

Protein degraded. Disease targeted. Lives transformed.



August 2023

Forward-looking Statements and Intellectual Property

Forward-looking Statements

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C4T is a Leader in Delivering on the Promise of Targeted Protein Degradation

Our Mission

To deliver on the promise of targeted protein degradation science to create a new generation of medicines that transform patients' lives





Robust Pipeline of Degrader Medicines Pursuing Multiple Targets in Oncology

Program	Target	Indications	Discovery	Pre-clinical	Early phase development	Late phase development	Rights
CFT7455	IKZF1/3	Multiple Myeloma & Non- Hodgkin's Lymphoma					
CFT8634	BRD9	Synovial Sarcoma & SMARCB1-null Cancers					
CFT1946	BRAF-V600	V600 Mutant Cancers					
CFT8919 ¹	EGFR L858R	Non-Small Cell Lung Cancer					
Chromatin Regulating Targets		Various Cancers					
Oncogenic Signaling Targets		Various Cancers					
Transcription Factor Targets		Various Cancers					

Two Clinical Readouts On Track by Year-End

CFT7455 IKZF1/3	Present Phase 1 dose escalation data from the Phase 1/2 trial 2H
CFT8634 BRD9	Present Phase 1 dose escalation data from the Phase 1/2 trial 2H
CFT1946 BRAF V600	 ✓ First patient dosed in the Phase 1/2 trial ✓ Present new preclinical data
CFT8919 EGFR L858R	 ✓ Secure China Partnership ✓ Achieved FDA clearance of IND
	Cash Runway into 2H 2025 ¹

CFT7455 Targeting IKZF1/3

Multiple Myeloma (MM) & Non-Hodgkin's Lymphoma (NHL)



CFT7455 Phase 1/2 Trial Progressing through Dose Escalation

Arm A: R/R Multiple Myeloma & Non-Hodgkin's Lymphoma Monotherapy¹ N = 5

> Schedule: 28-day cycle/7-days off

> > <u>Status</u>: Complete

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1. 28-day cycle /7 days off dose limiting toxicity (DLT) window

2. Combination therapy arms will open once the selected CFT7455 dose level has been cleared for safety

6-12 patient food effect enrichment arm also included during escalation, not pictured in schema

Single Agent CFT7455 Arm A Data Demonstrated Potent On Target Degradation; Dosing Schedule Modified to Improve Therapeutic Index



0 Patient 1 Percent Change dFLC Patient 2 -25-Patient 3 -50 -75 -100 20 10 15 n 5 Day 15 AUC0-24h (hr*ng/mL) Modified Dosing Schedule with 14 Days Off 10-· 100 - 75

CFT7455 Concentration (ng/mL) IKZF3 Degradation % 50 = issue 0.1 LLOQ Concentration 0.01 at which stasis 14 21 observed in 28 H929 xenoaraft Time (days) Simulated 50 µg Plasma PK Simulated IKZF3 Degradation 50 µg

Meaningful Reductions in dFLC

Multiple Paths to Success for CFT7455 across Evolving Multiple Myeloma Landscape

Potential for CFT7455 to replace other IKFZ1/3 degraders and become a backbone therapy

Potential to combine CFT7455 with nextgeneration therapies

IKZF1/3 degrader competitive landscape

1L	2L	3L	4 L	
 Lenolidomic dexametho approved 	de + Isone	• Pomc dexai	alidomide + methasone approved	
 Iberdomide + dexamethasone in development 		 Mezig dexai devel 	gdomide + methasone in lopment	
	CF	T7455 -		

CFT7455, as one molecule, has the potential to be utilized across all lines of therapy with or without dexamethasone

Bi-specific T-Cell Engagers	CAR T-Cell Therapies
Antibody-Drug Conjugates	BCMA-CD38
Monoclonal Antibodies	Small Molecule Inhibitors/ Modulators



CFT8634 Targeting BRD9

Synovial Sarcoma & SMARCB1-Null Solid Tumors



Oncogenic SS18-SSX Fusion Leads to BRD9 Dependency in Synovial Sarcoma



cBAF, canonical BAF; ncBAF, noncanonical BAF; pBAF, polybromoBAF.

Incorporation of SS18-SSX fusion results in eviction of SMARCB1

- cBAF complex compromised
- Oncogenic state

Inactivation of SMARCB1 leads to dependency on ncBAF complex

- BRD9 is uniquely present in ncBAF
- Synthetic lethal dependency on BRD9 in synovial sarcoma and other SMARCB1-deficient cancers

CFT8634 Phase 1/2 Trial Progressing through Dose Escalation



BRD9 Previously Considered an Undruggable Target where Inhibitors are Ineffective for Synovial Sarcoma

Unmet Need

No approved therapies specifically for synovial sarcoma

Current treatment options offer limited benefit:

- **PFS of ~7 months**¹ in the front-line setting
- PFS ~5 months² in the relapsed refractory setting

Degrader Rationale

Oncogenicity of BRD9 depends on protein function not addressed by traditional inhibitors



Key Properties of CFT8634

- Orally bioavailable
- Potent
- Selective



Sources:
 Wang BC, et al. Front Oncol. 2021;11:76228.
 Sleijfer S, et al. J Clin Oncol. 2009;27(19):3126-3132.
 Progression Free Survival (PFS)

CFT8634 Pharmacokinetic and Pharmacodynamic Data Supportive of Proof of Mechanism



Effective Degradation of a Previously Undruggable Target



Recent Publications Support Potential Additional Indications for BRD9



Interferonopathies

BRD9 is a druggable component of interferon-stimulated gene expression and antiviral activity

EMBOpress



Interferonopathies

BRD9 regulates interferon-stimulated genes during macrophage activation via cooperation with BET protein BRD4

PNAS

Ovarian Cancer



The bromodomain containing protein BRD-9 orchestrates RAD51–RAD54 complex formation and regulates homologous recombination-mediated repair

Nature Communications



Clear Cell Renal Cell Carcinoma

Aberrant activation of m6A demethylase FTO renders HIF2a^{low/-} clear cell renal cell carcinoma sensitive to BRD9 inhibitors

Science Translational Medicine



Prostate Cancer

BRD9 Is a Critical Regulator of Androgen Receptor Signaling and Prostate Cancer Progression

AACR Journals



Multiple Myeloma

BRD9 Is Essential for Ribosome Biogenesis and the Survival of Multiple Myeloma Cells

ASH Annual Meeting 2022

Currently Evaluating Opportunities for Indication Expansion



CFT1946 Targeting BRAF V600

Melanoma, Colorectal (CRC) & Non-Small Cell Lung Cancer (NSCLC)



Current Standard of Care BRAF Inhibitors Lead to Resistance

Unmet Need

Resistance to approved BRAF inhibitors results in a median PFS of less than 15 months²

Toxicity associated with inhibiting wild-type BRAF

Key Properties of CFT1946

• Orally bioavailable

Therapeutics

- Potent and selective against BRAF V600 mutant targets while sparing wild-type activity
- Preclinical activity in setting of resistance to BRAF inhibitors



Degrader prevents dimer formation and avoids paradoxical activation

Sources: 1. 2022 Market size from

2022 Market size from EvaluatePharma.

NIH SEER Database, Primary Literature Consensus. https://www.ncbi.nlm.nih.gov/pmc/articles/MC5931274/, https://pubmed.ncbi.nlm.nih.gov/26980021/

CFT1946 Shows Superior Efficacy Compared to Approved BRAF Inhibitor in Preclinical Models



CFT1946 is the First Clinical BRAF V600 Degrader



1. 28-day cycle/ dose limiting toxicity (DLT) window

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2. BRAF inhibitor naïve expansion arm to be aligned across global health authorities

Colorectal cancer (CRC); Recommended Phase 2 dose (RP2D); Non-Small Cell Lung Cancer (NSCLC)

CFT1946 Has the Potential to Address Multiple Tumor Types with BRAF V600 Mutations





::: C4

- 1. ACS Figures & Facts 2022: https://www.cancer.org/research/cancer-facts-statistics/all-cancer-facts-figures/cancer-factsfigures-2022.htm
- Owsley J, et al. Experimental Biology and Medicine. 2021;246(1):31-39
- heropeutics Paik, P. K., et al. Journal of clinical oncology: official journal of the American Society of Clinical Oncology. 2011; 29(15), © 2023 C4 Therapeutics, Inc. 20 2046-2051
 - Bylsma, L. C., et al. Cancer medicine. 2020;9(3), 1044–1057.

CFT8919 Targeting EGFR L858R

EGFR L858R + Non-Small Cell Lung Cancer (NSCLC)



Potential for CFT8919 to Improve Outcomes for NSCLC Patients with EGFR L858R Mutations

Osimertinib and other inhibitors provide suboptimal response for NSCLC patients with L858R mutation





Key Properties of CFT8919

- Orally bioavailable
- Potent and selective against L858R, regardless of secondary mutations
- Allosteric binding

Sources:
 Soria, J.-C. et al. NEJM 378, 113–125 (2018),
 Sher, T. et al, Mayo Clin. Proc. 83, 355-367 (2008),
 2022 market size from EvaluatePharma.

Mutant EGFR (mEGFR); Non-small cell lung cancer (NSCLC); Tryosine Kinase Inhibitor (TKI)

Strategic Partnership with Betta Pharmaceuticals in Greater China

Expert Partner in	Expedite
Greater China	Development
 Betta has a proven track record of developing NSCLC medicines including EGFR inhibitors 	 High prevalence of EGFR L858R driven NSCLC in Greater China allows for faster clinical development in target population

Expected Next Steps:

- Betta to file a CTA with the NMPA
- Phase 1 dose escalation to begin in Greater China



Deal Terms:

- \$10 million upfront and \$25 million equity investment
- C4T eligible to receive up to \$357 million in potential milestones and low to middouble-digit percent royalties on net sales in the licensed territories



EGFR L858R Driven NSCLC has Higher Prevalence in Asia

Non-Small Cell Lung Cancer (NSCLC)



~200K Patients in the US diagnosed in 2022



~693K Patients in Asia diagnosed in 2020

~10 – 15% Analysis of NSCLC patients have mutant EGFR (mEGFR) in the U.S. population

~40%

~40[%] of mEGFR NSCLC patients have the L858R activating mutation

Sources

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- Recondo 2018 Nature Rev Clin Oncol <u>https://www.nature.com/articles/s41571-018-0081-4</u>
- Globacan 2020
- American Cancer Society. Facts & Figures 2023. American Cancer Society.
- SEER Cancer Stat Facts: Lung and bronchus cancer. National Cancer Institute, <u>https://seer.cancer.gov/statfacts/html/lungb.html</u>.

• Mao 2021 Pathol Oncol Res (https://pubmed.ncbi.nlm.nih.gov/34257561/)

Melosky 2021 Mol Diagnosis Ther (https://link.springer.com/article/10.1007/s40291-021-00563-1)

CFT8919 is Selective for EGFR L858R and Active in a Setting of Osimertinib Resistance in Preclinical Models

Specific for EGFR Exon 21 Mutants



Active in setting of EGFR C797S





Thank You!



