



# Monofunctional Degradation Activating Compounds: From New Targets to the Clinic

TPD Summit, October 2022

Chris Nasveschuk on behalf of the C4 Team



# Forward-looking Statements and Intellectual Property

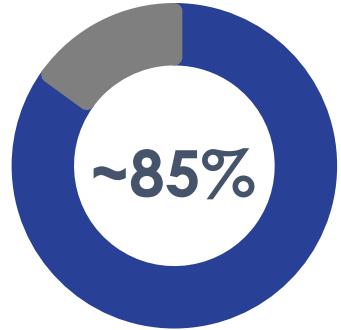
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## Intellectual Property

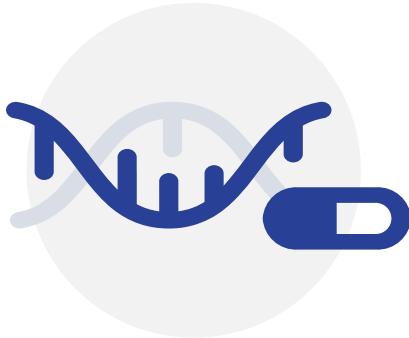
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# Targeted Protein Degradation Has the Potential to Transform Treatment of Disease



## TPD Has an Expansive Target Landscape

85% of proteins are currently undruggable or poorly drugged



## TPD Offers a Powerful Modality

Benefits of genetic knockdown with a small molecule approach

**C4T's TORPEDO platform creates therapeutic candidates that have the potential to improve patient care**



Overcome Resistance



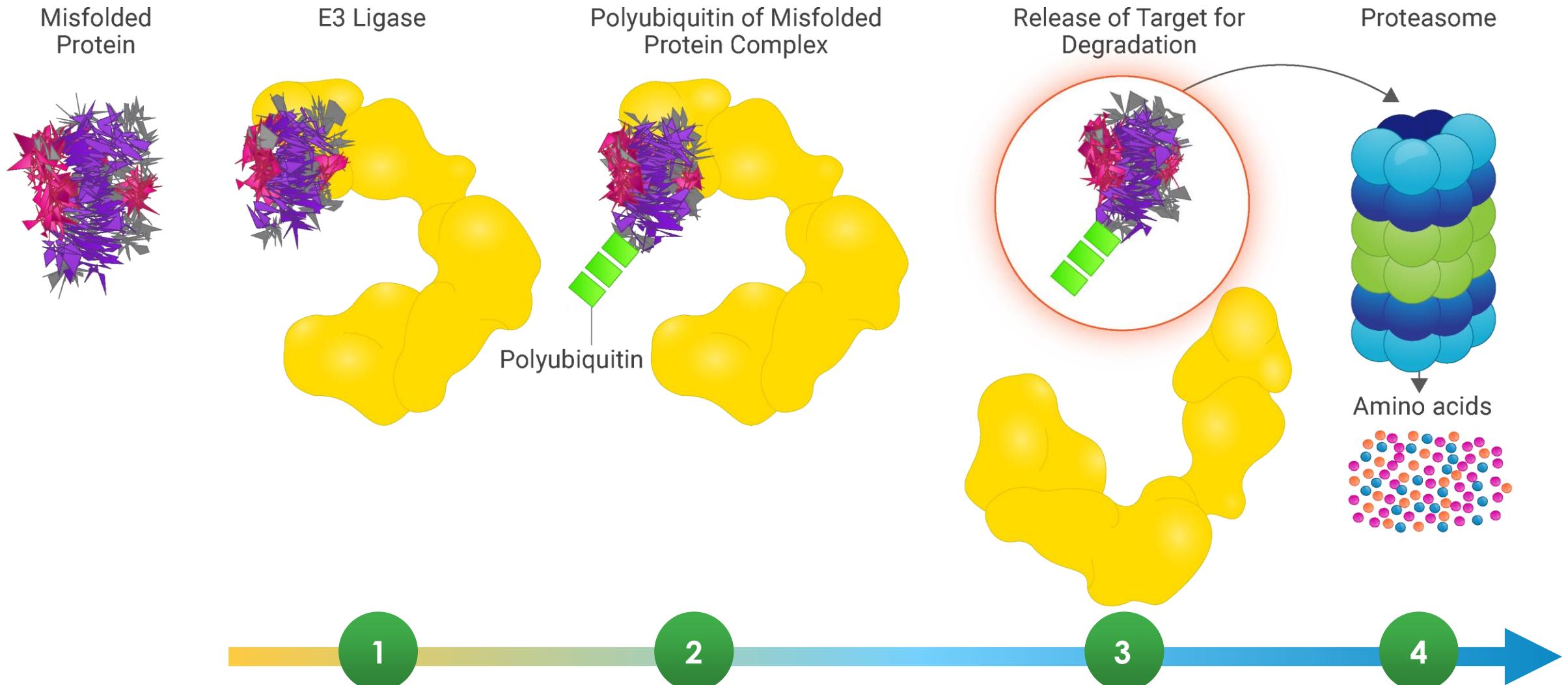
Drug Undruggable Targets



Improve Treatment Options

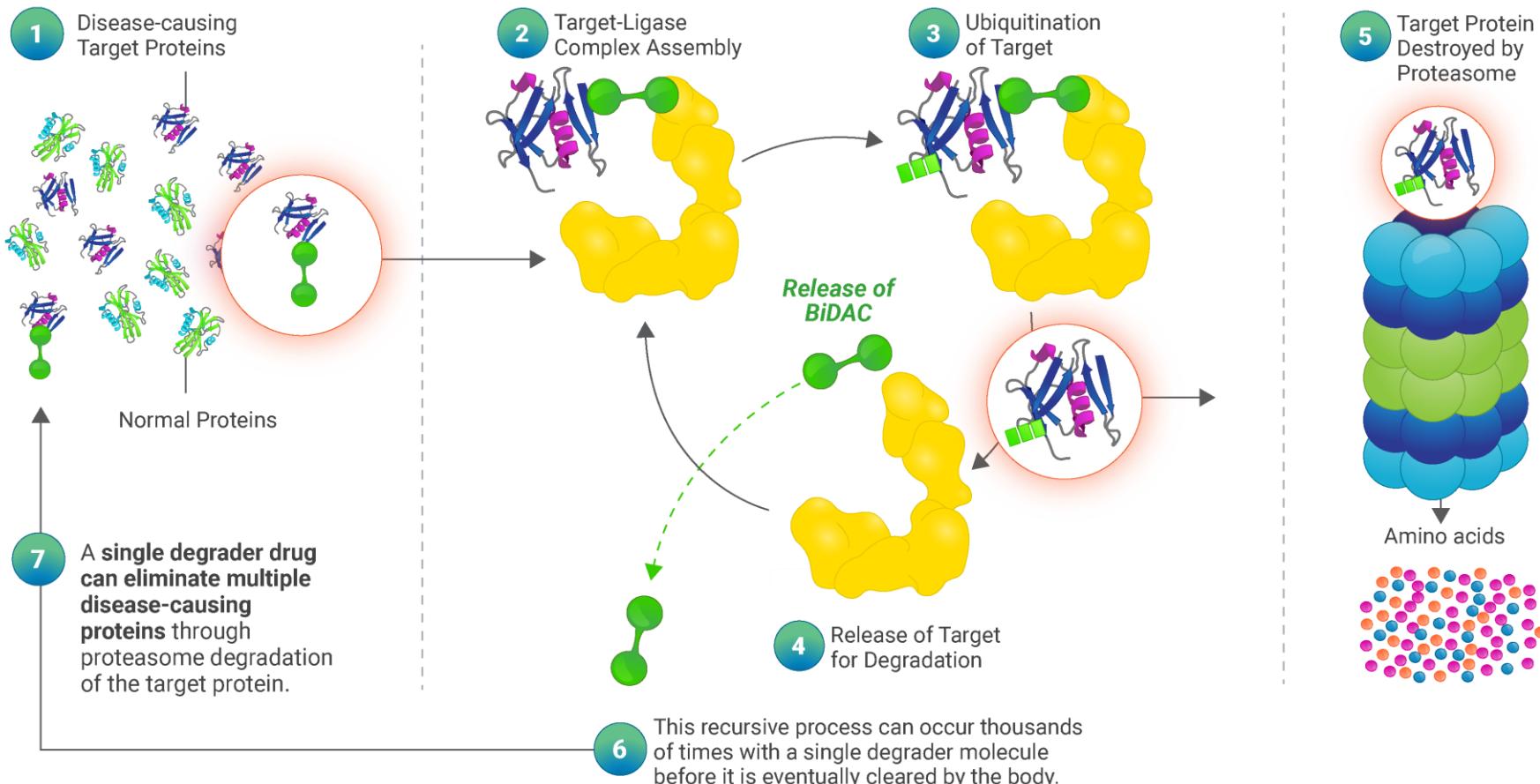
Source: Hopkins, A., Groom, C. The druggable genome. *Nat Rev Drug Discov* 1, 727–730 (2002).

# The Human Body Has A Natural Process to Destroy Unwanted Proteins



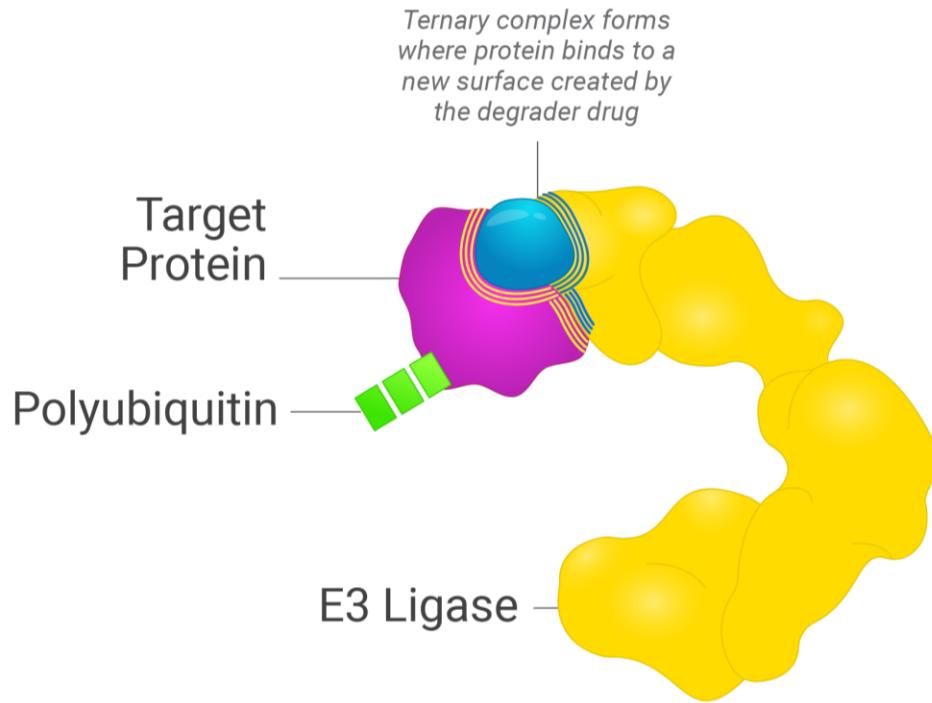
# Targeted Protein Degradation Leverages the Body's Natural Process to Destroy Disease-Causing Proteins

## Focus on Overall Catalytic Degradation

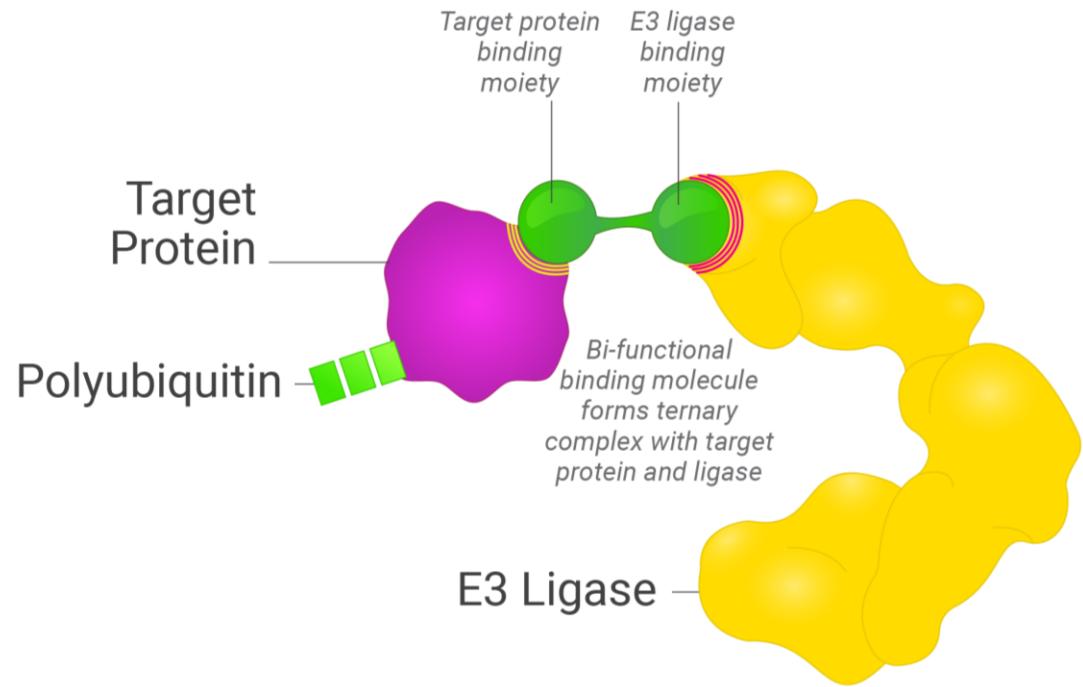


# TORPEDO Platform Offers Flexibility to Design MonoDAC and BiDAC Degraders

## MonoDAC Degrader

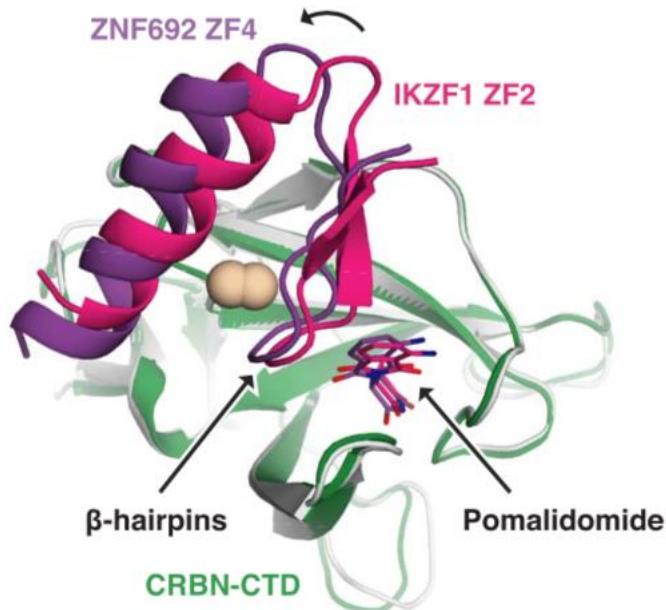


## BiDAC Degrader

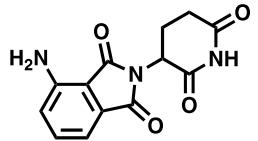


Flexibility to Address Different Targets with Tailored Approach

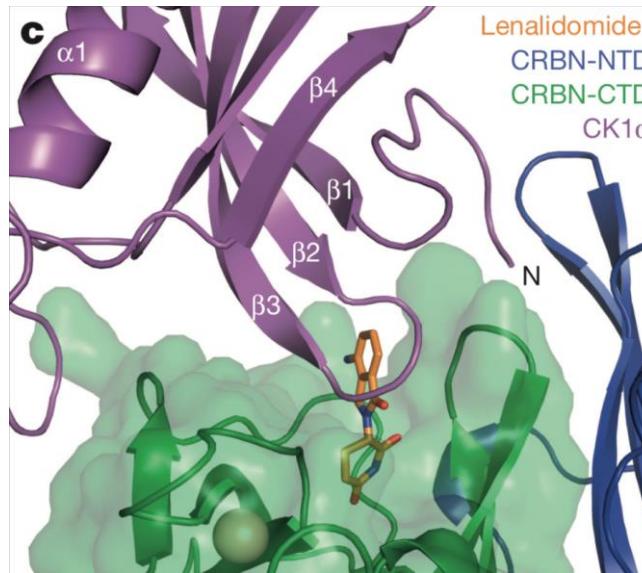
# Molecular Glue Degraders (MGDs) in the Literature



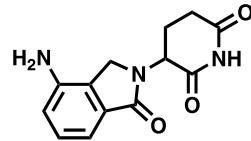
Thalidomide-related Zinc Finger degraders



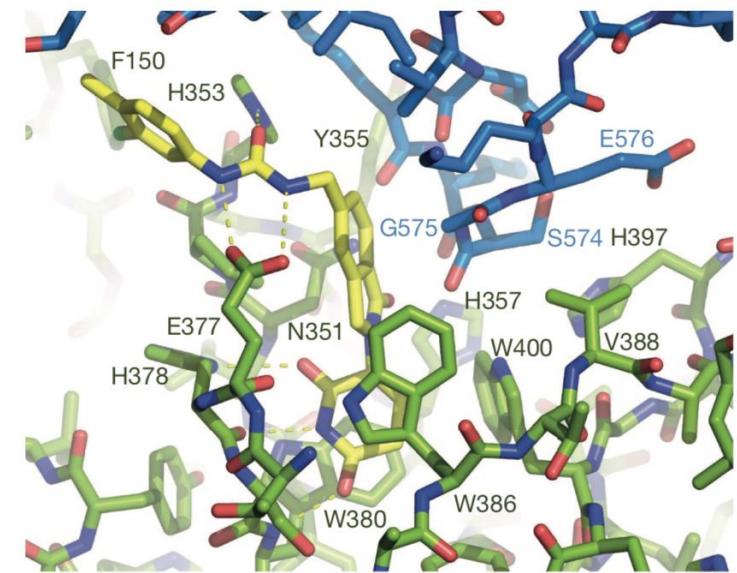
Science 2018, 362(6414), eaat0572



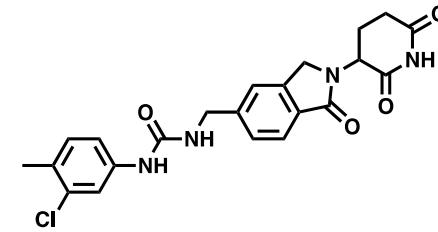
Lenalidomide, CK1- $\alpha$  degrader



Nature 2016, 532(7597), 127

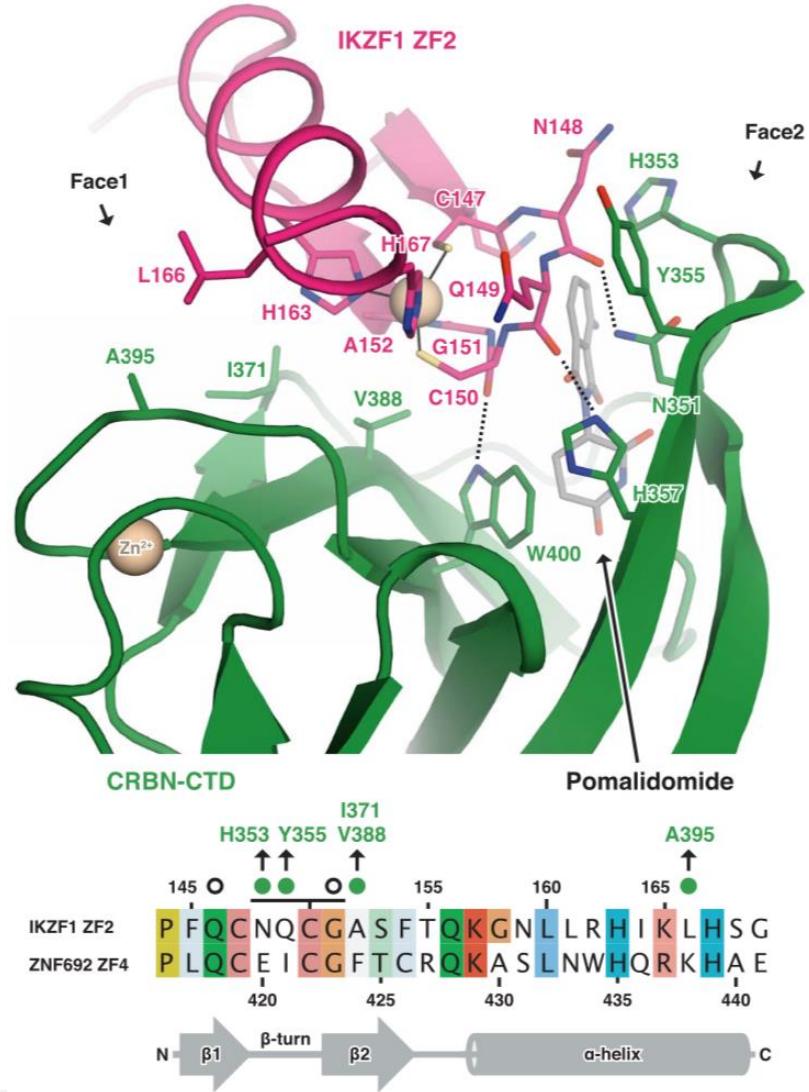
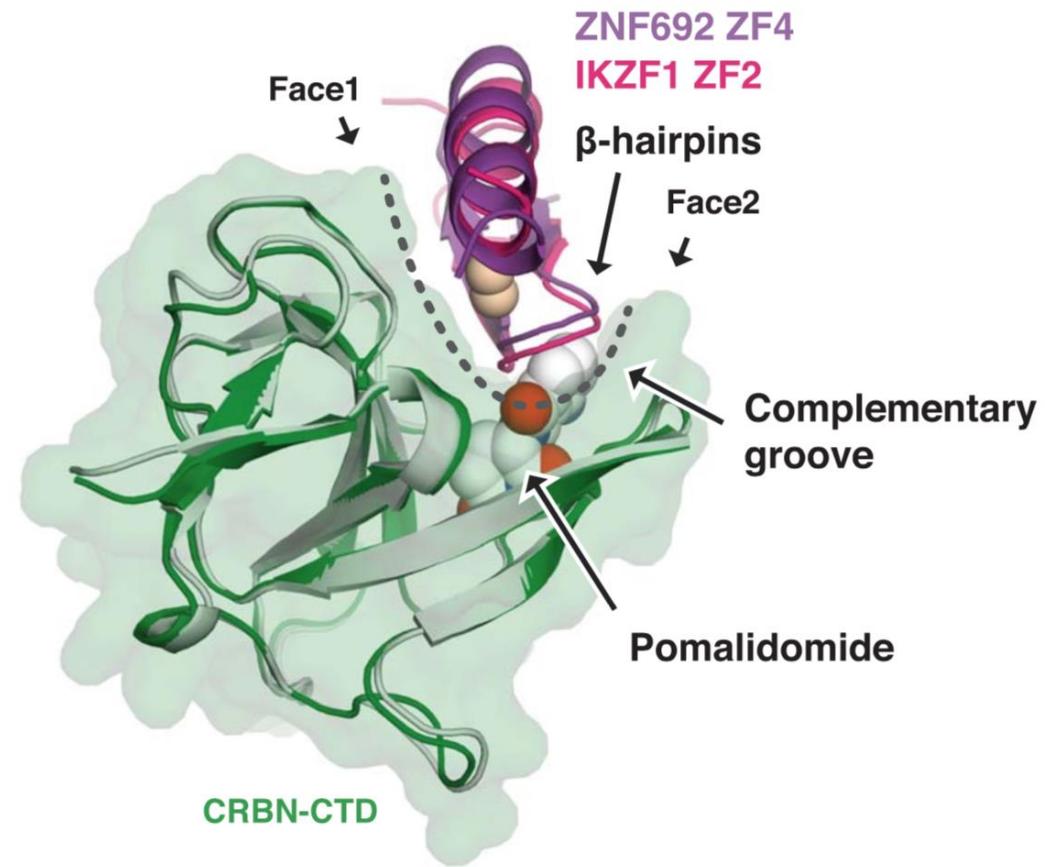


CC-885/90009 GSPT1 degraders



Nature 2016, 535(7611), 252

# MGDs Literature in the Literature



β-turn is the primary recognition element, secondary interactions play a role in activity and selectivity

Science 2018, 362(6414), eaat0572

# Why MonoDAC Degraders and Molecular Glue Degraders?

## Disease Target and Potential for New Medicines

- Access to Not-Yet-Drugged/or Unligandable Targets
- MonoDAC degraders and MGDs exhibit degradation-only pharmacology

## Drug Properties and Performance

- Access to degrader drug candidates within more traditional Rule of 5 physicochemical property space

MonoDAC Degraders and MGDs should be aligned to the right target and clinical opportunities



C4 Therapeutics

# Identification and Characterization of MonoDAC CFT7455

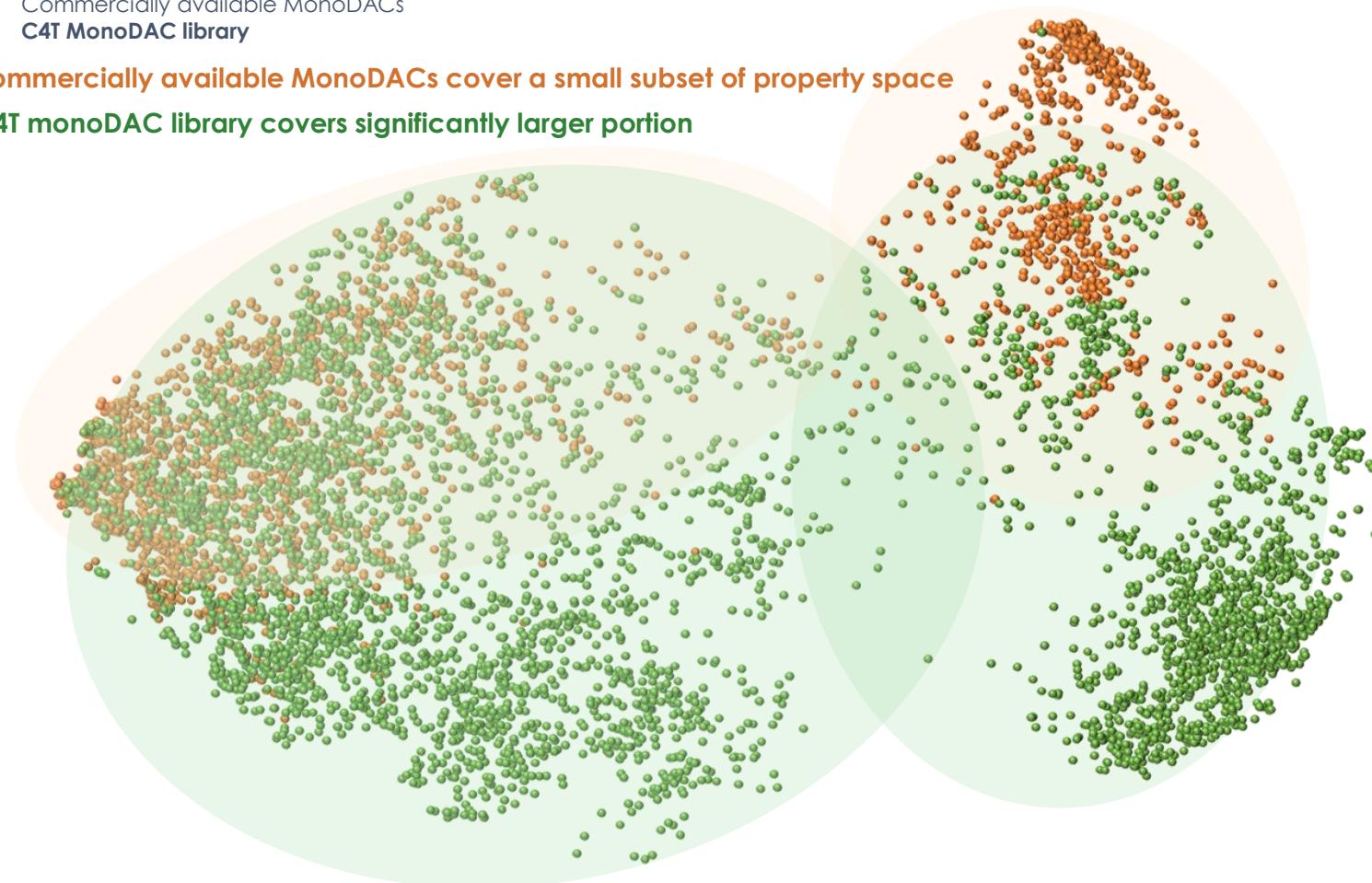


# C4T MonoDAC Library: Expanding the Cereblon Toolbox

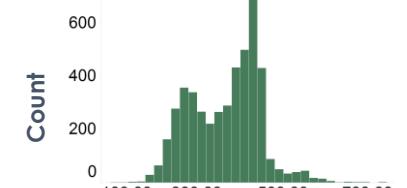
- Commercially available MonoDacs
- C4T MonoDAC library

Commercially available MonoDacs cover a small subset of property space

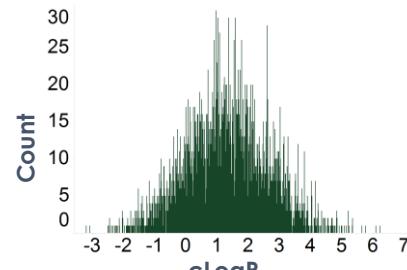
C4T monoDAC library covers significantly larger portion



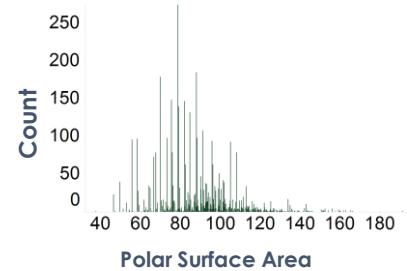
Drug-like property space



Molecular Weight



cLogP



Polar Surface Area

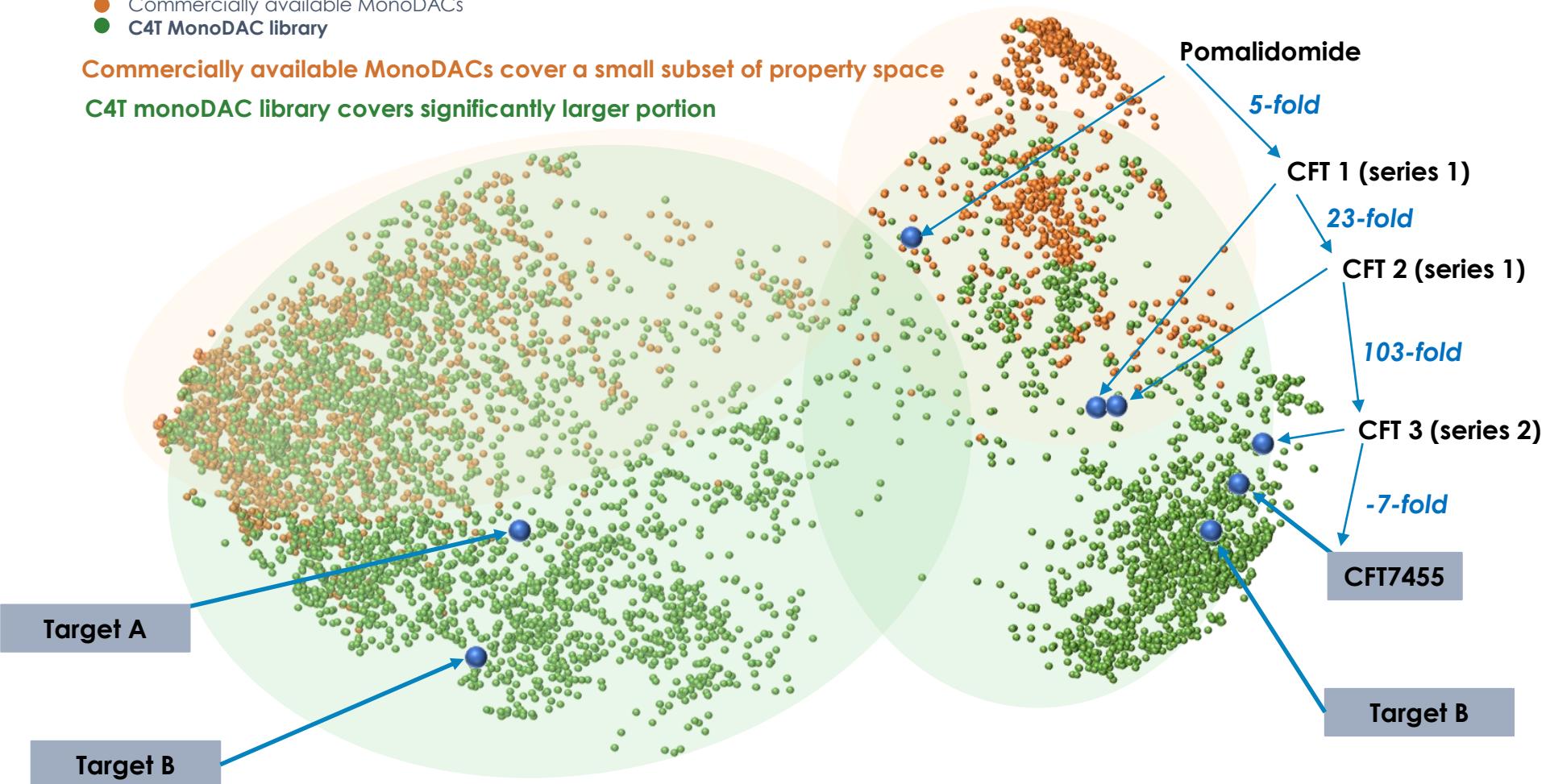
>5,000 membered library constructed from >200 unique scaffolds to maximize MonoDAC structural diversity and CRBN surface remodeling

# C4T MonoDAC Library in Action

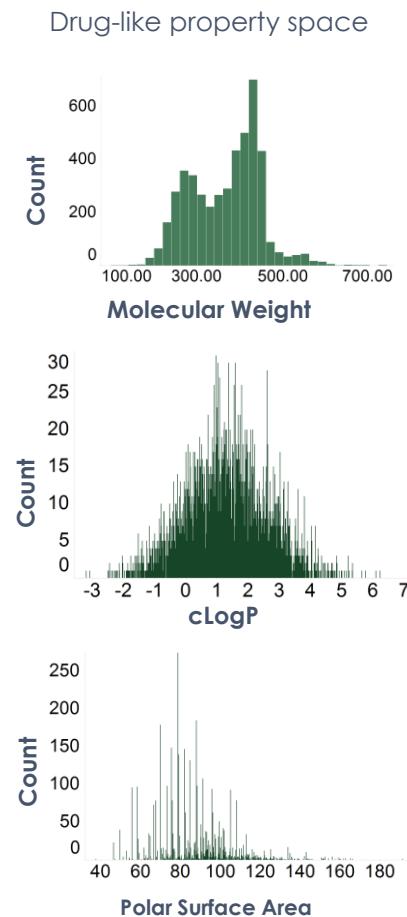
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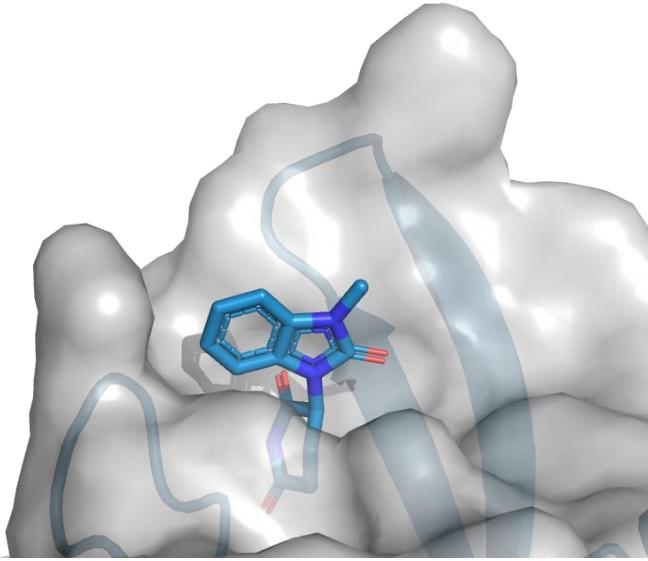
C4T MonoDAC Library has produced hits to novel MonoDAC Targets and a development candidate



# Deep Structural Design Expertise of MonoDAC Degraders

## Benzoimidazolone

C4T unpublished  
1.17 Å

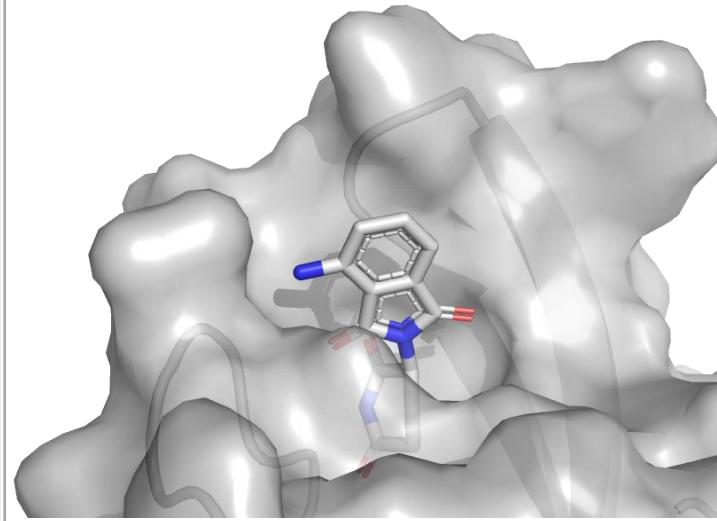


Compound 1  
CRBN FP  $K_D$  = 830 nM



## Pomalidomide

PDB 6h0f



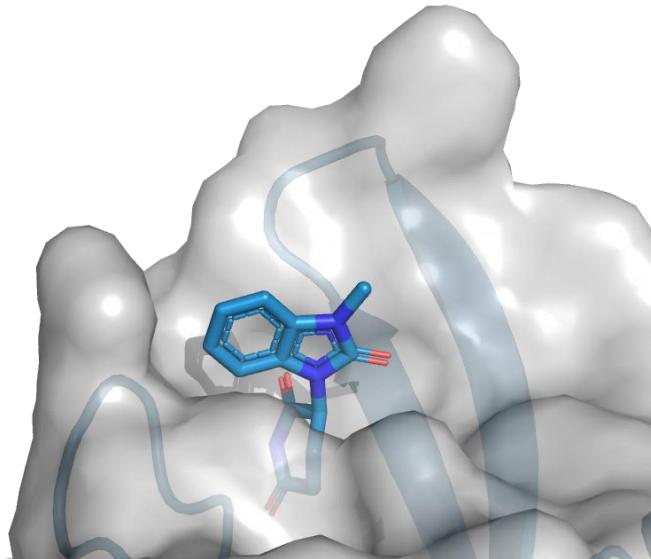
Pomalidomide  
 $K_D$  = 1600 nM



# Deep Structural Design Expertise of MonoDAC Degraders

## Benzoimidazolone

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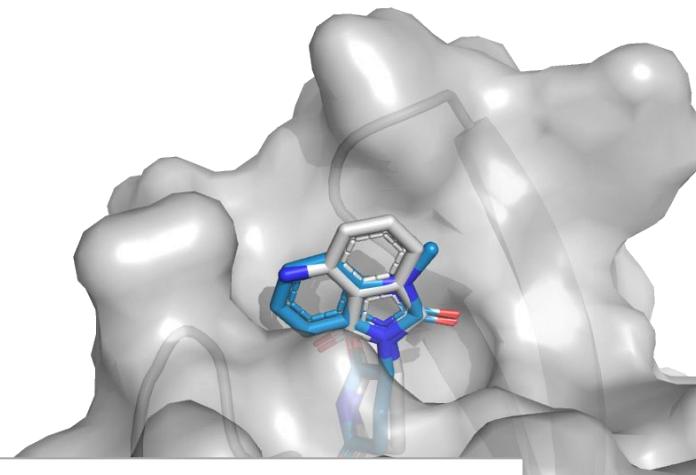
Compound 1  
CRBN FP  $K_D$  = 830 nM



CRBN, cereblon; FP, fluorescence polarization; IKZF1, Ikaros family zinc finger protein 1.

## Pomalidomide

PDB 6h0f



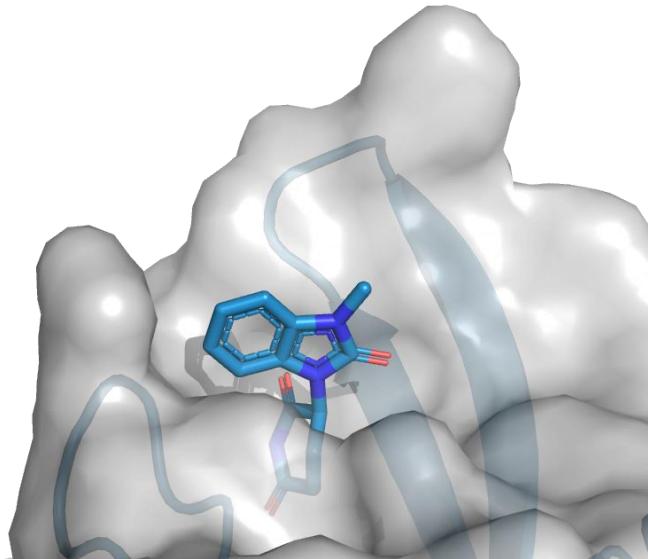
Pomalidomide  
 $K_D$  = 1600 nM



# Deep Structural Design Expertise of MonoDAC Degraders

## Benzoimidazolone

C4T unpublished  
1.17 Å



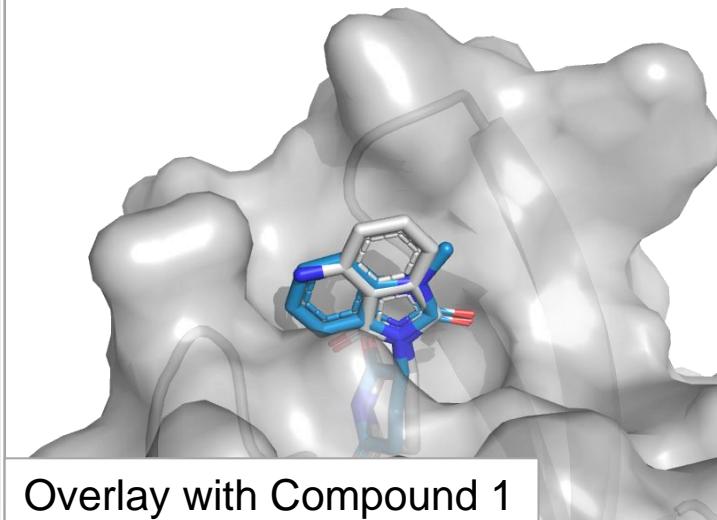
Compound 1  
CRBN FP  $K_D$  = 830 nM



CRBN, cereblon; FP, fluorescence polarization; IKZF1, Ikaros family zinc finger protein 1.

## Pomalidomide

PDB 6h0f



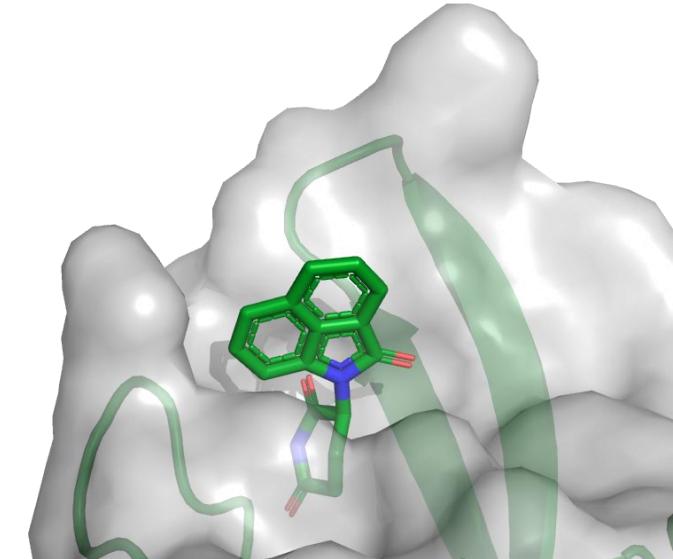
Overlay with Compound 1

Pomalidomide  
 $K_D$  = 1600 nM



## Benzoisoindolinone

C4T unpublished  
1.06 Å



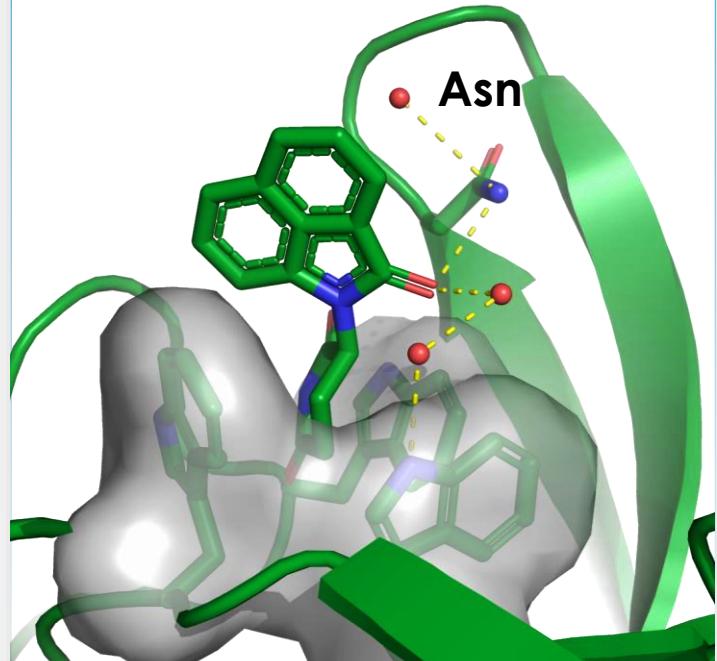
Compound 2  
 $K_D$  = 34 nM



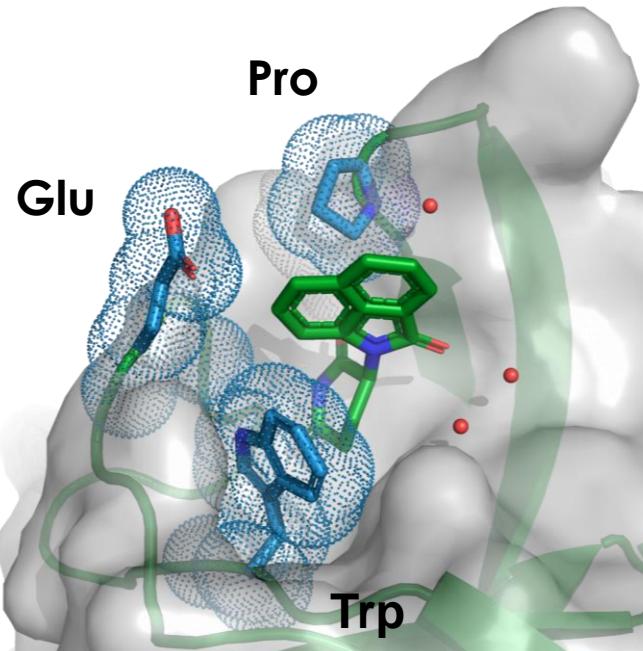
50-fold  
affinity  
increase

# Exploring CRBN Interactions with the Potent Tricyclic Core

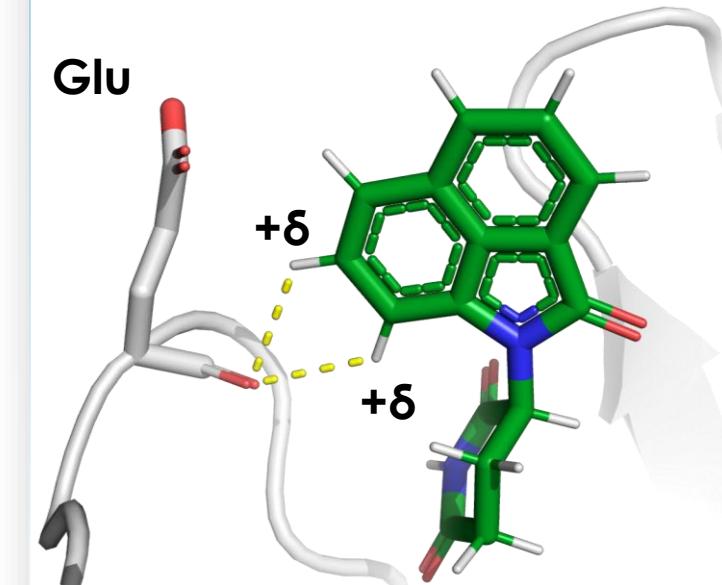
Tri-Trp Pocket Interactions



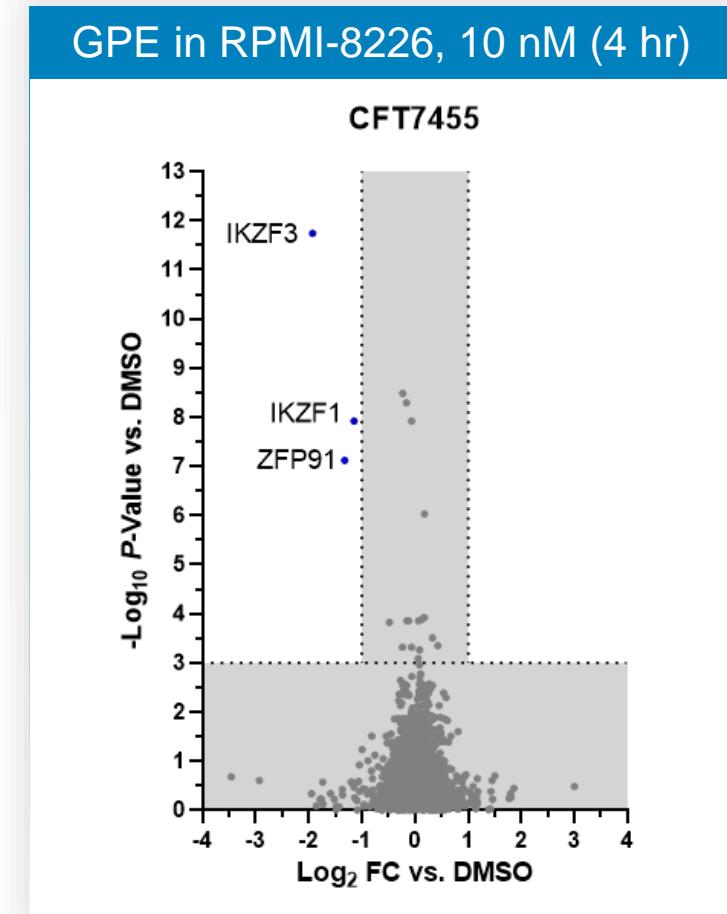
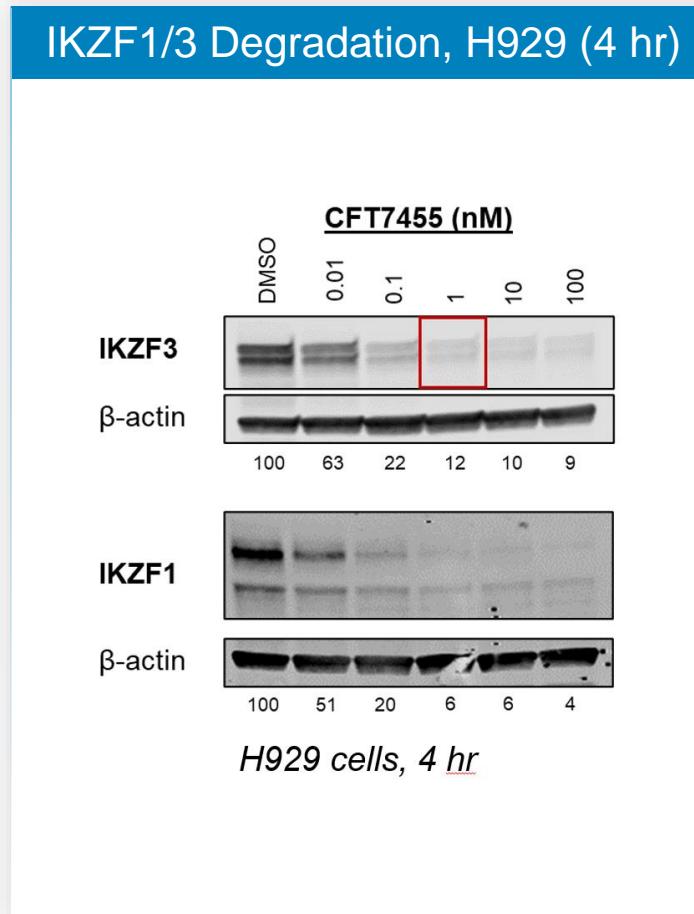
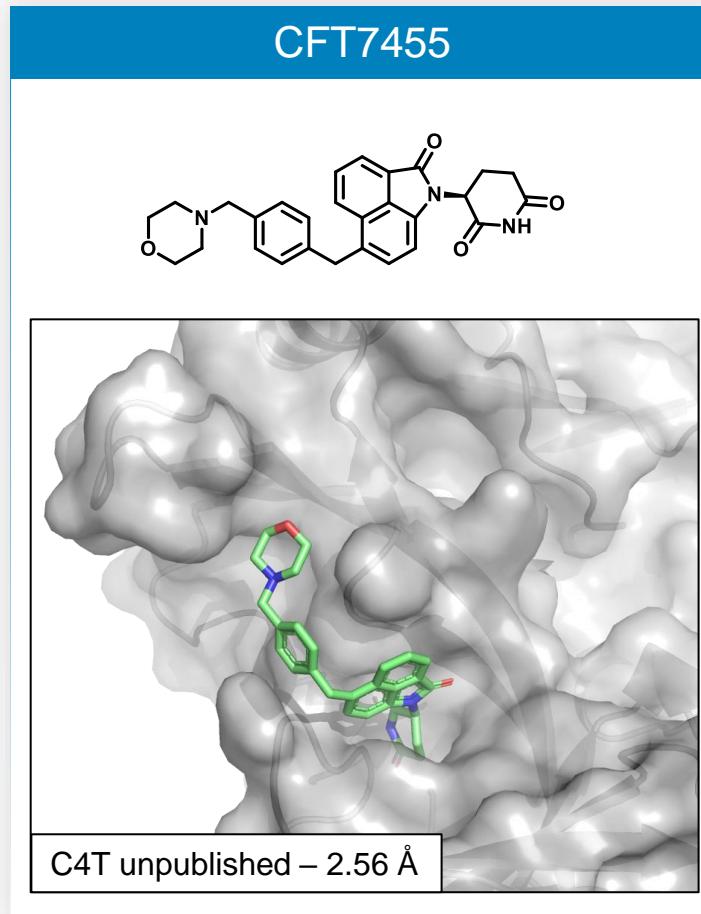
Increased Hydrophobic Contacts with CRBN



+δ Aromatic C-H Interactions with Backbone Carbonyl

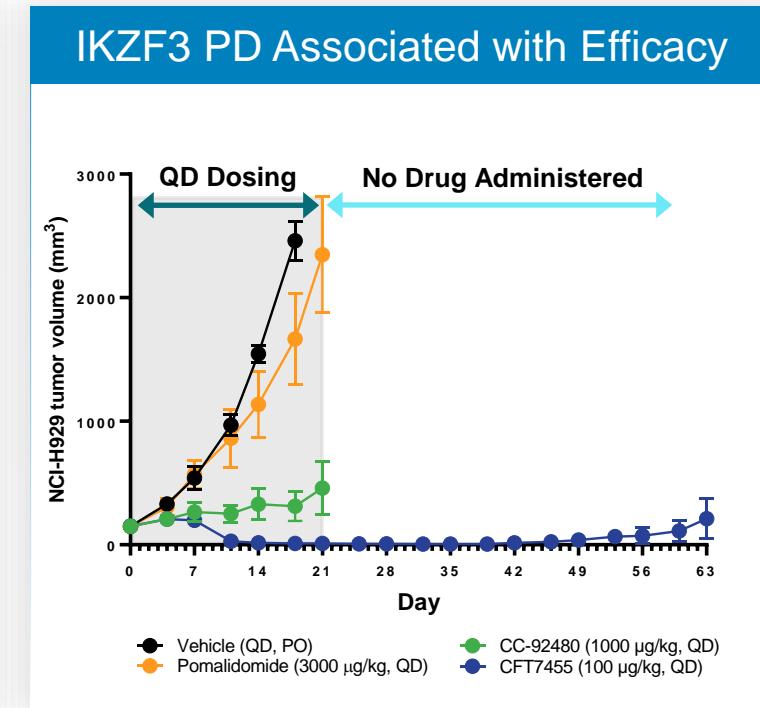
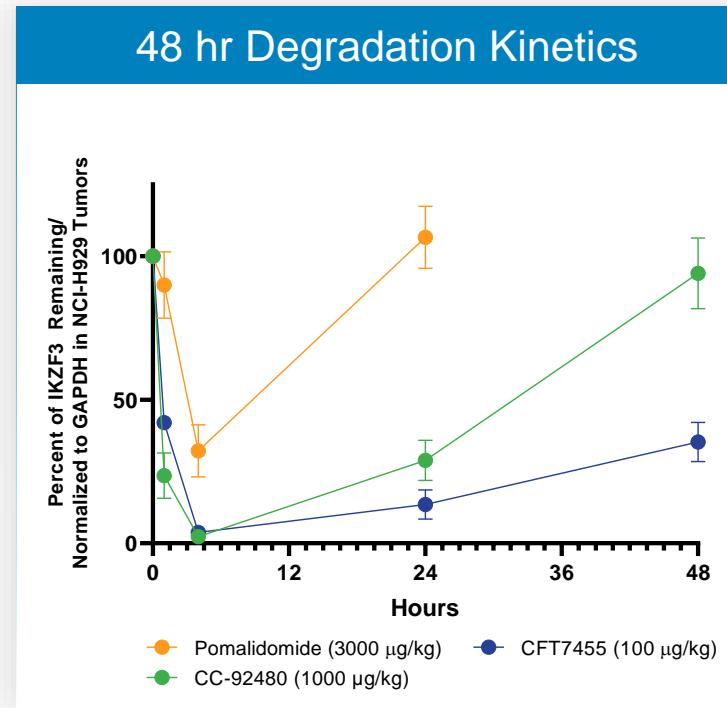
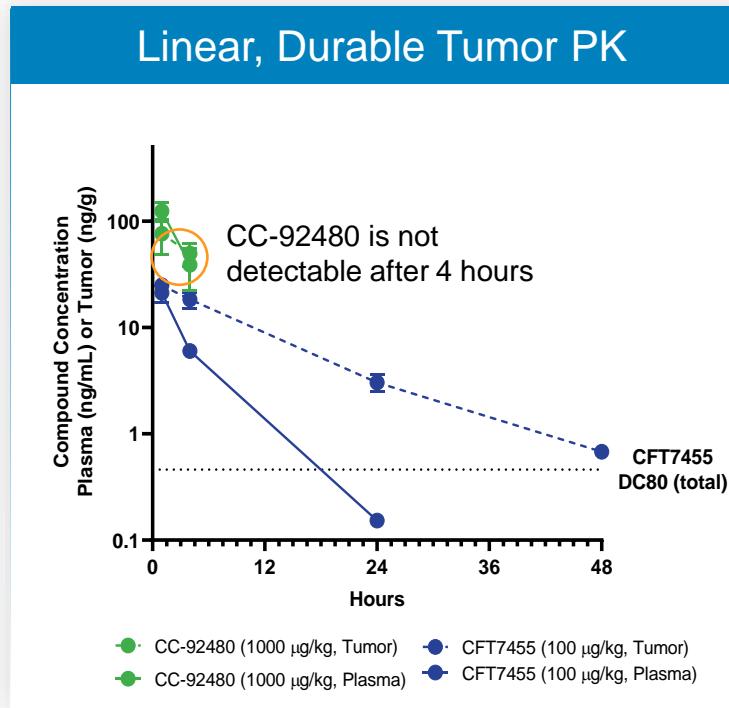


# CFT7455, A Highly Potent and Selective MonoDAC Degrader



GPE, global proteomics experiment; IKZF1/3, Ikaros family zinc finger proteins 1 and 3.  
C4 Therapeutics data on file.

# CFT7455, A Highly Potent and Selective MonoDAC Degrader



CFT7455 displays linear and durable tumor PK translating into deep IKZF3 degradation and regression in MM xenograft models

IKZF3, Ikaros family zinc finger protein 3; MM, multiple myeloma; PD, pharmacodynamics; PK, pharmacokinetics, QD; once daily.  
C4 Therapeutics data on file.



Finding MonoDAC Hits  
to Novel Neosubstrates



# Strategic Approach to MonoDAC Hit Identification

## In Silico Ternary Complex Design

- Identify G-loop containing proteins across the proteome
- Generate ternary complex models that inspire new monoDAC design
- Approach does not necessarily identify productive degraders

## Ternary Complex Identification

- Develop high-throughput biochemical methods to identify ternary complex formation
- Develop high-throughput cellular assay methods to identify ternary complex formation
- Approaches do not necessarily identify productive degraders

## Cellular Degradation

- HiBiT assay – mechanism-informed reduction of target protein levels
- Off-target/off-mechanism and off-target/on-mechanism activity could confound hit identification

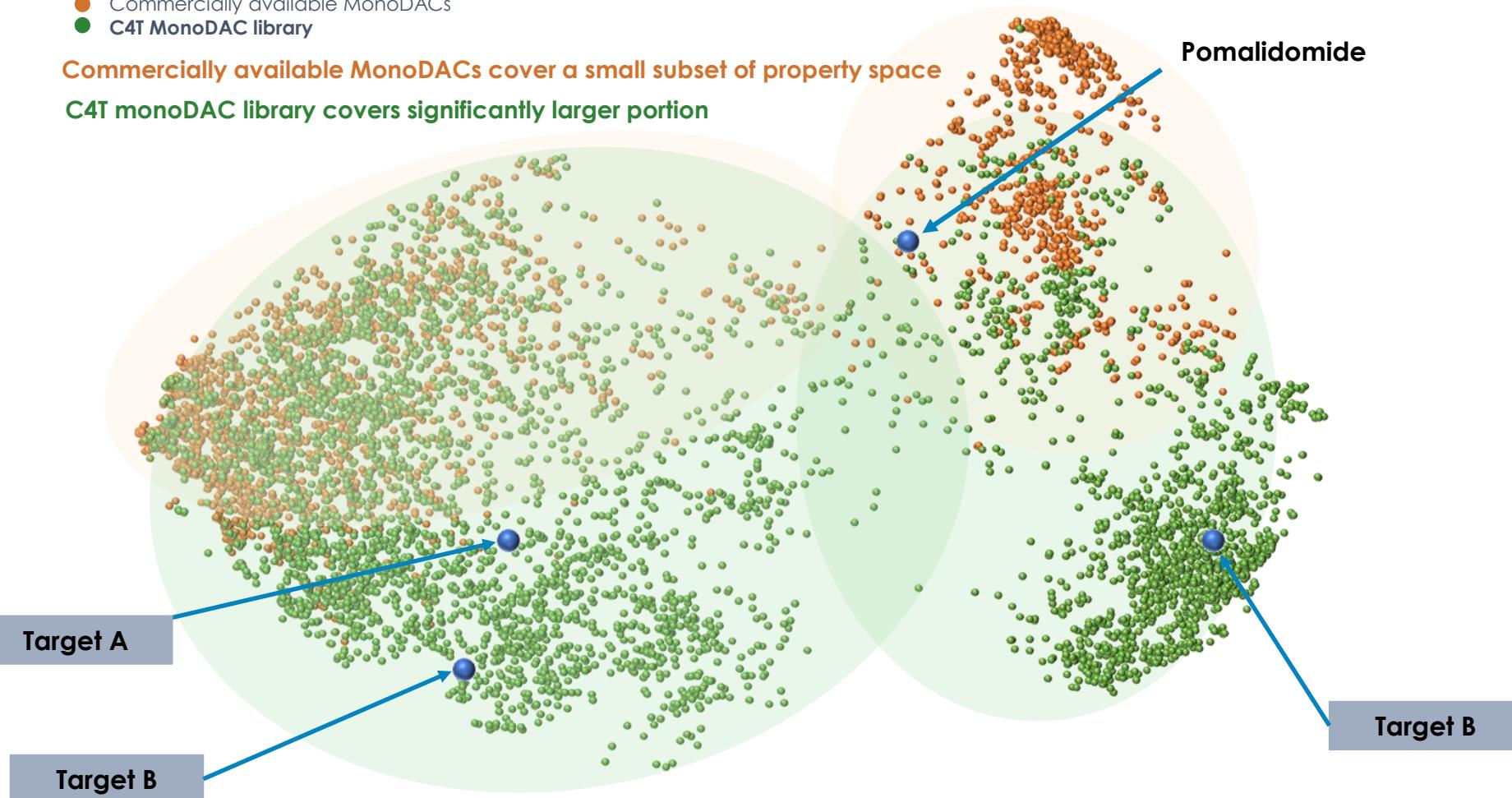
A comprehensive approach that will also identify and expand MonoDAC degrons

# C4T MonoDAC Library in Action – Cellular Degradation of Specific Target

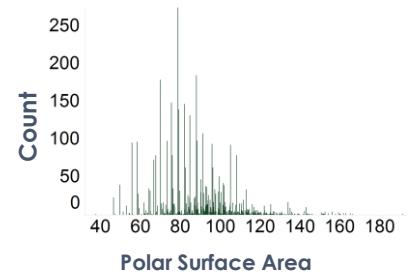
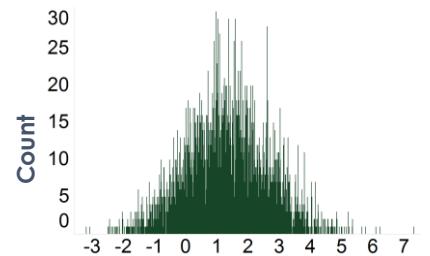
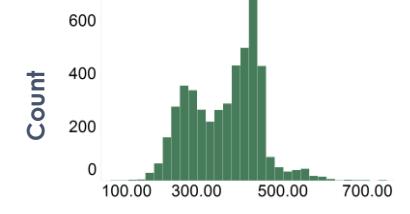
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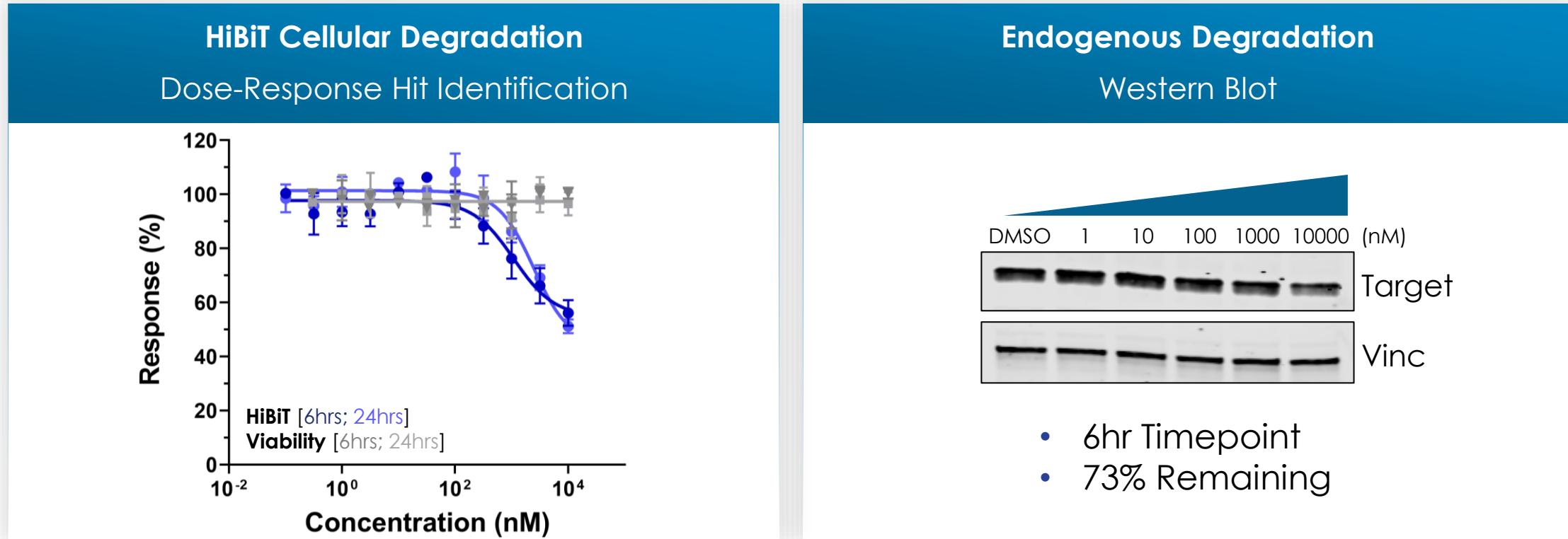
C4T monoDAC library covers significantly larger portion



Drug-like property space



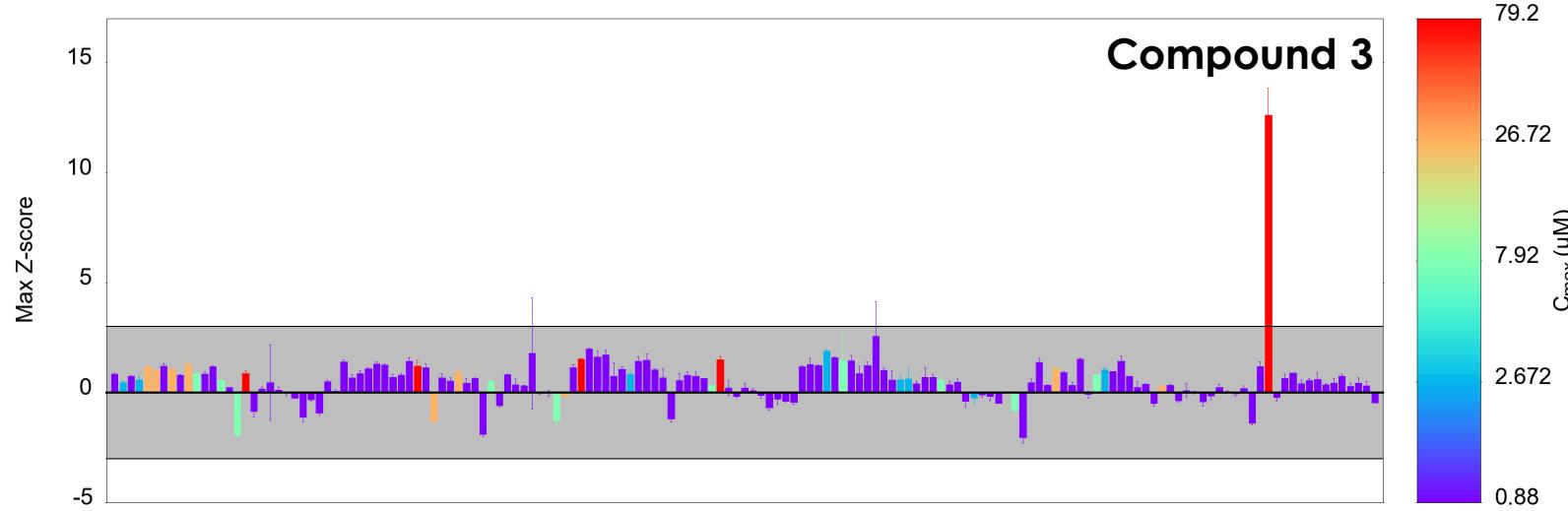
# C4T MonoDAC Library in Action – Cellular Degradation of Target A



Identified hit reduces tagged and endogenous Target A protein levels

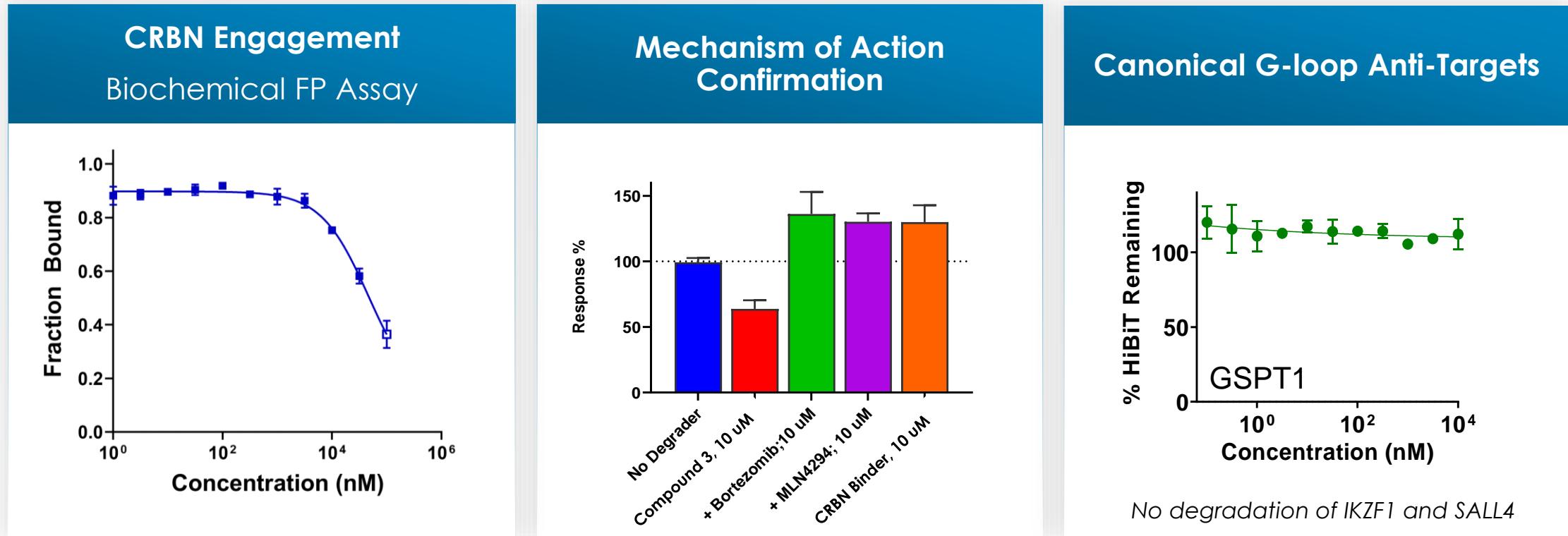
# C4T MonoDAC Library in Action – Biochemical Ternary Complex

- AlphaLISA assay format
- Ternary complex between Target A and CRBN
- Analysis of library subset shown



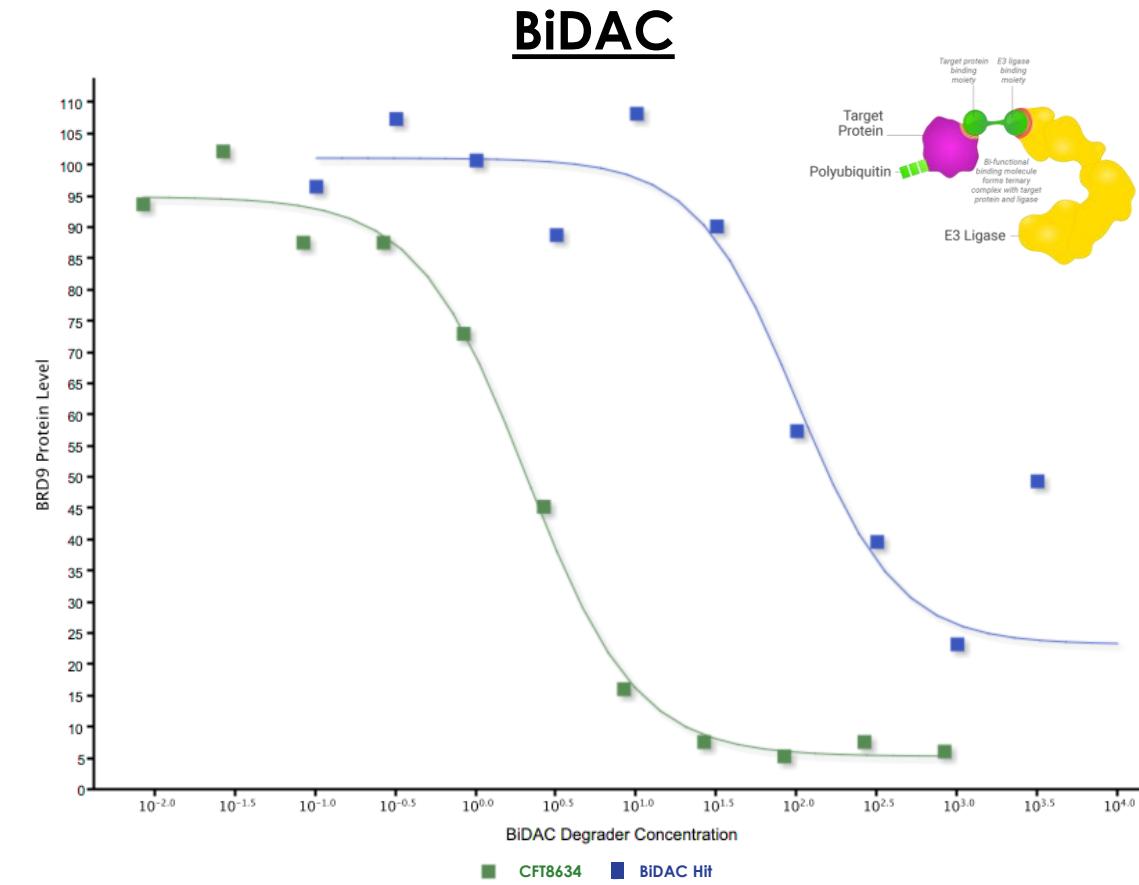
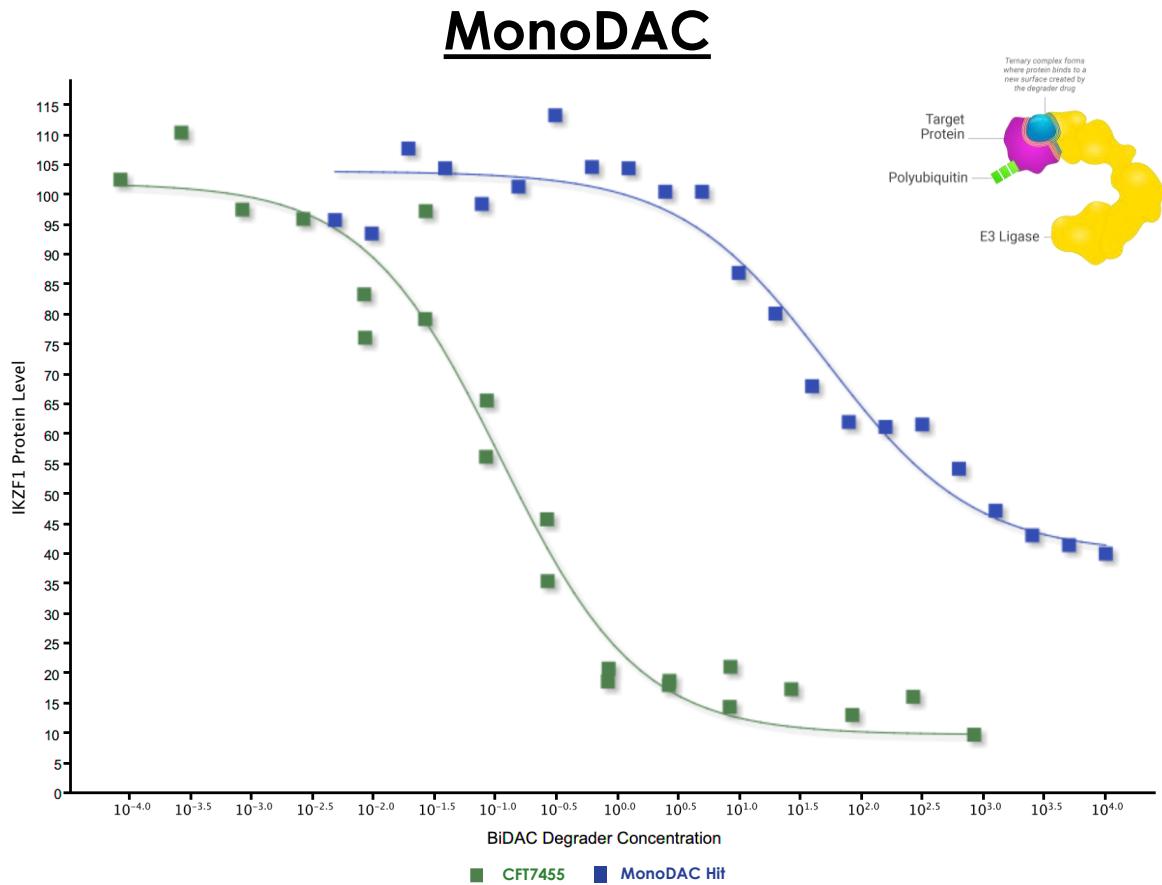
Biochemical screen of the MonoDAC library identifies the same hit compound

# C4T MonoDAC Library – Hit Validation



Compound 3 demonstrates selective on-target, on-mechanism degradation of Target A

# MonoDAC and BiDAC: Target-Tailored Approach



C4T's Platform has produced both MonoDAC and BiDAC Development Candidates

# Thank You C4T Team

