

bridgebio

hope through
rigorous science

R&D Day

October 12, 2021



Today's agenda

Introduction	Grace Rauh V.P. Communications, BridgeBio Pharma
Genetic Basis of Disease	Richard Scheller, Ph.D. Chairman of R&D, BridgeBio Pharma
BridgeBio's Endless Summer	Neil Kumar, Ph.D. Founder and CEO, BridgeBio Pharma
Precision Cardiorenal Introduction	Cameron Turtle, D. Phil. Chief Strategy Officer, BridgeBio Pharma
Acoramidis: TTR Stabilizer for ATTR	Jonathan Fox, M.D., Ph.D. Chief Medical Officer, BridgeBio Cardiorenal
Encaleret: CaSR Inhibitor for ADH1	Mary Scott Roberts, M.D. Sr. Director, Clinical Development, BridgeBio Cardiorenal
Gene Therapy Platform	Eric David, M.D., J.D. CEO, BridgeBio Gene Therapy
Mendelian Programs: PH1, LGMD2i, RDEB	Uma Sinha, Ph.D. Chief Scientific Officer, BridgeBio Pharma
Precision Oncology Programs: KRAS, SHP2	Eli Wallace, Ph.D. Chief Scientific Officer, BridgeBio Oncology
BridgeBioX	Charles Homcy, M.D. Chairman of Pharmaceuticals, BridgeBio Pharma
Conclusion	Neil Kumar, Ph.D. Founder and CEO, BridgeBio Pharma

Q&A

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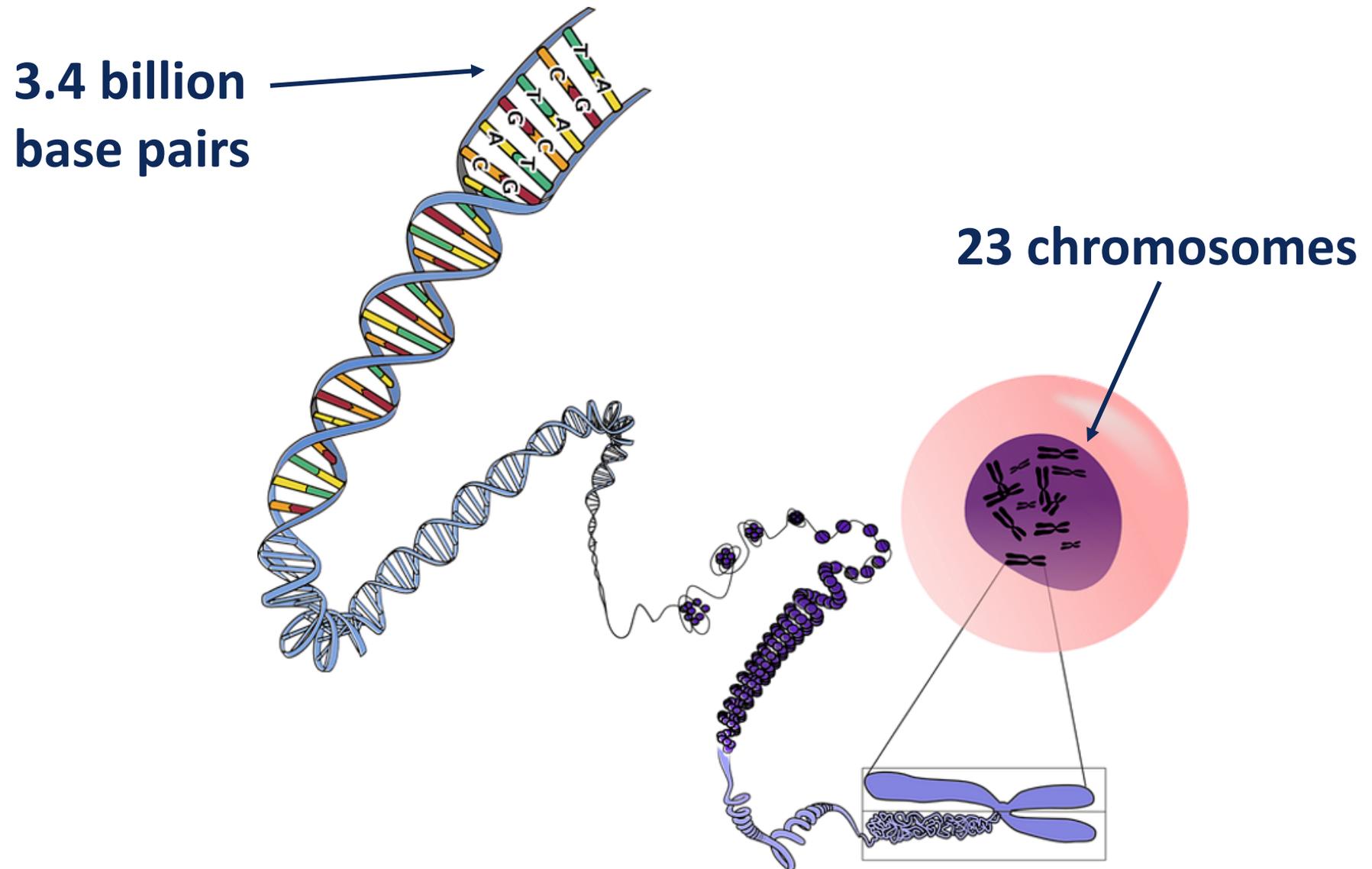
Genetic basis of disease

Richard Scheller, Ph.D.

Chairman of R&D



DNA structure

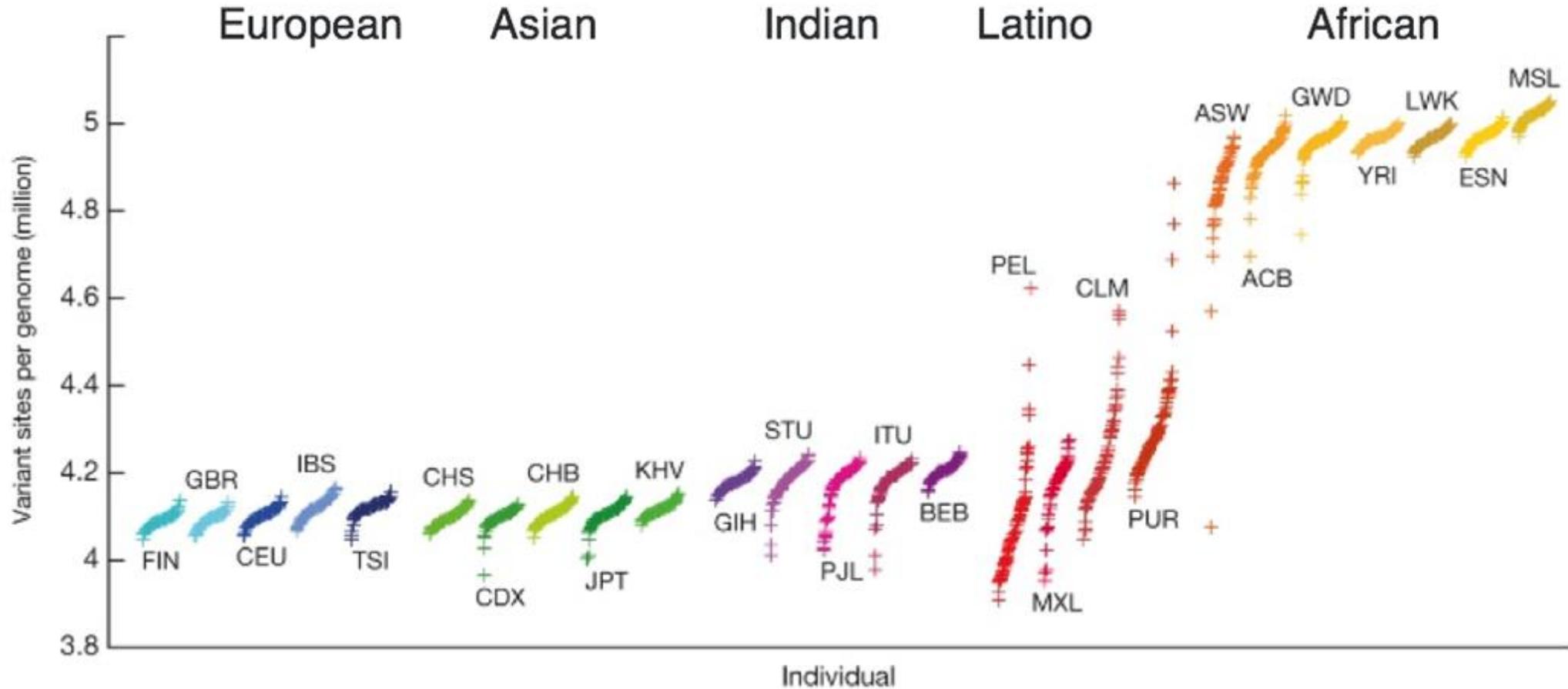


The human genome differs from our closest animal relatives by approximately 50M changes

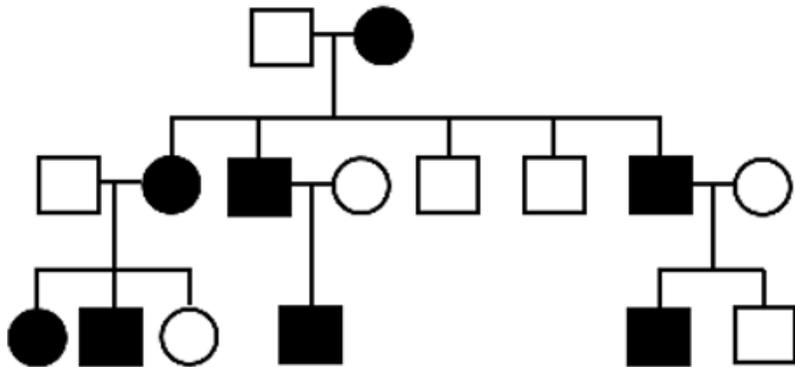




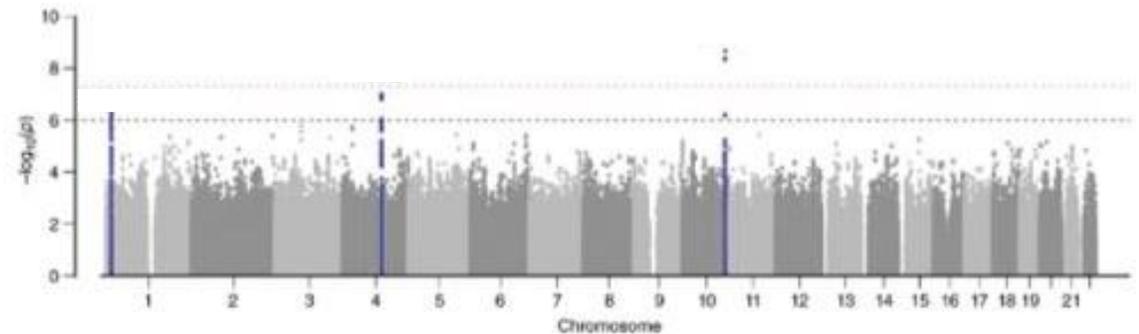
We differ from each other by 4-5 million variants - African genomes are the most diverse



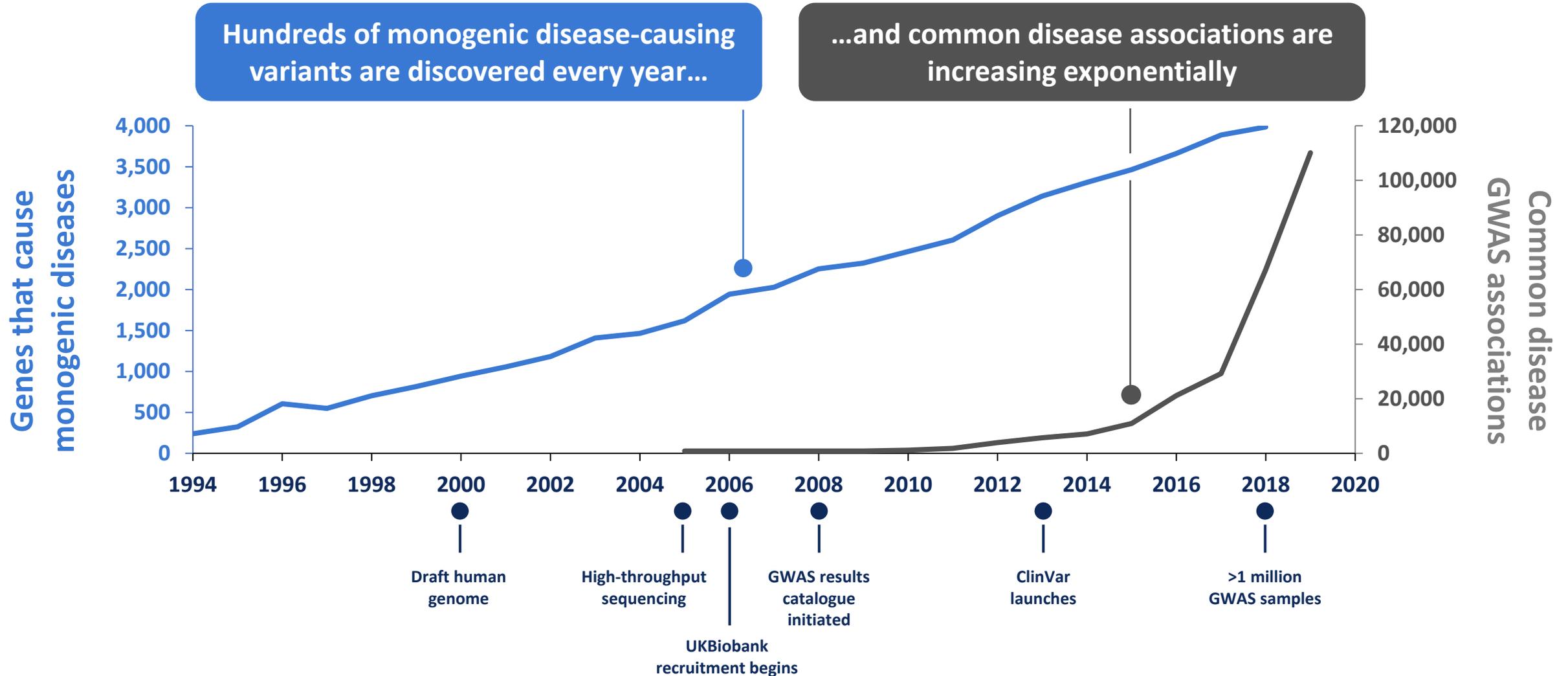
Monogenic target identification & validation



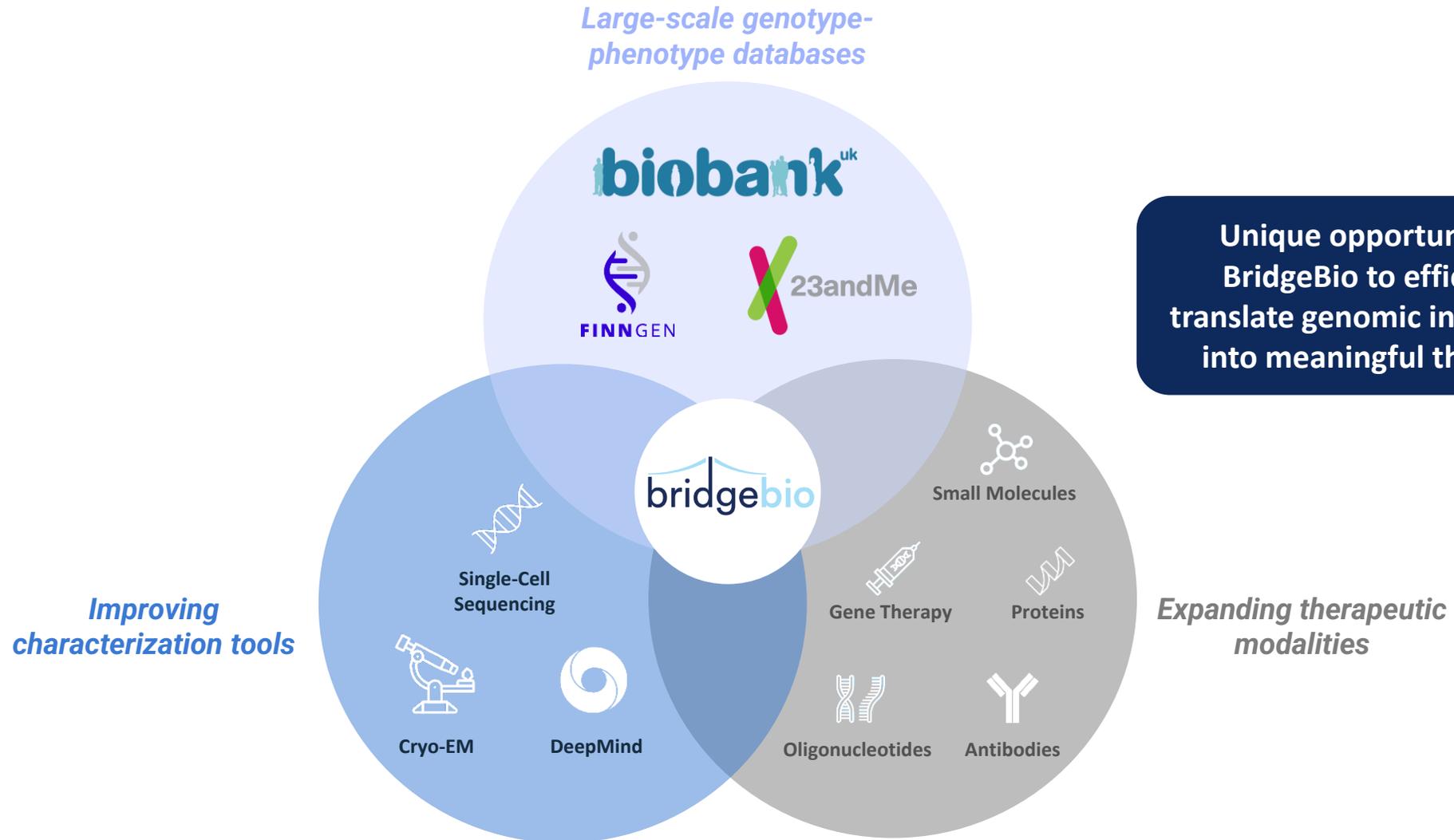
Indication expansion into common diseases



Single gene variants can cause or contribute to disease



Convergence of genetic innovation driving drug development



BridgeBio's endless summer

Neil Kumar, Ph.D.
Founder and CEO



Currently, few examples of sustainable innovation engines for genetic medicines

Big pharma R&D destroys value in aggregate

Big pharma R&D IRR



- R&D IRR is less than cost of capital for big pharma

Biotech companies have expectations that can't be met

The biotech market requires constant and significant innovation to create long term stable ROIC

- Currently, biotech EV is ~\$1.4 trillion
 - Assume – One wants to grow market cap by 12% YoY
 - Roughly, capital leaving the system by dividends + M&A = capital raised by IPOs + follow-ons
- If 70% of the value comes from new drugs, biotech would need to generate drugs worth ~\$2 trillion over the next 10 years, or approvals with aggregate **~\$40 billion peak year sales every year**

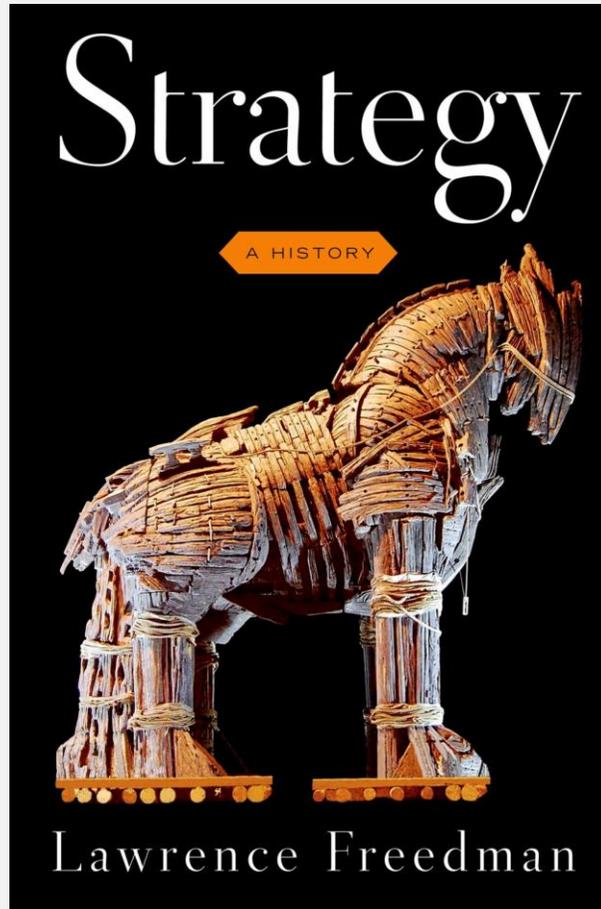
What does a sustainable genetic medicine innovation ecosystem look like? Criteria #1

Criteria #1: Need to solve for diseconomies of scale early, and economies of scale late



What does a sustainable genetic medicine innovation ecosystem look like? Criteria #2

Criteria #2: Each program needs to be NPV positive and supported by beautiful science



Each program is NPV positive

Realistic market size estimates

Only 2.3% of brands today >\$2 Bn

Capital efficient

IND cost < \$15 Mn for small molecule

Beautiful science

High POTS programs

More like engineering, less biology

Product market fit

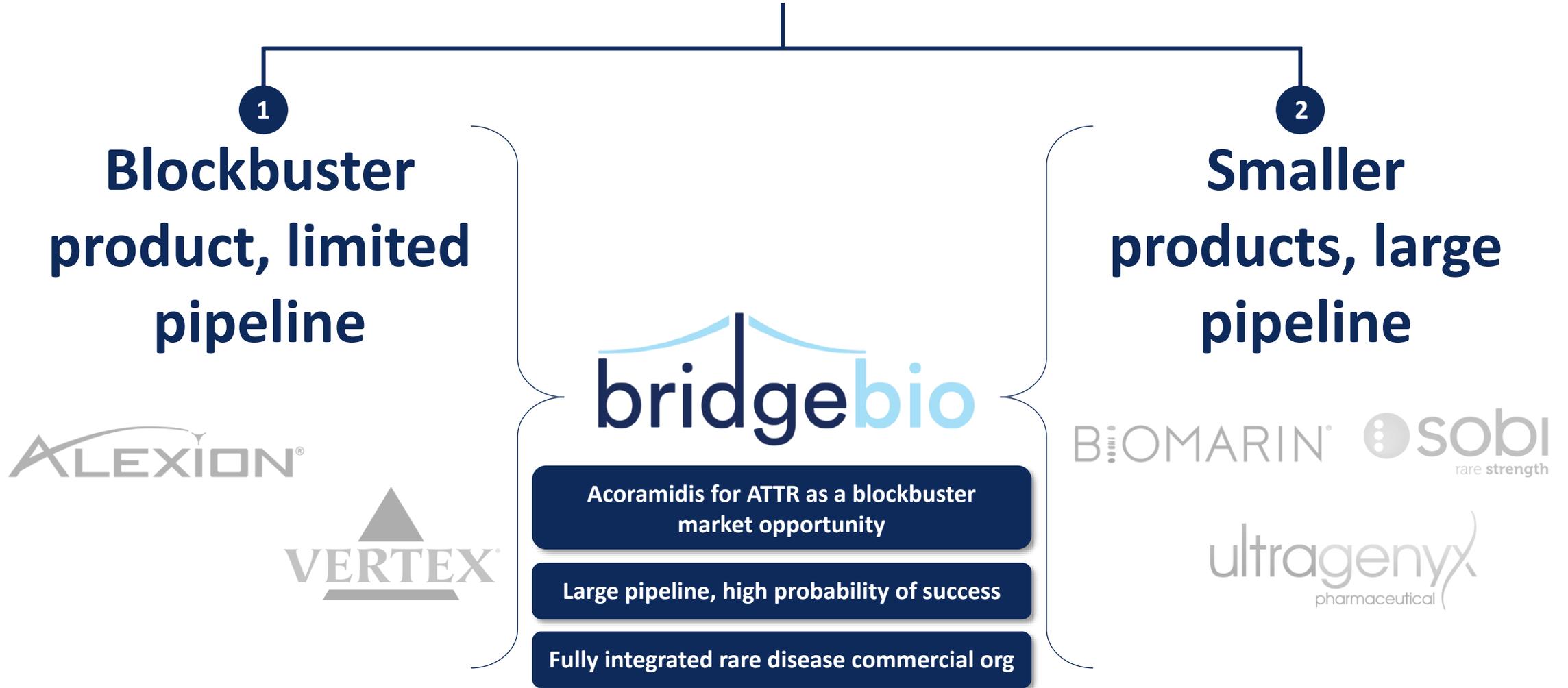
Therapies which match patient need

BridgeBio satisfies the criteria of a sustainable genetic medicine innovation engine

		Key attributes of BridgeBio
Key criteria of a sustainable genetic medicine innovation engine	Criteria #1	▪ The willingness and scale to fail and to re-allocate capital, within a decentralized company model
		▪ Focus at the level of individual diseases and assets. Drug R&D is a game of details
	Criteria #2	▪ Distinctive early-stage asset selection, based on a deep understanding of clinical unmet need, genetics, and underlying molecular pathophysiology
		▪ Efficient corporate structure that cuts no corners on science and medicine, but limits G&A, infrastructure and needless management
	People	▪ Experienced, product-focused R&D leadership that can define go / no-go's, required product attributes, and can drive programs through the clinic efficiently

Putting it all together: The opportunity to build the next great genetic medicine company

Traditional genetic medicines players fall into 1 of 2 archetypes



Fingerprints of hope – #1 BridgeBio's distinctive productivity

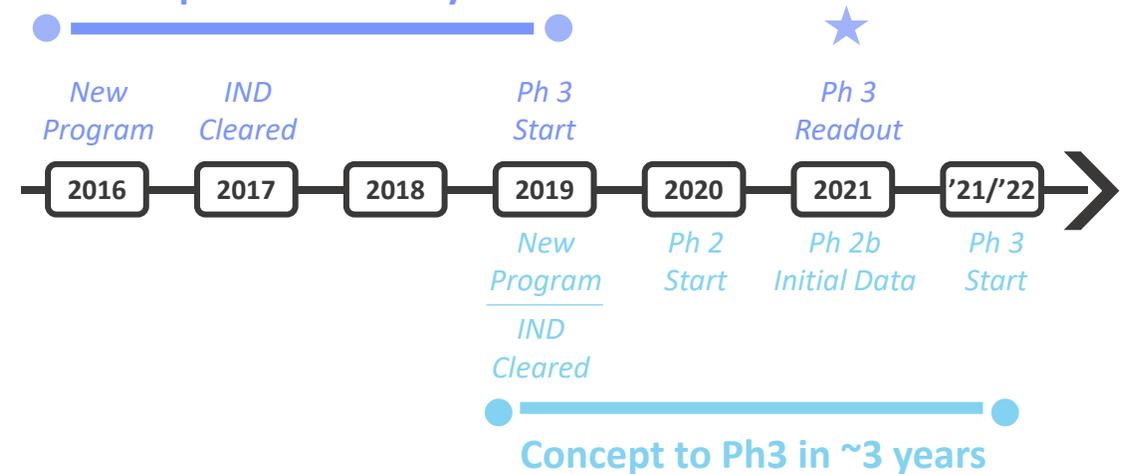
In less than 6 years since inception, BridgeBio has delivered...



Select Programs:

ATTR

Concept to Ph3 in ~3 years



ADH1

Concept to Ph3 in ~3 years

...building the framework for efficient, repeatable results

Fingerprints of hope – #2 BridgeBio's unique product platform

Recent Additions

- 4 new databases
- Bayesian methods for precise disease prevalence estimates
- 14 new university partnerships
- >5,000 new rare variants, >100 new causal genes discovered
- NMR spectroscopy for new drug targets
- AI for deciphering new protein structures
- Phenotypic screening for largest genetic diseases
- ASO screens for haploinsufficiency diseases
- 4 new clinical trials
- Activated 62 new sites in 11 countries
- Telperian partnership for ML empowered precision analytics
- Science 37 partnership for agile, decentralized clinical trials
- Two commercial launches (MoCD Type A, 2L CCA)
- 95% of lives covered in 6mths of NULIBRY launch
- Established a PAP to provide qualified patient's free access
- European office open, LATAM office upcoming

DISCOVER



Computational genomics, systemic disease mapping, broad network of academic partnerships

CREATE



Molecular dynamics assisted chemistry, gene therapy, therapeutic proteins, antisense oligos

TEST



20 ongoing trials across >450 sites and 26 countries, central operations toolkit and analytics

DELIVER



Global infrastructure, diagnostics, patient support, disease state awareness

Fingerprints of hope – #3 BridgeBio's great drug developers

Scientific insight and judgment from industry leaders with a proven track record



Charles Homcy, MD
Founder and Chairman of
Pharmaceuticals



Frank McCormick, PhD
Founder and Chairman of
Oncology



Richard Scheller, PhD
Chairman of R&D



Len Post, PhD
Advisor



Phil Reilly, MD, JD
Advisor



Experienced team of R&D operators responsible for 100+ INDs and 20+ approved products

Mendelian / Cardiorenal



Uma Sinha, PhD
Chief Scientific Officer



Robert Zamboni, PhD
Chemistry



Jonathan Fox, MD, PhD
Chief Medical Officer,
Cardiorenal



Eli Wallace, PhD
Chief Scientific Officer, Oncology



Pedro Beltran, PhD
SVP, Oncology



Fingerprints of hope – #4 BridgeBio's pipeline, including potential best-in-class candidates

	Indication	Drug Mechanism	Pt. pop. (US+EU)	Discovery	Pre-IND	Phase 1	Phase 2	Phase 3	Approved	Partner
Mendelian	MoCD type A	NULIBRY™ (Synthetic cPMP, fosdenopterin)	100							MEDISON
	Achondroplasia	Low-dose FGFRi (infigratinib)	55k							
	LGMD2i	Glycosylation substrate (ribitol)	7k							
	RDEB	Recombinant COL7 (BBP-589)	2k							
	PKAN / organic acidemia	Pank activator (BBP-671)	7k							
	VM / LM	Topical PI3K inhibitor (BBP-681)	117k							
	Netherton	Topical KLK inhibitor (BBP-561)	11k							
	PTEN autism	PI3Kb inhibitor (BBP-472)	120k							
	4 undisclosed small molecule programs		>500k							
	4 undisclosed antisense oligonucleotide programs		>300k							
Precision Cardiorenal	ATTR amyloidosis	TTR stabilizer (acoramidis)	>400k							AstraZeneca
	ADH1	CaSR antagonist (encaleret)	12k ¹							
	PH1 / frequent stone formers	GO1 inhibitor (BBP-711)	5k / 1.5m							
	Undisclosed DCM small molecule program		>250k							
	Undisclosed DCM AAV gene therapy program		>250k							
Precision Oncology	FGFR2+ cholangiocarcinoma (2L)	TRUSELTIQ™ (FGFRi, infigratinib)	4k							
	FGFR2+ cholangiocarcinoma (1L)	FGFRi (infigratinib)								HELFINN
	FGFR3+ adjuvant urothelial	FGFRi (infigratinib)	21k							
	FGFR1-3+ tumor agnostic	FGFRi (infigratinib)	24k							
	FGFR1-3+ gastric cancer	FGFRi (infigratinib)	41k ²							
	MAPK / RAS-driven cancer	SHP2i monotherapy (BBP-398)	>500k							
		SHP2i combo therapy (BBP-398)								
	KRAS-driven cancer	KRAS G12C dual inhibitor	>500k							
	PI3Kα:RAS Breaker									
	KRAS G12Di	>500k								
Solid tumors	GPX4i	>500k								
Gene Therapy	CAH	AAV5 gene therapy (BBP-631)	>75k							
	Canavan	AAV9 gene therapy (BBP-812)	1k							
	TMC1 hearing loss	AAV gene therapy (BBP-815)	2k							
	Galactosemia	AAV gene therapy (BBP-818)	>7k							
	TSC1/2	AAV gene therapy	>100k							
	Cystinuria	AAV gene therapy	20k							
	3 capsid discovery collaborations									

¹US carriers

²China + Japan patient population

BridgeBio's endless summer



MoCD Type A

2L CCA

ATTR-CM/PN

ADH1

Achon

CAH

PKAN/OA

PH1/FSF, VM

SH2P2, UC, RDEB

LGMD2i, Canavan

KRAS

ALS, Autism

CF, A1AT, GALT

TMC1, TSC1/2

Presentations to come



- **Precision Cardiorenal:**
 - **Introduction – Cameron Turtle**
 - **Acoramidis, a TTR stabilizer, for ATTR – Jonathan Fox**
 - **Encaleret, a CaSRi, for ADH1 – Mary Scott Roberts**
- **Gene therapy platform – Eric David**
- **Wave 3 Mendelian programs – Uma Sinha**
- **Precision Oncology – Eli Wallace**
- **BridgeBioX – Charles Homcy**

Precision cardiorenal

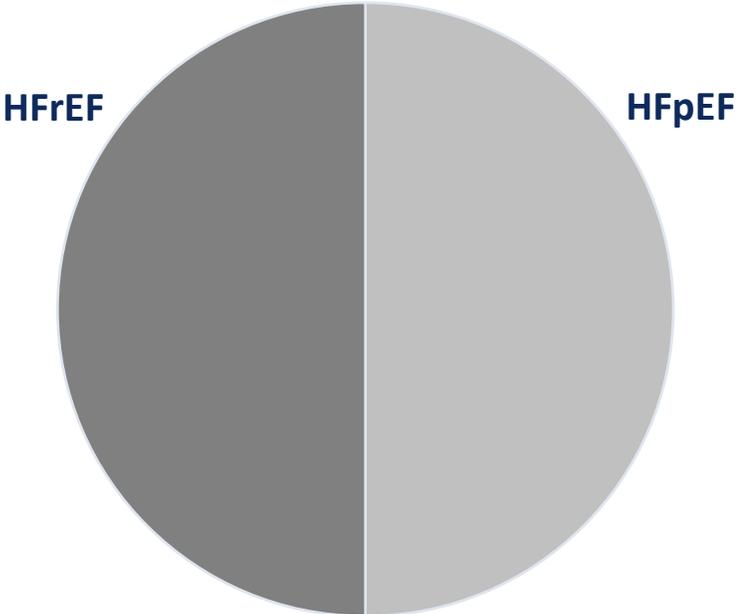
Cameron Turtle, D. Phil.

Chief Strategy Officer

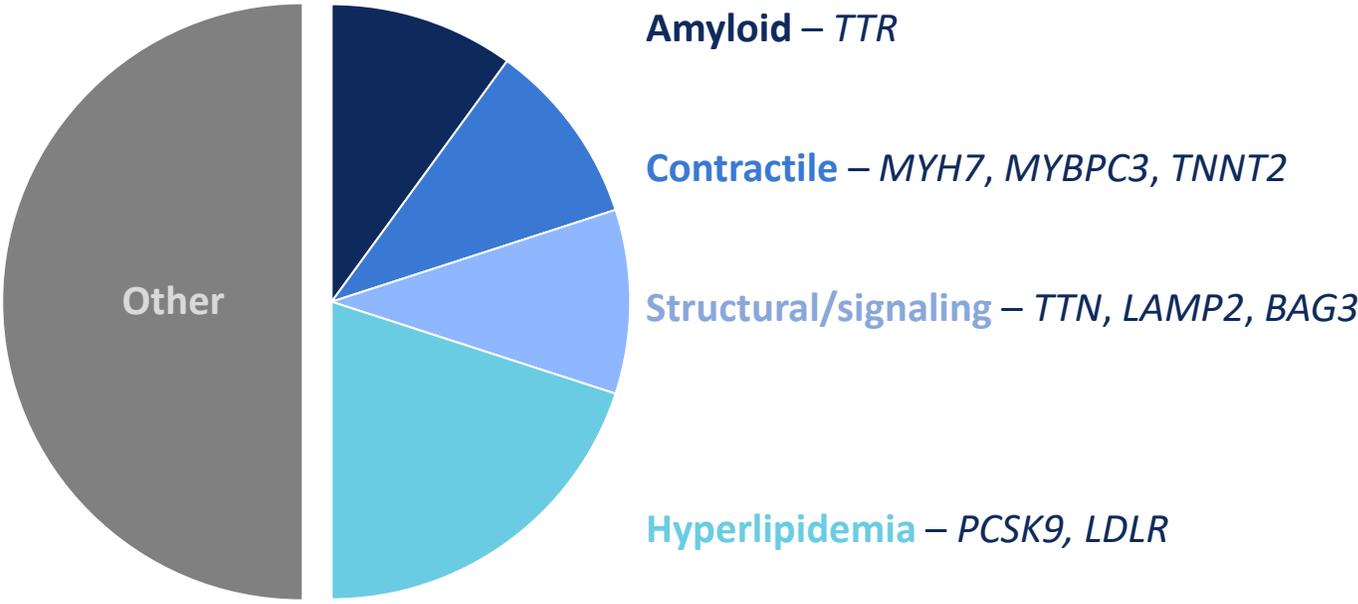


Genetic drivers of cardiac disease are unlocking precision medicine targets

Diagnosis by phenotype



Diagnosis by mechanism – genetic associations

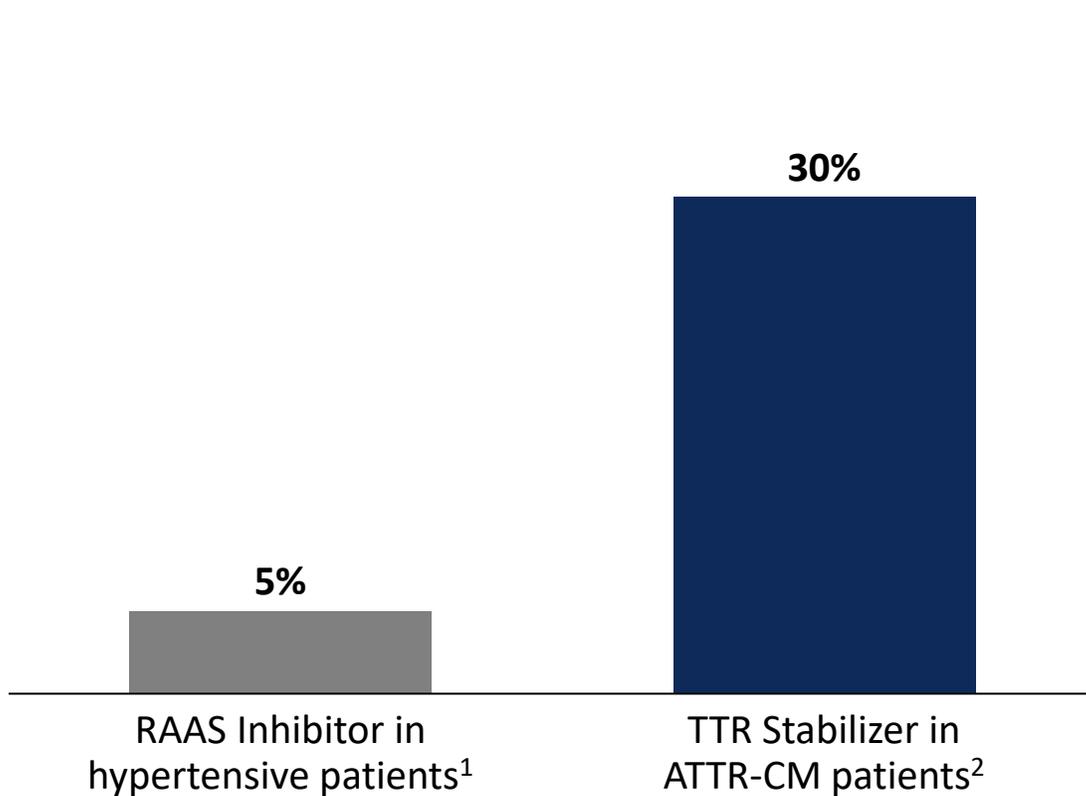


HFrEF = heart failure reduced ejection failure; HFpEF = heart failure preserved ejection failure
Note: Proportions not to scale, genetic associations not exhaustive

Precision medicines have delivered increased treatment effect sizes

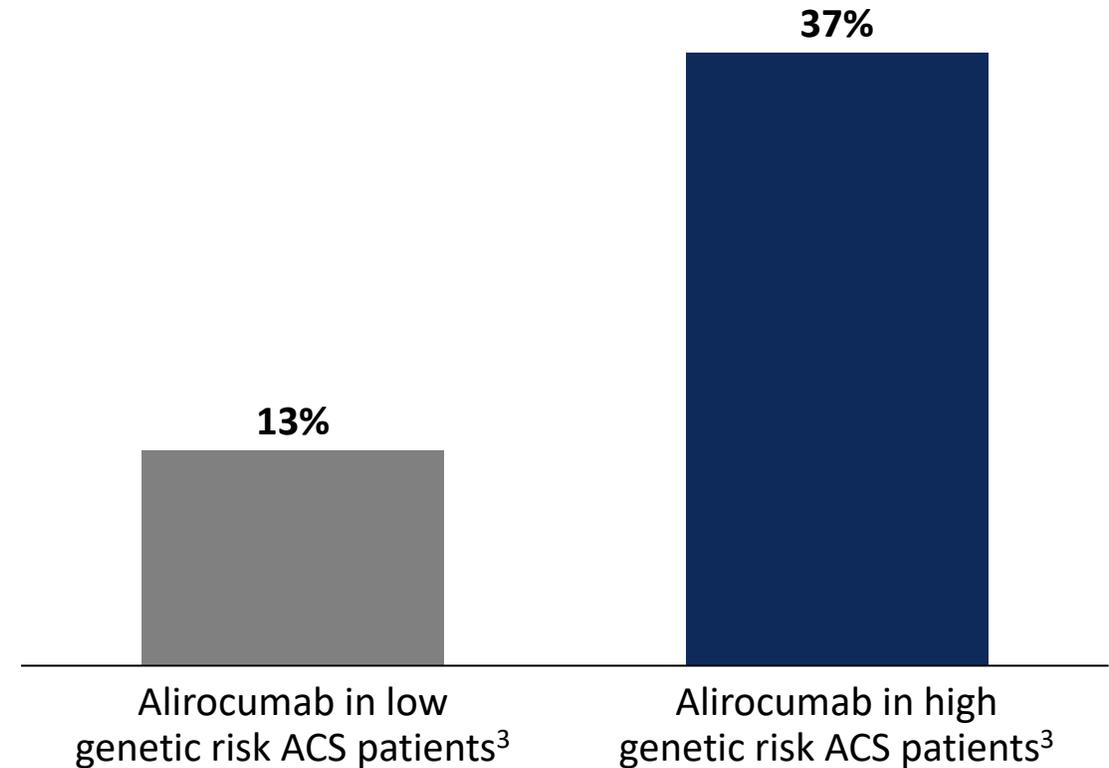
Mendelian disease

% reduction in mortality



High genetic risk in broader diseases

% reduction in mortality



Cardiorenal pipeline overview

	Indication	Drug Mechanism	Pt. pop. (US+EU)	Discovery	Pre-IND	Phase 1	Phase 2	Phase 3
Precision Cardiorenal	ATTR amyloidosis	TTR stabilizer (acoramidis)	>400k	█	█	█	█	█
	ADH1	CaSR antagonist (encaleret)	12k ¹	█	█	█	█	
	PH1 / frequent stone formers	GO1 inhibitor (BBP-711)	5k / 1.5m	█	█	█		
	Undisclosed DCM small molecule program		>250k	█				
	Undisclosed DCM AAV gene therapy program			█				

█ Featured Programs

¹US carriers

Acoramidis: TTR stabilizer for ATTR

Jonathan Fox, M.D., Ph.D.

Chief Medical Officer, Cardiorenal



Acoramidis for transthyretin (TTR) amyloidosis (ATTR)



Len
Living with ATTR-CM

Prevalence
400k+
Worldwide

Pathophysiology
Systemic disease most commonly presenting as cardiomyopathy or peripheral neuropathy

Genetic Driver

Destabilized TTR leading to amyloid accumulation

Therapeutic Hypothesis

TTR stabilizer designed to mimic protective T119M mutation

Design Criteria for Optimal Therapy

-  Near-complete stabilization of TTR
-  Preservation of TTR tetramer
-  Oral Dosing

ATTR is a systemic disease with multiple manifestations

Clinical Presentation

Diagnostic Approach

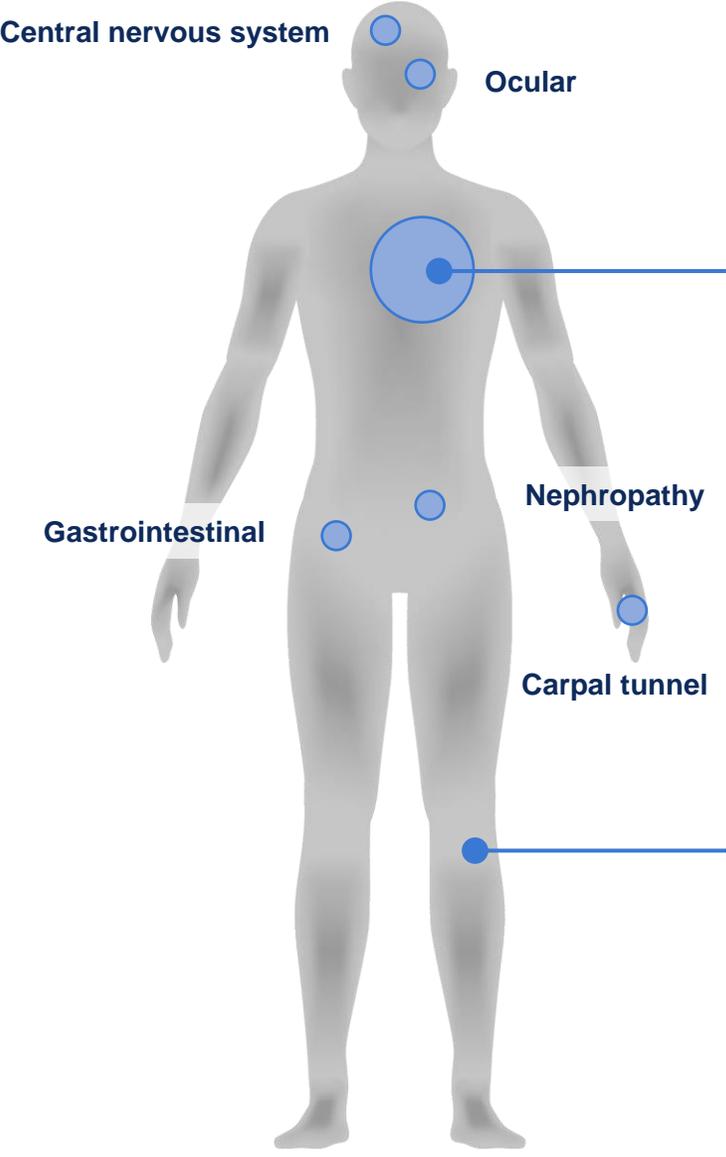
Therapeutic Approach

Published Data

Pivotal Trial Design

6MWD Importance

Upcoming Milestones



Cardiomyopathy (ATTRwt-CM or ATTRv-CM)

- Deposition of wild-type or variant (e.g., V122I) TTR amyloid in the heart, leading to predominantly diastolic heart failure
- Likely affects 400K+ worldwide, majority undiagnosed
- Late onset (age 50+), progressive and fatal with median survival of 3-5 years from diagnosis

Polyneuropathy (ATTR-PN)

- Affects ~10K worldwide, primarily in EU and Japan
- Exclusively caused by variant TTR (e.g., V30M)
- Onset between ages of 30 and 50, progressive and fatal with median survival of 5-10 years from diagnosis

Source: Grogan, M et al. JACC 2016, 68:1014-20; Planté-Bordeneuve, V. and Said, G. Lancet Neurol 2011, 10:1086-97

Rapid increase in patient finding driven by non-invasive diagnosis techniques

Clinical Presentation

Diagnostic Approach

Therapeutic Approach

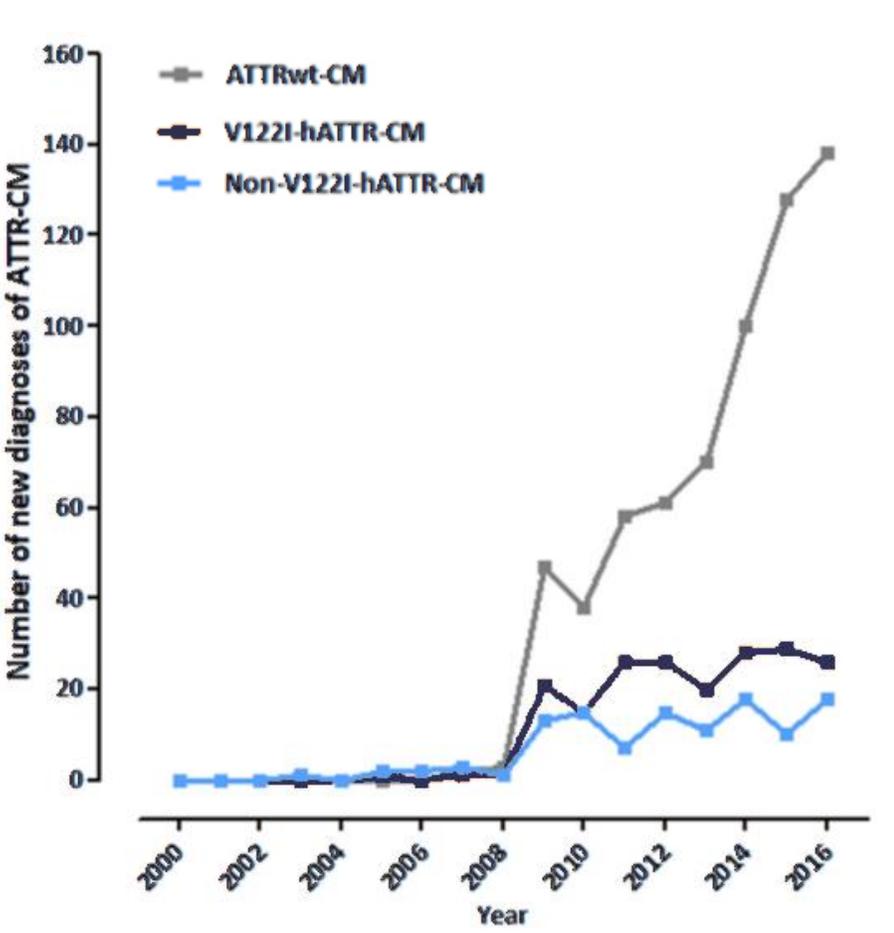
Published Data

Pivotal Trial Design

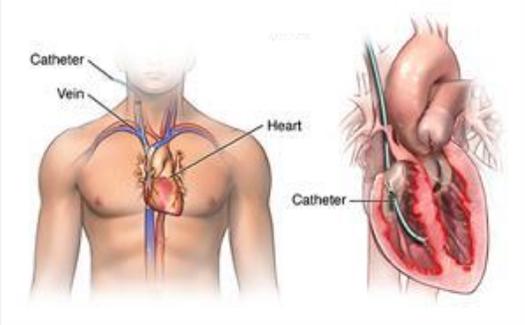
6MWD Importance

Upcoming Milestones

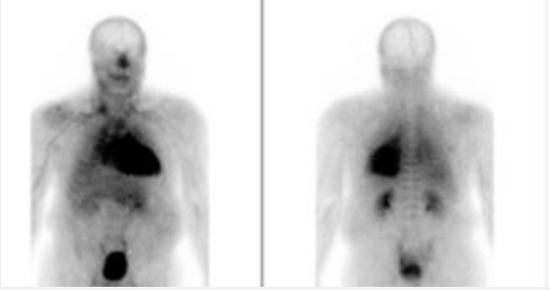
New ATTR-CM diagnoses per year at single site



Invasive heart biopsies



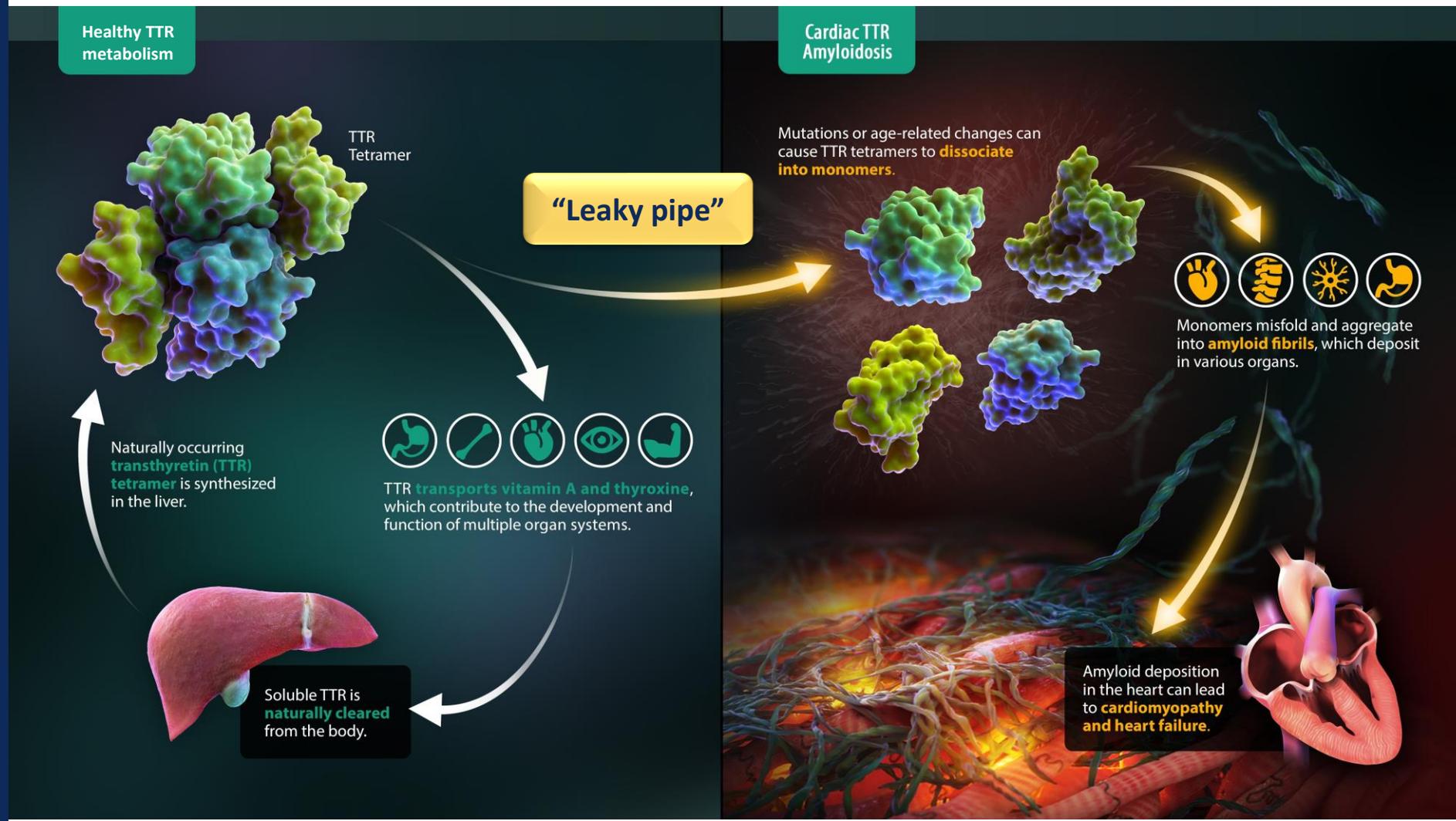
Non-invasive technetium scans



Source: Lane et al., 2019; Johns Hopkins Medicine; American College of Cardiology

TTR plays a physiological role in the body

- Clinical Presentation
- Diagnostic Approach
- Therapeutic Approach
- Published Data
- Pivotal Trial Design
- 6MWD Importance
- Upcoming Milestones



Targeting diseases at their source optimizes safety and efficacy

Clinical Presentation

Diagnostic Approach

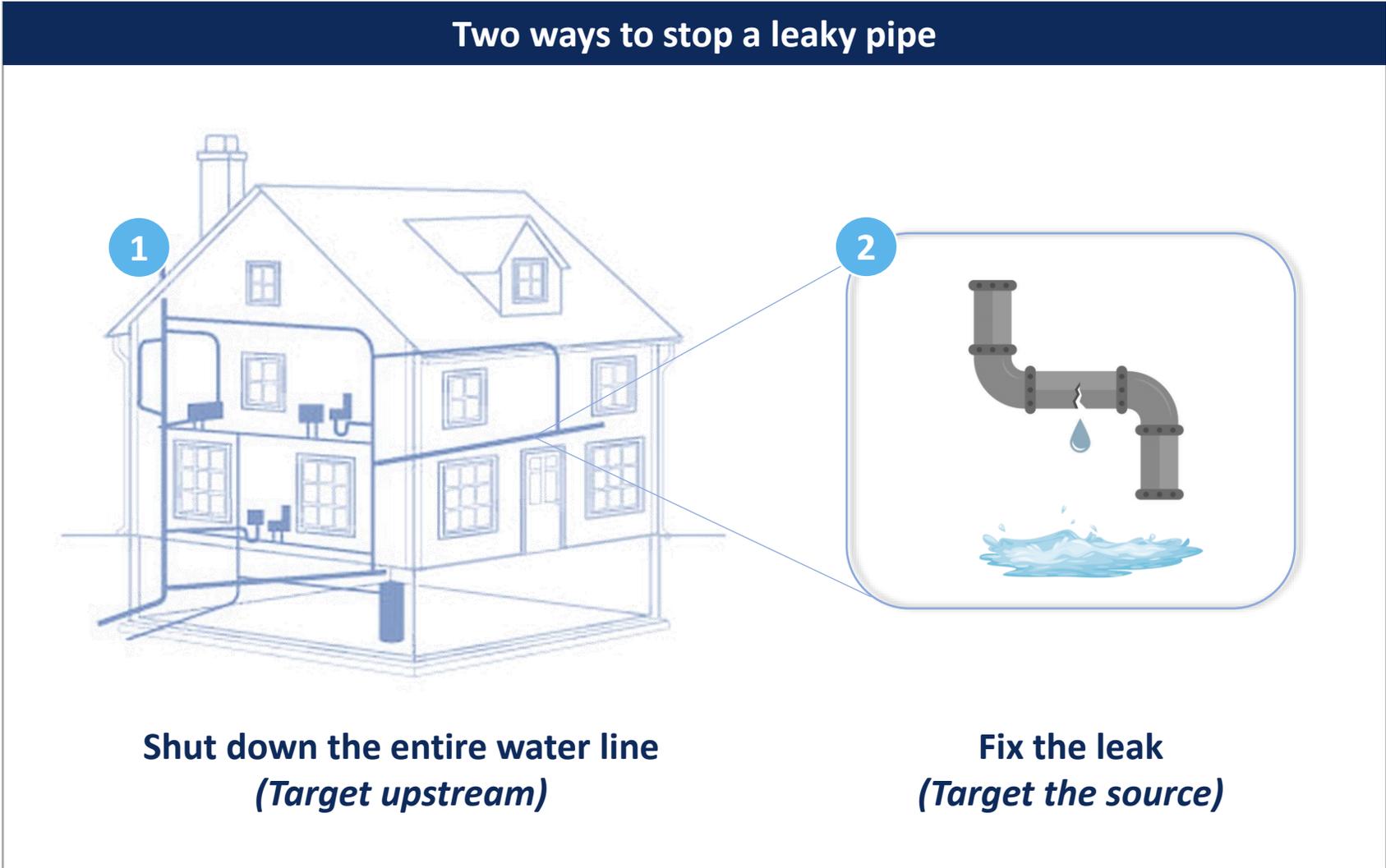
Therapeutic Approach

Published Data

Pivotal Trial Design

6MWD Importance

Upcoming Milestones



Acoramidis was designed to treat ATTR at its source

Clinical Presentation

Diagnostic Approach

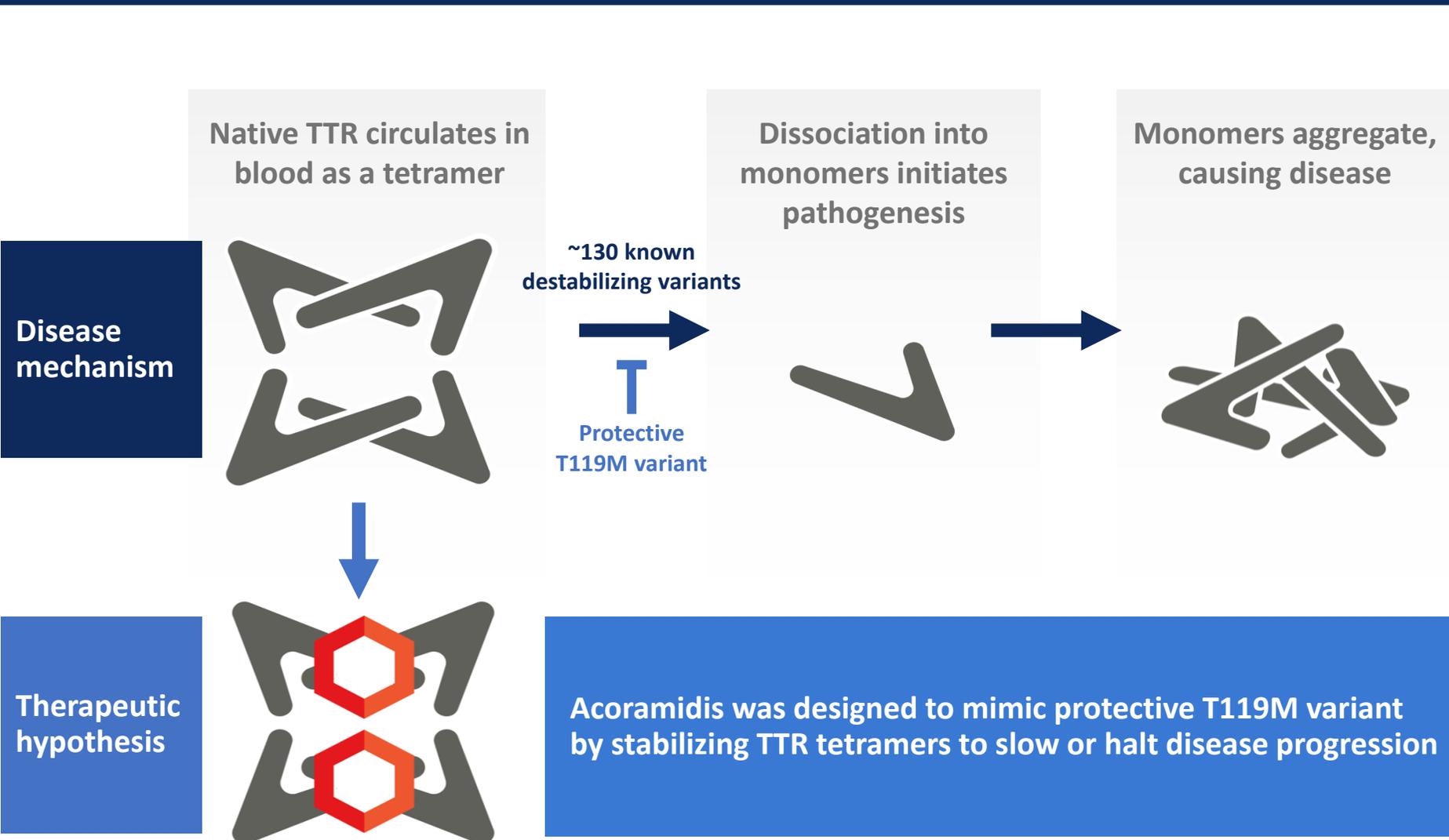
Therapeutic Approach

Published Data

Pivotal Trial Design

6MWD Importance

Upcoming Milestones



Human genetics suggest TTR instability is associated with disease severity

Clinical Presentation

Diagnostic Approach

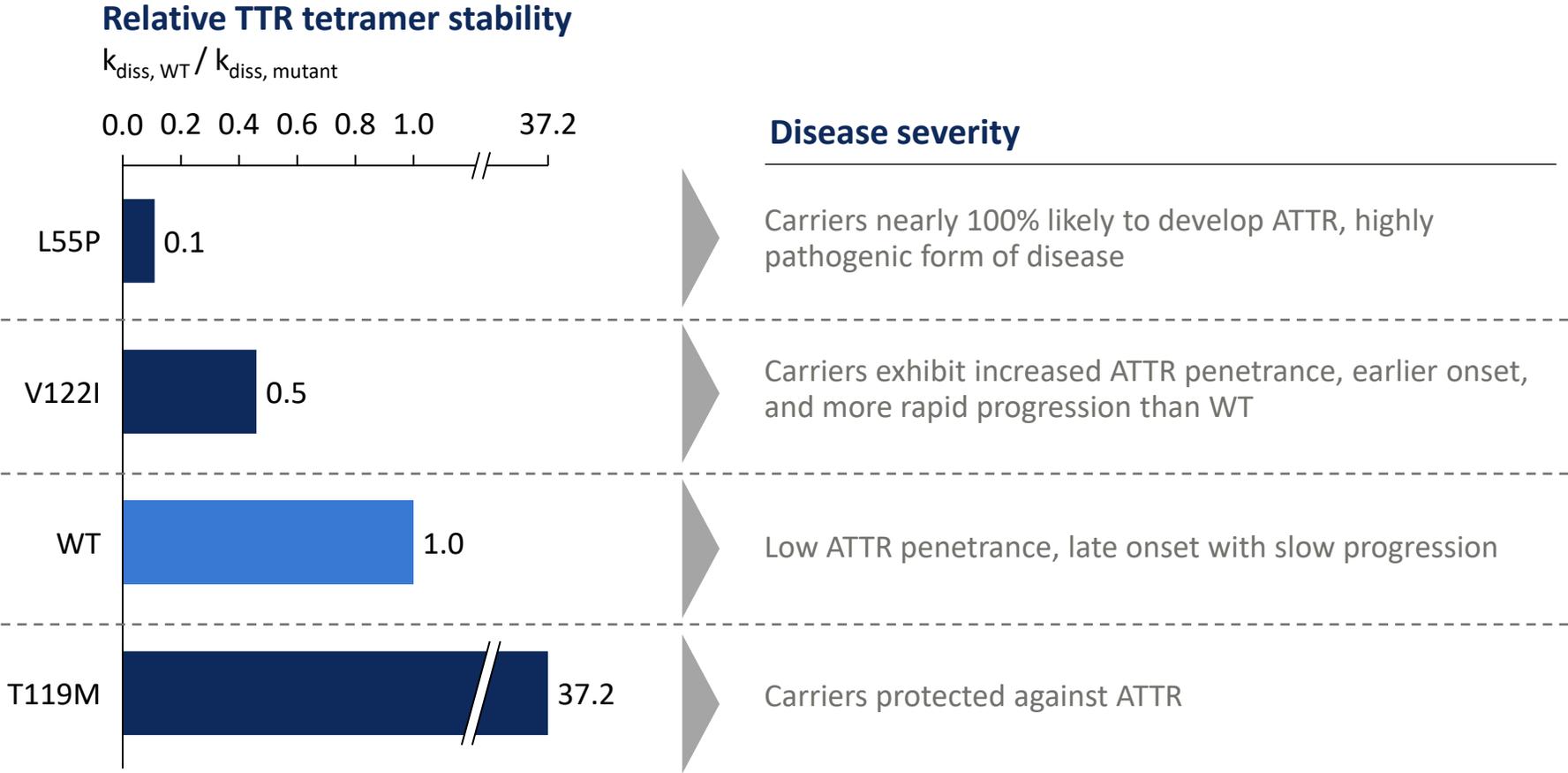
Therapeutic Approach

Published Data

Pivotal Trial Design

6MWD Importance

Upcoming Milestones



- Greater TTR destabilization correlates with earlier disease onset, increased disease severity
- ATTR-protective mutations stabilize TTR tetramer, preventing dissociation

k_{diss} = dissociation constant
Source: Hammarstrom, P. et al. PNAS 2002, 99:16427-16432

Higher dose of tafamidis increased stabilization and improved clinical benefit

Clinical Presentation

Diagnostic Approach

Therapeutic Approach

Published Data

Pivotal Trial Design

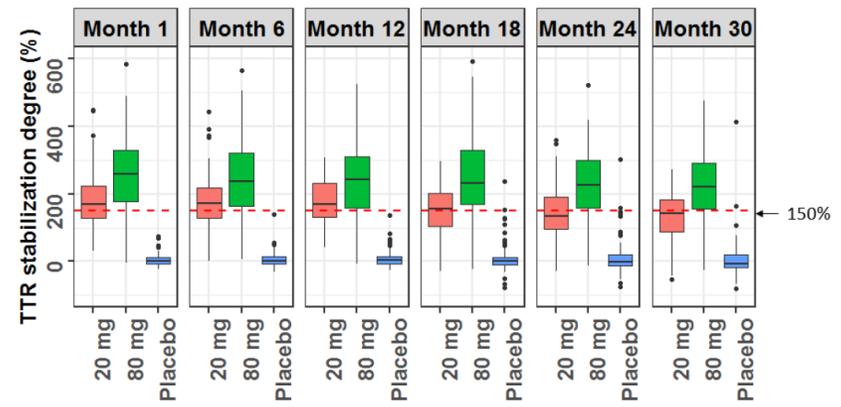
6MWD Importance

Upcoming Milestones

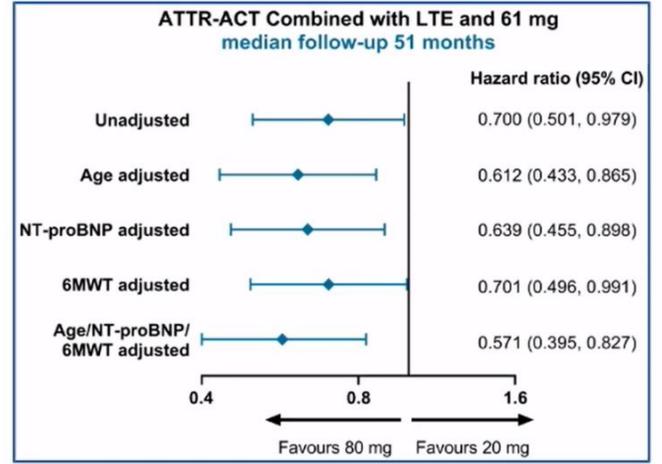
Phase 3 ATTR-ACT study tested two doses of tafamidis (20 mg & 80 mg) vs. placebo

- Participants receiving 80 mg of tafamidis (vs. 20 mg) exhibited greater TTR stabilization¹
- Benefit of tafamidis 80 mg vs. 20 mg was evident on all-cause mortality in analysis of ATTR-ACT combined with long-term extension (LTE)²

TTR stabilization¹



All-cause mortality²



¹FDA CDER Clinical Pharmacology and Biopharmaceutics, Clinical Review (Vyndaqel/Vyndamax), 2019
²Damy T, et al. Eur J Heart Fail. 2020. "Efficacy and safety of tafamidis doses in the Tafamidis in Transthyretin Cardiomyopathy Clinical Trial (ATTR-ACT) and long-term extension study"

Acoramidis demonstrates near-complete stabilization of TTR

Clinical Presentation

Diagnostic Approach

Therapeutic Approach

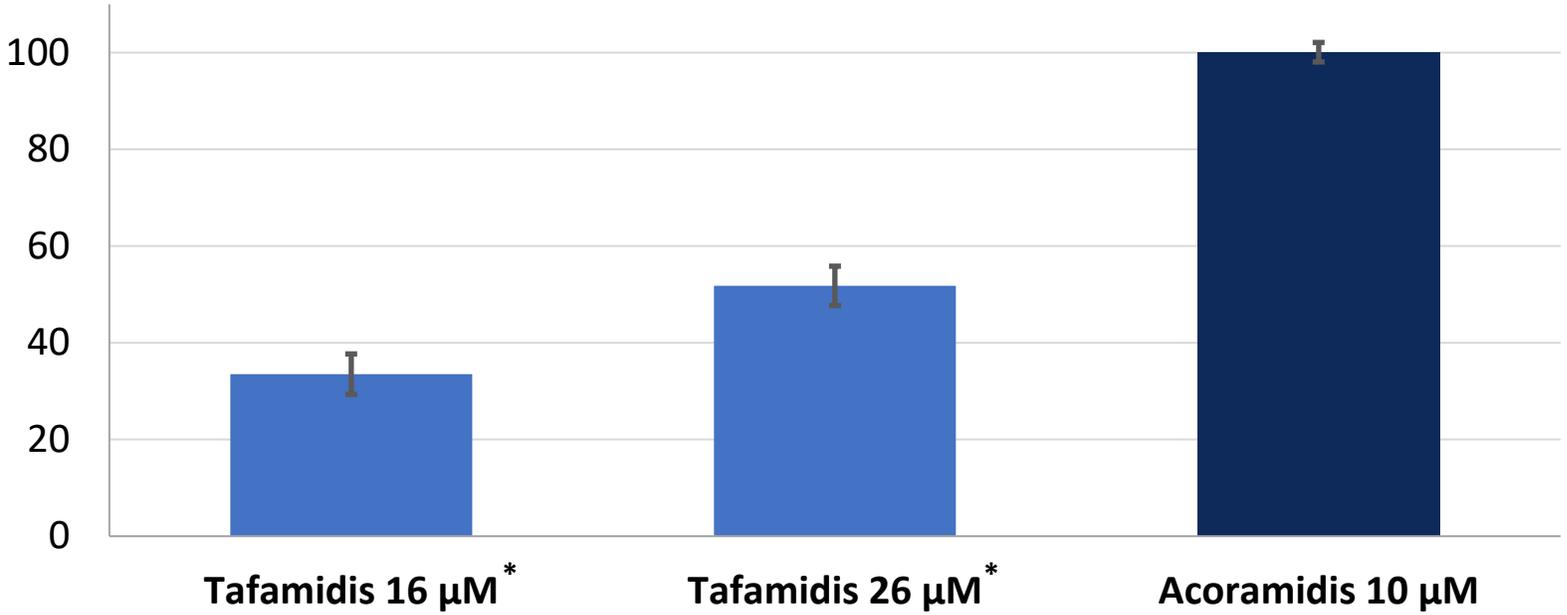
Published Data

Pivotal Trial Design

6MWD Importance

Upcoming Milestones

TTR target site occupancy by FPE assay¹
%, mean +/- SD



Acoramidis demonstrated near-complete TTR stabilization in vitro at clinical concentrations

Source: Ji, A.X., et al. Differential Transthyretin Binding, Kinetic Stability and Additive Ex Vivo Stabilization by AG10 Compared to Tafamidis. American Heart Association Scientific Sessions 2019
¹FPE characterization of TTR binding site occupancy in serum incubated with stabilizer, n = 12
*Tafamidis 80 mg mean C_{min} and C_{max} per FDA review of tafamidis meglumine ATTR-CM NDA

Phase 2 ATTR-CM trial provided randomized 28-day and 15-month OLE data

Clinical Presentation

Diagnostic Approach

Therapeutic Approach

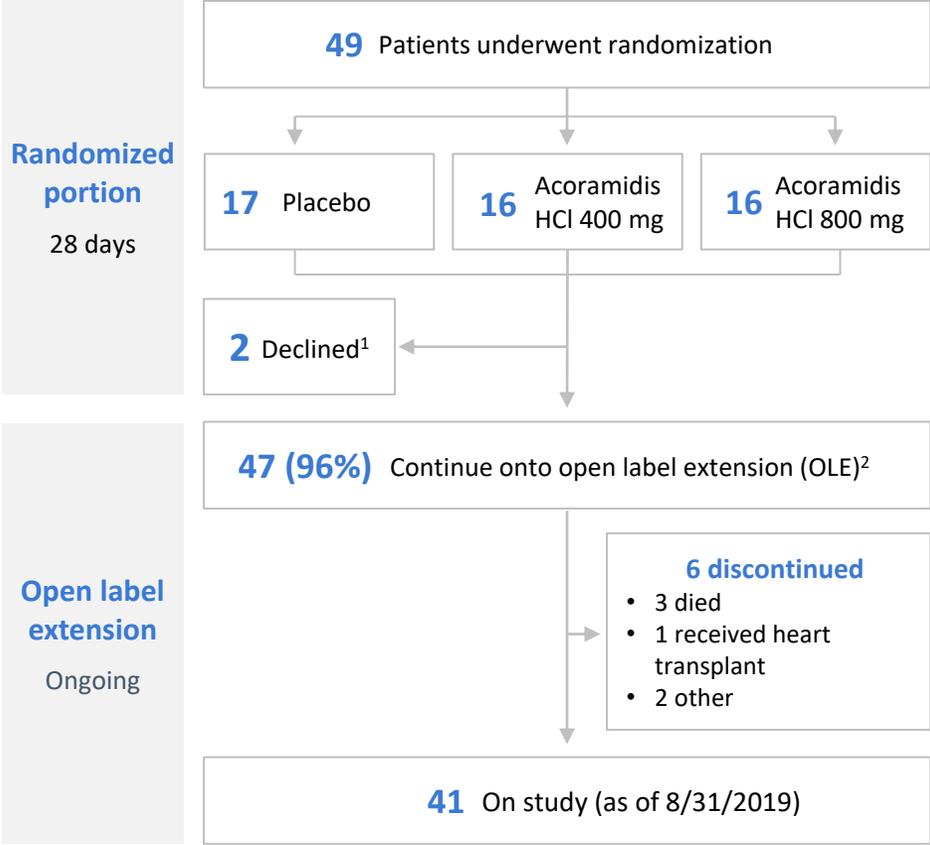
Published Data

Pivotal Trial Design

6MWD Importance

Upcoming Milestones

Schematic of acoramidis Phase 2 studies⁴



Outcomes

Primary

Safety and tolerability

- Adverse events
- Clinical events and vital signs
- Clinical laboratory parameters

Secondary and Exploratory

- Pharmacokinetics
- Pharmacodynamics
- Echocardiographic parameters

Publications to date:

- Randomized 28-day study³
- Open label extension analysis as of 8/31/2019, median 65 weeks from acoramidis 201 initiation (randomized), median 53 weeks on acoramidis⁴

¹Both declined participation due to geographical constraints regarding study visits
²Median rollover period of 72 days (range 41-152 days)
³Judge, D.P. et al. JACC Vol. 74, No. 3, 2019:285 – 95
⁴Judge, D.P. et al. American Heart Association 2019

Acoramidis has been well-tolerated and demonstrated near-complete TTR stabilization in preclinical, Ph 1, and Ph 2 studies

Clinical Presentation

Diagnostic Approach

Therapeutic Approach

Published Data

Pivotal Trial Design

6MWD Importance

Upcoming Milestones

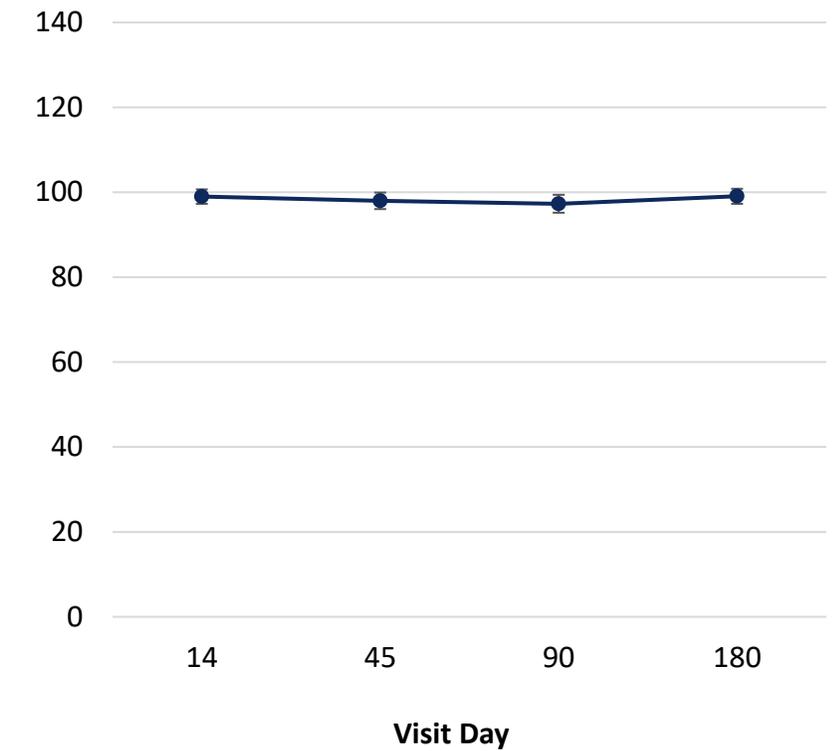
Phase 2 safety summary¹

Randomized portion	Placebo N = 17	Acoramidis (pooled doses) N = 32
Any Adverse Event	15 (88%)	21 (66%)
Mild	6 (35%)	11 (34%)
Moderate	8 (47%)	9 (28%)
Severe	1 (6%)	1 (3%)
Any SAE	2 (12%)	1 (3%)
AF and CHF	1 (6%) ¹	0
Leg cellulitis	1 (6%)	0
Dyspnea	0	1 (3%)

In long-term OLE, acoramidis was generally well tolerated with a pattern of adverse events consistent with underlying disease severity, concurrent illnesses, and age of participants

Phase 2 OLE TTR stabilization²

TTR stabilization at steady-state trough level
%, mean ± SEM

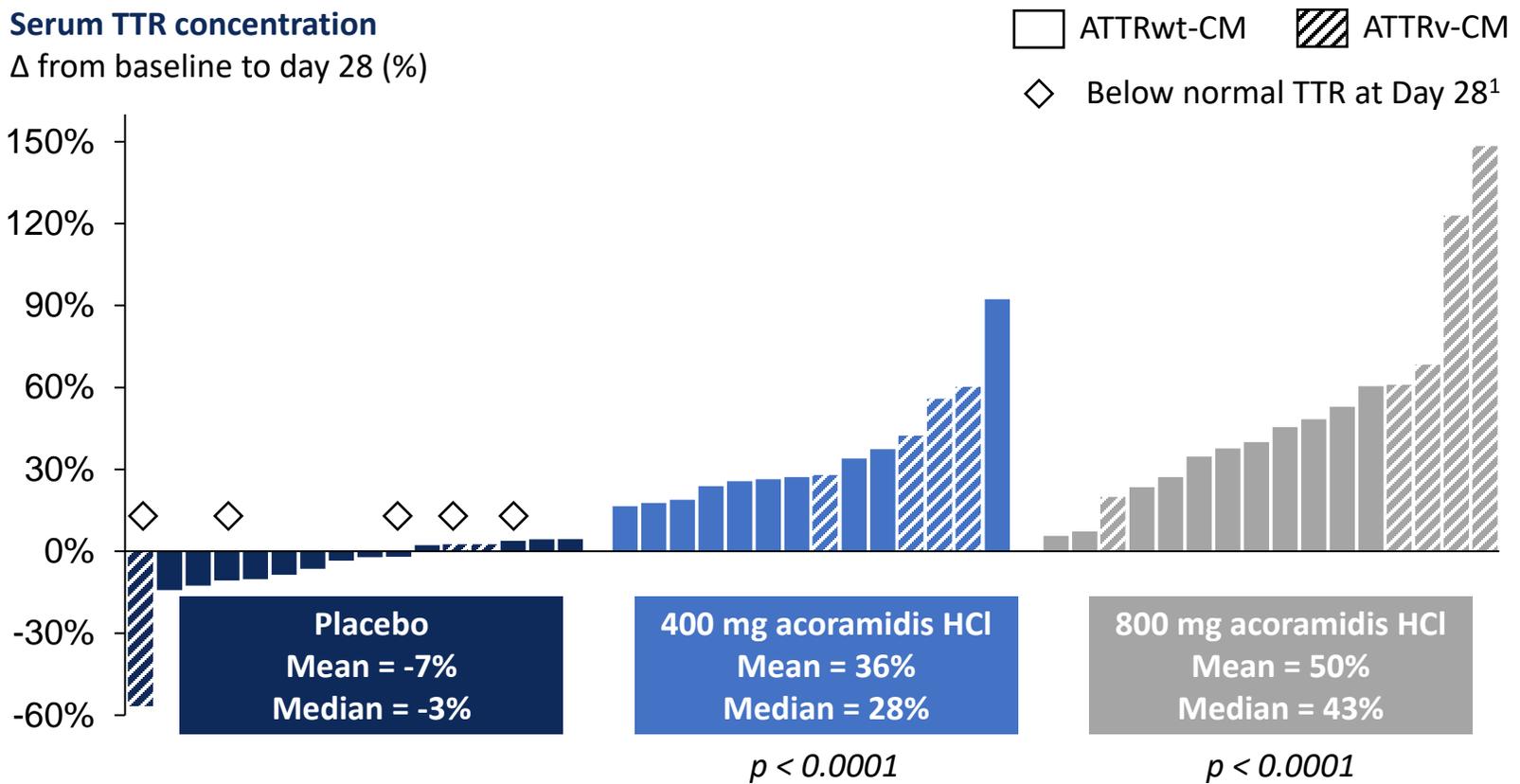


¹Judge, D.P. et al. JACC Vol. 74, No. 3, 2019:285 – 95

²Judge, D.P. et al. American Heart Association 2019

Acoramidis increased serum TTR concentrations in a dose-dependent manner

- Clinical Presentation
- Diagnostic Approach
- Therapeutic Approach
- Published Data
- Pivotal Trial Design
- 6MWD Importance
- Upcoming Milestones



- Dose-dependent increase in serum TTR concentrations in acoramidis-treated subjects
- Greater effect observed in ATTRv subjects – final TTR concentrations normalized to comparable levels as in treated ATTRwt subjects

Note: Serum TTR concentrations not available at baseline for one 400 mg subject and at Day 28 for one 400 mg and one placebo subject
 Source: Judge, D.P. et al. J Am Coll Cardiol. 2019;74(3):285-295.
¹Normal reference range for serum TTR 20-40 mg/dL (3.6-7.3 μ M)

NT-proBNP and TnI remained stable in acoramidis-treated participants throughout OLE

Clinical Presentation

Diagnostic Approach

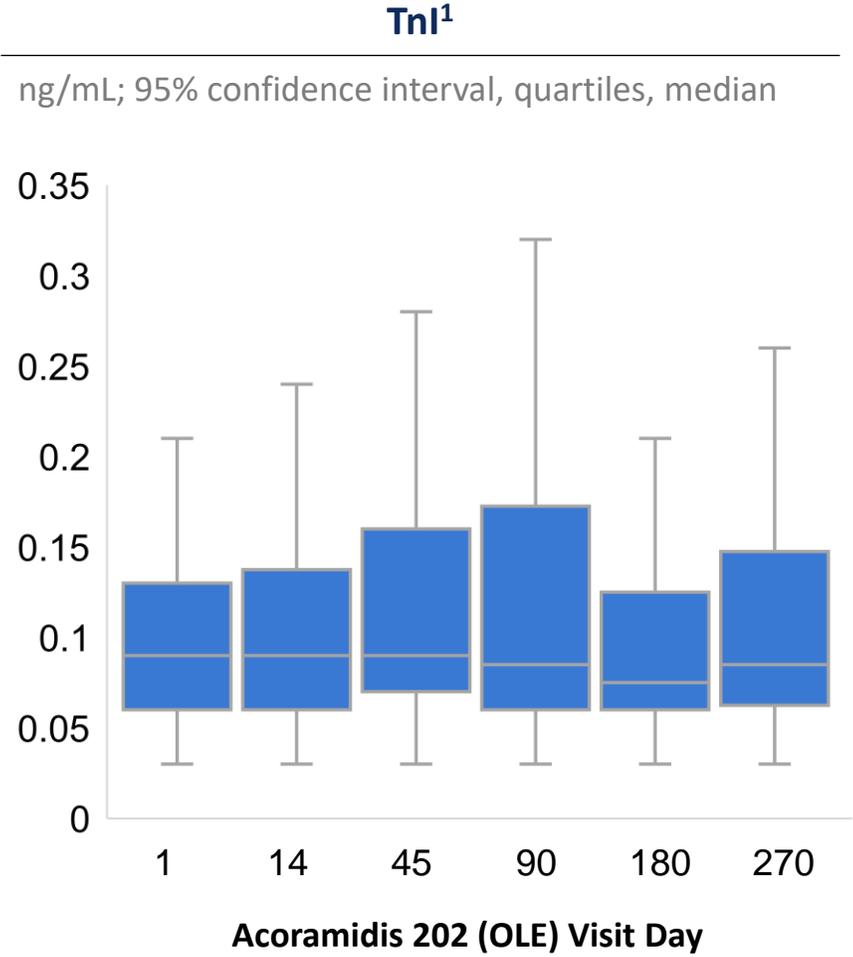
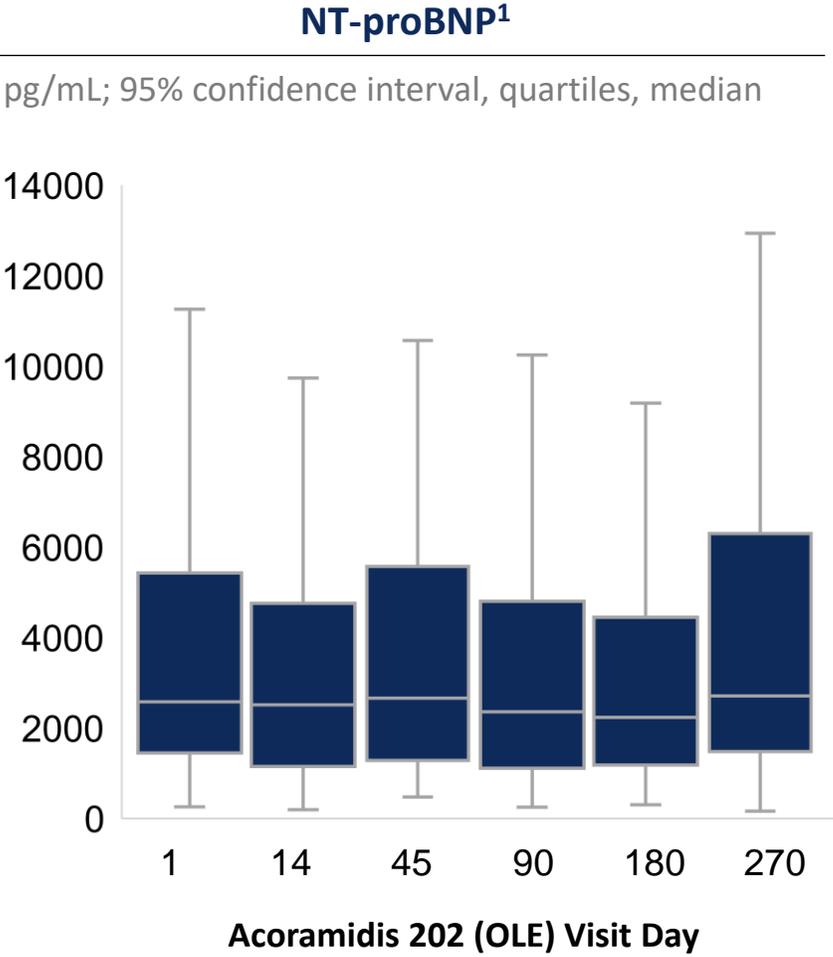
Therapeutic Approach

Published Data

Pivotal Trial Design

6MWD Importance

Upcoming Milestones



¹Judge, D.P. et al. American Heart Association 2019. NT-proBNP = N-Terminal pro B-type Natriuretic Peptide. TnI = troponin I

Echocardiography parameters remained stable in acoramidis-treated participants throughout OLE

Clinical Presentation

Diagnostic Approach

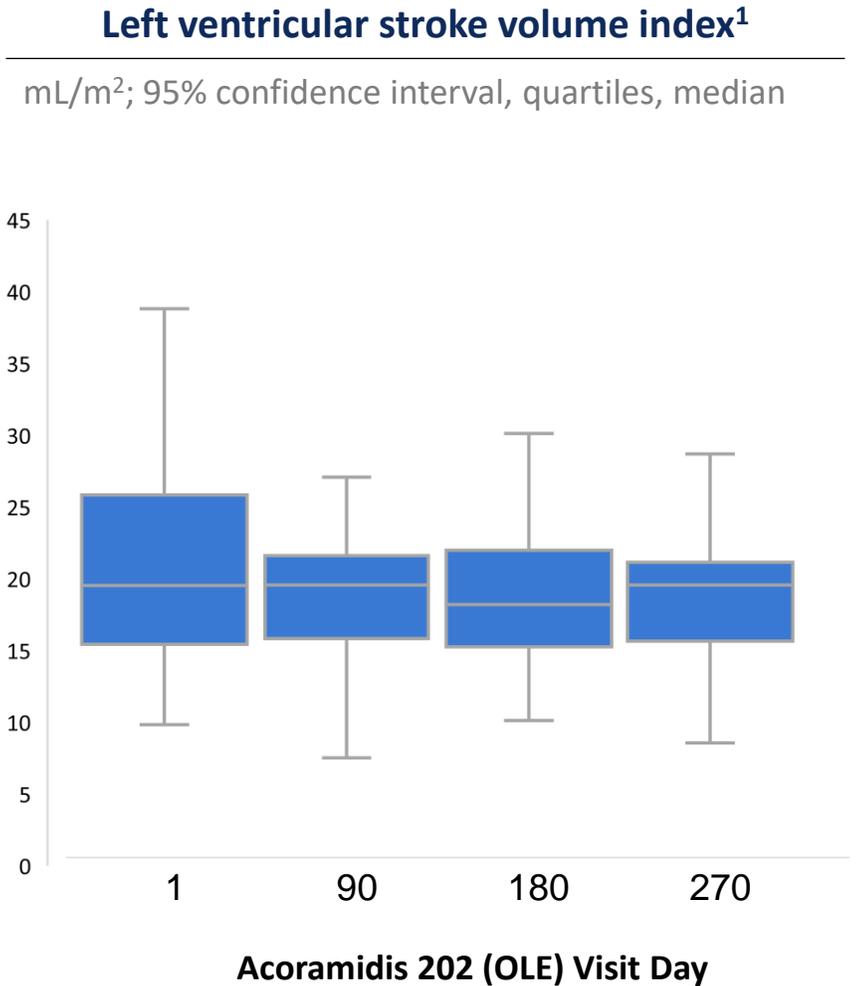
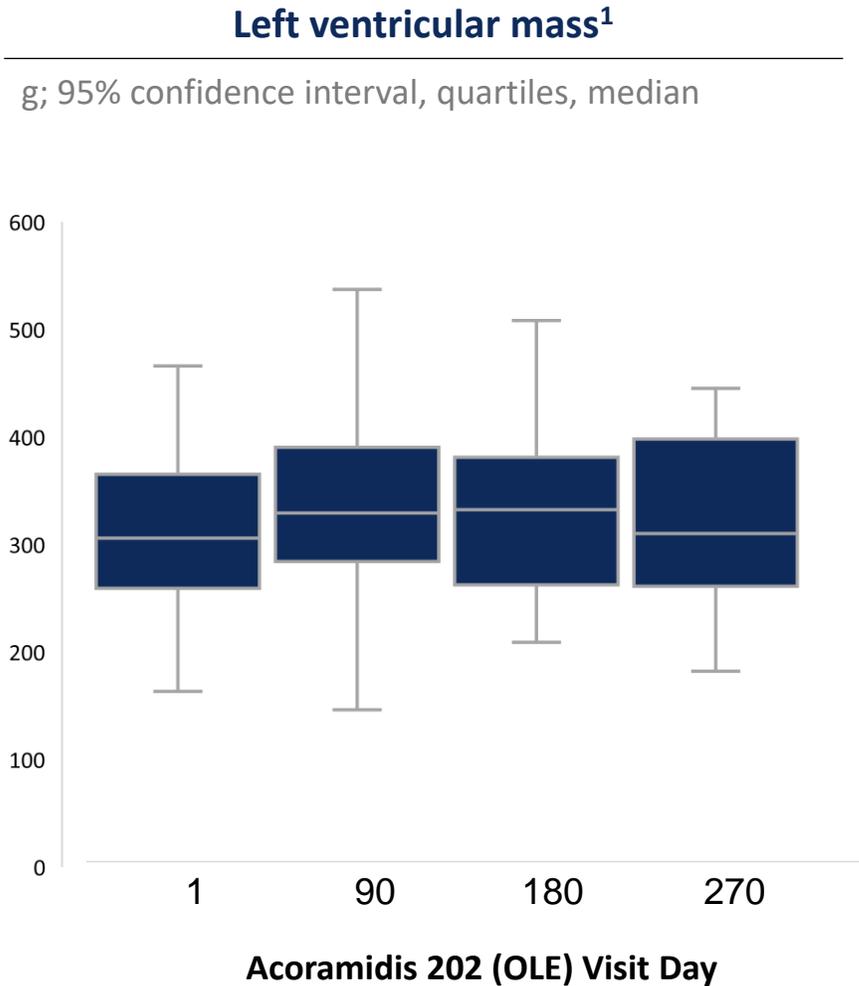
Therapeutic Approach

Published Data

Pivotal Trial Design

6MWD Importance

Upcoming Milestones



¹Judge, D.P. et al. American Heart Association 2019

Embedded Ph3 design includes 12-month and 30-month primary endpoints

Clinical Presentation

Diagnostic Approach

Therapeutic Approach

Published Data

Pivotal Trial Design

6MWD Importance

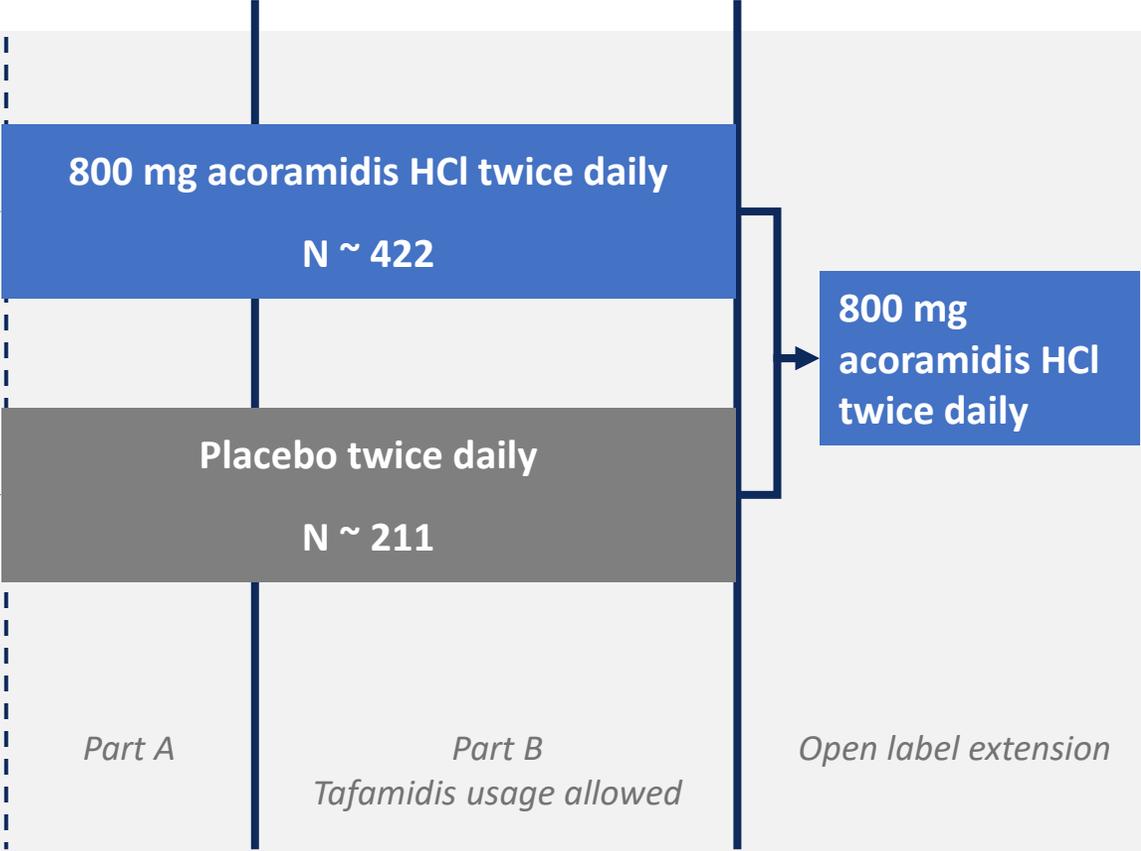
Upcoming Milestones



- Subjects with diagnosed ATTR-CM (WT or variant)
- NYHA Class I-III
- ATTR-positive biopsy or ^{99m}Tc scan
- Light chain amyloidosis excluded if diagnosis by ^{99m}Tc

12-month primary endpoint:
Change in 6MWD

30-month primary endpoint:
Hierarchical composite



Secondary endpoints include: Kansas City Cardiomyopathy Questionnaire, serum TTR, TTR stabilization
 6MWD = Six minute walk distance; NYHA = New York Heart Association;
^{99m}Tc = Technetium labeled pyrophosphate (PYP) or bisphosphonate (e.g., DPD); dx = diagnosis;
 CV hosp = cardiovascular-related hospitalizations

6-Minute Walk Test is a clinically meaningful, treatment-responsive endpoint

Clinical Presentation

Diagnostic Approach

Therapeutic Approach

Published Data

Pivotal Trial Design

6MWD Importance

Upcoming Milestones



Simple, sub-maximal exercise test to assess aerobic capacity and endurance



Demonstrated to measure treatment benefit in heart failure, COPD, and pulmonary arterial hypertension



Higher rates of mortality observed with lower 6MWD in multiple cardiopulmonary diseases^{1,2}

“The 6-minute walk test (6MWT), a measure of functional capacity, was identified as a **predictor of overall survival** in patients with ATTR-CM.”
- Maurer et al., 2020

1 Ingle, L. et al., Biomed Res Int 2014
2 Lane, T. et al., Circulation 2019

Rapid functional decline in untreated ATTR-CM patients provides opportunity to demonstrate robust clinical benefit

Clinical Presentation

Diagnostic Approach

Therapeutic Approach

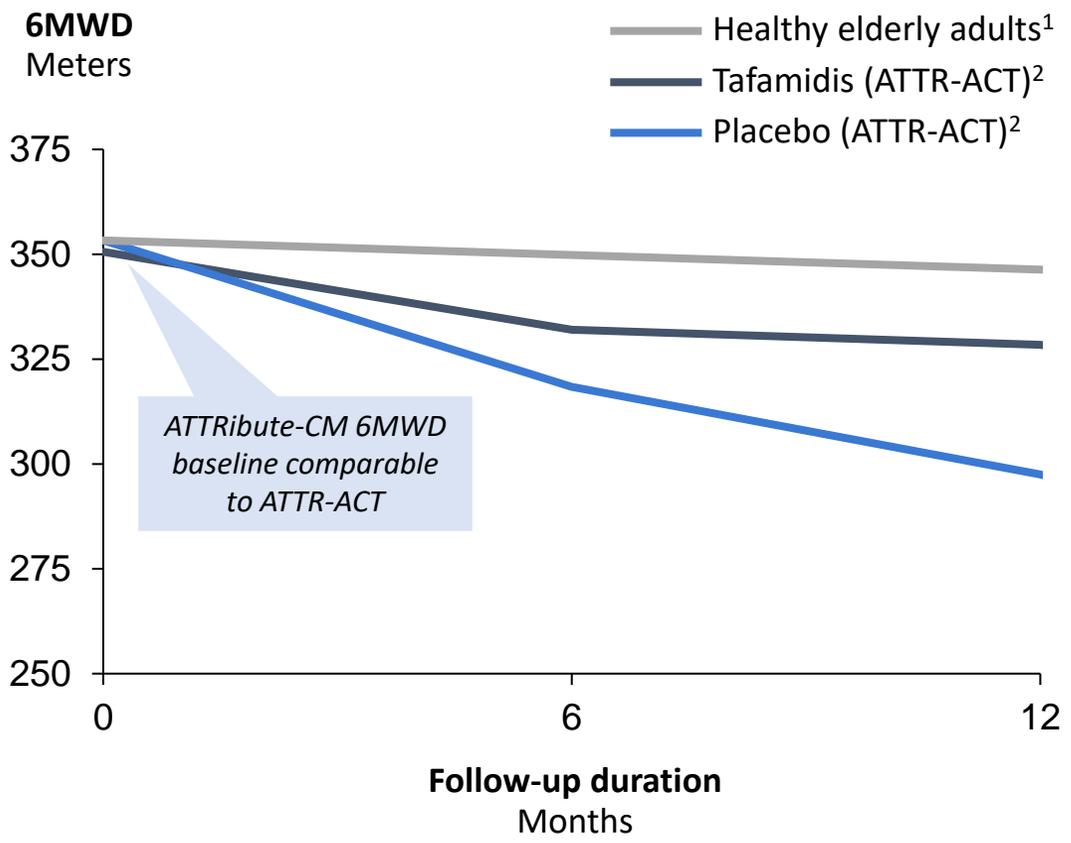
Published Data

Pivotal Trial Design

6MWD Importance

Upcoming Milestones

Summary of 6MWD data in ATTR and healthy cohorts



12-month decline:

- Healthy elderly adult: -7m
- ATTR-ACT (tafamidis): -25m

Hypothesis: near-complete stabilization of TTR by acoramidis may slow or halt functional decline in 6MWD

¹Enright, P.L. et al. Chest 2003. N = 3333 healthy elderly adults, baseline set to match ATTR-ACT placebo group
²Maurer, M.S. et al. NEJM 2018. N = 264 (tafamidis), N = 177 (placebo) ATTR-CM trial participants

Timeline of upcoming milestones

Clinical Presentation

Diagnostic Approach

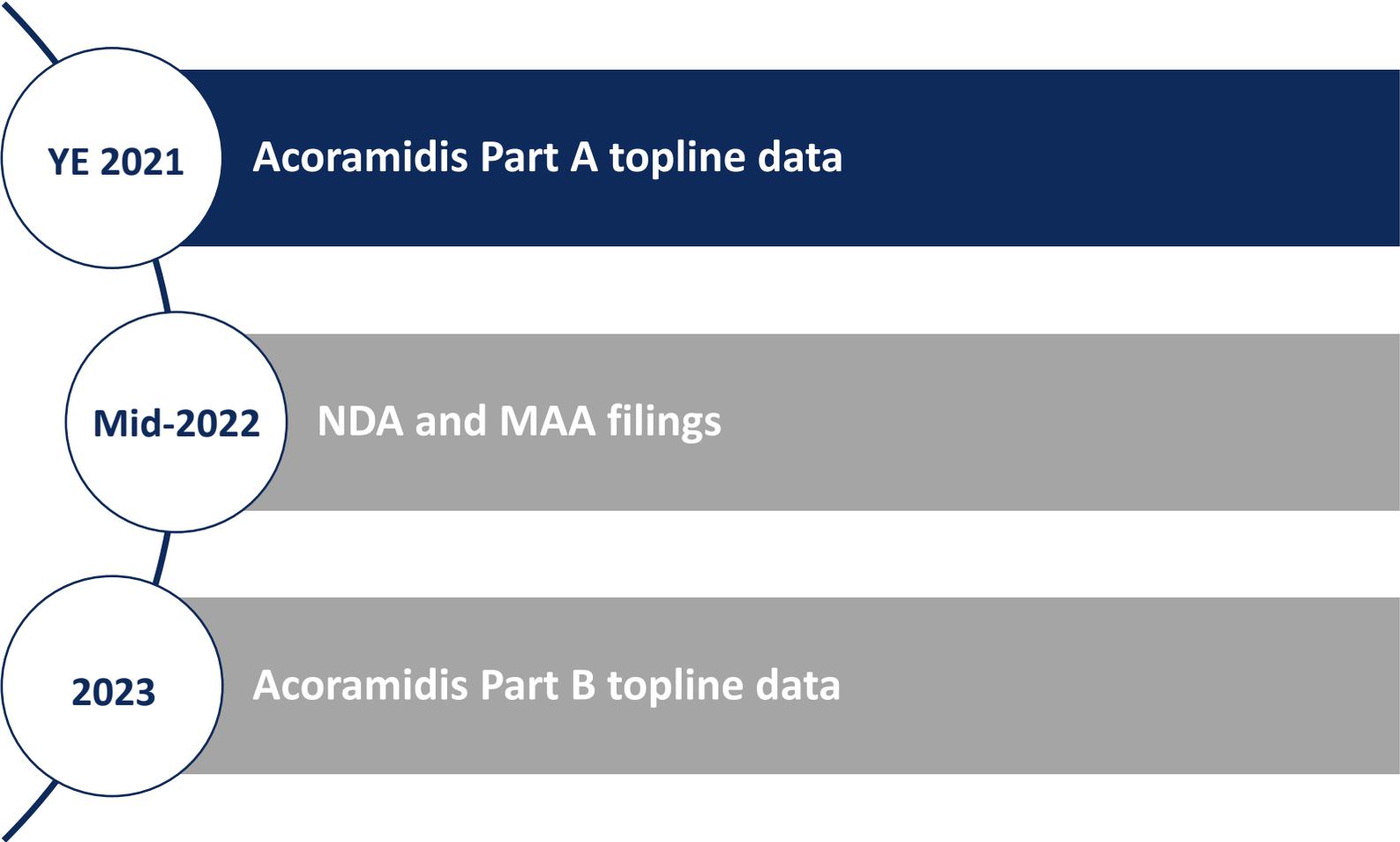
Therapeutic Approach

Published Data

Pivotal Trial Design

6MWD Importance

Upcoming Milestones



Note: NDA/MAA filings to occur upon receipt of positive Part A

Encaleret: CaSR inhibitor for ADH1

Mary Scott Roberts, M.D.
Sr. Director, Clinical Development,
Cardiorenal



Encaleret for autosomal dominant hypocalcemia type 1 (ADH1)



**Alexis and Jackson
Living with ADH1**

Prevalence

12k+

US

Pathophysiology

Decreased blood calcium, elevated urine calcium, and lower parathyroid hormone secretion

Genetic Driver



Hyperactivation of calcium-sensing receptor (CaSR)

Therapeutic Hypothesis



Selectively antagonize CaSR to normalize downstream effects

Design Criteria for Optimal Therapy

- ✓ Directly target CaSR to potentially resolve key symptoms
- ✓ Phase 2 data suggests potential to normalize blood Ca and urine Ca
- ✓ Oral Dosing

ADH1-causing variants hyperactivate the CaSR and disrupt calcium homeostasis leading to potentially life-threatening symptoms

Disease Overview

Therapeutic Approach

Phase 2 Design

Phase 2 Clinical Data

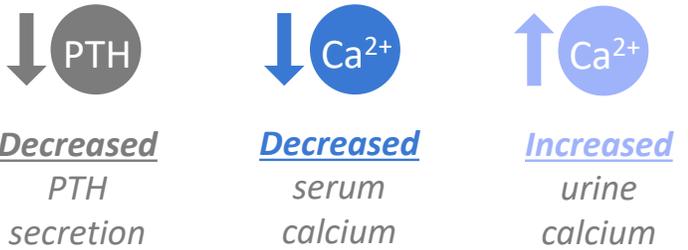
Summary & Next Steps

Disease Mechanism

Normal CaSR senses and regulates serum Ca levels to maintain calcium homeostasis



Hyperactive CaSR causes dysregulation of calcium homeostasis



Clinical Manifestation

Presenting symptoms

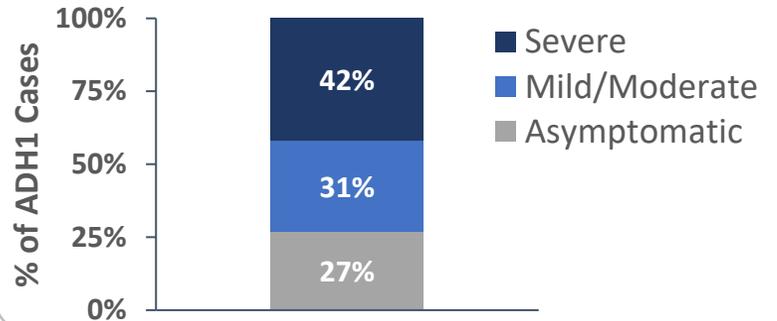
- Hypocalcemic seizures
- Paresthesia
- Tetany
- Muscle cramps

Long-term complications

- Nephrocalcinosis
- Nephrolithiasis
- Chronic kidney disease

Median age of ADH1 dx¹: 25 (0-77) years

Symptom presentation¹



¹Roszko, et al., ASBMR Annual Meeting, 2021. Abbreviations: dx = diagnosis. Age of dx presented as median (range)

ADH1 symptom severity is associated with blood calcium levels and current treatment inadequately addresses symptom burden

Disease Overview

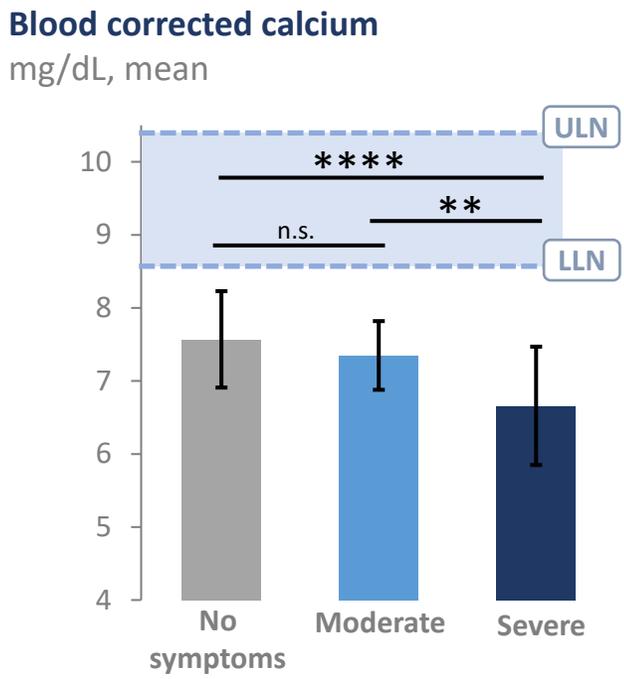
Therapeutic Approach

Phase 2 Design

Phase 2 Clinical Data

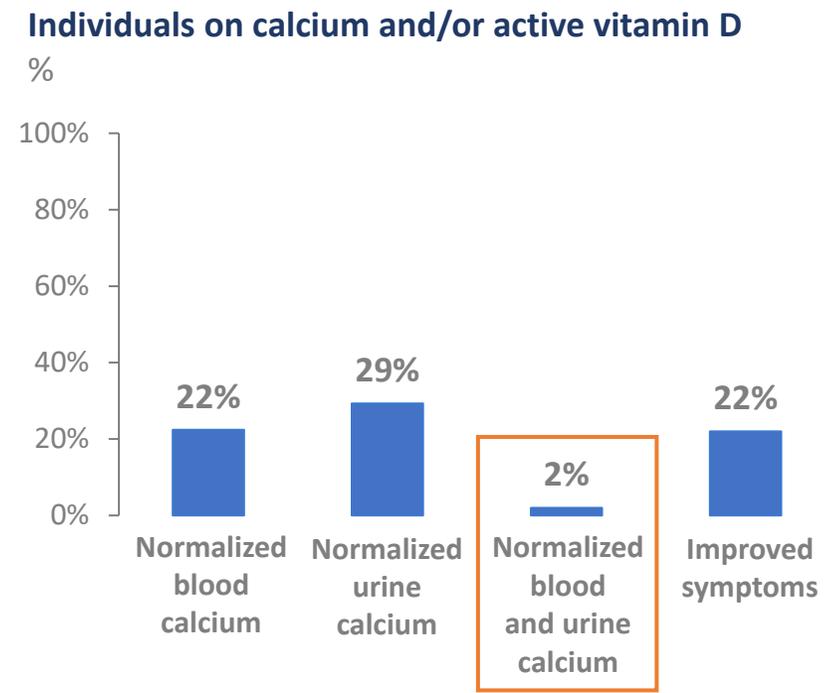
Summary & Next Steps

Blood calcium at clinical presentation



Severely symptomatic individuals exhibited significantly lower blood calcium compared to asymptomatic and moderately symptomatic¹

ADH1 medical intervention



Only 2% of individuals normalized both blood and urine calcium, and only 22% reported symptom improvement on-treatment¹

ULN = upper limit of normal, LLN = lower limit of normal. ** p-value < 0.01. **** p-value < 0.0001. n.s. = not statistically significant

¹Roszko, et al., ASBMR Annual Meeting, 2021

Successful CaSR antagonism would increase PTH secretion and renal calcium reabsorption

Disease Overview

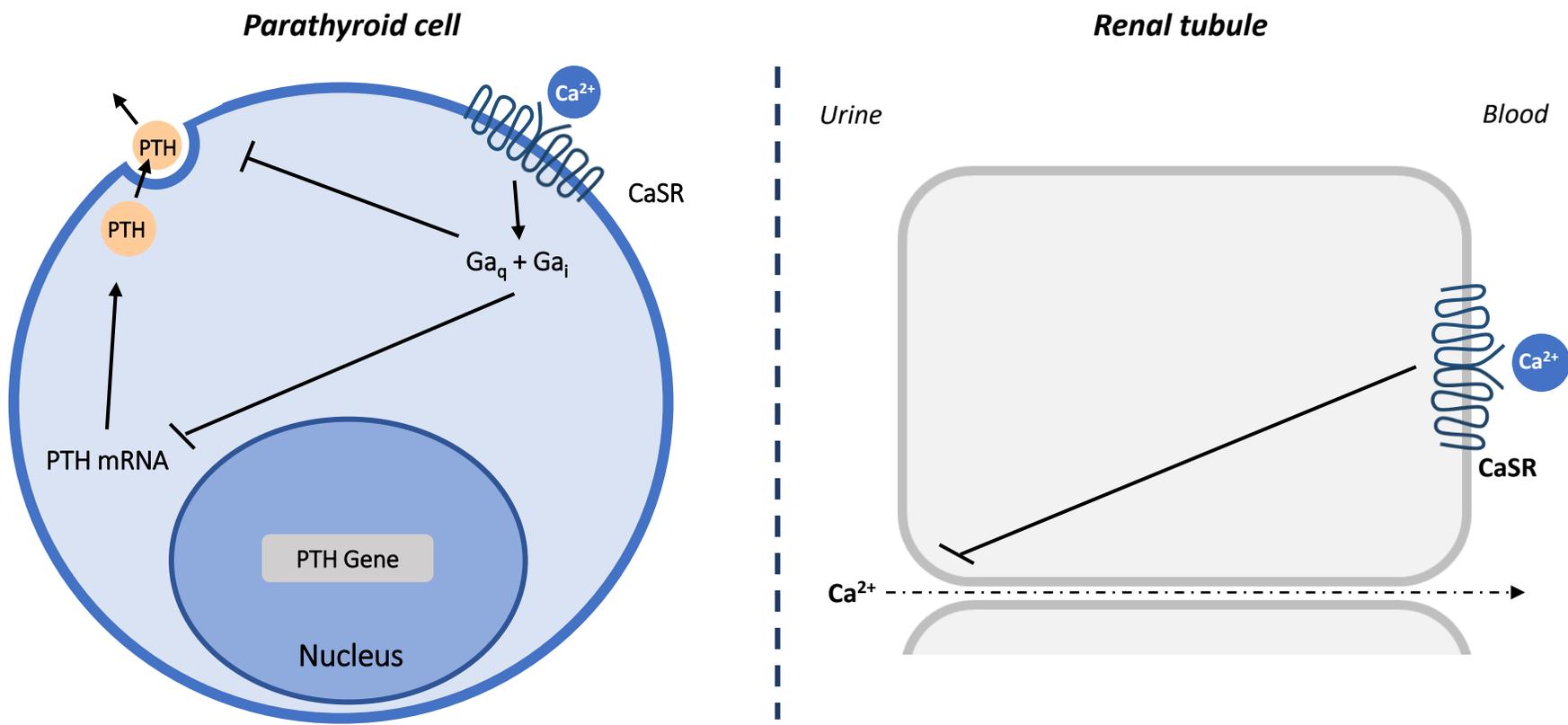
Therapeutic Approach

Phase 2 Design

Phase 2 Clinical Data

Summary & Next Steps

Parathyroid and renal CaSR antagonist action



Encaleret is designed to antagonize the CaSR to increase PTH release, increase blood calcium, and reduce urine calcium

Figures adapted from: ¹Berne and Levy Physiology, 6th ed. Chapter 39; ²Toka, H.R., et al. Physiology. 2015

Encaleret is designed to address the underlying disease mechanism and simultaneously normalize blood calcium and urine calcium

Disease Overview

Therapeutic Approach

Phase 2 Design

Phase 2 Clinical Data

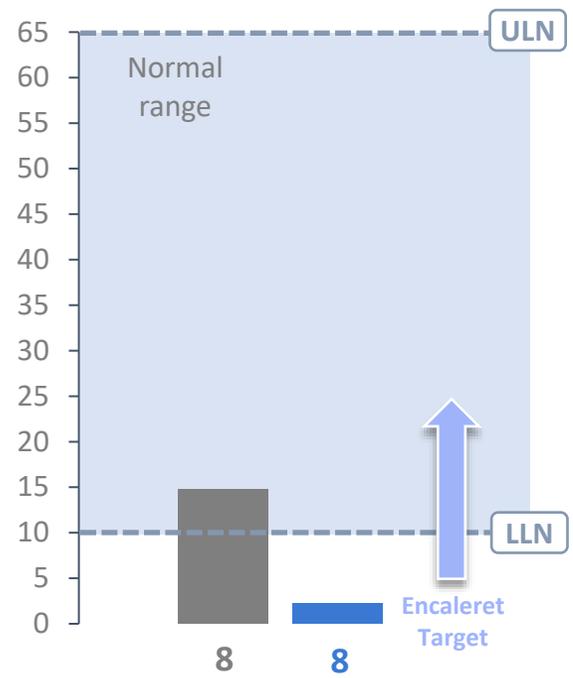
Summary & Next Steps

Summary of key disease measures in ADH1 patients with and without supplementation

■ Without supplementation ■ With supplementation

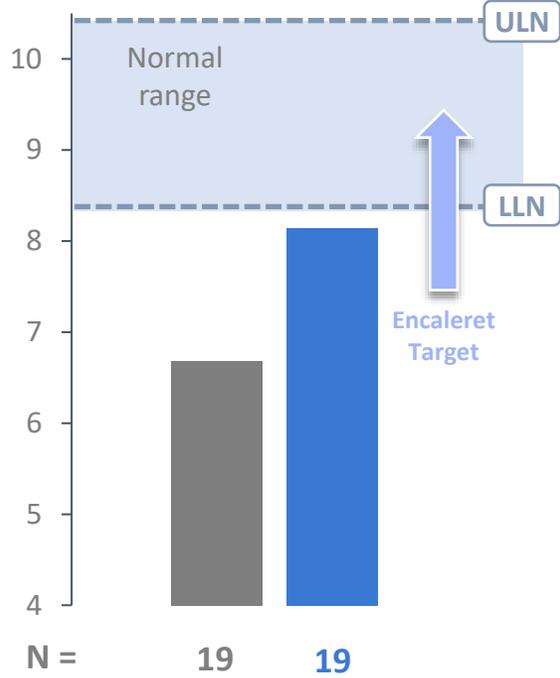
Blood parathyroid hormone

pg/mL, mean



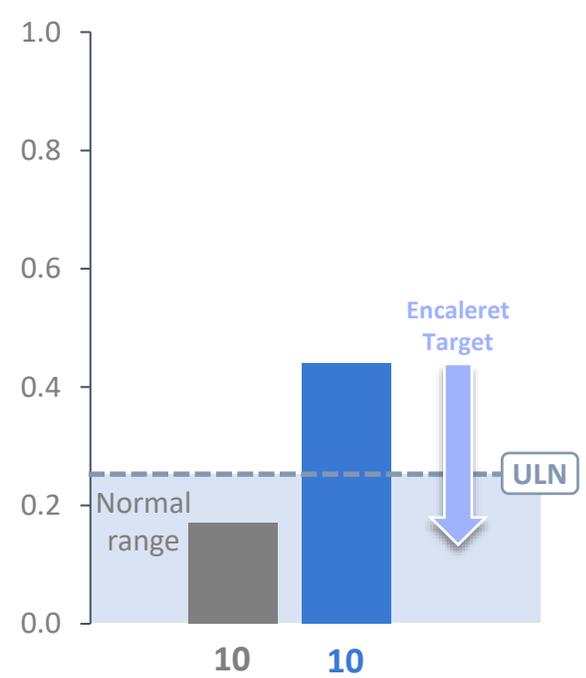
Blood corrected calcium

mg/dL, mean



Urine calcium

Calcium: Creatinine ratio, mean



ULN = upper limit of normal, LLN = lower limit of normal
Source: Pearce et al., NEJM 1996. PTH values reported as below detection limit or undetectable were recorded as "0"

Encaleret Phase 2 study design

Disease Overview

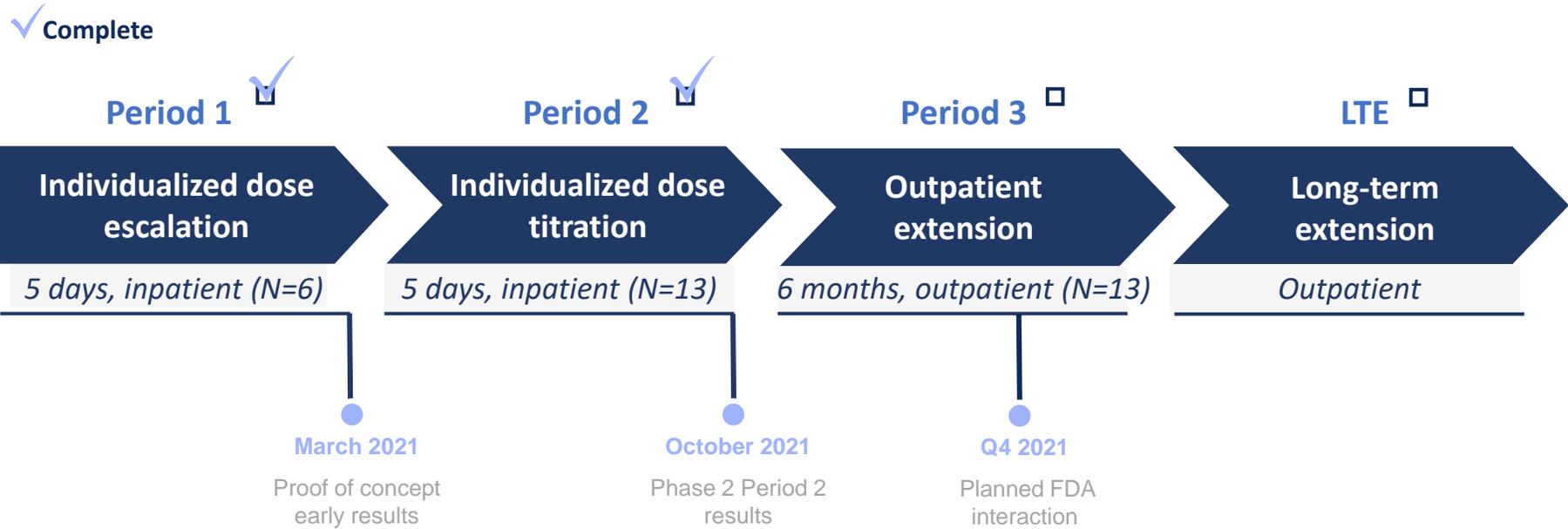
Therapeutic Approach

Phase 2 Design

Phase 2 Clinical Data

Summary & Next Steps

Program Overview



Key study objectives:

- Safety and tolerability
- Blood calcium concentration
- Urine calcium concentration
- Intact parathyroid hormone concentration

Additional measures:

- Blood 1,25-(OH)₂ Vitamin D, magnesium, and phosphate
- Urine creatinine, cAMP, citrate, phosphate, sodium, magnesium
- Bone turnover markers (serum collagen C-telopeptide, serum procollagen Type 1 N-propeptide)

Period 2 individualized dose titration phase resulted in a lower Day 5 mean encaleret dose as compared to Period 1

Disease Overview

Therapeutic Approach

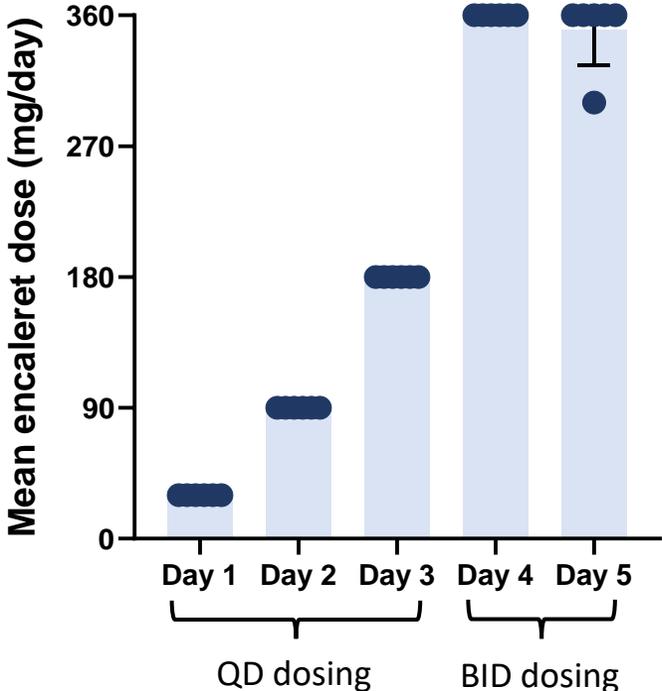
Phase 2 Design

Phase 2 Clinical Data

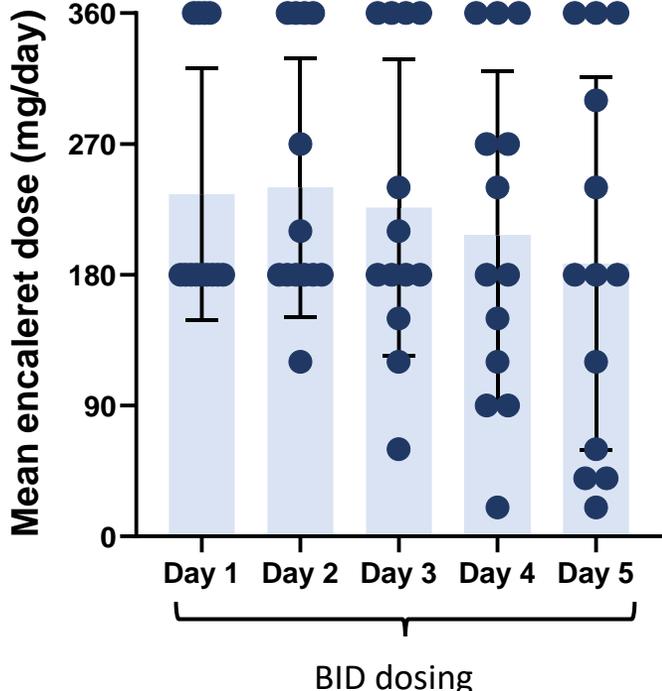
Summary & Next Steps

Period 1 and Period 2 encaleret dosing summary

Period 1 Dosing
Defined dose escalation
Day 5 Mean: 350.0±22.4 mg/day



Period 2 Dosing
Individualized dose titration
Day 5 Mean: 187.7±128.2 mg/day



Data reported as mean±SD.

Study participants exhibited hypocalcemia, elevated urine calcium, suppressed PTH, and elevated phosphate at baseline

Disease Overview

Therapeutic Approach

Phase 2 Design

Phase 2 Clinical Data

Summary & Next Steps

Baseline characteristics

Characteristic	Study Population N = 13	Normal Range
Age, mean, yr (range)	39 (22-60)	
Female, n (%)	8 (62%)	
Nephrocalcinosis, n (%)	10 (77%)	
ECG QT _c B (msec)	452 ± 16	< 440
Corrected Calcium (mg/dL)*	8.0 ± 0.7	8.4 – 10.2
Intact PTH (pg/mL)*	2.8 ± 3.4	15 – 65
Phosphate (mg/dL)*	5.1 ± 1.1	2.3 – 4.7
Magnesium (mg/dL)*	1.8 ± 0.1	1.6 – 2.6
24h Urine Calcium (mg/24h)	441 ± 258	< 250-300
Supplements		
Elemental Calcium (mg/day) [mean (range)]	2628 (750-4800)	
Calcitriol (µg/day) [mean (range)]	0.8 (0.2-2.0)	
CASR Variants	C131Y (2), P221L (2), E604K (1), A840V (3), F788C (1), T151M (1), Q245R (1), I692F (1), E228K (1)	

Data reported as mean±SD. ECG QT_cB = electrocardiogram Bazett-corrected Q-T interval. *Measurements taken pre-dose Day 1 in Period 1 or Period 2

Encaleret continues to be generally well-tolerated with no serious adverse events reported¹

Disease Overview

Therapeutic Approach

Phase 2 Design

Phase 2 Clinical Data

Summary & Next Steps

Summary of Period 1 and Period 2 safety measures

	Period 1 N = 6	Period 2 N=13
Number of subjects experiencing any Serious Adverse Event	0 (0%)	0 (0%)
Number of subjects experiencing any Adverse Event	6 (100%)	10 (77%)
Mild	6 (100%)	10 (77%)
Moderate	1 (17%)	0 (0%)
Severe	0 (0%)	0 (0%)
Number of Adverse Events Reported	19	12
Mild	18 (95%)	12 (100%)
Moderate	1 (5%)	0 (0%)
Severe	0 (0%)	0 (0%)
Treatment-related Adverse Events²	3 (16%)	8 (67%)
Hypocalcemia	1 (33%)	0 (0%)
Hypophosphatemia	2 (67%)	7 (88%)
Hypercalcemia	0 (0%)	1 (12%)

¹Data as of September 3, 2021. ²Treatment-related adverse events were transient and resolved with dose-adjustment. Treatment-related AEs were counted as the number of events per period and are presented as a percentage of the total number of AEs. The most common AEs (≥ 2 subjects) were hypophosphatemia, hypocalcemia, and headache

Encaleret treatment increased blood calcium and parathyroid hormone and decreased urine calcium in ADH1 participants during Period 1

Disease Overview

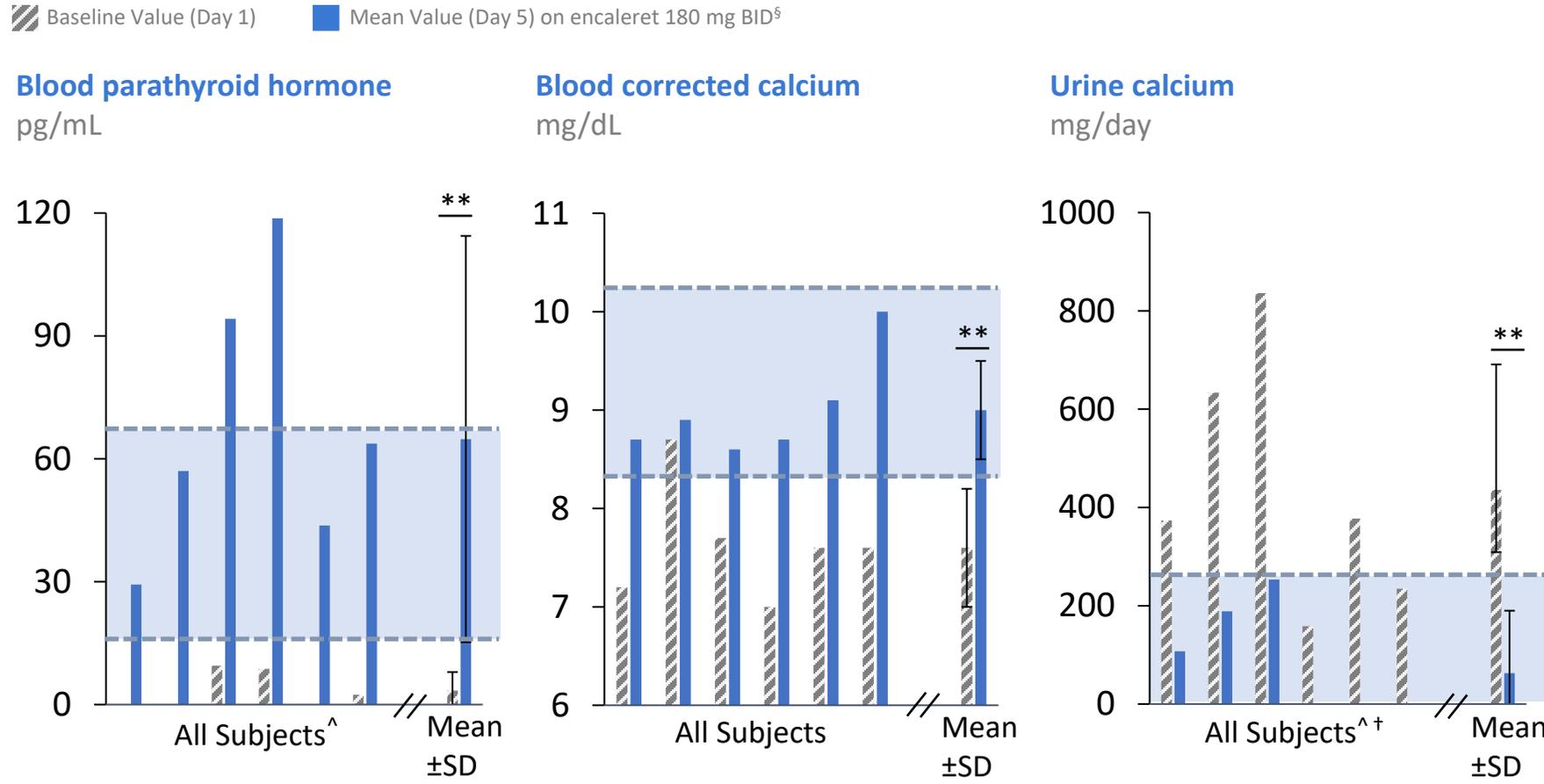
Therapeutic Approach

Phase 2 Design

Phase 2 Clinical Data

Summary & Next Steps

Individual and mean responses on Day 1 and 5 of Period 1 (N=6)



[§] Encaleret dose adjusted to 180/120 in 1 subject on Day 5. [^] Values below limit of assay quantitation recorded as "0". [†] Day 4 values used in two subjects given Day 5 values unavailable. Gray shading reflects normal range. ** p-value < 0.01.

Blood phosphate and magnesium levels also normalized over 5-day dose escalation, on average. Data not plotted above.

Encaleret treatment normalized mean blood and urine calcium during Period 2

Disease Overview

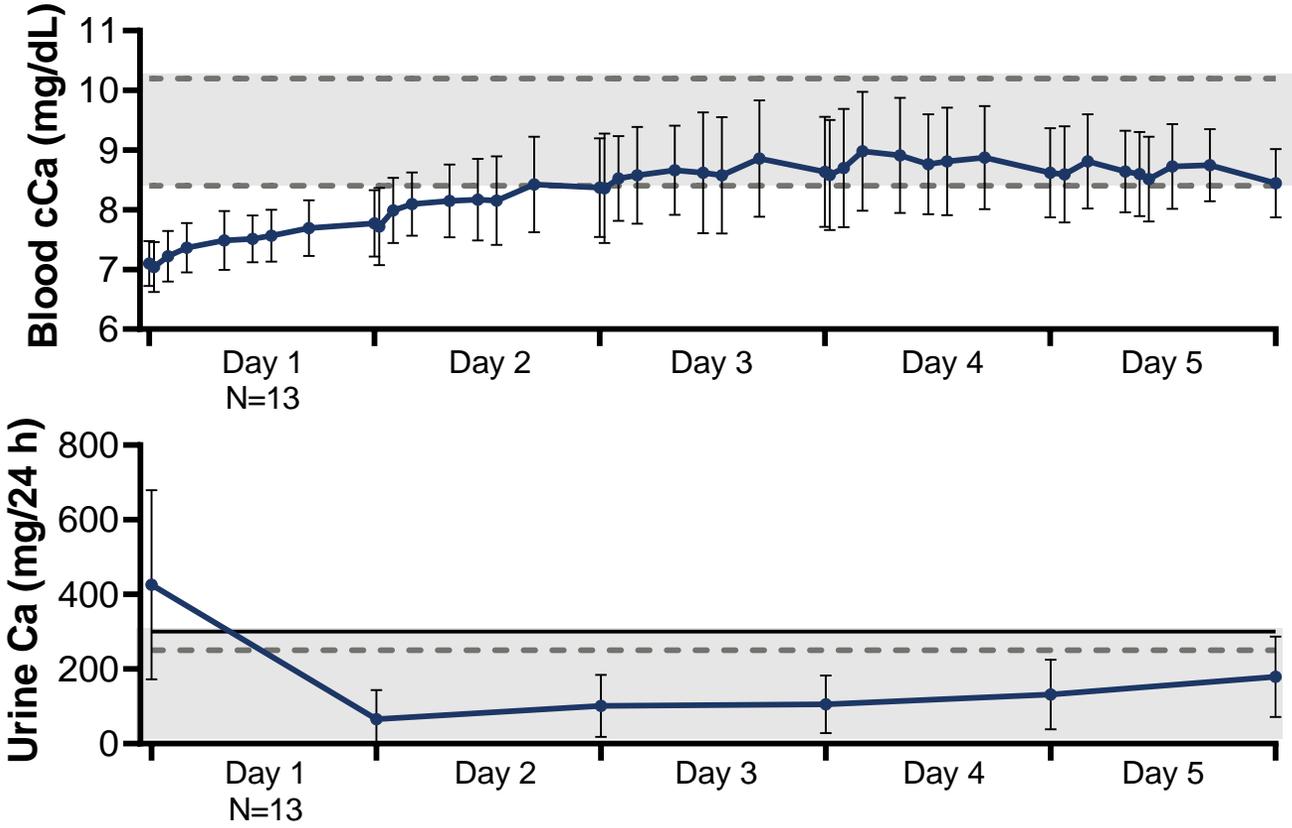
Therapeutic Approach

Phase 2 Design

Phase 2 Clinical Data

Summary & Next Steps

Mean responses on Day 1 through Day 5 in Period 2 (N=13)



Increasing urine calcium is likely due to both increasing corrected calcium and decreasing encaleret dose

Data reported as mean±SD. Values below limit of assay quantitation recorded as "0". Gray shading reflects normal range. Solid line for urine calcium reflects the upper limit for men and dashed line reflects upper limit for women

Encaleret increased PTH and decreased mean blood phosphate during Period 2

Disease Overview

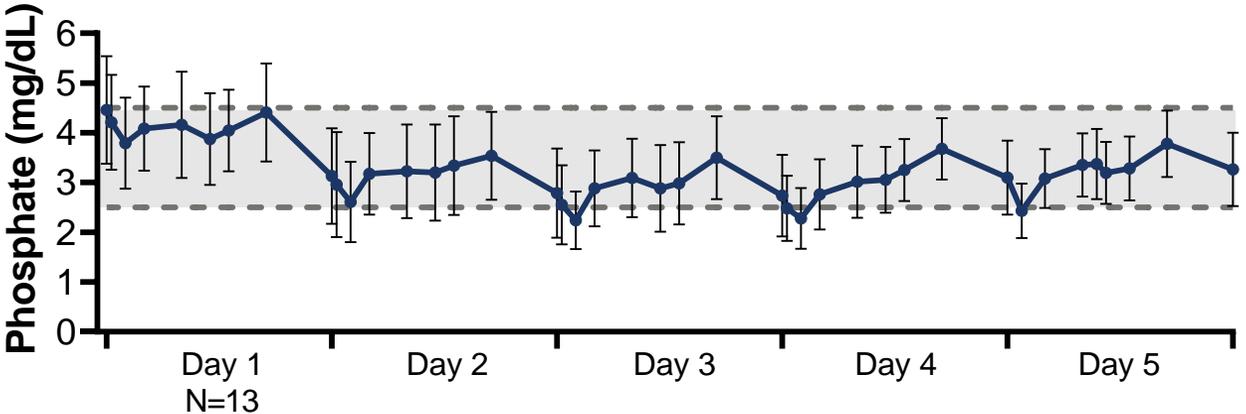
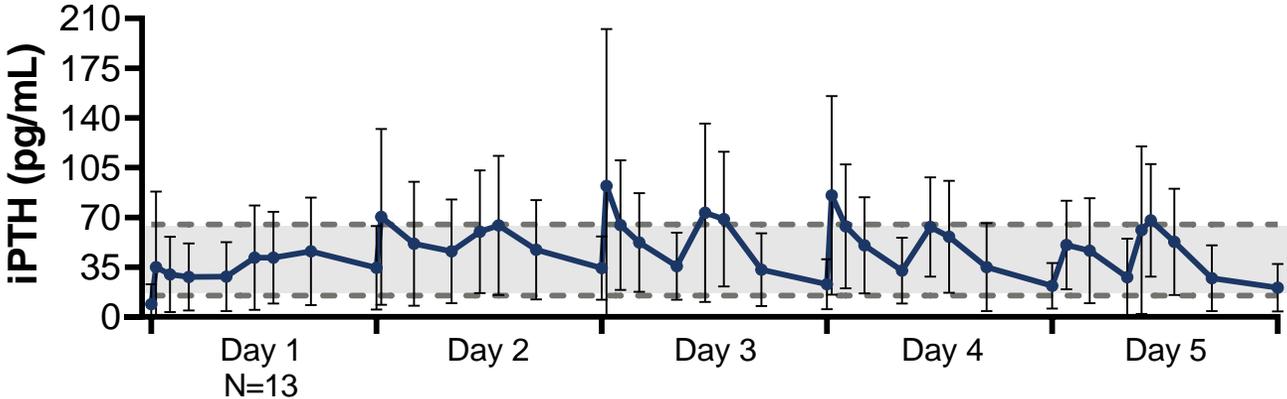
Therapeutic Approach

Phase 2 Design

Phase 2 Clinical Data

Summary & Next Steps

Mean responses on Day 1 through Day 5 in Period 2 (N=13)



Data reported as mean±SD. Values below limit of assay quantitation recorded as "0". Gray shading reflects normal range

Summary reported Phase 2 data and next steps

Disease Overview

Therapeutic Approach

Phase 2 Design

Phase 2 Clinical Data

Summary & Next Steps

Summary of encaleret development program

- ✓ In 13 participants, encaleret normalized mean blood calcium and 24-hour urine calcium excretion, increased PTH, and decreased phosphate into the normal range during both Periods 1 and 2
- ✓ Individualized BID dosing in Period 2 resulted in a decrease in the mean Day 5 encaleret dose as compared to Period 1
- ✓ Encaleret was well-tolerated when administered once or twice daily over 5 days, with no serious adverse events reported
- ✓ Consistent improvements in mineral homeostasis suggest encaleret may become an effective treatment for ADH1
- ✓ Granted Fast Track Designation and Orphan Drug Designation by the FDA

Next 12 months

- Interact with FDA
- Present complete Phase 2 data
- Initiate Phase 3 registrational study

Planned activities

- Pediatric development program in ADH1
- Evaluation of encaleret in non-genetic hypoparathyroidism

BridgeBio Gene Therapy

Eric David, M.D., J.D.

CEO, BBGT



Gene therapy pipeline overview

	Indication	Drug Mechanism	Pt. pop. (US+EU)	Discovery	Pre-IND	Phase 1	Phase 2	Phase 3
Gene Therapy	CAH	AAV5 gene therapy (BBP-631)	>75k					
	Canavan	AAV9 gene therapy (BBP-812)	1k					
	TMC1 hearing loss	AAV gene therapy (BBP-815)	2k					
	Galactosemia	AAV gene therapy (BBP-818)	>7k					
	Tuberous sclerosis complex 1	AAV gene therapy	>100k					
	Tuberous sclerosis complex 2	AAV gene therapy						
	Cystinuria	AAV gene therapy	20k					
	Undisclosed DCM gene therapy program	AAV gene therapy						
	3 capsid discovery collaborations							

 Featured Programs

Research and manufacturing capabilities



Facility | 20,000 sq ft lab space in Raleigh, NC

People | 60+ gene therapy employees (>50% in research or CMC)

Capabilities | Vector development, optimization, analytical development, and production (200L)

External Manufacturing | Dedicated GMP manufacturing suite at Catalent

BBGT program updates

Program	Status Update	Next Catalyst
Congenital Adrenal Hyperplasia (CAH)	Trial enrollment underway	Initial Phase 1/2 data
Canavan	Trial enrollment underway	Initial Phase 1 biomarker data
Transmembrane Channel Protein 1 (TMC1)	Proof-of-concept established in multiple disease models	IND enabling studies
NEW Galactosemia	Proof-of-concept established in disease model	IND enabling studies

BBP-631: AAV5 gene therapy for congenital adrenal hyperplasia (CAH)



Maddie
Living with CAH

Prevalence

>75k

US & EU

Pathophysiology

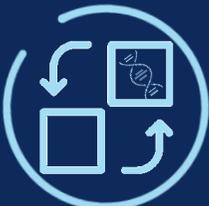
Inability to produce cortisol causes need for supraphysiologic doses of synthetic steroids, 3-4x increase in mortality risk, hirsutism, Cushingoid symptoms

Genetic Driver



Loss of function of 21-hydroxylase (21-OH)

Therapeutic Hypothesis



AAV5 gene therapy to provide 21-OH

Design Criteria for Optimal Therapy

- ✓ Only approach designed to induce endogenous cortisol and mineralocorticoid production
- ✓ Durable transgene delivery to the adrenal gland of NHPs
- ✓ Low threshold to correct phenotype

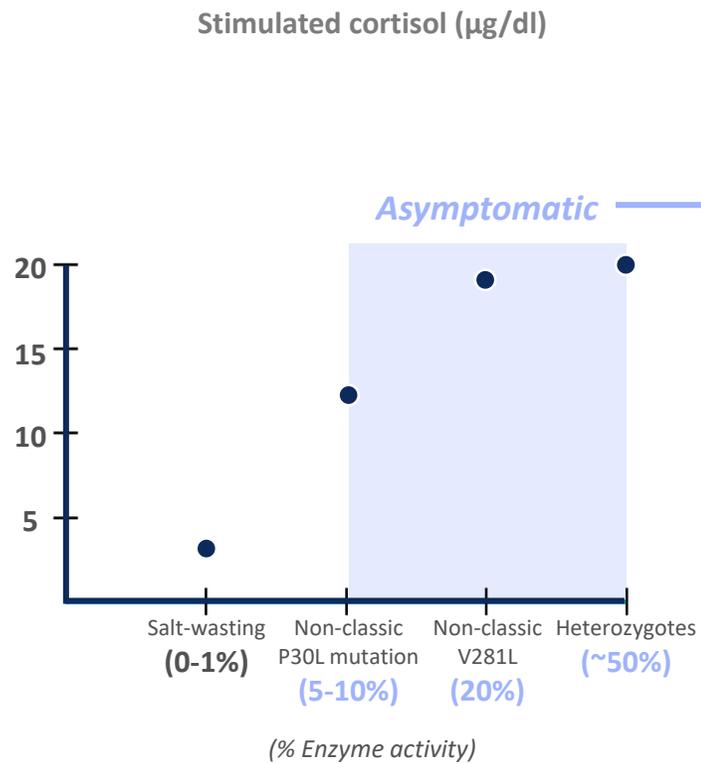
5-10% of WT enzyme may be sufficient for clinical impact

Preclinical Data

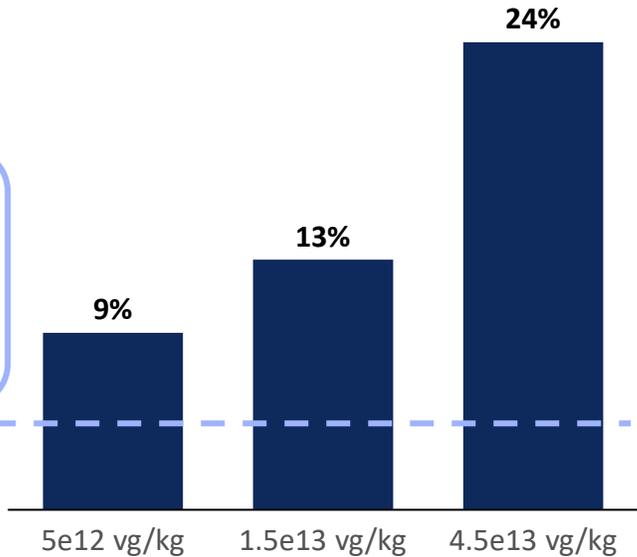
Phase 1/2 Design

Genotype-phenotype studies show that >5-10% of enzyme activity results in nonclassical CAH

NHP protein data suggests potentially therapeutic levels of 21-hydroxylase enzyme



Human 21-hydroxylase protein as a % of NHP 21-hydroxylase protein (Mass Spec quantification)



- Mass-spec methods to quantify protein expression by identifying differential peptides between human and NHP 21-OH
- Data suggest dose-dependent enzyme expression in the adrenal cortex from 9%-24% of WT levels

Source: Perdomini, Gene Therapy 2017; ESGCT 2019; data on file

Phase 1/2 first-in-human trial design

Preclinical Data

Phase 1/2 Design

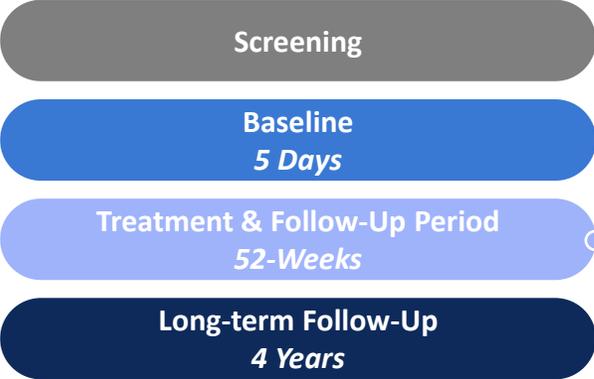
Status

- Trial enrollment underway

Eligibility

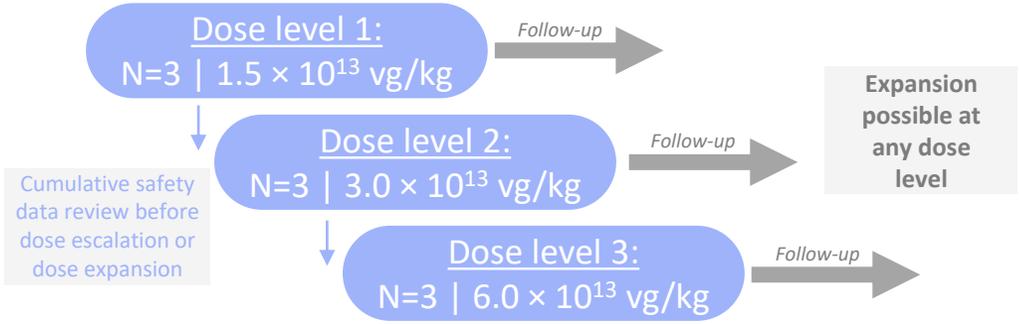
- Age >18 years with classic CAH (simple virilizing or salt-wasting) due to 21-Hydroxylase Deficiency (21-OHD)
- Screening/baseline 17-OHP levels > 5-10 × ULN

FIH Trial Design



Dose Escalation Design

Three dose levels of BBP-631 are planned for the study



Primary Objectives

- Evaluate safety
- Levels of endogenous cortisol (pre- and post-ACTH stimulation)
- Quality-of-life assessment

BBP-812: AAV9 gene therapy for Canavan disease



Prevalence

~1k

US & EU

Pathophysiology

Dysregulated amino acid metabolism leads to neurodegeneration and loss of motor function. Most children do not live past age 10

Genetic Driver



Loss of function of aspartoacylase (ASPA)

Therapeutic Hypothesis



AAV9 gene therapy to provide ASPA

Design Criteria for Optimal Therapy



Profound phenotypic rescue observed in mouse models by Dr. Guangping Gao



IV delivery enables widespread delivery across CNS



Clear PD biomarkers to measure effect of gene therapy (NAA)

Tobin
Child with Canavan

CANAspire, our FIH trial for BBP-812, will assess two doses before expanding

Phase 1/2 Design

Status

- Trial enrollment underway

Eligibility

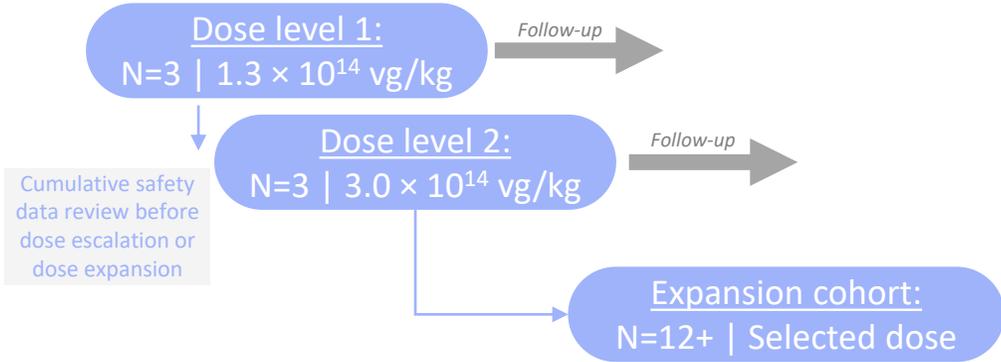
- Age ≤ 30 months with Canavan disease
- Stable health with no acute or chronic renal and liver disease
- Negative for AAV9 total antibodies

FIH Trial Design



Dose Escalation and Expansion Design

Two dose levels of BBP-812 are planned for the study



Primary Objectives

- Evaluate safety and tolerability
- Levels of NAA (CSF, urine)
- Developmental milestones (e.g., TIMPSI, GMFM-88, Canavan Disease Rating Scale)

BBP-815: AAV gene therapy for TMC1 genetic hearing loss

Addressable Prevalence

~2k

US & EU

Pathophysiology

Lack of functional TMC1 protein in the inner and outer hair cells of the cochlea leads to severe-to-profound bilateral hearing loss

Genetic Driver



Loss of function of transmembrane channel protein 1 (TMC1)

Therapeutic Hypothesis



Intracochlear delivery of AAV gene therapy to provide TMC1 and enable natural hearing processes

Design Criteria for Optimal Intervention



Only approach designed to enable natural hearing



Dose-dependent, durable rescue of the hearing phenotype in profoundly deaf mice



Near complete transduction of inner and outer hair cells in NHPs

Gene therapy is the only modality designed to address TMC1 hearing loss at its source and allow for endogenous production of the TMC1 protein

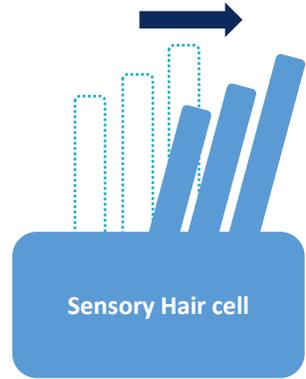
Therapeutic Approach

Preclinical Data

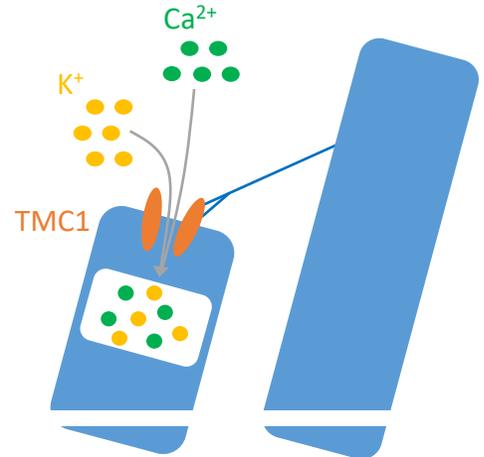
BBP-815 is the only intervention designed to enable natural hearing

TMC1 = transmembrane channel protein 1

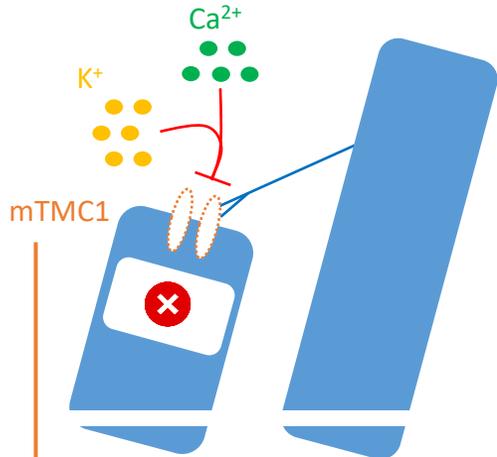
Sound-evoked displacement of hair cell stereocilia



Functional TMC1 enables hair cell depolarization



Mutated TMC1 disrupts electrical response to sound



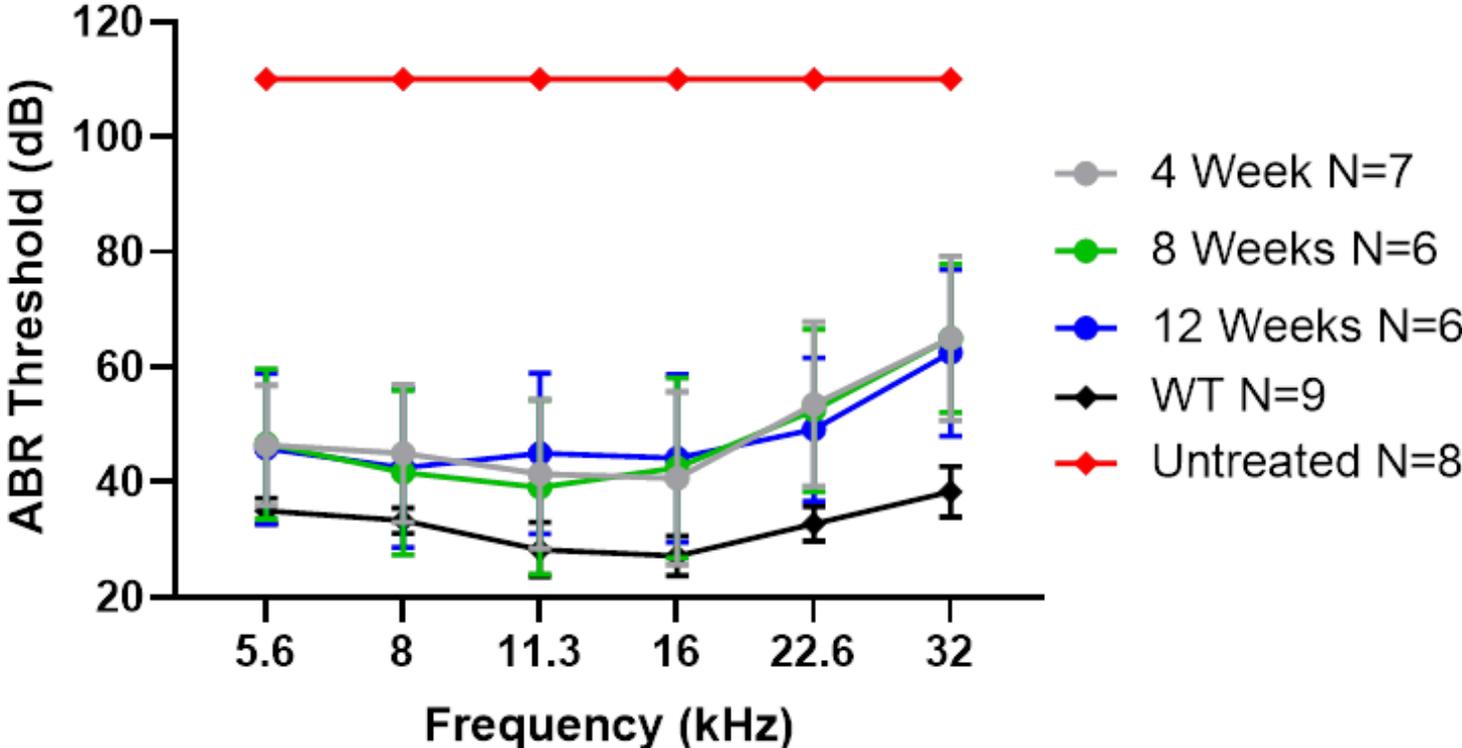
BBP-815 is designed to produce the TMC1 protein directly in hair cells of the cochlea and enable mechanotransduction of sound

BBP-815 durably rescues hearing in profoundly-deaf mice

Therapeutic Approach

Preclinical Data

Durability of Auditory Brainstem Response (ABR) in TMC1-N193I mice dosed with BBP-815



Source: BridgeBio Gene Therapy data on file

BBP-818: AAV gene therapy for classic galactosemia type I

Prevalence

7k+

US & EU

Pathophysiology

Inability to metabolize galactose leads to impaired speech, developmental delays, impaired motor function, primary ovarian insufficiency, and osteopenia

Genetic Driver



Loss of function of galactose-1-phosphate uridylyltransferase (GALT)

Therapeutic Hypothesis



AAV gene therapy to provide GALT enzyme and enable galactose metabolism

Design Criteria for Optimal Therapy



Only approach designed to enable galactose metabolism, reduce gal-1p, and restore galactosylation of lipids and proteins



Dose-dependent, durable restoration of the GALT enzyme in liver and CNS



Low threshold to correct phenotype

Classic galactosemia is a slowly progressive disease that impacts development of the CNS, ovaries, and bone despite strict dietary restrictions

Unmet Need

Therapeutic Approach

Preclinical Data



Toddlers/Children

- Speech/ language disorders
- Developmental delays
- Impaired growth
- Cataracts
- Learning delays
- Neurological Impairments, including gait, balance, fine motor tremors
- Behavioral and emotional issues
- Tremor



Teens

- Puberty and fertility problems (females)
- Primary ovarian insufficiency (POI)
- Anxiety
- Growth delays
- Social challenges
- Learning difficulties
- Tremor



Adults

- Tremor
- Seizures
- Anxiety
- Depression
- ADHD
- Cataracts
- Early-onset dementia

Top 3 Lifelong Complications (despite dietary restriction):

- **85% had impaired CNS development**
- **80% had impaired ovarian development**
- **26.5% had impaired bone development**

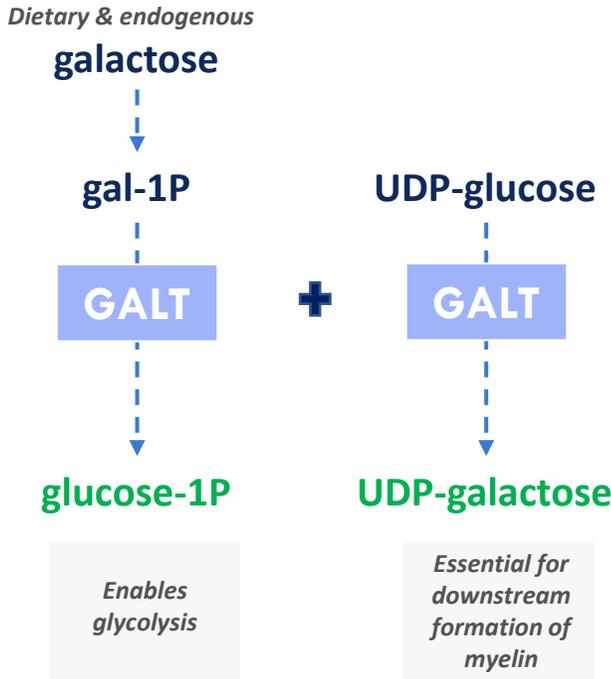
Gene therapy is the only therapy designed to treat galactosemia at its source and allow for endogenous production of the GALT enzyme

Unmet Need

Therapeutic Approach

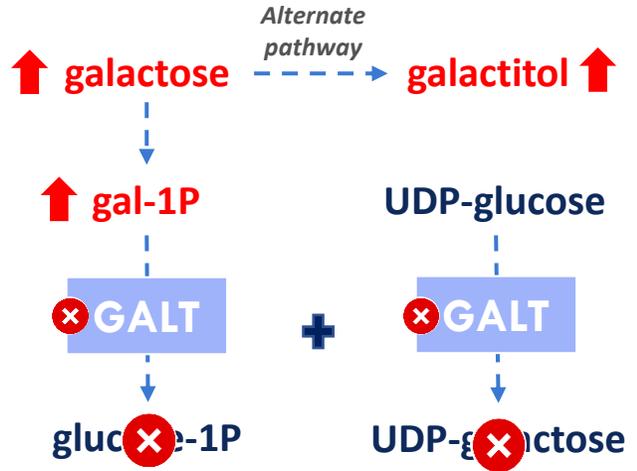
Preclinical Data

Functional galactose metabolism (Leloir Pathway)



- In typical galactose metabolism, galactose from both diet and endogenous production is processed by the GALT enzyme into downstream glucose-1P and UDP-galactose

Dysregulation of galactose metabolism in type I galactosemia



- In type I galactosemia, gal-1P is unable to be processed, leading to buildup of gal-1P, galactose, and galactitol, all of which are toxic.
- UDP-glucose is unable to be converted into UDP-galactose, which is essential for myelination
- Key symptoms include speech impairment, ataxia, significant developmental delays, primary ovarian insufficiency and osteopenia

GALT = Galactose-1-phosphate-uridylyltransferase; gal-1P = galactose-1-phosphate
 Source: Berry. Classic Galactosemia and Clinical Variant Galactosemia. 2021

BBP-818 achieves significant, dose-dependent production of the GALT enzyme in GALT-KO mouse brain; these levels may be sufficient for clinical impact

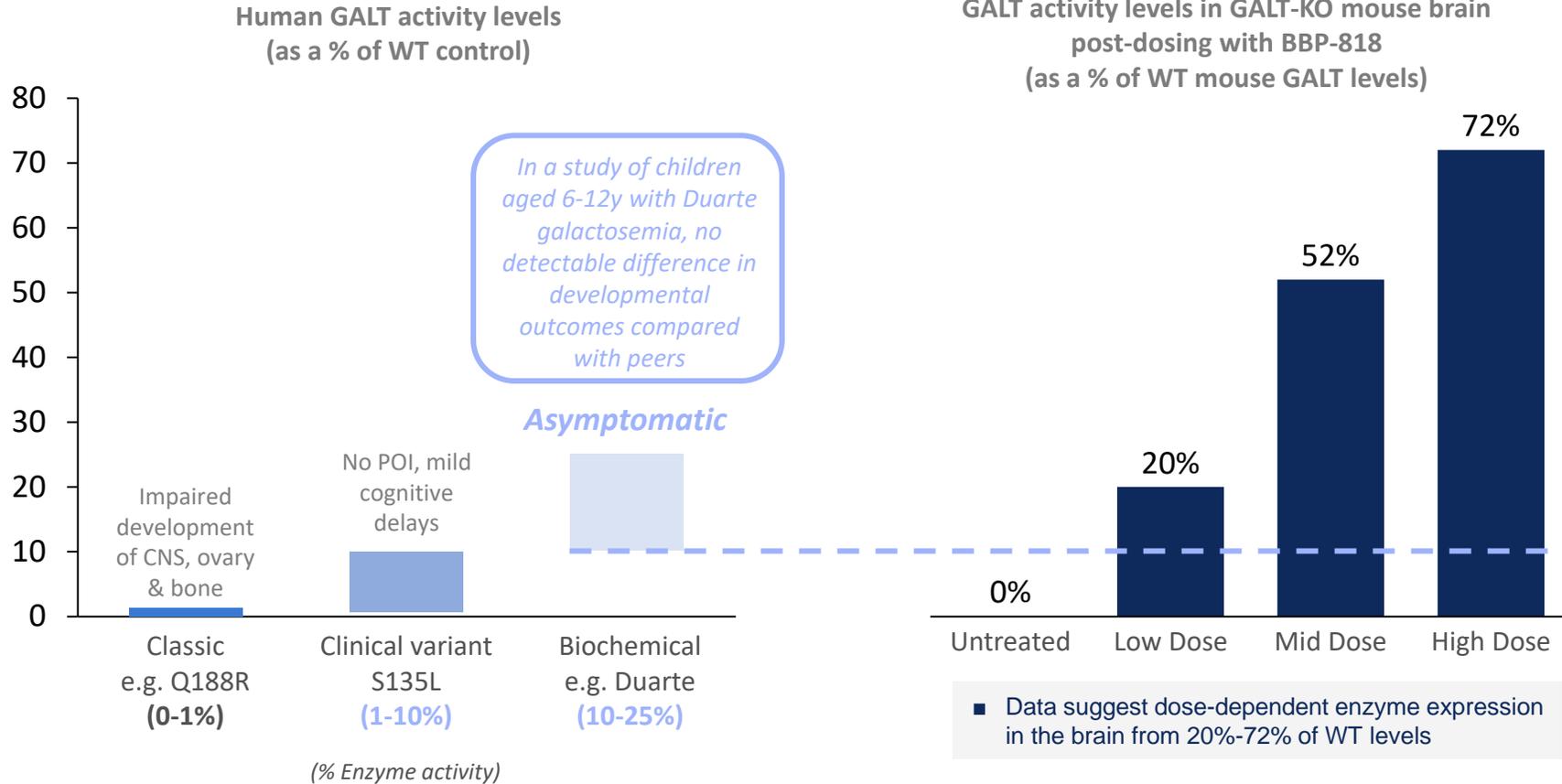
Unmet Need

Therapeutic Approach

Preclinical Data

Genotype-phenotype studies show patients with >10% of GALT activity tend to be asymptomatic, and those with >1% tend to have much milder phenotype

GALT-KO mouse data suggests potentially therapeutic levels of GALT enzyme production



Source: BridgeBio Gene Therapy data on file, Berry 2021, Fridovich-Keil 2014

BridgeBio Gene Therapy

EXPERIENCED GENE THERAPY TEAM



Senior leadership team of industry veterans



FLEXIBLE MANUFACTURING MODEL



In-house capabilities & flexible facility build-out



20,000 sq ft lab space in Raleigh, NC

GROWING PIPELINE



Robust pipeline with clinical readouts in 2022

2 clinical programs

2 pre-IND programs

4 discovery programs

3 capsid discovery collaborations

Mendelian Wave 3 Programs

Uma Sinha, Ph.D.
Chief Scientific Officer



Mendelian pipeline overview

	Indication	Drug Mechanism	Pt. pop. (US+EU)	Discovery	Pre-IND	Phase 1	Phase 2	Phase 3	Approved
Mendelian	MoCD type A	NULIBRY™ (synthetic cPMP, fosdenopterin)	100						
	Achondroplasia	Low-dose FGFRi (infigratinib)	55k						
	LGMD2i	Glycosylation substrate (ribitol)	7k						
	RDEB	Recombinant COL7 (BBP-589)	2k						
	PKAN / organic acidemia	Pank activator (BBP-671)	7k						
	VM / LM	Topical PI3Ki (BBP-681)	117k						
	Netherton	Topical KLKi (BBP-561)	11k						
	PTEN autism	PI3Kb inhibitor (BBP-472)	120k						
	4 undisclosed small molecule programs		>500k						
	4 undisclosed antisense oligonucleotide programs		>300k						
Precision Cardiorenal	ATTR amyloidosis	TTR stabilizer (acoramidis)	>400k						
	ADH1	CaSR antagonist (encaleret)	12k ¹						
	PH1 / frequent stone formers	GO1 inhibitor (BBP-711)	5k / 1.5m						
	2 undisclosed DCM programs		>250k						

 Featured Programs

¹US carriers

Primary Hyperoxaluria Type 1 (PH1)



Violet
Living with PH1

Prevalence

5k

US & EU

Pathophysiology

Excess oxalate result in the buildup of oxalate stones in the kidneys, urinary tract, and other vital organ

Genetic Driver



Loss of function of AGXT gene causing excess oxalate

Therapeutic Hypothesis



GO1 inhibitor to reduce oxalate production

Design Criteria for Optimal Therapy

 Highest pharmacodynamic response (max inhibition of GO)

 Oral dosing

GO = glycolate oxidase

GO1 inhibitor (BBP-711) is designed to treat PH1 at its genetic source

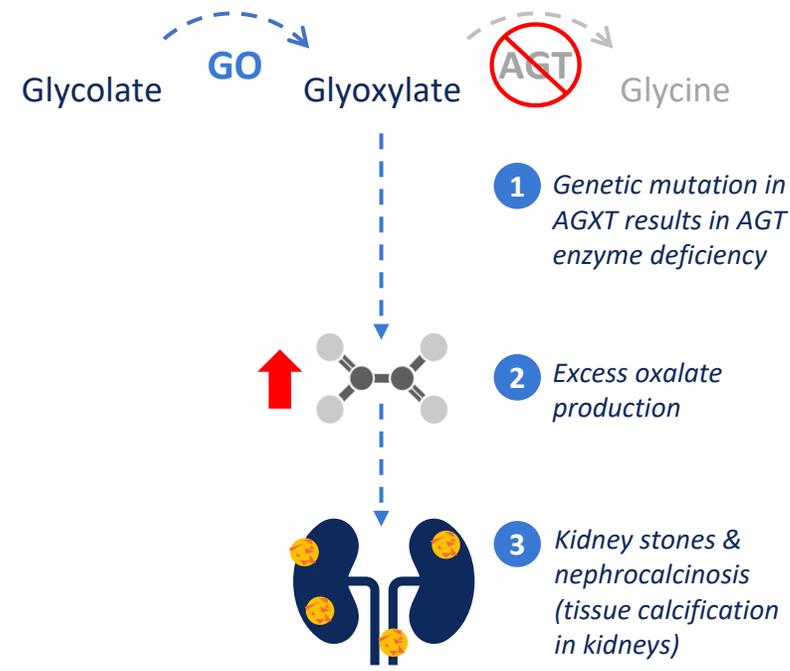
PH1 Program Overview

Phase 1 Clinical Data

Expansion opportunity

Disease Mechanism

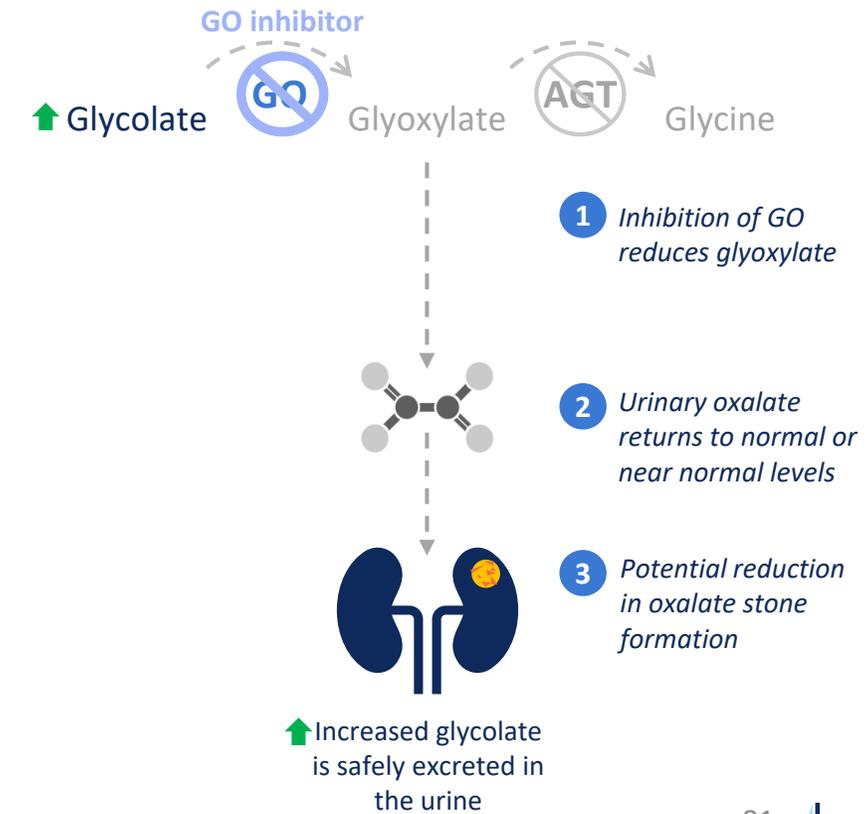
- ✓ Functional AGXT metabolizes glyoxylate into glycine resulting in normal oxalate production
- ✗ Loss of function mutation in AGXT results in excess oxalate production, causing oxalate stone formation



Hyper - oxal - uria = Excess - oxalate - in the urine

Therapeutic Approach

- 💡 Inhibition of GO reduces the precursor substrate (glyoxylate) for oxalate production



Well tolerated in Phase 1 with dose-dependent increases in plasma glycolate

PH1 Program Overview

Phase 1 Clinical Data

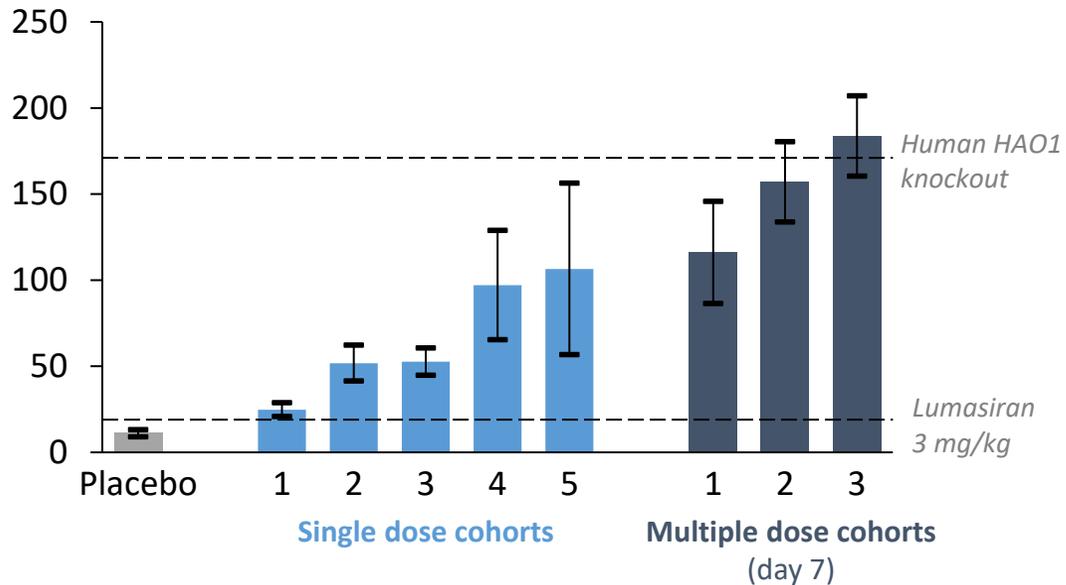
Expansion opportunity

Oral doses in healthy adult volunteers

Mean maximal plasma glycolate concentration

Mean \pm SD (μ M)

Preliminary and interim data



Well tolerated

- No safety signals of clinical concern
- All AEs were mild or moderate

Favorable PK/PD profile

- Potential to maximally inhibit GO with once-daily dosing
- Largest glycolate response observed to date by targeting GO

Timeline

2022

Phase 1 data & Phase 2/3 start

Key Endpoints

- Increase in plasma glycolate
- Safety and tolerability
- Pharmacokinetic profile

Expansion opportunity in recurrent stone formers with hyperoxaluria

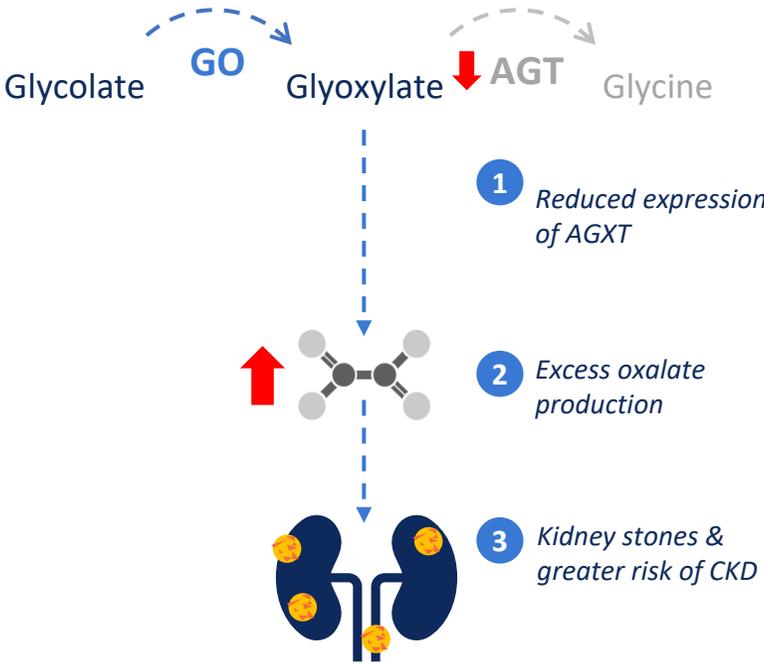
PH1 Program Overview

Phase 1 Clinical Data

Expansion opportunity

Disease Mechanism

✘ Epigenomic changes in AGXT results in excess oxalate production, causing oxalate stone formation



Prevalence

~1.5M

US & EU

Epigenomic Driver



Hypermethylation and downregulation of AGXT causing excess oxalate

Gianmoena, Kathrin, et al. "Epigenomic and transcriptional profiling identifies impaired glyoxylate detoxification in NAFLD as a risk factor for hyperoxaluria." *Cell Reports* 36.8 (2021): 109526.

Limb-Girdle Muscular Dystrophy Type 2i (LGMD2i)



Seamus
Living with LGMD2i

Prevalence¹
7k
US & EU

Pathophysiology
Progressive muscle weakness resulting in the loss of ability to perform routine daily functions

Genetic Driver

Loss of function of FKR gene

Therapeutic Hypothesis

Add glycosylation substrate to drive residual enzyme activity

Design Criteria for Optimal Therapy

-  Naturally occurring compound with strong safety profile
-  First potential disease targeting therapy
-  Oral dosing

¹Includes all patients with potentially treatable mutations

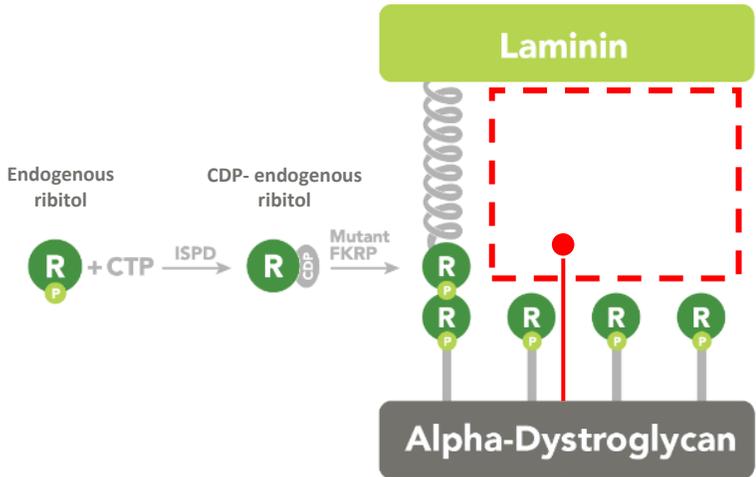
Ribitol (BBP-418) is being investigated as an upstream substrate to drive residual activity of the mutant FKRP enzyme

LGMD2i Program Overview

Update / Next Steps

Disease Mechanism

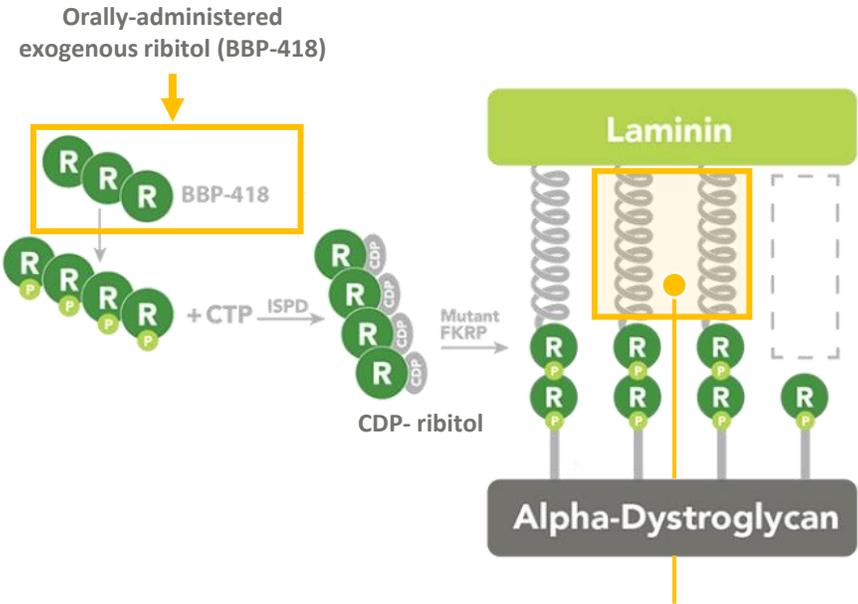
- ✔ Functional FKRP fully glycosylates alpha-dystroglycan (α -DG) which stabilizes cells by binding extracellular ligands
- ✘ Partial loss of function mutation in FKRP result in dysfunctional, hypo-glycosylated α -DG in muscle cells which increases cell susceptibility to damage



Mutations in FKRP prevent addition of CDP-ribitol to alpha-dystroglycan (hypo-glycosylated α -DG) limiting α -DG's ability to function as a "shock absorber" for muscle fibers

Therapeutic Approach

- 💡 Supply supraphysiological levels of ribitol upstream to drive residual activity of mutant FKRP enzyme and increase α -DG glycosylation levels



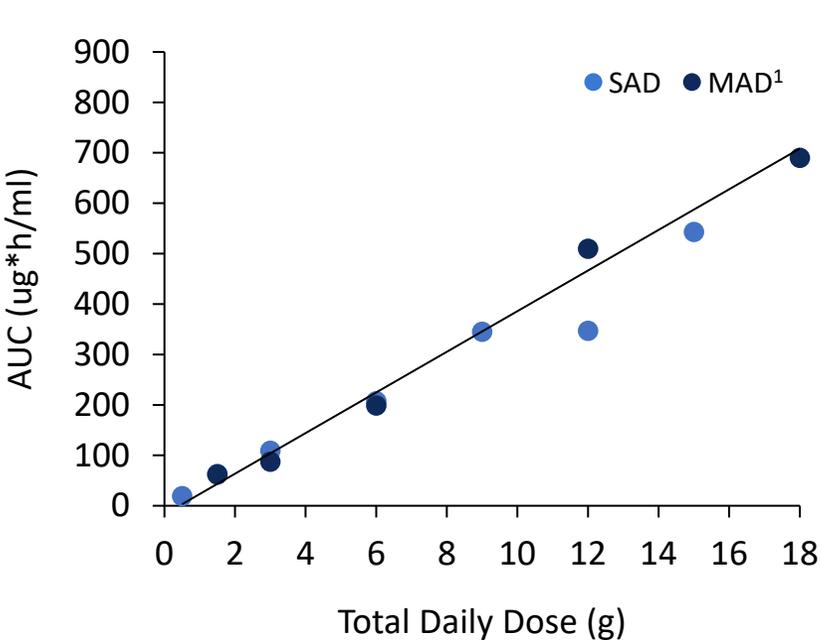
Potential partial restoration of α -DG glycosylation

Strong safety profile in Phase 1 with first LGMD2i patient dosed in 1Q21

LGMD2i Program
Overview

Update / Next Steps

Phase 1 findings: Exposure by total daily dose for healthy volunteers



- ✓ *Strong safety profile with all doses well-tolerated*
- ✓ *Dose-proportional exposures up to the highest dose tested*
- 💡 *Informed Phase 2 dosing selection*
- 💡 *Phase 2 ongoing*

Timeline

2022

Phase 2 Data

Key Endpoints

- Safety and tolerability
- Functional clinical assessments
- Glycosylated α -DG
- Creatine Kinase (marker of muscle breakdown)

¹MAD cohorts include 1.5g QD, 3g QD, 3g BID, 6g BID, 9g BID

Recessive Dystrophic Epidermolysis Bullosa (RDEB)



Child Living with RDEB

Prevalence
2k
US & EU

Pathophysiology
Loss of structural adhesion causes debilitating blistering, tearing and scarring of the skin and impacts oral and GI

Genetic Driver

Loss of function of collagen type VII protein (C7)

Therapeutic Hypothesis

Systemic treatment by providing C7 replacement therapy through IV infusion

- Design Criteria for Optimal Therapy**
-  Only systemic treatment targeting RDEB
 -  Potential to provide GI & oral benefits while also proactively improving the quality of wounds
 -  Convenient dosing schedule for patient and families

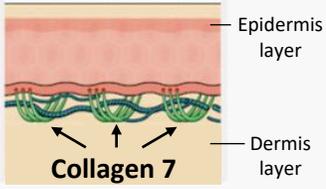
PTR-01 is currently the only systemic treatment targeting RDEB at its source

RDEB Program Overview

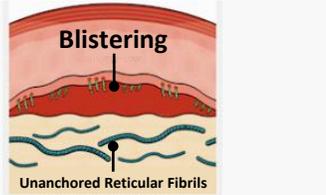
Update / Next Steps

Disease Mechanism

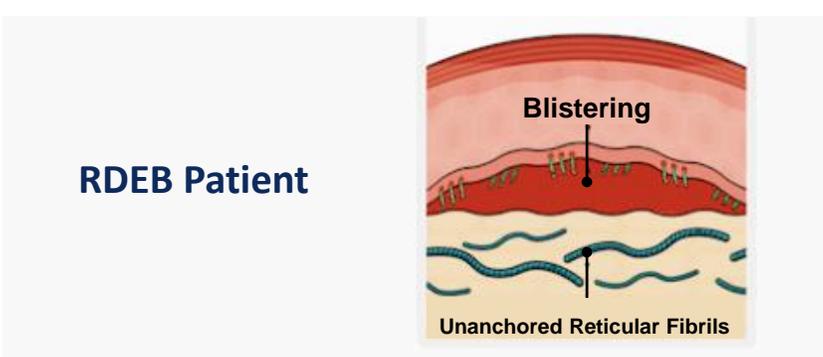
✓ *Functional C7 forms anchoring fibrils (AFs) to secure the epidermis to the dermis in healthy skin*



✗ *Mutated C7 causes severe blistering due to loss of structural adhesion*

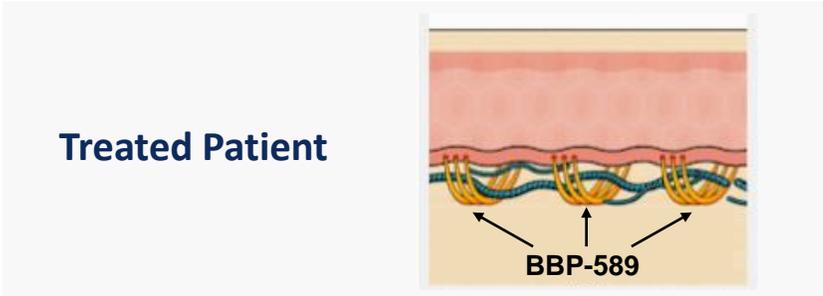


Therapeutic Approach



RDEB Patient

BBP-589 deposits in the skin, assembles into functional AFs, and reconstitutes DEJ adhesion in patients with RDEB



Treated Patient

Symptoms

Localized skin

Universal skin

Systemic tissues



- Severe blistering
- Wounding
- Extensive scarring



- Dystrophy and loss of nails
- Mitten deformity of hand and feet
- Higher risk for aggressive squamous-cell carcinoma



- Joint contractures
- Oral and dental malformations
- Severe impact to upper GI track
- Nutritional deficits and anemia

DEJ: dermal-epidermal junction

Well tolerated in Phase 1 with dose-dependent increase in C7 skin deposition

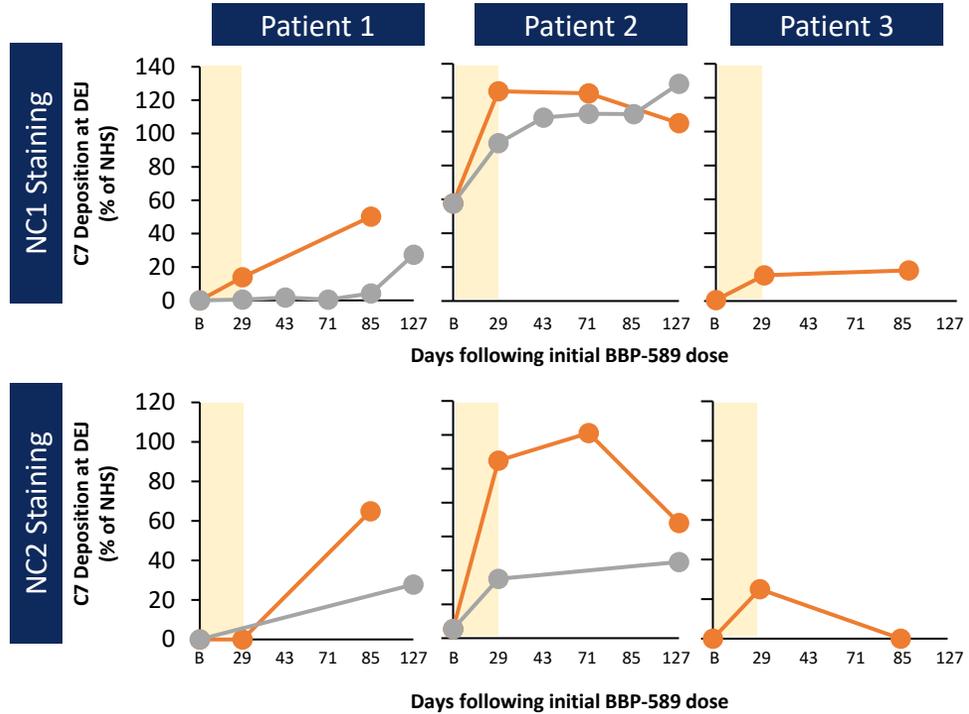
RDEB Program Overview

Update / Next Steps

Phase 1: Immunofluorescence C7 Deposition

Treatment window: Pts received 3 doses of BBP-589 (3mg/kg) during this time

● 3 mg/kg ● 0.3 mg/kg



All 3 patients receiving 3mg/kg dose exhibited increased C7 deposition at the DEJ

Timeline

1H22

Phase 2 data

Key Endpoints

- Wound improvement
- Deposition of C7 in skin biopsy
- Safety and tolerability

Patients 1 and 2 participated in the 3mg/kg and 0.3mg/kg cohorts. DEJ: dermal-epidermal junction

Mendelian program summary

15+ Mendelian Disease Programs

1 FDA approval

MoCD Type A



5 mid/late-stage programs

ATTR, Achondroplasia, ADH1, RDEB, LGMD2i

3 early clinical programs

PH1, PKAN, VM/LM

10+ preclinical programs

Catalysts (YE21 / 2022)

Acoramidis – Part A readout

4Q 2021

Achondroplasia, RDEB, LGMD2i, PH1

2022

Precision Oncology: Program Updates

Eli Wallace, Ph.D.

Chief Scientific Officer,
Oncology



Precision oncology pipeline overview

	Indication	Drug Mechanism	Pt. pop. (US+EU)	Discovery	Pre-IND	Phase 1	Phase 2	Phase 3	Approved	
Precision Oncology	FGFR2+ cholangiocarcinoma (2L)	TRUSELTIQ™ (FGFRi, infigratinib)	4k							
	FGFR2+ cholangiocarcinoma (1L)	FGFRi (infigratinib)	4k							
	FGFR3+ adjuvant urothelial	FGFRi (infigratinib)	21k							
	FGFR1-3+ tumor agnostic	FGFRi (infigratinib)	24k							
	FGFR1-3+ gastric cancer	FGFRi (infigratinib)	41k ¹							
	MAPK / RAS-driven cancer	SHP2i monotherapy (BBP-398)	>500k							
		SHP2i combo therapy (BBP-398)	>500k							
	KRAS-driven cancer	KRAS G12C dual inhibitor	>500k							
		PI3Kα:RAS Breaker	>500k							
		KRAS G12Di	>500k							
	Solid tumors	GPX4i	>500k							



 Featured Programs

¹China + Japan patient population

KRAS mutant-driven cancers



Basia
Living with pancreatic cancer (>90% KRAS-driven)

Prevalence
>500k
US & EU

Pathophysiology
RAS is the most frequently mutated oncogene, leading to abnormal cell proliferation and survival

Program Highlights

- G12C dual inhibitor**
✓ MOA: first to directly bind and inhibit both GTP (active) and GDP (inactive) states of KRAS^{G12C}
- PI3K α :RAS Breaker**
✓ MOA: first to block RAS-driven PI3K α activation with the potential to avoid adverse effects on glucose metabolism
- G12D inhibitor**
✓ MOA: directly bind and inhibit KRAS^{G12D} - the single most prevalent KRAS mutant

MOA = mechanism of action

Partnerships afford us exceptional collaborators and resources

Research Capabilities

G12C Dual Inhibitor

PI3K α :RAS Breaker



- Partnership with the National RAS Initiative, including **60 of the world's foremost academic RAS researchers**
- Cutting edge RAS **structural biology expertise**
- Utilization of **cutting-edge instrumentation and techniques**, as well as the **expertise** to lead experiments

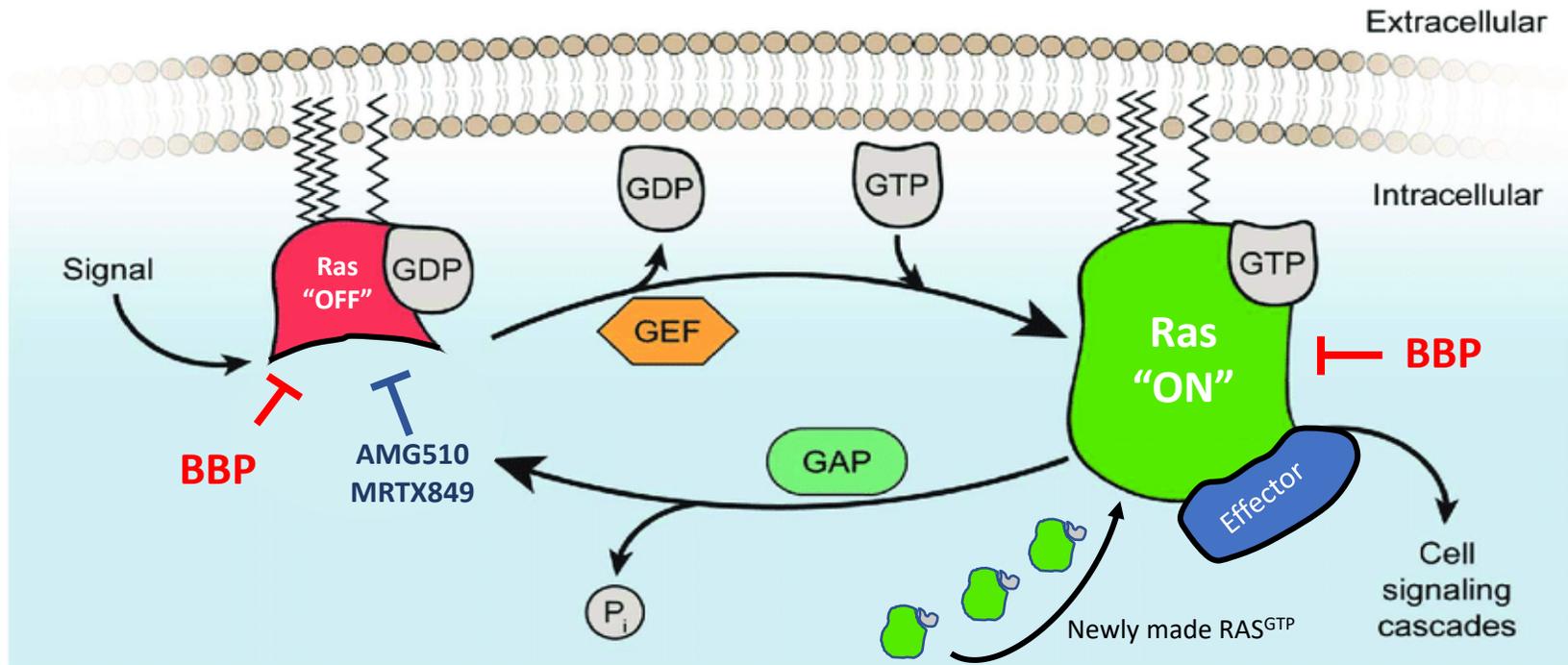
- Home to Sierra: the **world's 3rd fastest computing system**
- Enables **multi-microsecond molecular dynamics simulations** of protein complexes, and highly efficient in silico docking simulations
- This computing power, combined with RAS structural biology expertise at the NCI, delivers **unique insights that fuel our drug design**

We hypothesize that a compound that inhibits both GTP (active) and GDP (inactive) forms of KRAS^{G12C} will be superior to one that only inhibits the latter

Research Capabilities

G12C Dual Inhibitor

PI3K α :RAS Breaker



	GTP (active) / GDP (inactive) dual inhibitor e.g. BBP compounds	GDP (inactive) inhibitors e.g. AMG510, MRTX849
1	Blocks oncogenic signaling from KRAS ^{G12C} GTP (active) ✓	
2	Prevents KRAS ^{G12C} GDP (inactive) from cycling to KRAS ^{G12C} GTP (active) ✓	✓
3	Prevents resistance from residual KRAS ^{G12C} GTP (active) signaling ✓	

Source: Adapted from Schoneborn & Heumann, IJMS, 2018. Note: Conclusions based on preclinical models

BridgeBio G12C inhibitors modify both GTP (active) and GDP (inactive) forms of KRAS^{G12C}

Research Capabilities

G12C Dual Inhibitor

PI3K α :RAS Breaker

					
			BBP	AMG510	MRTX849
% modified	KRAS ^{G12C} GTP (active)	15'	100	0	0
		120'	100	0	0
	KRAS ^{G12C} GDP (inactive)	15'	100	80	73
		120'	100	83	80
KRAS ^{G12C} : RAF1 Effector Binding IC ₅₀ (nM)			35	>100,000	20,000
H358 pERK IC ₅₀ @ 30' (nM)			8	50	310

Multiple series of dual inhibitors progressing to identify development candidate

Note: Conclusions based on preclinical models

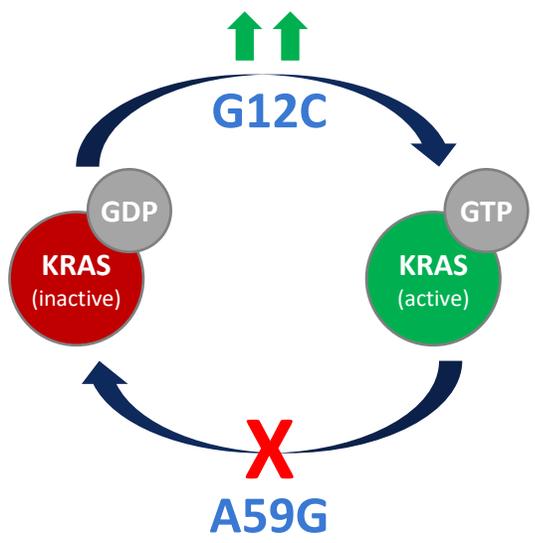
RAS-GTP “locked” mutant A59G, provides strong evidence for cellular GTP-state inhibitor activity

Research Capabilities

G12C Dual Inhibitor

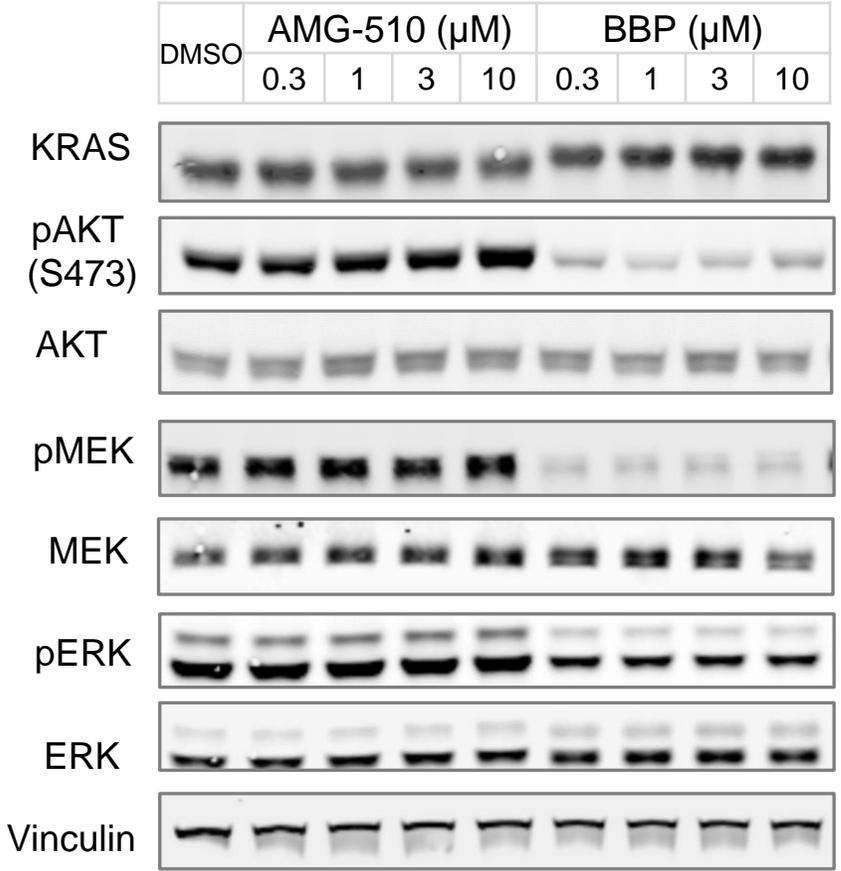
PI3K α :RAS Breaker

Impact of KRAS Mutations on Nucleotide Turnover



A59G is a 'transition state' mutant that abrogates GTPase activity and locks KRAS in GTP-state

KRAS^{G12C/A59G}



Strong pAKT, pMek and pERK inhibition observed with BBP KRAS-GTP/GDP dual inhibitor

Note: Conclusions based on preclinical models

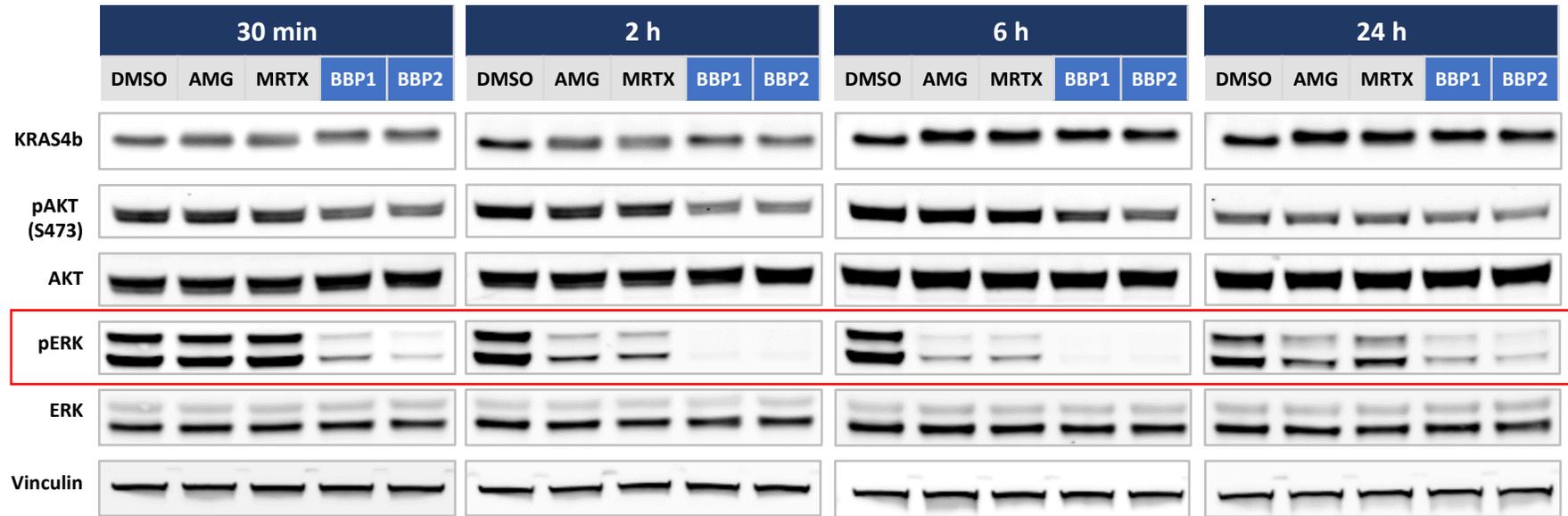
BridgeBio G12C dual inhibitors engage KRAS^{G12C} more quickly and potently than inhibitors that only target the GDP (inactive) form

Research Capabilities

G12C Dual Inhibitor

PI3K α :RAS Breaker

KRAS^{G12C} homozygous, p53 mutant CRC



GTP/GDP dual inhibitors:

- ✓ Quickly engage the target because they do not depend on nucleotide cycling to reveal the substrate
- ✓ Show faster and greater inhibition of pERK and pAKT than GDP (inactive) inhibitors

Note: Conclusions based on preclinical models

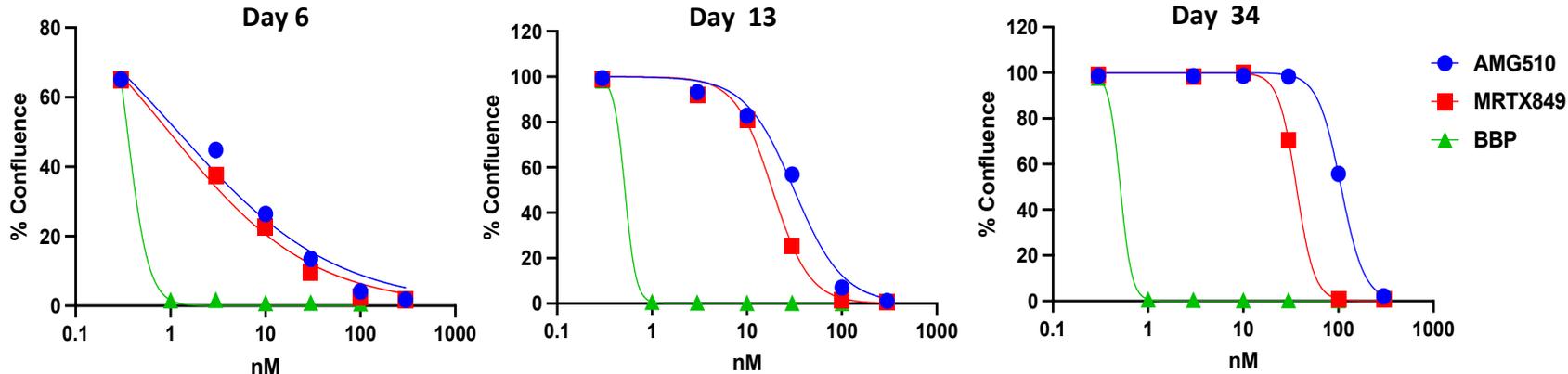
BridgeBio G12C dual inhibitors are more potent and retain activity compared to inhibitors that only target the GDP (inactive) form

Research Capabilities

G12C Dual Inhibitor

PI3K α :RAS Breaker

Clonogenic Assay



% Confluence (IC₅₀, nM)

	BBP	AMG510	MRTX849
Day 6	< 1	7	5
Day 13	< 1	32	19
Day 34	< 1	107	36

GTP/GDP dual inhibitors:

- ✓ Potently inhibit colony formation
- ✓ Retain potent activity suggesting that inhibiting both states of mutant KRAS reduces or delays development of resistance

Note: Conclusions based on preclinical models

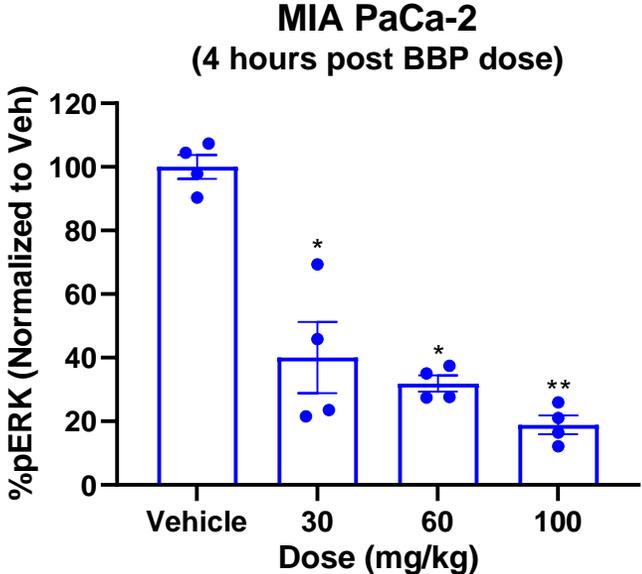
BridgeBio dual GTP/GDP inhibitors show potent and sustained inhibition of pERK in vivo

Research Capabilities

G12C Dual Inhibitor

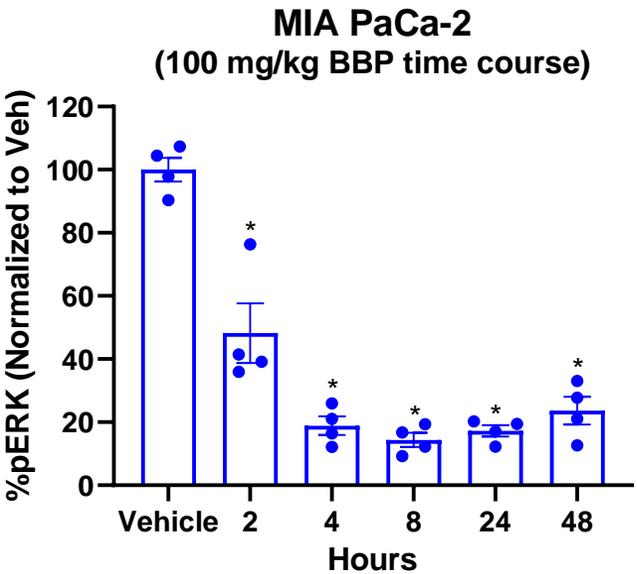
PI3K α :RAS Breaker

Dose Response



One-way ANOVA with Tukey's test vs vehicle *p<0.001, **p<0.0001

Time Response



One-way ANOVA with Tukey's test vs vehicle *p<0.0001

GTP/GDP dual inhibitors:

- ✓ Dose dependently inhibit pERK
- ✓ Provide sustained pERK inhibition over 48 hours after a single dose

Note: Conclusions based on preclinical models

BBP induces tumor regressions and is well tolerated in the MIA PaCa-2 CDX model

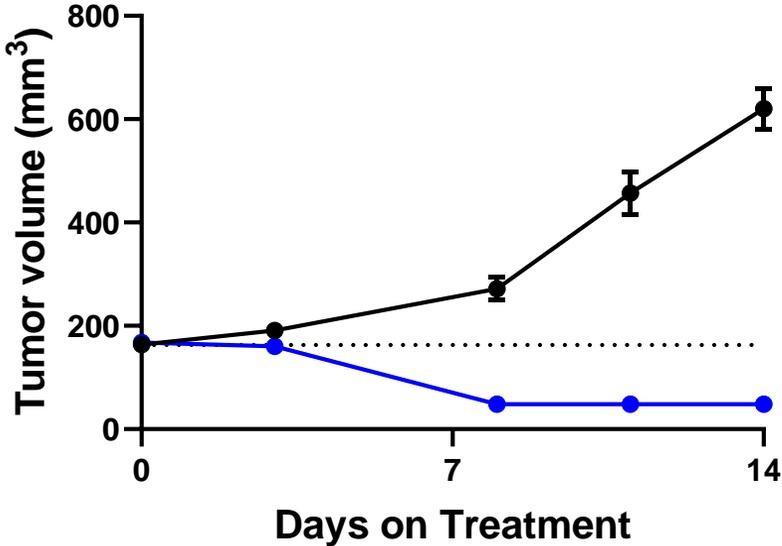
Research Capabilities

G12C Dual Inhibitor

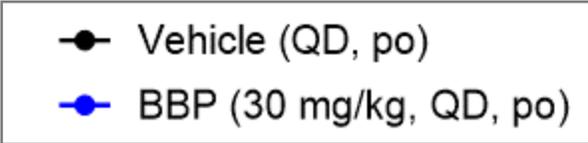
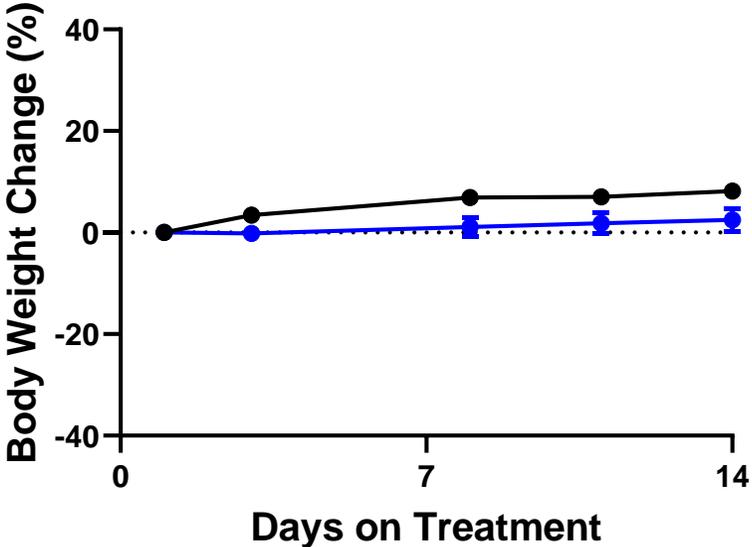
PI3K α :RAS Breaker

MIA PaCa-2 pancreatic CDX (KRAS^{G12C})

Tumor volume



Body weight % change



Note: Conclusions based on preclinical models

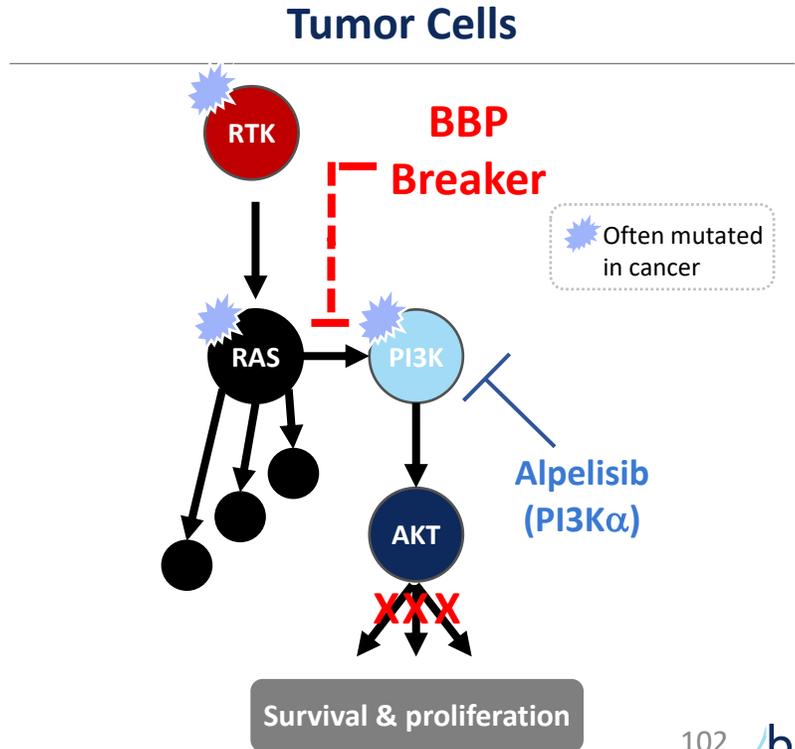
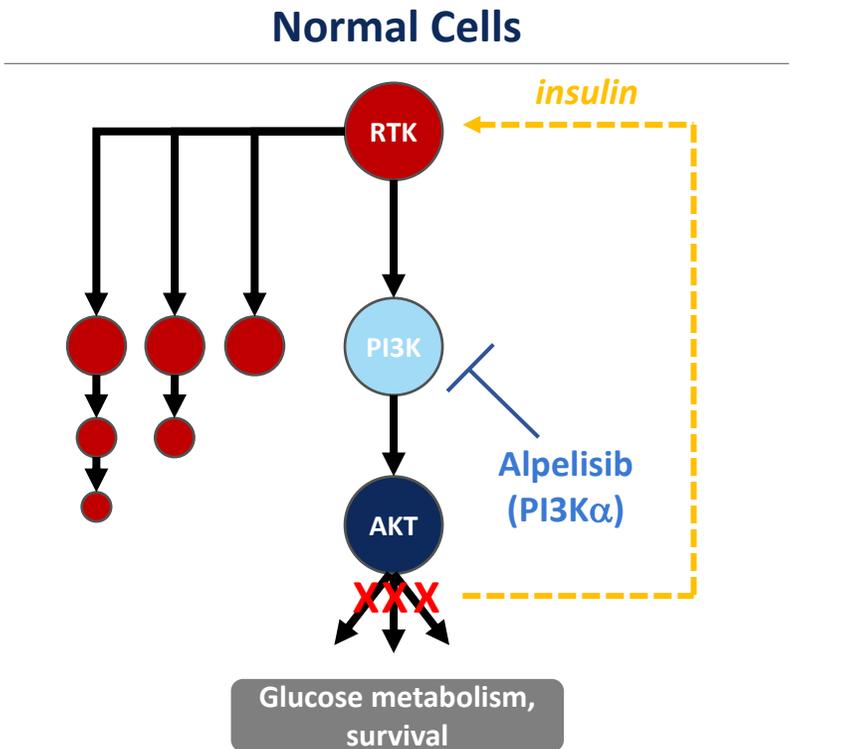
Novel approach to target PI3K α is tumor cell specific and differentiates from kinase inhibitors

Research Capabilities

G12C Dual Inhibitor

PI3K α :RAS Breaker

- PI3K α kinase inhibitors *block normal cell signaling* as well as RAS-driven PI3K α pathway activation in tumor cells, resulting in *dose-limiting hyperglycemia and insulin-driven resistance*
- Our novel approach of inhibiting PI3K α :RAS PPI with a “*PI3K α Breaker*” should avoid hyperglycemia and insulin-driven resistance by specifically targeting tumor cells and may provide multiple therapeutic opportunities:
 - *Tumors with RAS or PI3K α helical mutations and RTK mutant/amplified drivers*
 - *Potential combination with ERK pathway inhibition (BRAFi, MEKi, ERKi, KRAS^{G12C}i)*



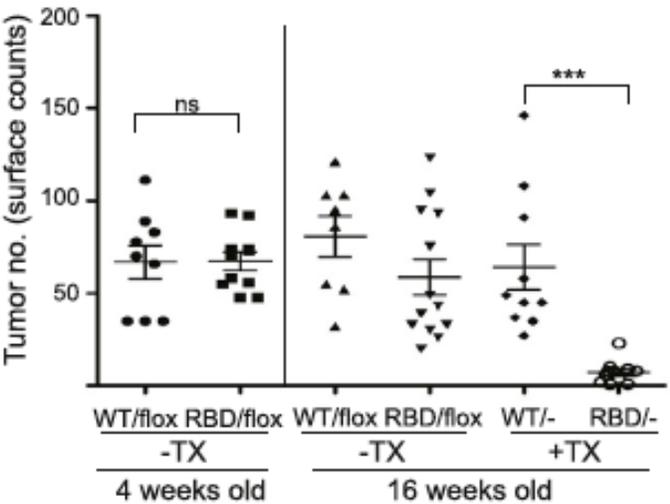
Impairing the PI3K α :RAS interaction blocks oncogene-driven NSCLC tumor growth in vivo

Research Capabilities

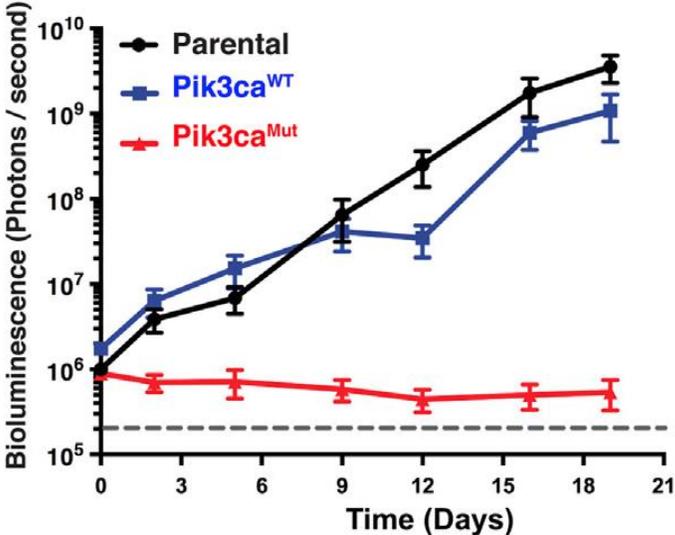
G12C Dual Inhibitor

PI3K α :RAS Breaker

KRAS^{G12D}-driven tumor growth is inhibited in mice with T208D and K227A mutations in the RAS-Binding Domain (RBD) of PIK3CA



Mutating the RBD of PIK3CA in the EGFR^{T790M/L858R} mutant NSCLC model H1975 abrogates xenograft tumor growth



- Mutating the RBD of PIK3CA in NSCLC models has profound effects on EGFR- and KRAS^{G12D}-driven tumorigenesis
- RBD mutant mice have normal glucose metabolism

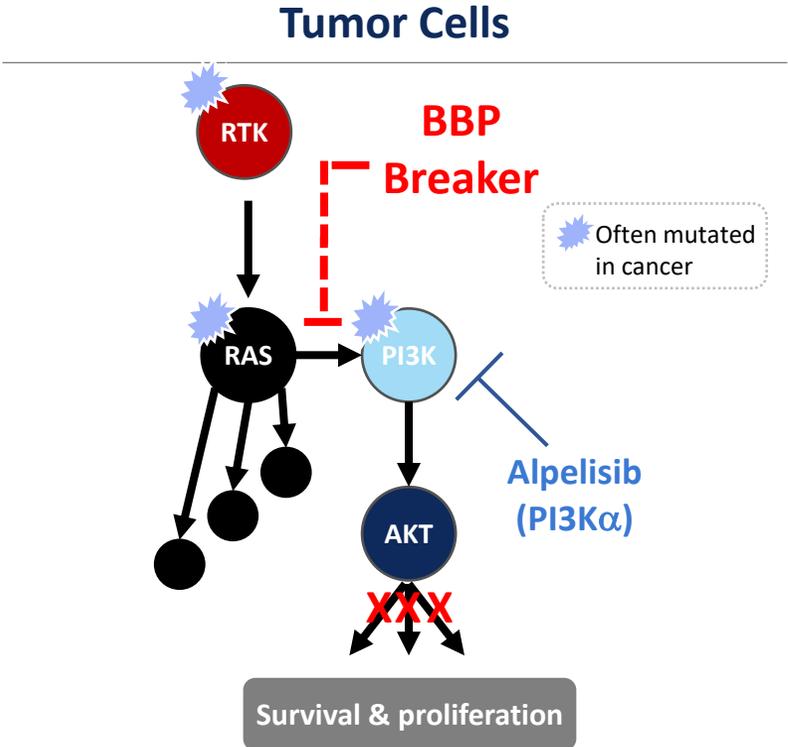
Sources: Gupta et al Cell 2007; Castellano et al Cancer Cell 2013; Murillo et al., Cell Reports 2018

BridgeBio has discovered potent and selective PI3K α :RAS breakers

Research Capabilities

G12C Dual Inhibitor

PI3K α :RAS Breaker



- Structural insights provide a novel approach to develop PI3K α :RAS breakers
- PI3K α :RAS breakers selectively bind to PI3K α
 - PI3K α amino acid sequence in the region of the binding pocket is unique amongst all the isoforms
 - No binding affinity to KRAS
- PI3K α :RAS breakers do not affect kinase activity of PI3K α

	BBP	Alpelisib
PI3K α Binding (IC ₅₀ , nM)	177	N/A
pAKT (IC ₅₀ , nM)	21	169
Cell Viability	134	744

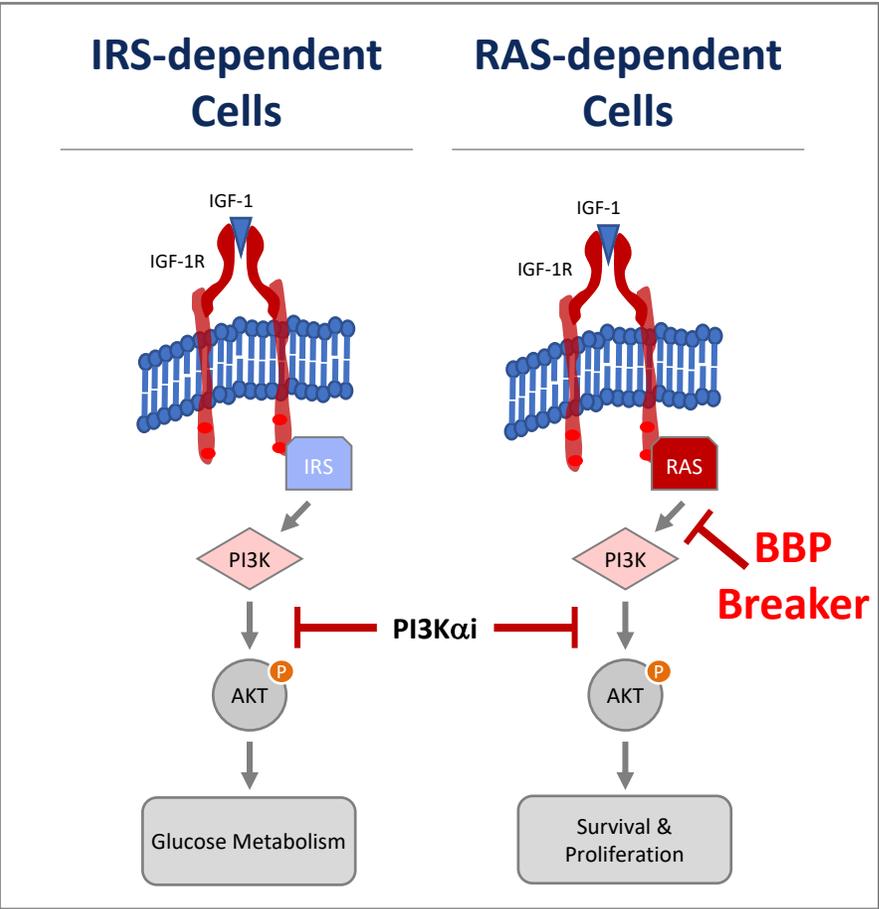
Multiple series of compounds have been identified as potent PI3K α :RAS PPI inhibitors

Cellular experiments show that only PI3K α breaker differentiates between RAS and IRS-driven pAKT activation

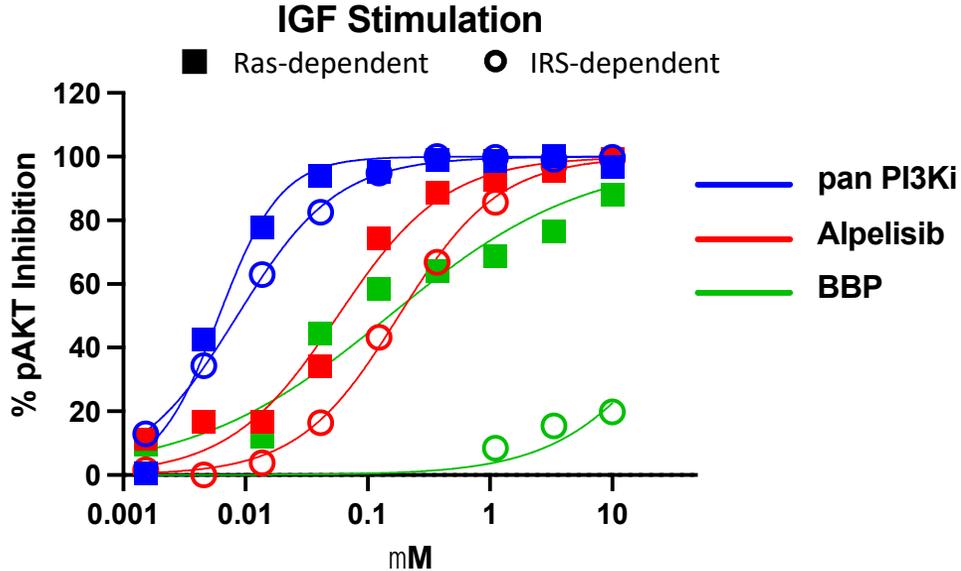
Research Capabilities

G12C Dual Inhibitor

PI3K α :RAS Breaker



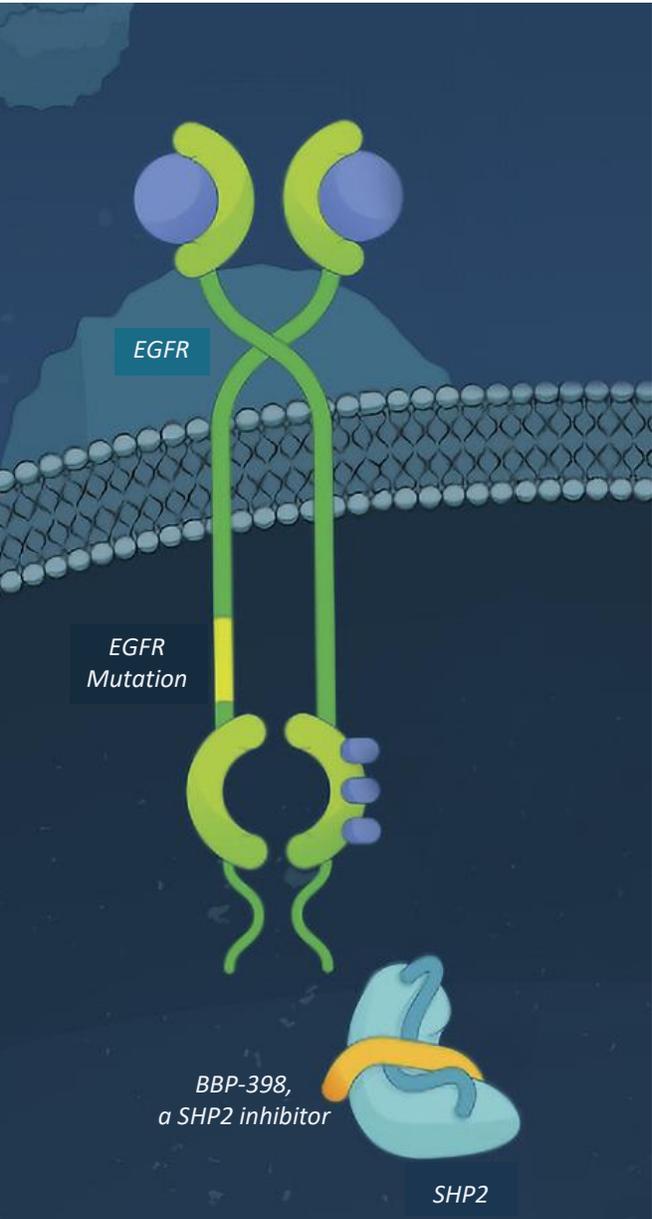
% AKT inhibition in cells treated with IGF



These data suggest that PI3K α breakers may avoid the on-target hyperglycemia associated with PI3K α kinase inhibitors

IRS – insulin receptor substrate; IGF-1 – insulin-like growth factor-1

BBP-398: SHP2 inhibitor for treatment resistant cancer



Prevalence
>500k
US & EU

Pathophysiology
SHP2 acts upstream of RAS/ERK in RTK and cytokine signaling to regulate cell proliferation, survival, adhesion, and migration

Program Highlights

- ✓ BBP-398 is a selective, orally bioavailable, allosteric SHP2 inhibitor
- ✓ Potential to be best-in-class based on optimal PK profile that may enable tolerable once-daily dosing
- ✓ Monotherapy dose escalation is ongoing with plans to initiate combination studies next year

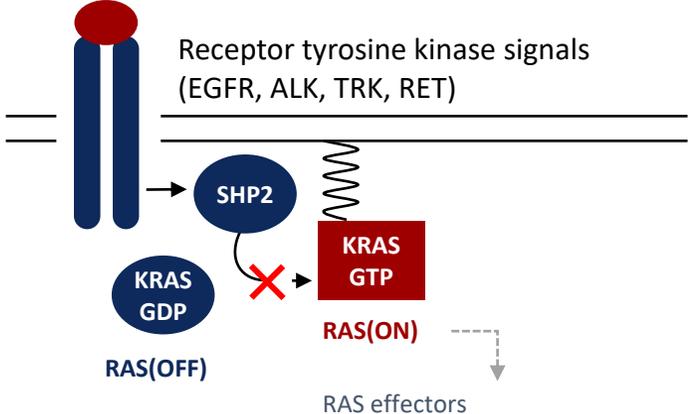
BBP-398 shows best-in-class potential in a large cancer market

Overview

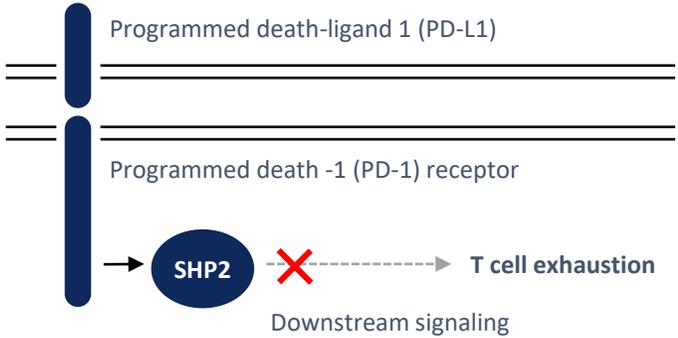
Preclinical Data

Phase 1 Dose Escalation

Our SHP2i blocks downstream MAPK signaling and abrogates T cell exhaustion



Tumor cell proliferation and survival



We believe BBP-398 has the ideal properties for combination with a multitude of other therapeutic classes

- ✓ Human half life: ~10-15 hours
- ✓ Optimal PK profile which may enable better tolerability in combination

Initial clinical combinations of focus based on SHP2i preclinical data

Combination Agent	Patient Population ¹
KRAS G12Ci	70,000
EGFRi  (LianBio)	150,000
PD-1  Bristol Myers Squibb™	700,000

¹US incidence estimated from SEER, TCGA; all scaled for WW incidence

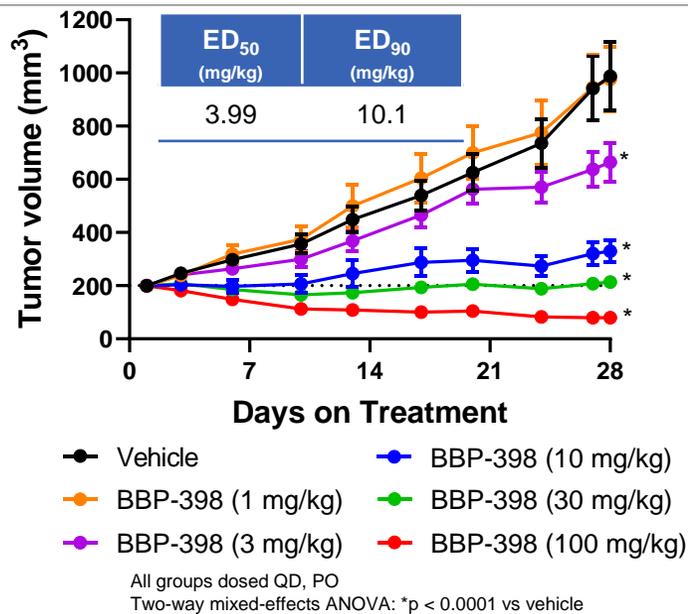
Predicted BBP-398 pharmacokinetics support once daily oral administration to achieve target coverage

Overview

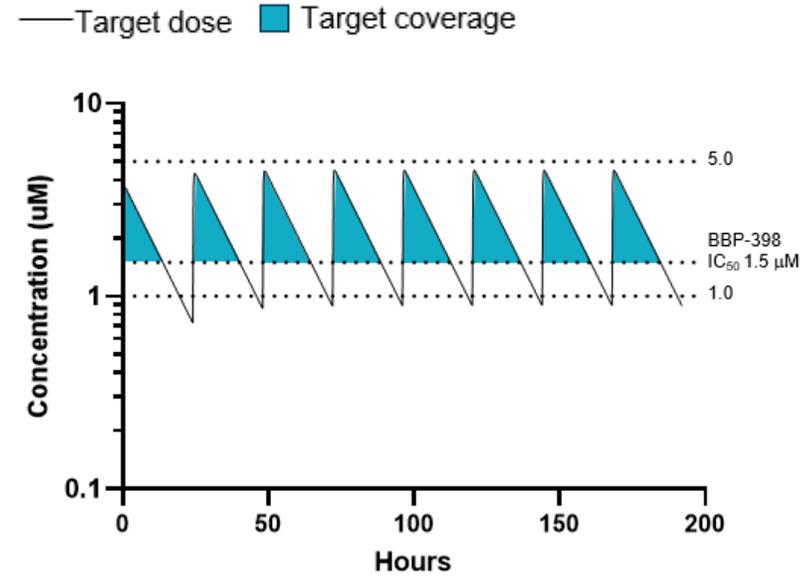
Preclinical Data

Phase 1 Dose Escalation

HCC827 (EGFR^{ex19del} & EGFR^{amp})
- NSCLC CDX



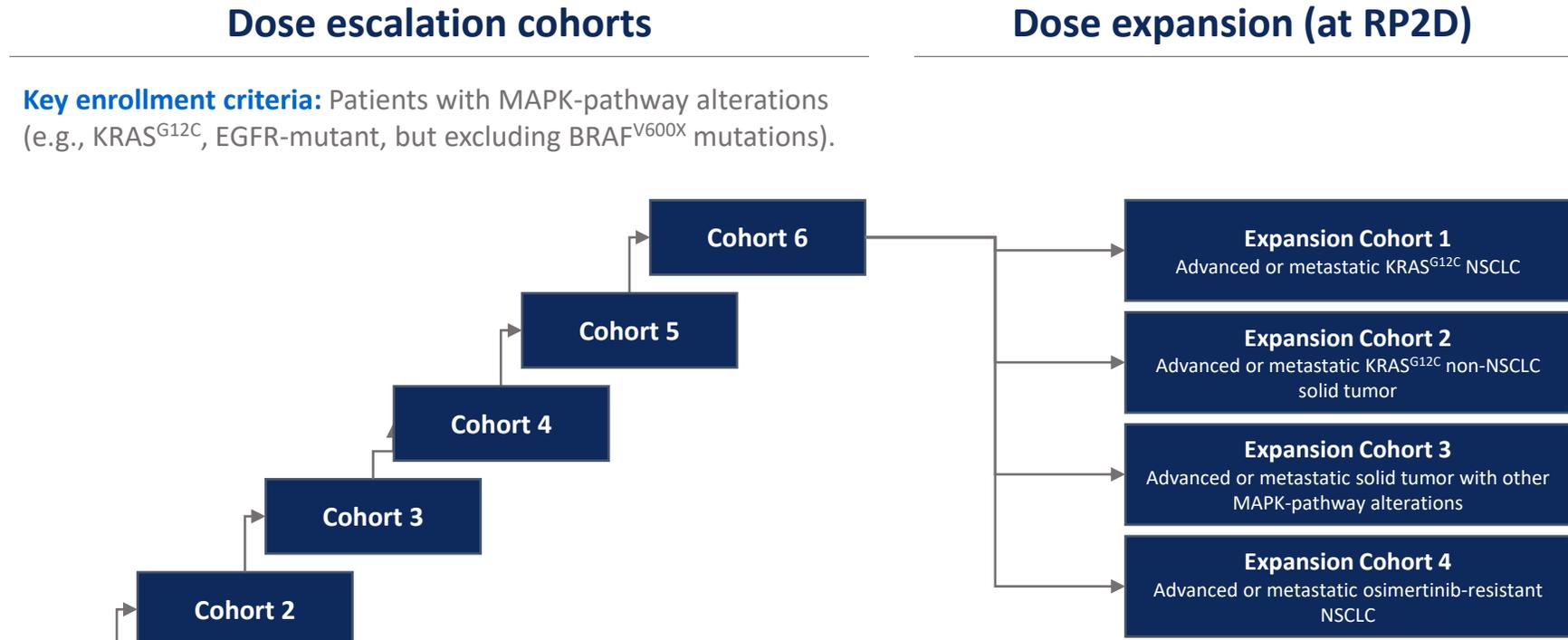
BBP-398 steady-state PK simulation for optimal efficacy



Predicted clinical exposure supports coverage of efficacy target in patients may be achieved with continuous once daily dosing

Phase 1 dose escalation with BBP-398 is ongoing: Observed PK/PD is in-line with preclinical predictions

- Overview
- Preclinical Data
- Phase 1 Dose Escalation**



BBP-398 dose escalation ongoing
Dose expansion at RP2D and combinations with Nivolumab and G12Ci planned for 2022

Precision oncology summary

BridgeBio Oncology

- Infigratinib approved for 2nd line FGFR2 fusion cholangiocarcinoma with multiple late-stage studies ongoing
- Identified multiple series of differentiated novel KRAS^{G12C} GTP/GDP inhibitors
- Identified multiple series of differentiated novel PI3K α :RAS Breakers
- Progressing potentially best-in-class SHP2 inhibitor BBP-398 with differentiated pharmacokinetic profile that may enable once-daily dosing in combination studies

2022 Targets

- RAS development candidate
- Present BBP-398 Phase 1 monotherapy data
- Initiate BBP-398 combination studies (KRAS G12Ci, IO, EGFRi)

BridgeBioX

Charles Homcy, M.D.
Chairman of Pharmaceuticals



bridgebio lab at Stanford

Our discovery lab is located in the **Stanford** Life Sciences District – we aim to create an academic/industry hybrid environment, and foster a culture driven by **intellectual curiosity** and a dedication to **patient impact**

