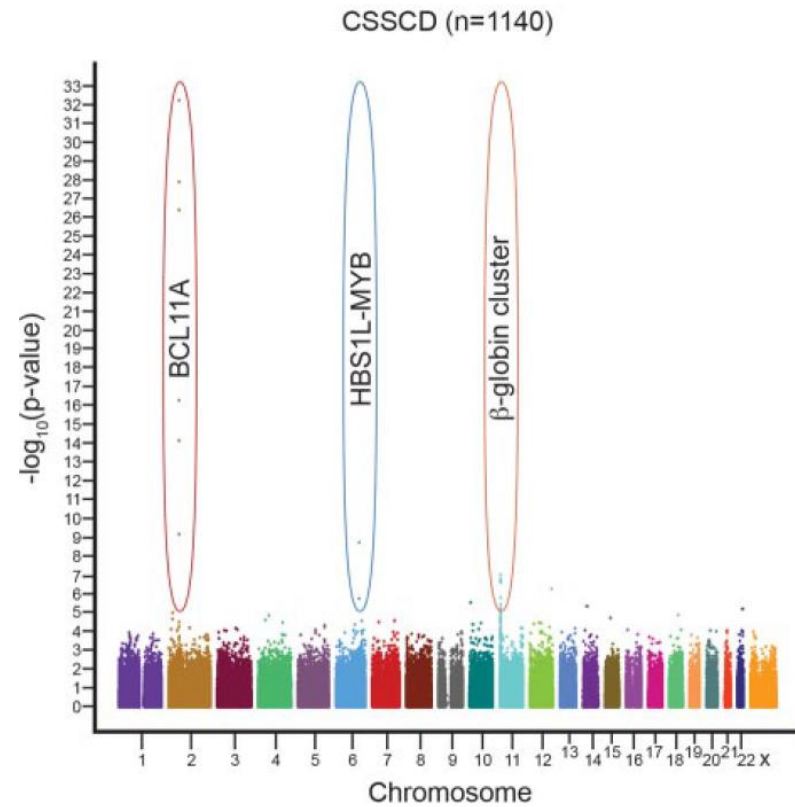
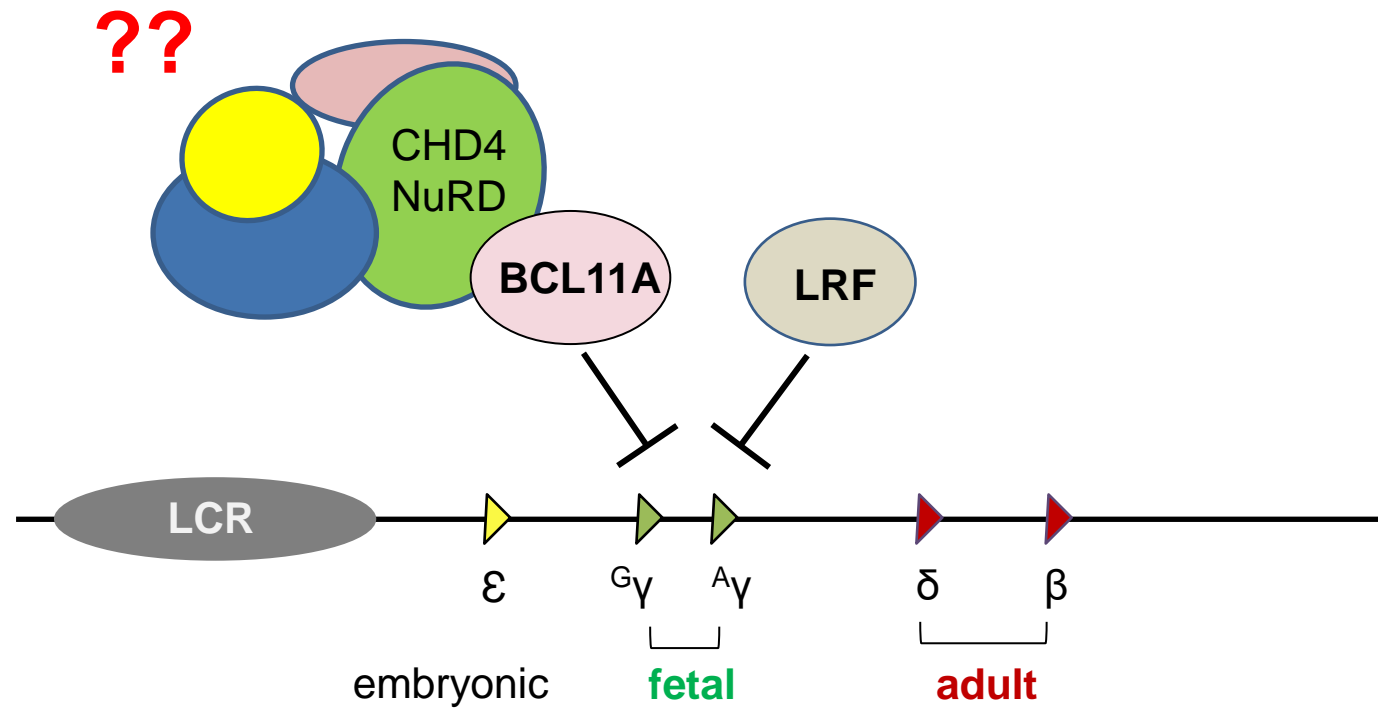


Figure courtesy of Guillaume Lettre



What controls the fetal-to-adult switch?

(More regulators? Novel therapeutic opportunities?)

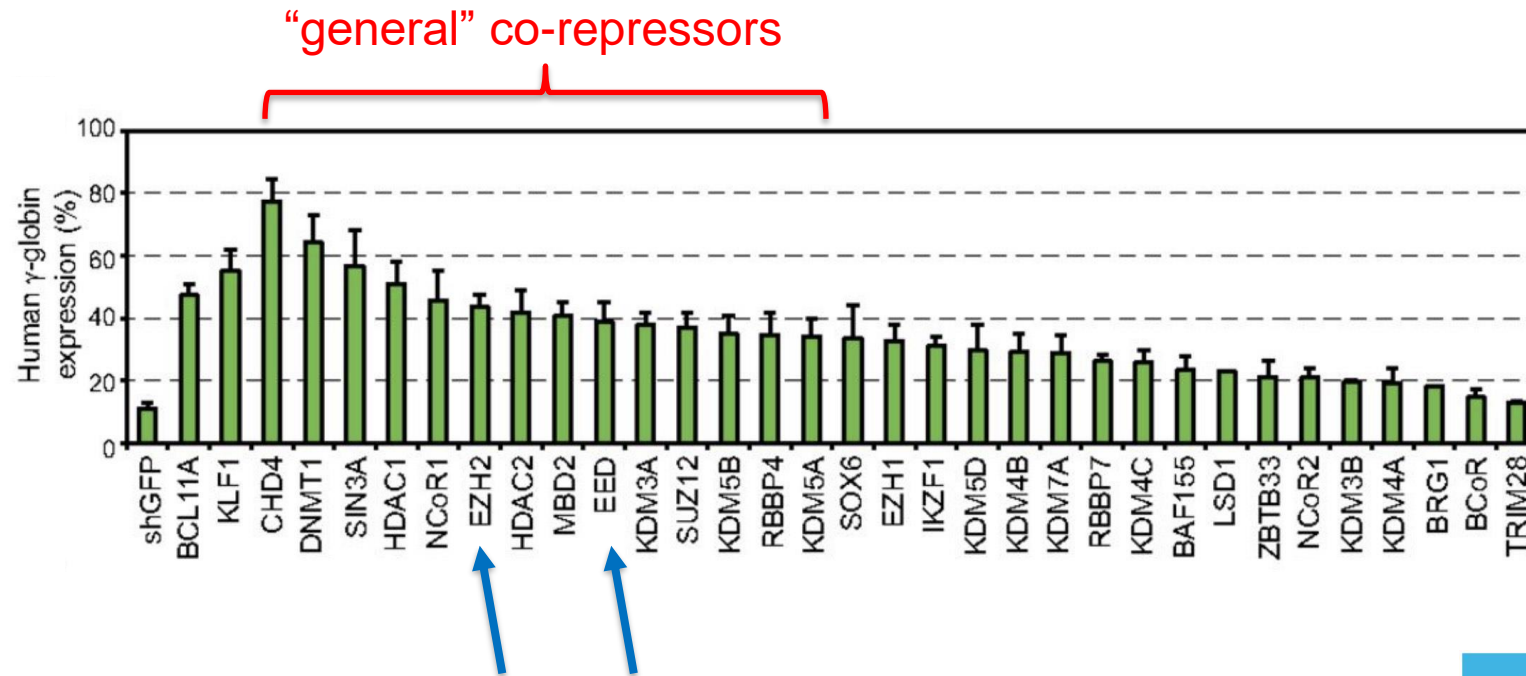


Hemoglobin switching involves widely expressed co-regulators

Corepressor-dependent silencing of fetal hemoglobin expression by BCL11A

Jian Xu^a, Daniel E. Bauer^a, Marc A. Kerényi^b, Thuy D. Vo^a, Serena Hou^a, Yu-Jung Hsu^a, Huilan Yao^b, Jennifer J. Trowbridge^a, Gail Mandel^b, and Stuart H. Orkin^{a,c,1}

PNAS 2013

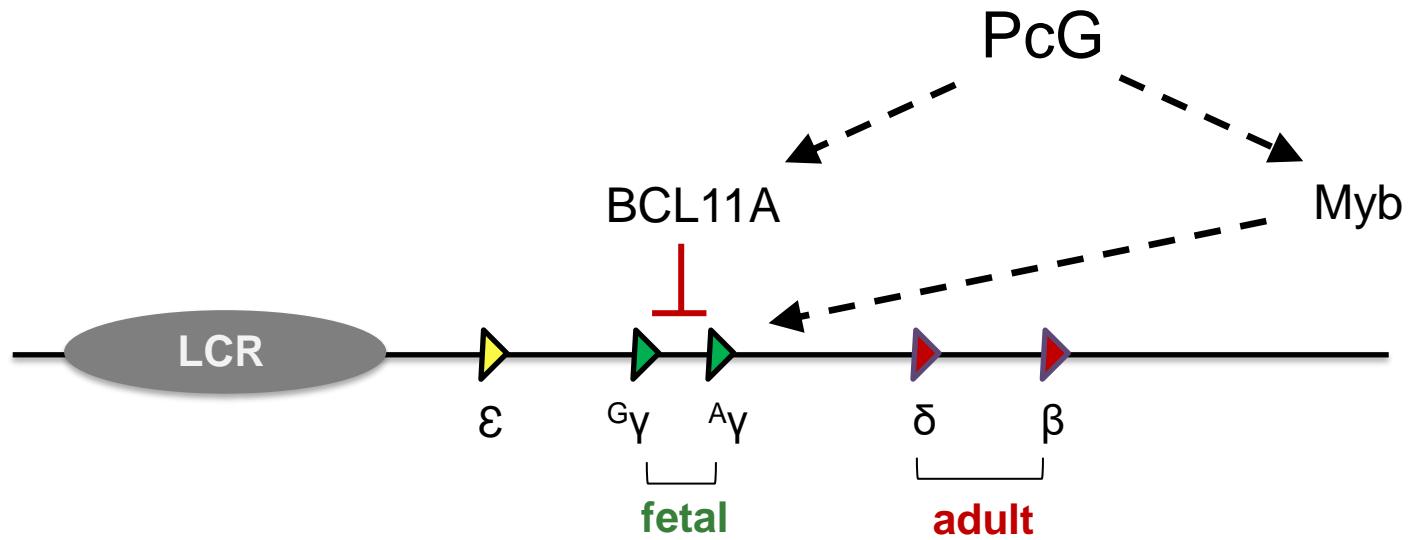


Hemoglobin switching involves widely expressed co-regulators

Our screens also identified several PRC2 components as fetal hemoglobin regulators, providing independent validation of PcG as targets.



Polycomb influences BCL11A and Myb levels



FTX-6058 Preclinical Pharmacology and Mechanistic Insights

Judith Dunn, PhD

President, Research & Development



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Sickle Cell Disease (SCD) is Caused by a Mutation in the β -globin Gene Resulting in Sickled Red Blood Cells, VOCs, and Anemia

THE DISEASE

Genetic disorder of red blood cells (RBC) caused by mutation in Hemoglobin-Beta (HBB) gene



PATIENT EXPERIENCE

- Vaso-occlusive crises (VOCs)
- Anemia
- Other complications, including stroke and organ damage



morbidity and mortality

TREATMENT OPTIONS

Short of SCT, current therapies are unable to address broad SCD symptomatology

- Hydroxyurea (current SOC) offers limited benefit and is only effective in a subset of individuals with SCD
- Newly approved therapies **address only a subset of SCD symptomatology** (i.e., anemia or VOCs)



only addressing subsets

GLOBAL IMPACT

Sickle cell disease is prevalent globally



Existing Therapeutic Agents Do Not Effectively Address the Unmet Need in SCD, and Underscores the Need for Novel Therapies




Hydroxyurea

Current Standard of Care




-  Potential to ameliorate broader SCD disease pathology
-  Significant non-responder population and waning efficacy over time
-  Potential safety risks (i.e., myelosuppression) and tolerability issues

HbS Polymerization Inhibitors

Increasing Total Hemoglobin

-  Addresses anemia (i.e., low total hemoglobin) by increasing levels of sickled hemoglobin (HbS)
-  Increasing HbS has not translated to clinical outcomes, such as VOCs
-  Approach does not address broad SCD disease pathology

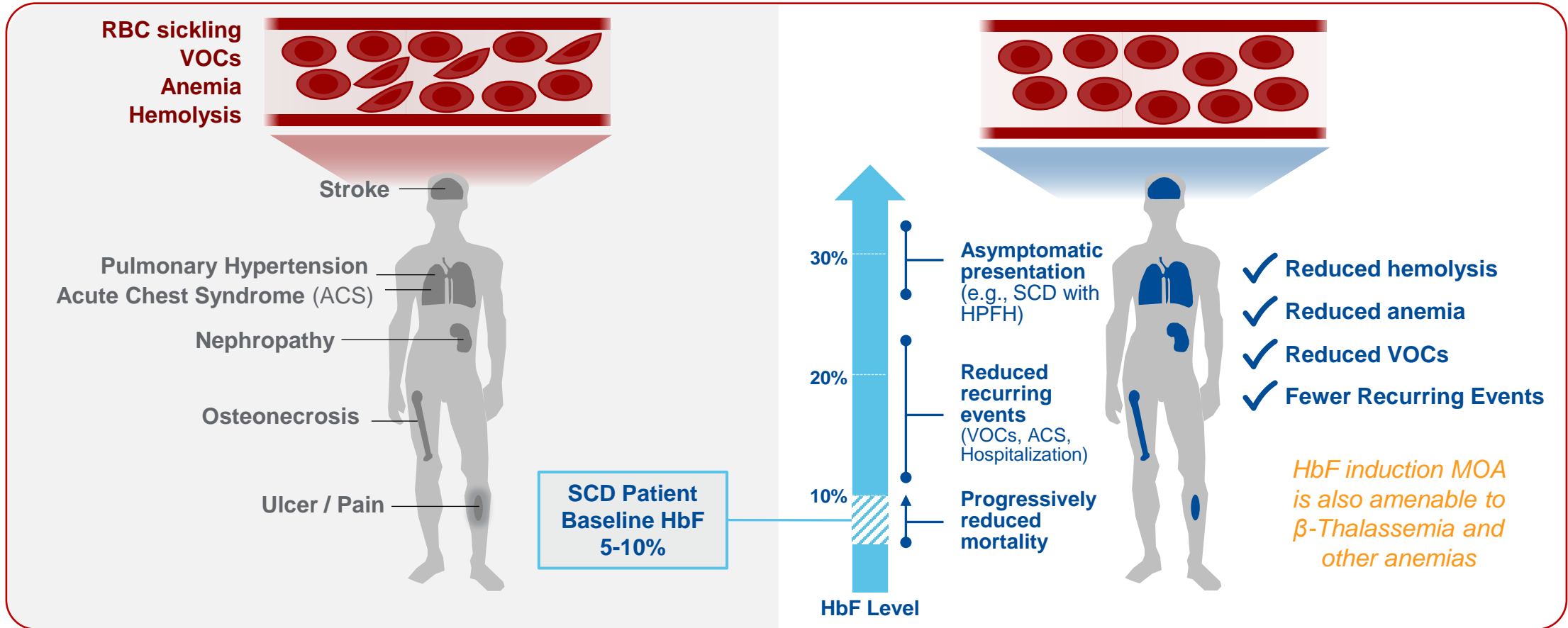
P-Selectin Inhibitors

-  Demonstrated ability to reduce the number of VOCs
-  Approach does not address broad SCD disease pathology
-  IV administration impacts patient convenience

Increasing Fetal Hemoglobin (HbF) is the Only Mechanism Shown to Broadly Improve Outcomes in SCD

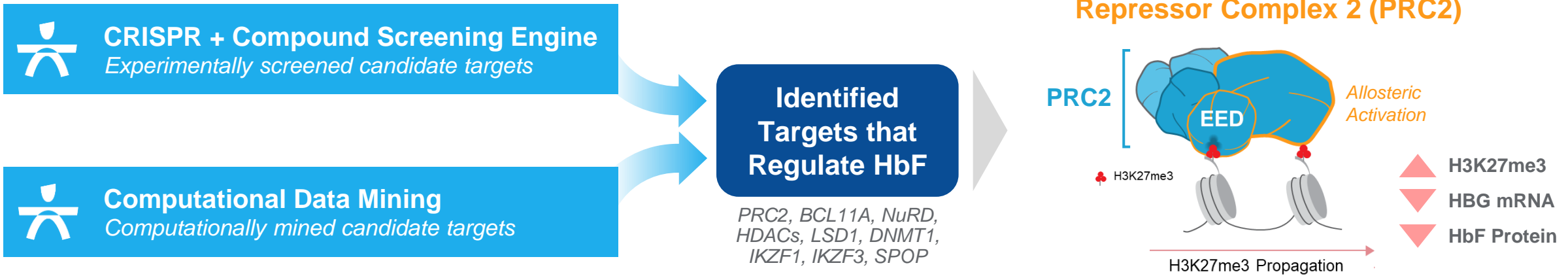
SCD Patient

SCD Patient with High Fetal Hemoglobin (HbF)



SCD individuals can have additional mutations that cause a condition known as hereditary persistence of fetal hemoglobin (HPFH), which leads to reduced or no symptoms in patients with SCD and β -thalassemia

FulcrumSeek Identified Embryonic Ectoderm Development (EED) as a Target for HbF Induction



Internal Medicinal Chemistry Led to FTX-6058, a Potent and Selective EED Inhibitor



FTX-6058 Induced HbF 2 – 3 Fold Across Multiple Preclinical Models with Strong Correlation between mRNA and Protein

Extensive Preclinical Validation

✓ HUDEP-2 Cells (Human)

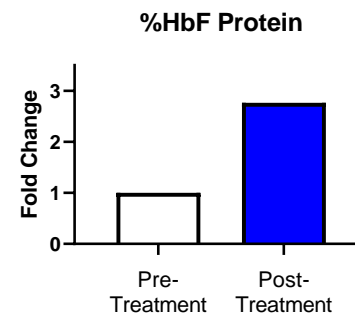
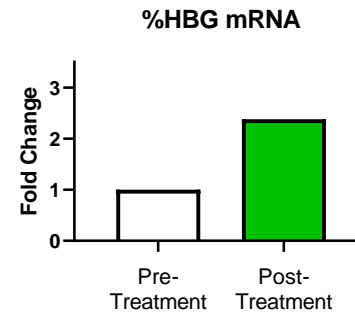
✓ Healthy CD34+ Cells (Human)

✓ SCD CD34+ Cells (Human)

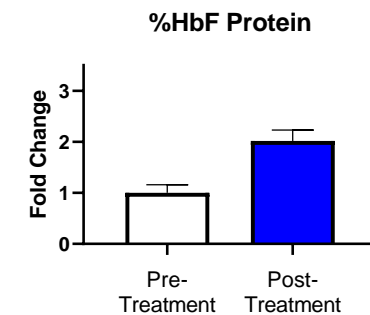
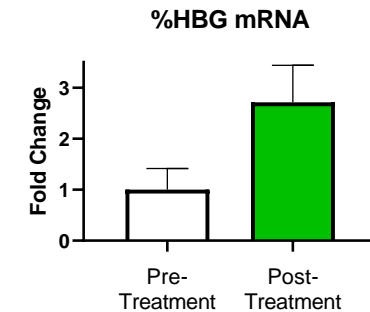
✓ Wild-type Mouse*

✓ SCD Townes Mouse

Healthy CD34+ Cells



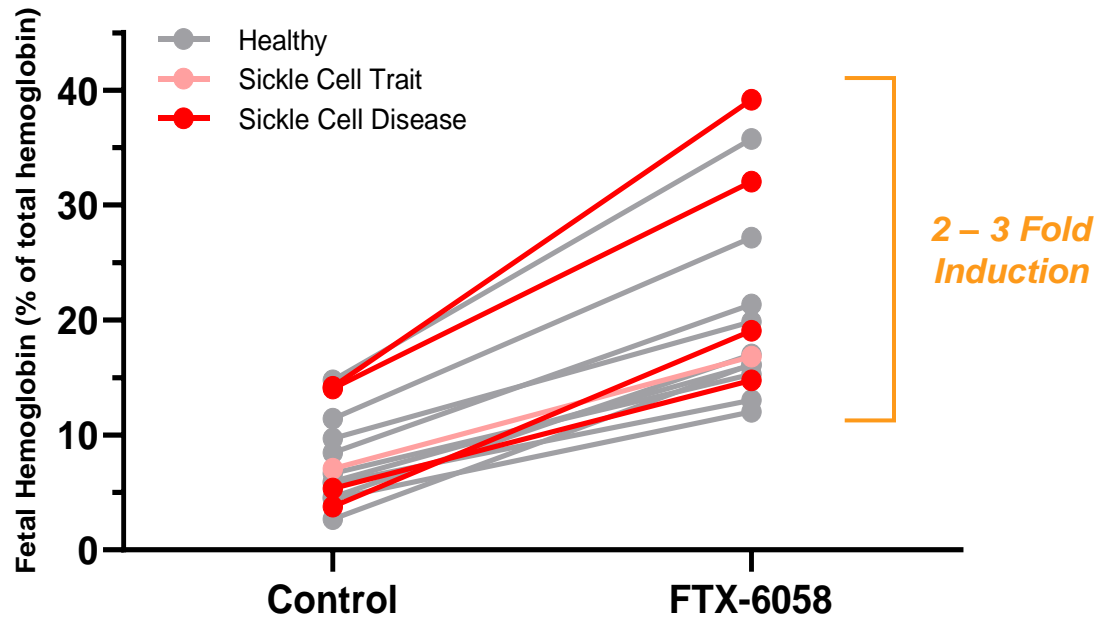
SCD Townes Mouse



Consistent 2 – 3 fold induction and strong correlation between fetal hemoglobin mRNA and protein observed both *in vitro* and *in vivo*

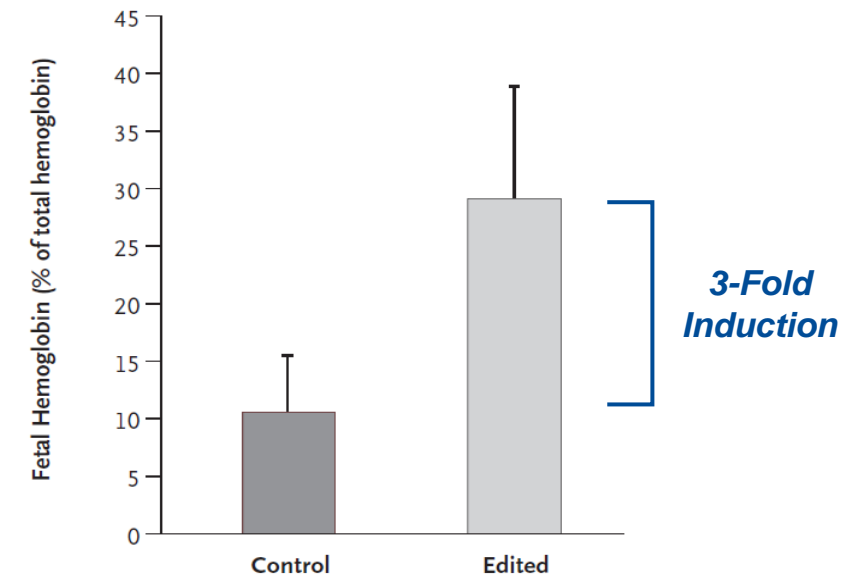
In Preclinical Studies, FTX-6058 Demonstrated HbF Induction Comparable to BCL11A Gene Editing

FTX-6058 HbF Induction in Healthy and SCD CD34+ Donors




- FTX-6058 achieved an absolute 8 – 25% increase in HbF, which has the potential to broadly address SCD symptomatology
- Demonstrated ability to achieve potentially “curative” HbF levels (e.g., 25 – 35% HbF) associated with asymptomatic disease

HbF Induction with BCL11A Gene Editing (CTX001) in Healthy CD34+ Donors¹



- CTX001, a gene editing therapy targeting BCL11A, achieved ~3-fold HbF induction in healthy donor CD34+ cells
- Robust preclinical HbF induction has translated to the clinic, achieving “curative” HbF levels and asymptomatic disease

HbF Fold Induction from CD34+ Cell Assay is Highly Translatable to the Clinic, with Initial HbF Induction Observed After ~1 Month of Treatment

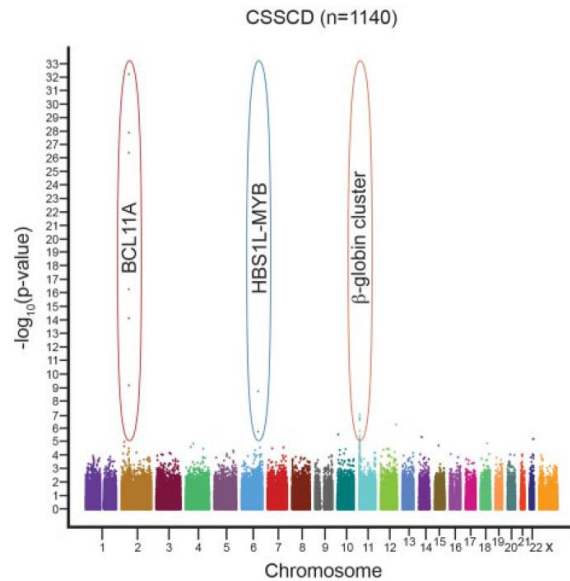
Mechanism / Asset	BCL11A KO	DNMT1i	HU	PDE9i	 FTX-6058
Preclinical HbF Fold Induction¹	3-fold	1.5 – 2 fold	1.1 – 1.2 fold	No Change	2 – 3 fold
SCD Clinical HbF Fold Induction²	~10-fold ³	2 – 2.5 fold ⁴	~1.7-fold ⁵	<1.1 fold ⁶	TBD <i>(Phase 1b)</i>
Time to Maximal HbF Induction	~5 months	>2 months ⁷	Up to 6 months	4 – 6 months	~3 months <i>(projection)</i>

Therapeutic goal is to induce HbF 2 – 3 fold over baseline levels

FULCRUM THERAPEUTICS ¹ Fulcrum generated data in erythroid cells derived from healthy CD34+ cells; ² Calculated as fold change over baseline %HbF (as a % of total hemoglobin) ³ CRISPR Therapeutics EHA 2021 virtual congress (SCD); ⁴ Molokie, R. PLoS Med. 2017; utilizing cohort 5 data (0.16 mg/kg data); ⁵ Data from the Multicenter Study of Hydroxyurea (MSH); Steinberg, MH. Blood. 1997; ⁶ Data from IMR-687 monotherapy in parent study provided in Imara 2021 EHA investor event; ⁷ based on 8-week treatment period 26

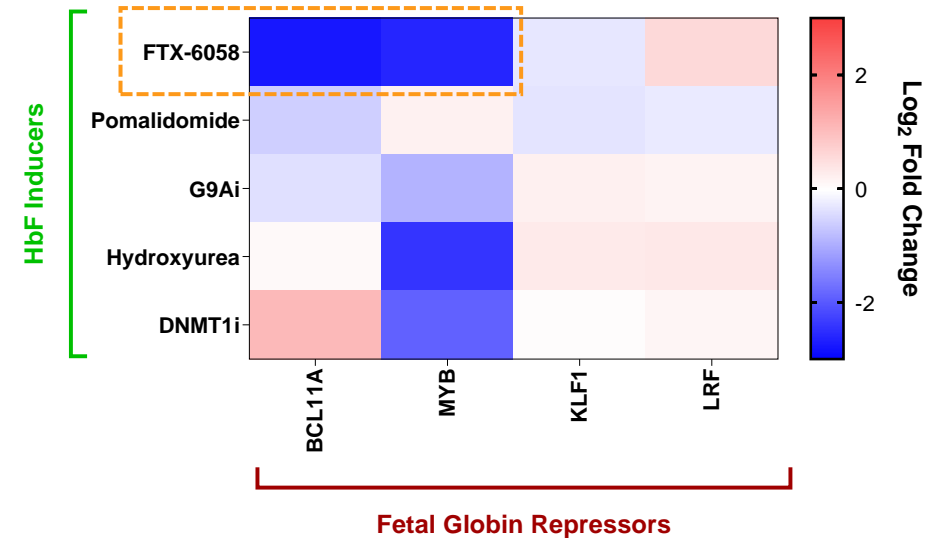
FTX-6058 Demonstrates Potent Downregulation of Key Fetal Globin Repressors, including BCL11A and MYB

Genome-wide Association Studies in SCD Have Identified Key Fetal Hemoglobin Repressors



- HPFH results from polymorphisms in BCL11A, the intergenic region of HSB1L-MYB, and the β -globin cluster
 - Downregulation of BCL11A and MYB have been demonstrated to induce fetal hemoglobin
 - Increased HbF levels also result from mutations in the β -globin gene

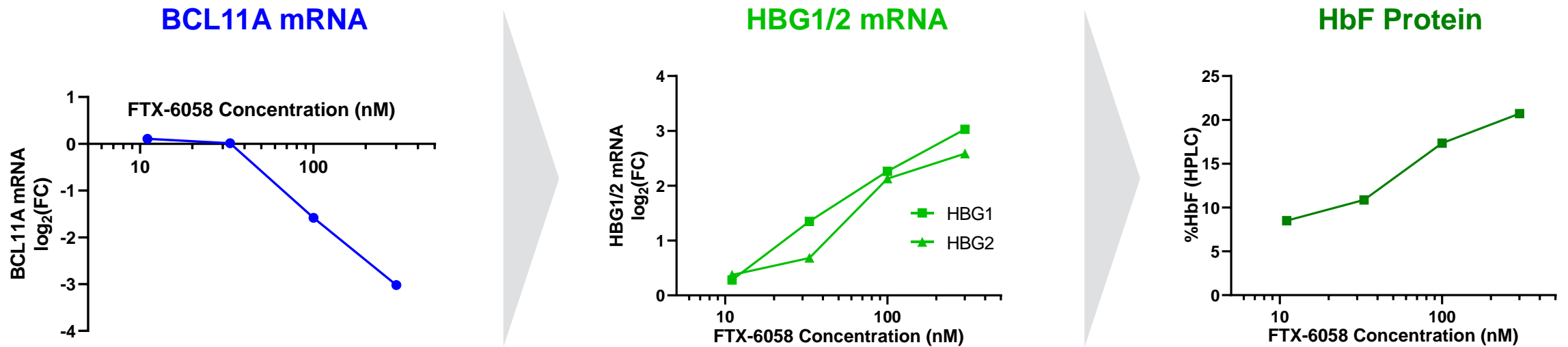
FTX-6058 Potently Downregulates BCL11A and MYB in Erythroid Cultures Derived from CD34+ Cells



- FTX-6058 potently downregulates master fetal globin repressors BCL11A and MYB
- Robust reduction of BCL11A mRNA only observed with FTX-6058
- No effect on other validated HbF repressors, such as KLF1 or LRF

FTX-6058 Demonstrates Robust Relationship Between BCL11A Downregulation and Fetal Hemoglobin Induction in Erythroid Cells

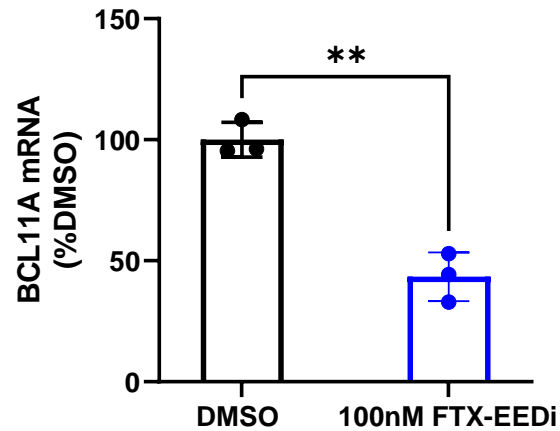
EEDi Modulation of BCL11A and HBG1/2 mRNA Translates to Robust HbF Protein Induction



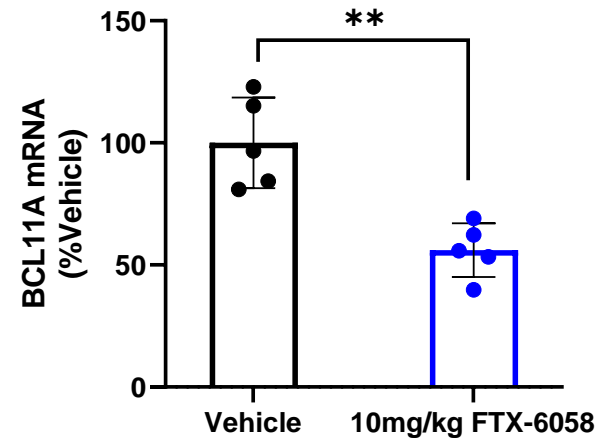
- FTX-6058 treatment results in dose-dependent BCL11A downregulation, and subsequent HBG1/2 upregulation and HbF induction
- Observe 2 – 3 fold HbF protein induction when BCL11A expression is reduced >50%
- Continue to establish relationship between MYB downregulation and HBG mRNA / HbF protein induction

In Vitro BCL11A Downregulation Observed with EED Inhibition Translates In Vivo

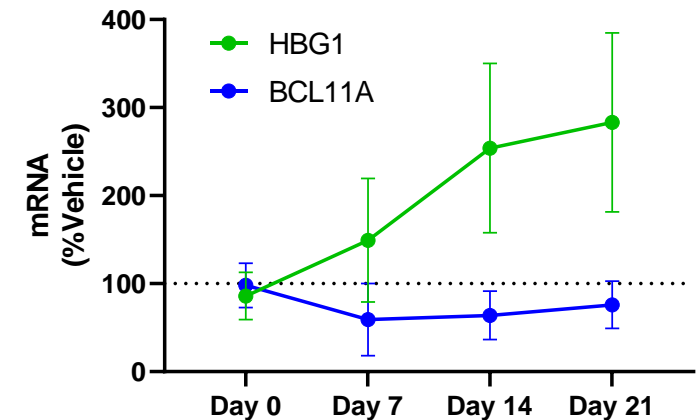
CD34+ Cell Derived Erythroid Culture



Wild-Type Mouse



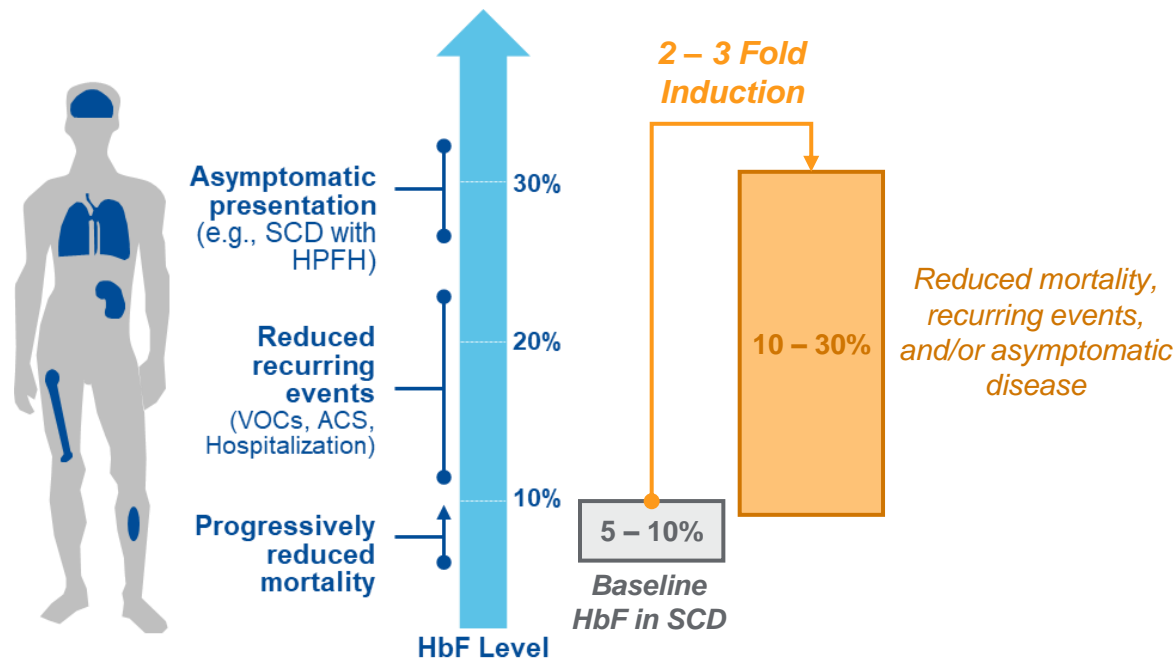
Townes SCD Mouse



- Observe consistent ~50% reduction in BCL11A across *in vitro* and *in vivo* studies
- In WT and Townes mouse, achieve ~50% reduction in BCL11A by day 5 – 7 of FTX-6058 treatment
- Durable BCL11A reduction achieved in Townes mouse model translates to robust 2 – 3 fold HBG1 induction

Oral HbF Inducer has the Potential to be the Preferred SCD Treatment Option for Patients and Providers

HbF Induction has the Potential to Provide Broad Clinical Benefits in SCD



FTX-6058 has Significant Advantages Over Gene Editing / Therapy Approaches

Oral HbF Inducer (FTX-6058)	<ul style="list-style-type: none"> • Convenient once daily oral pill with the potential to address broad SCD symptomology • Achieving 25 – 35% HbF levels has the potential to provide a functional cure • Potential for improved safety and tolerability compared to myeloablative conditioning and stem cell transplant • Oral pill is favorable for global distribution
Gene Editing / Therapy	<ul style="list-style-type: none"> • Potential for a functional cure, though requires a highly invasive procedure • Myeloablation safety risks and unknown efficacy durability • Likely significant barriers to access

FTX-6058 Phase 1 Healthy Volunteer Update

Christopher Morabito, MD

Chief Medical Officer

*****Results are from ongoing, blinded Phase 1 clinical trial*****



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FTX-6058 Phase 1 Healthy Volunteer Trial Design

Phase 1 Design and Endpoints

Primary

- Safety and tolerability

Secondary

- Pharmacokinetic measurements (bioavailability and half-life measurements)

Exploratory

- Target engagement
- HBG (fetal hemoglobin) mRNA
- F-reticulocytes (i.e., reticulocytes containing HbF protein)

Opportunistically assessing HBG mRNA and F-reticulocytes in whole blood after FTX-6058 treatment

Overview of Phase 1 SAD / MAD Cohorts

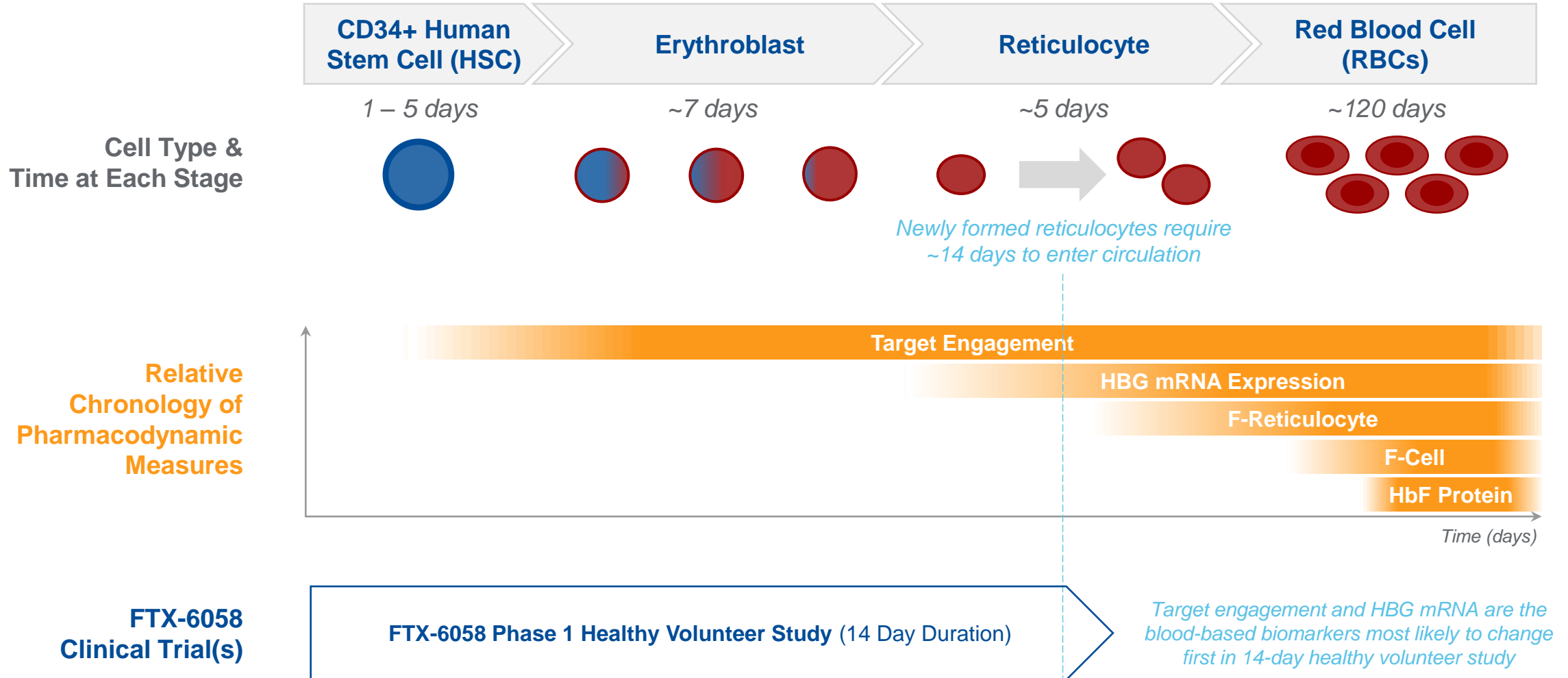
SAD Cohorts	Dose (mg)	MAD Cohorts (QD, 14d)
Cohort 1 ✓	2	Cohort 1 ✓
Cohort 2 ✓	4	
	6	Cohort 2 ✓
Cohort 3 ✓	10	Cohort 3 ✓
Cohort 4 ✓	20	Cohort 4 ✓
Cohort 5 ✓	30	Cohort 5 ✓
Cohort 6 ✓	40	
Cohort 7 ✓	60	
	90	

SCD Cohort (6 mg)
Ongoing

✓ Completed cohorts

- Each MAD cohort has 8 subjects (6 on study drug and 2 on placebo)
- Predicted human dose from PK/PD modeling is 4mg, and supports QD dosing
- The 6, 10, and 20mg doses were projected to achieve maximal target engagement and HbF induction

Erythropoiesis in Healthy Volunteers Influenced Biomarker Selection of 14-day Ph 1 Study



FTX-6058 has been Generally Well-Tolerated

Summary of Related TEAEs

SAD								
	Placebo	FTX-6058						Blinded
Adverse Event	N = 14	2mg N = 3	4mg N = 3	10mg N = 5	20mg N = 5	30mg N = 5	40mg N = 5	60mg N=5
Eosinophilia Count Increased	1 (7%)	0	0	0	0	0	0	0
Leukopenia	0	0	0	0	1 (20%)	0	0	0
Headache	0	0	0	0	0	0	0	1 (20%) ^a

^a Blinded data from SAD 60mg cohort

Food Effect	
	FTX-6058
Adverse Event	20mg ^c N = 10
Nausea	1

^c Cross-over study design

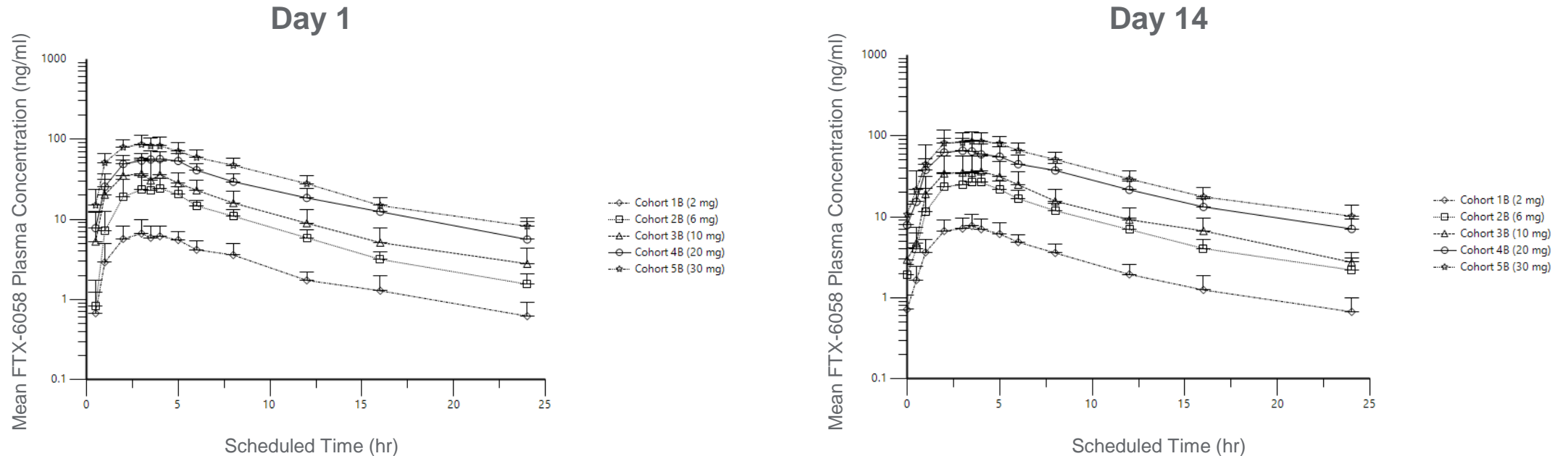
MAD						
	Placebo	FTX-6058				
Adverse Event	N =10	2mg N = 6	6mg N = 6	10mg N = 6	20 mg N=6	30 mg N=6
Diarrhoea (Loose Stool) ^b	1 (10%)	0	0	0	0	0
Dry Mouth	0	1 (17%)	0	0	0	0
Abnormal Stool	0	0	0	1 (17%)	0	0
Diarrhoea	1 (10%)	0	0	0	0	0
Neutrophil Count Decrease	0	0	0	1 (17%)	0	0
Headache	0	0	0	1 (17%)	0	0

^b Did not meet the WHO definition of diarrhea per protocol

- No SAEs reported to date and no discontinuation due to TEAE
- All treatment-emergent adverse events possibly related to FTX-6058 were Grade 1 or 2 severity per CTCAE criteria and resolved
- One Grade 3 and one Grade 4 TEAE: both unrelated to FTX-6058
 - Both TEAEs were asymptomatic, incidental creatine phosphokinase (CPK) increases detected at safety follow-up visit (i.e., 7 – 10 days post-treatment) in the 20mg food effect cohort and 10mg MAD cohort, respectively

FTX-6058 PK Profiles have Demonstrated Dose Proportionality in both SAD and MAD Cohorts

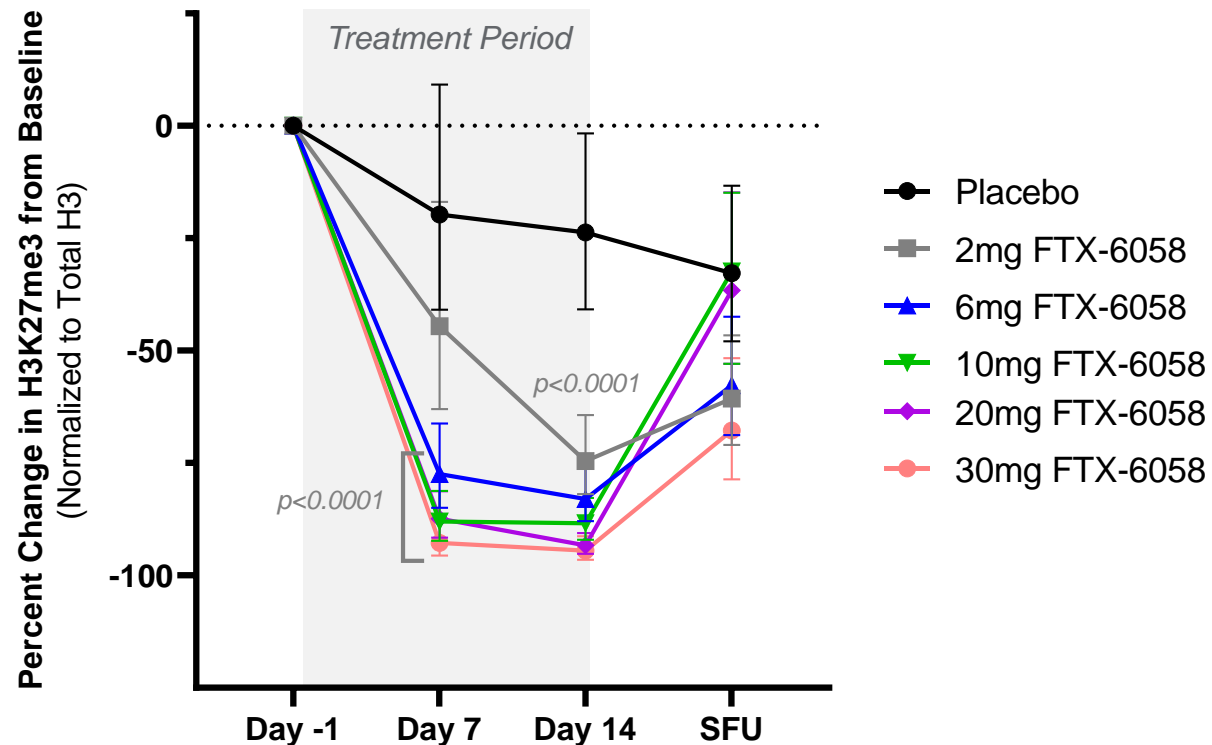
Plasma FTX-6058 Pharmacokinetics from MAD Cohorts



- Dose-proportional pharmacokinetics demonstrated across both SAD and MAD cohorts
- Mean half-life was approximately 6-7 hours in the MAD cohorts, and supports QD dosing
- No food effect observed with FTX-6058

All MAD Cohorts Demonstrate Approximately 75 – 95% Reduction in H3K27me3 Levels After 14 Days of FTX-6058 Treatment

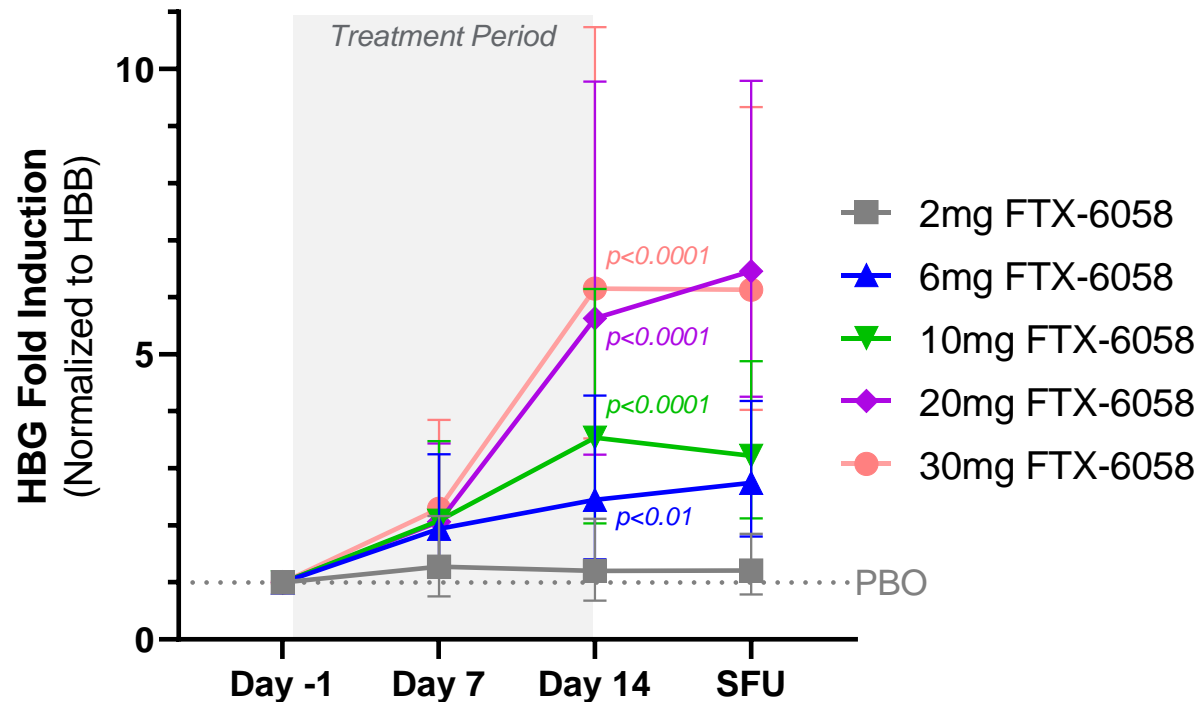
Mean Reduction (%) of H3K27me3 Levels



- Demonstrated proof-of-mechanism as evidenced by inhibition of H3K27me3 levels
- Robust target engagement observed in doses as low as 2mg
- Safety follow-up (SFU) samples were collected 7 – 10 days post 14-day treatment period

FTX-6058 Achieved Further HBG mRNA Induction in 20mg and 30mg Cohorts, With Up to a Mean 6.2-fold Induction

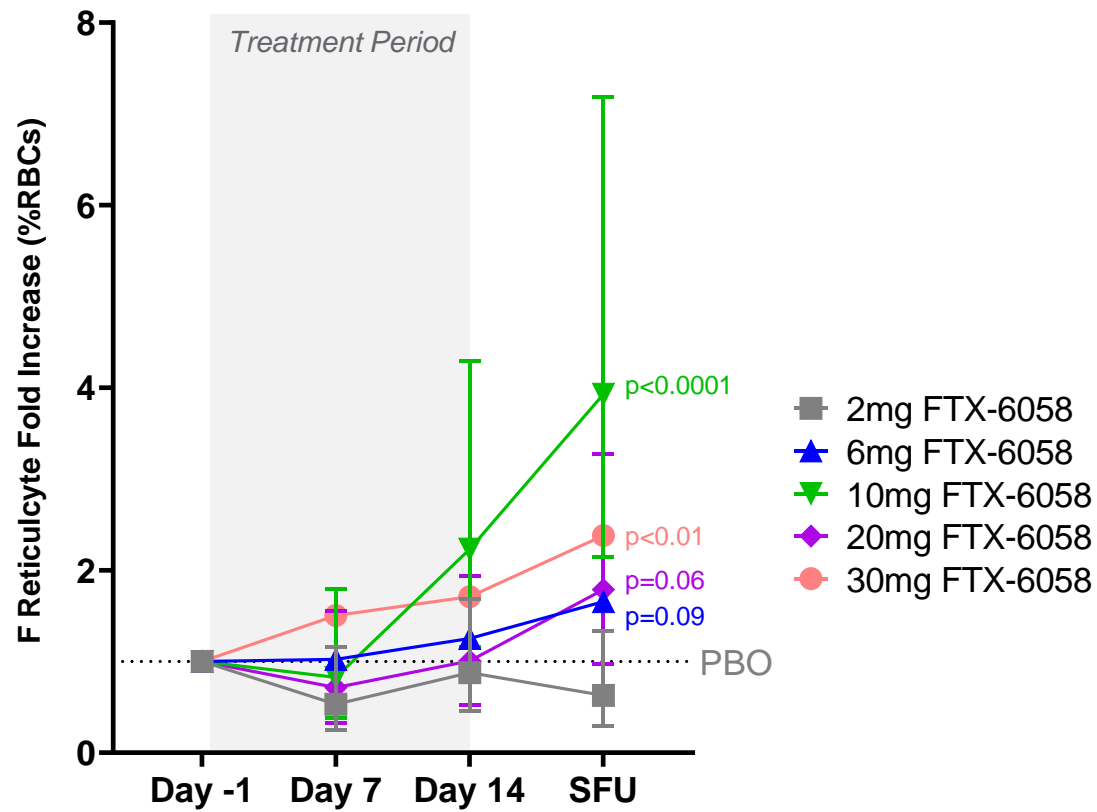
HBG mRNA Induction is both Time- and Dose-dependent in MAD Cohorts



- Demonstrated proof-of-biology as evidenced by HBG mRNA induction
- At Day 14, the 6 – 30mg FTX-6058 cohorts achieved a mean 2.4 – 6.2 fold HBG mRNA induction
- The maximal rate of HBG induction was observed in the 20mg and 30mg cohorts
- Maximal HBG induction has not yet been achieved
- Persistent and durable HBG induction observed 7 – 10 days after treatment (i.e., SFU)

Increases in F-Reticulocytes Provide Earliest Indication that HbF Protein Production is Beginning

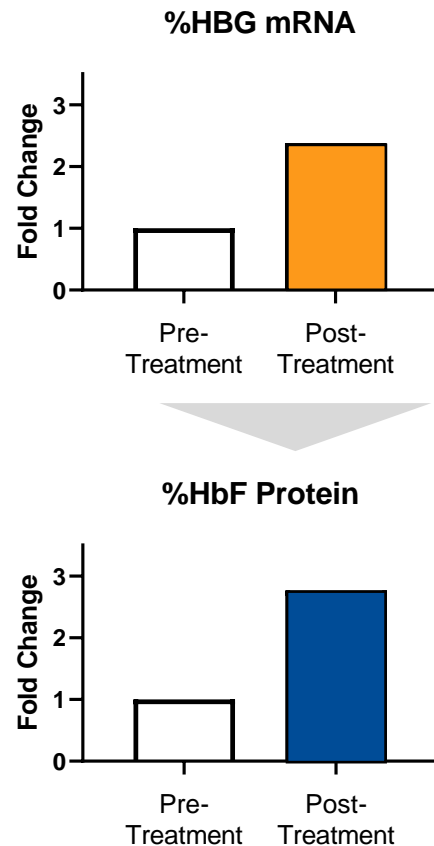
F-reticulocytes do not Predict HbF Fold Induction, but Demonstrate Translation of HbG to HbF



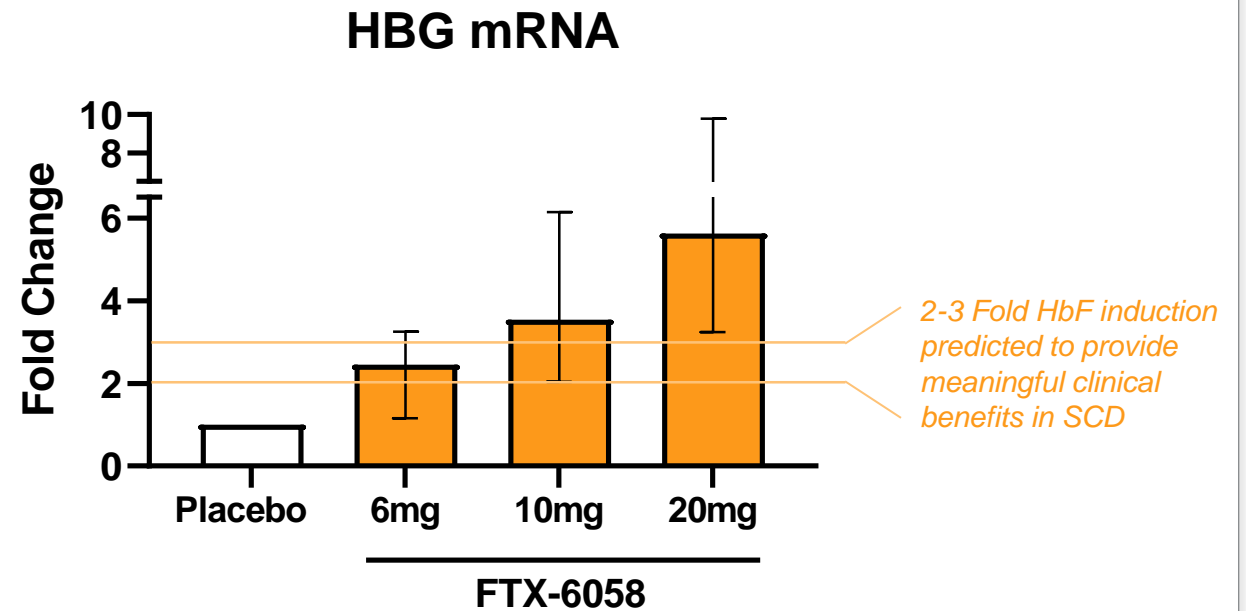
- Observed a mean **1.7 – 3.9 fold increase in F-reticulocytes** at the safety follow-up (SFU) visit with FTX-6058 doses $\geq 6\text{mg}$
- Increases in F-reticulocytes indicates that persistent HbG mRNA induction is beginning to translate to HbF production in newly formed reticulocytes
- Further increases in F-reticulocytes, F-cells, and HbF protein are projected with longer FTX-6058 treatment duration

FTX-6058 Clinical Results Achieve Induction Thresholds Predicted to Provide Meaningful Therapeutic Benefits and Potential Functional Cure to SCD Patients

Strong Correlation Between mRNA and Protein in Healthy CD34+ Cells

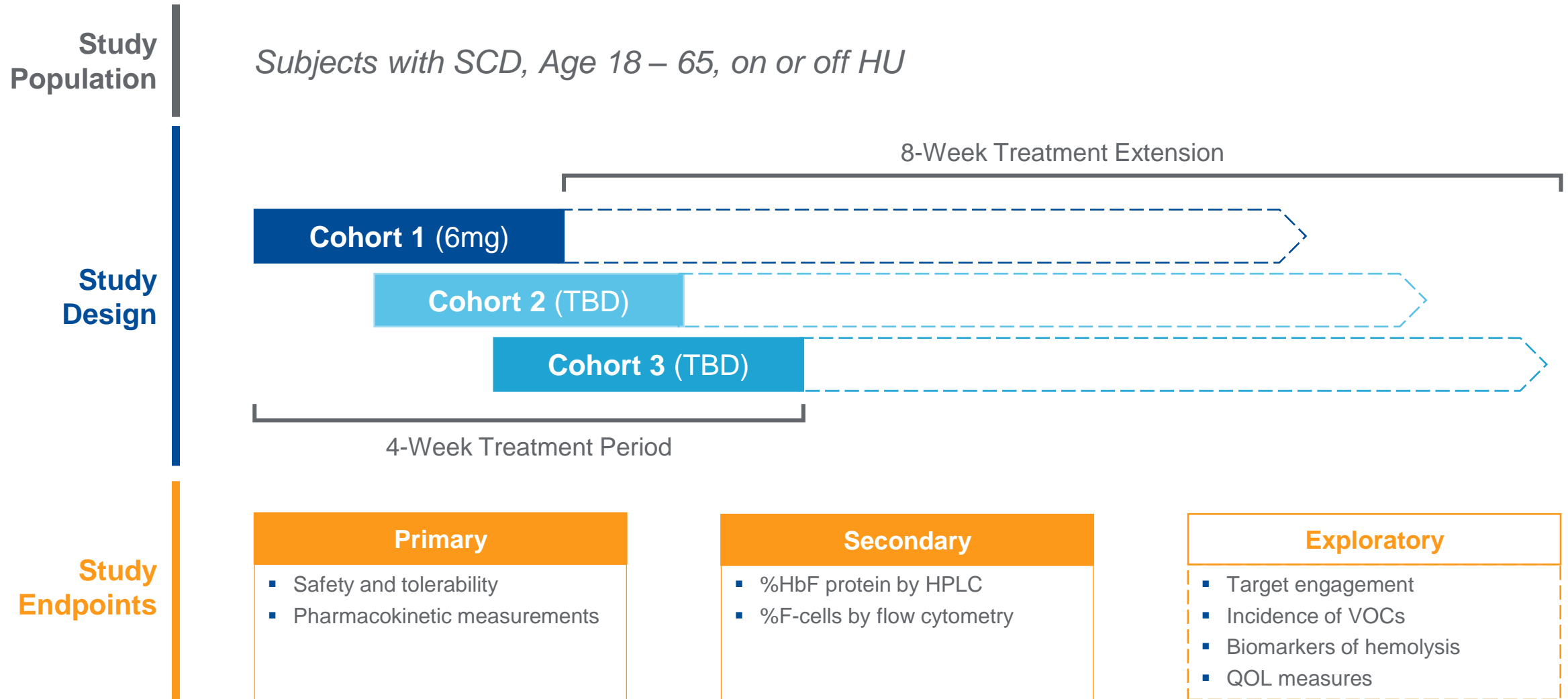


Clinical Data is Predictive of Meaningful HbF Induction

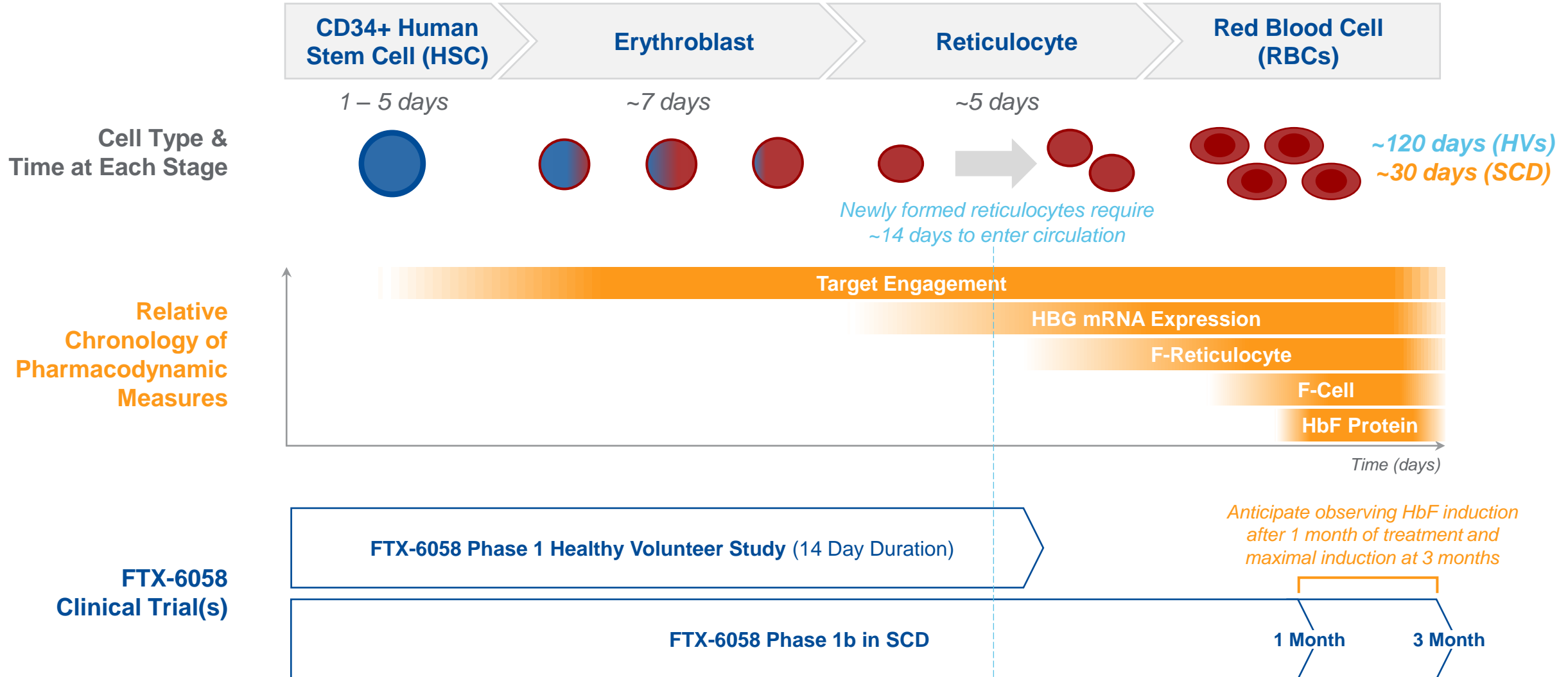


Anticipate similar HbF protein induction to translate in individuals with SCD

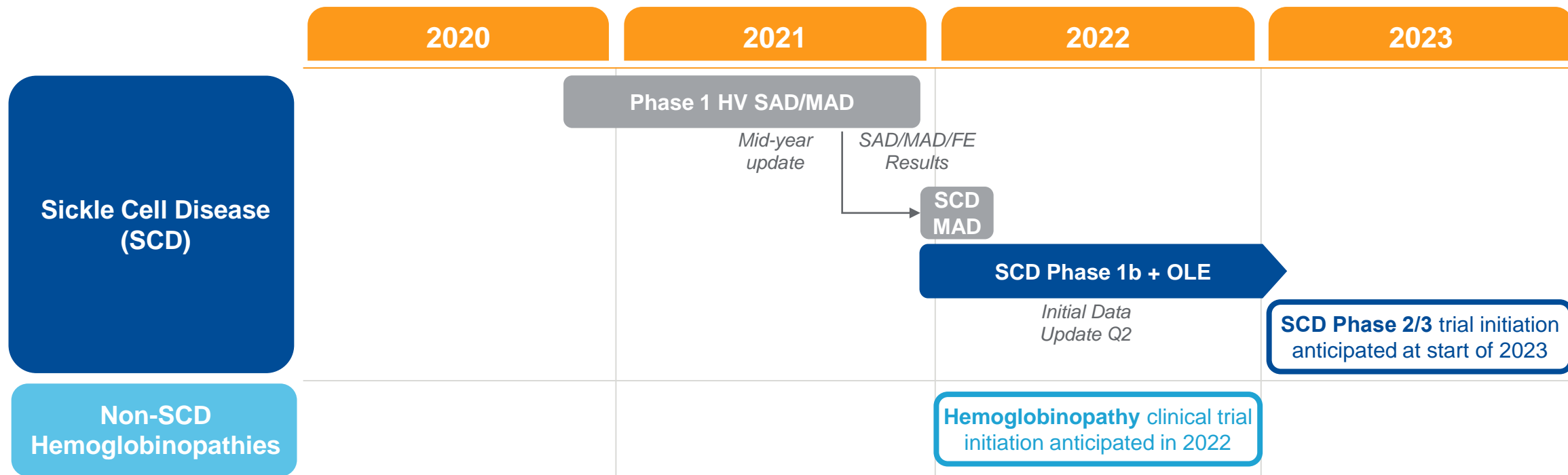
Phase 1b Clinical Trial Design



Increased Hemolysis and Short SCD RBC Lifespan Enables Quantitative Protein Measurements After 1 – 3 Months of Treatment in Phase 1b



FTX-6058 Anticipated Program Milestones



- Initiate Phase 1b SCD clinical trial by year-end
- Submit IND in additional hemoglobinopathies (e.g., β -Thalassemia) by year-end
- Report initial data from Phase 1b SCD clinical trial in Q2 2022
- Initiate Phase 2/3 SCD clinical trial in 2023



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Therapeutics

Q&A

