

Aurigene **PROTAC Research**

PROteolysis - TArgeting Chimeras (PROTACs) are hetero-bifunctional molecules that hijack the body's own natural protein disposal system to initiate selective degradation of the protein of interest (POI). Their novel mode of action offers the potential to target the "undruggable" proteome, which comprises of about 85% of human proteins.

The E3 ligand and POI ligand bind to their respective targets, to form a ternary complex. Once bound, an E3 and POI are in proximity, triggering the E3 to transfer multiple ubiquitin molecules from E2 to a lysine residue on the substrate of POI via the formation of a covalent bond.

Advantages of Protein Degradation

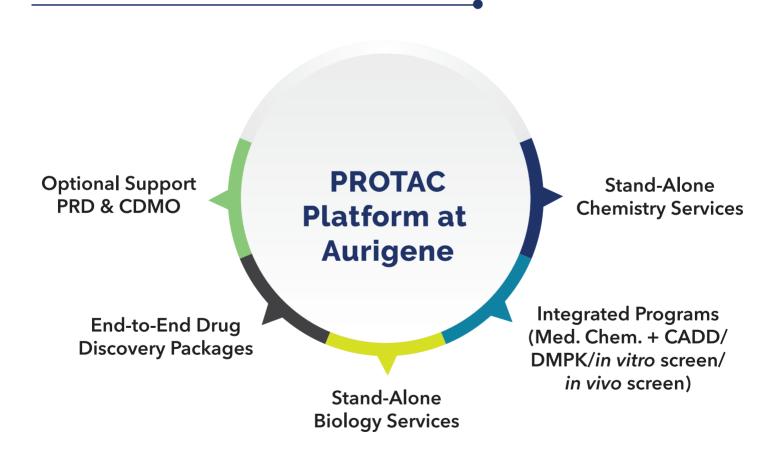
- Hit and run system, less toxic
- Less chance to develop resistance
- Robust, can degrade un-druggable targets
- Minimum amount to achieve efficacy



Why Aurigene?

- ~ 20 years of proven expertise in both Drug Discovery and Med. Chem. CRO
- Thorough understanding (~11 years) of PROTAC-related Med. Chem. space
- In-House library of Partial PROTACs (~500 no.) for rapid synthesis of analogues
- Synthetic capability to rapidly progress from mg to gm scale and beyond
- Experience with AUTACs, ATTECs, RIBOTACs etc.
- Experience with molecular glues

Flexible Business Models











PROTAC Discovery at Aurigene

PROTAC

Biology Understanding:

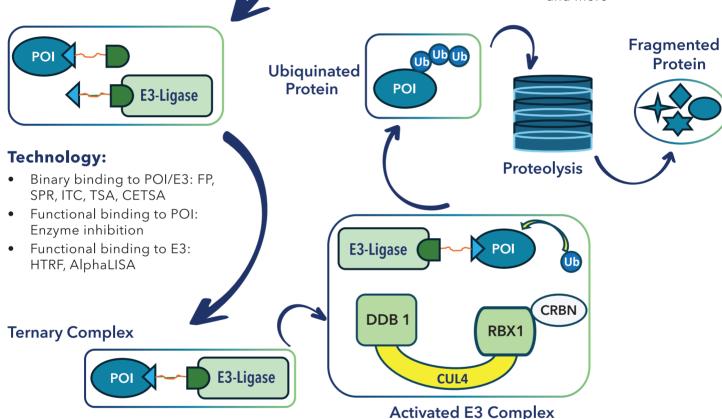
- Protein half-life
- Level and duration of knockdown
- Functional Isoform compensation
- Understand cellular E3 expression
- Understand effect of inhibition vs degradation

Technology and Design:

- CADD based ligand identification
- Linker design (flexible vs
- AI/ML based design
- MD simulation of POI:PROTAC: E3:PROTAC and POI:PROTAC:
 - E3 Ligase orientation space

PROTAC Synthesis

- > 10 years of expertise with PROTAC synthesis
- CRBN, VHL, MDM-2 and cIAP-based designs
- Synthesis of Molecular Glues
- In-House partial PROTAC library for rapid analoguing
- Synthesis of AUTACs, ATTECs, Oligo-PROTACs and more



Technology:

- Ternary binding to POI and E3: HTRF, NanoBRET, TR-FRET
- Ternary:Binary ratio
- Positive and negative co-operativity
- Half life of ternary complex
- Pull down in cellular context

Technology:

- Western / Jess
- Proteasome inhibitors
- AlphaLISA / HTRF



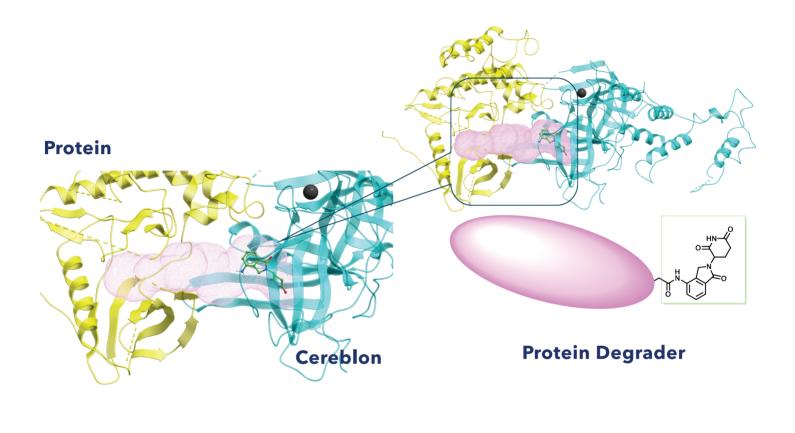






Novel CADD-Based TPD Designs

- Advance modelling tools predicted binary (POI and E3 ligase) and ternary complex (POI, E3 ligase and TPD)
- Deep Neural Network (DNN) model predicted degradation potential of DC50 <= 100 nM, Dmax >= 80%



Modeled Protein Degrader

POI





Cereblon ligand

E3 ligase -> Cereblon

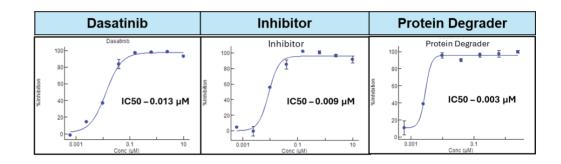
In vitro Biology: Inhibitor vs Degrader

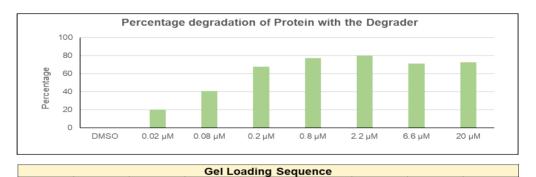
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		IC50 / DC50 (μM)
	ADP Glo	Western Blot	
Inhibitor	0.009	0.67	>20
Degrader	0.003	1.85	0.11

	% Inhibition / degradation at top conc.								
	ADP Glo	Western Blot							
Inhibitor	92	93	5						
Degrader	100	84	73						

- The inhibitor and protein degrader showed nanomolar potency in ADP Glo Assay.
- The small molecule showed IC50 of 670 nM in cell based NanoBRET Target Engagement Assay, where as the protein degrader demonstrated an IC50 of $1.85 \mu M$.
- The degrader showed dose dependent degradation of Protein in THP-1 cell line for 16 h. However, the small molecule was inactive. The tested concentrations are non cytotoxic (data supported by Cell Titer Glo assay).





0.2uM

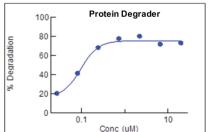
0.7uM

2.2uM

	1	2	3	4	5	6	7	8	9		
~180 ~130	=										
~100 ~70	100	_			hanni	Nessa	_	Luciani	-	Protein (62 kDa)	
~55	_										
~40	-	_								GAPDH (37 kDa)	
~35	-										
-											

0.02uM

0.08uM



6.6uM

20uM

Bottleneck

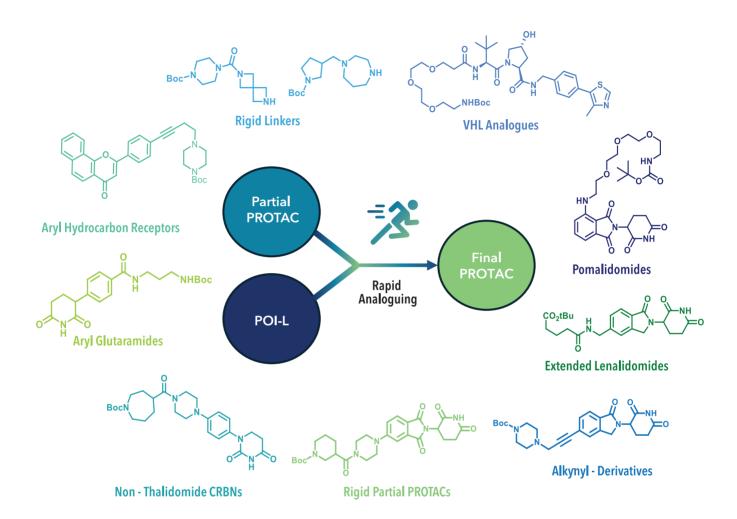
- Low permeability and metabolic stability
- Suboptimal PK parameters (Low AUC & T12/, High CI)

Ladder

DMSO

PROTAC Design using Aurigene's Partial PROTAC Library

Case Study: CADD-based design, synthesis and testing (in vitro, degradation, ADME & PK) of a small library in <2 months



- Linker-ligase library (~500 No)
- Rapid library synthesis
- Parallel purifications using Mass-based prep. HPLCs



Improvement of Physicochemical Parameters

Route	Dose (mg/Kg)	CO (ng/mL)	Cmax (ng/mL)	AUC last (h*ng/mL)	AUC INF_Obs (h*ng/mL)	T 1/2 (h)	MRT (h)	Cl_obs (mL/min/Kg)	% F
i.v.	1	559	-	59.5	60.1	0.15	0.09	278.7	-
SC	5	-	456	573	582	0.49	0.76		≥ 100

Route	Dose (mg/Kg)	CO (ng/mL)	C max (ng/mL)		AUC INF_Obs (h*ng/mL)	T 1/2 (h)	MRT (h)	Cl_obs (mL/min/Kg)	% F
i.v.	1	1236	-	276.5	280	1.54	0.65	59.9	-
SC	5		1348	1960	1990	5.72	2.57		≥ 100

Code	Metaboli	c Stability: 1	T 1/2 (min)	IC50	(µM)	Protein Degradation	
Code	MLM	RLM	HLM	ADP Glo	NanoBRET	% Degradation	
Protein Degrader	3.16	< 2.0	< 2.0	0.009	1.85	DC50 = 0.11 μM	
NB1-033-P	22.97	9.95	6.80	0.014	0.040	59% @ 0.078 µM, Hook Effect observed at higher concentrations	

Improved stability and PK parameters were observed in case of NB1-033-P. Further studies to improve the degradation potential are in progress.

The Aurigene Advantage

Truly Integrated Services

- Integrated discovery chemistry & Biology services (Small & large molecules)
- Early discovery to commercialization with Global regulatory & quality expertise
- Pre-formulation & formulation services, analytical & purification expertise
- Our R&D site is US FDA inspected
 - cGMP sites have multiple accreditations across globe
 - All our labs are GLP approved, animal facilities are AAALAC accredited

Intellectual Property Safety

Quality

Your IPs stay with you! All IP generated through projects belong to our clients

Information security

We are ISO27001 ISMS certified

Flexible Business Models

- Stand-Alone Chemistry Services, Mix-and Match programs (e.g. Med. Chem. + DMPK)
- End-to-End Drug Discovery Packages, Optional Support PRD & CDMO and more



Once with Aurigene, always with Aurigene! Our customers get a unique advantage of staying with us along the product lifecycle!

Thank You



For more information please visit

https://www.aurigeneservices.com/



To place an inquiry please visit

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contactapsl@aurigeneservices.com



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