

Discovery and characterization of potent degraders of IRAK4 and IMiD substrates for oncology indications

YMERA

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INVENTING NEW MEDICINES WITH TARGETED PROTEIN DEGRADATION

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Biology of Targeted Protein Degradation



Co-opting a Naturally Occurring Process to Regulate Protein Levels



E3 ligase recognizes protein



Ubiquitin chain transferred

3

Protein is marked for elimination

Broad Opportunity Only Binding Site Required

> Efficient Catalytic

Prolonged Impact Targeted Protein Degradation



Delivering on the Promise: Building an IRAK4 Franchise



KT-474 Selective heterobifunctional degrader of IRAK4

- First proof-of-mechanism for TPD in a randomized, placebo-controlled healthy volunteer study
- Demonstrated > 95% IRAK4 degradation in humans

KT-413

Dual-mechanism degrader of IRAK4 and IMiD substrates

• Phase 1 clinical trial in R/R B cell lymphomas ongoing

(the structures of KT-474 and KT-413 were not disclosed in this presentation)

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Degradation of IRAK4: Modulating Proinflammatory Cytokines and Cellular Proliferation

- IRAK4 is a key component of the myddosome and its function is dependent on both its kinase activity and on its scaffolding properties
- Activation of downstream pathways drive the scaffolding function of IRAK4 and are key drivers of cellular proliferation and proinflammatory cytokine and chemokine production



Degradation of IRAK4 and IMiD Substrates: Targeting Redundant Pro-survival Pathways in MYD88^{MT} DLBCL

- MYD88 L265P is a gain-of-function driver mutation which results in constitutive activation of the antiapoptotic NFkB signaling pathway
- Single-agent therapies that target activated NFκB signaling in DLBCL show limited activity in preclinical or clinical settings
- Redundant NFκB pathway activation and downregulation of Type 1 IFN is common in MYD88^{MT} lymphoma, supporting need to seek combination therapies
- Targeting simultaneous degradation of IRAK4 and IMiD substrates Ikaros and Aiolos shows synergistic activity in MYD88^{MT} models, supporting this targeted combination



IRAKIMiD: Functioning as a Heterobifunctional Degrader & a Molecular Glue



IRAK4 Ligand and Exit Vector Identification



Ternary Complex Modeling



X-ray: POM bound to CRBN-DDB1

X-rays and exploratory SAR inform design of ternary complex model



X-ray: KTX-733 bound to IRAK4



Ternary complex model to guide and inspire degrader design





Representative Motifs Explored

- Extensive exploration of IRAK4 ligand and linker required to probe SAR for both IRAK4 and IMiD substrate degradation
- Indazole scaffold became primary focus due to kinome selectivity, modularity, and ability to tune properties



Improving IRAK4 Degradation Potency





Parameter	KTX-671	KTX-315	Parameter	KTX-881	KTX-353
Linker	NH NH	NH - E	Linker		N N N
OCI-Ly10 Cell IRAK4 DC ₅₀ (nM)	> 1,000	22	OCI-Ly10 Cell IRAK4 DC ₅₀ (nM)	23	6.0

Linker modifications to alter POI exit vector directionality can improve IRAK4 degradation potency

Modulating Potency and Intrinsic Stability



	KTX-435	KTX-582	KTX-955
IRAK4 DC ₅₀ (nM)	18	4	5
lkaros DC ₅₀ (nM)	12	5	130
OCI-Ly10 CTG IC ₅₀ (nM)	270	28	1,800
HLM (μL/min/mg)	60	48	4

Linker modifications could enable an improvement on IRAK4 degradation efficiency and intrinsic stability but could significantly impact ability to degrade Ikaros and Aiolos

Merging Intrinsic Stability with IMiD Activity



	KTX-955	KTX-497	KTX-612
IRAK4 DC ₅₀ (nM)	5	3	7
lkaros DC ₅₀ (nM)	130	25	6
HLM (μL/min/mg)	4	1	3
RLM (μL/min/mg)	4	3	2

Understanding and Leveraging Permeability Data



Evaluation of IRAK4 degraders in rat PK studies: 10 mg/kg, PO

No correlation between oral absorption and passive permeability

Passive permeability trends with oral bioavailability

Number of HBDs Impacts Passive Permeability and Efflux



Hydrogen Bond Donors

 Reduction in hydrogen bond donors (HBDs) trends toward improved passive permeability and reduced MDR1mediated efflux



	KTX-088	KTX-109
IRAK4 DC ₅₀ (nM)	54	8
Hydrogen bond donors (#)	4	3
MDCK P _{app} (10 ⁻⁶ cm/s)	0.4	1.0
MDCK-MDR1 efflux ratio	8	0.3
R _{gyr} (Å)	9.2	9.2
Rat IV CL (mL/min/kg)	67	30
Rat PO PK (10 mg/kg) %F AUC (μM*hr)	0 0	4.2 0.13

Efflux Has Critical Impact on Oral Bioavailability

• In vivo MoA study indicated efflux has critical impact on oral bioavailability

Machine learning analysis to determine the most effective ER threshold to achieve oral F>10%



Within this specific series of degraders, an MDR1 efflux ratio of 16 was identified as the threshold above which oral absorption was significantly impacted

Efflux Ratio Impact on Oral Bioavailability



	KTX-037	KTX-045
IRAK4 DC ₅₀ (nM)	7	9
#HBD	4	4
P _{app} (10 ⁻⁶ cm/s)	19	4.9
Efflux ratio	35	10
R _{gyr} (Å)	5.8	6.1
RLM (µL/min/mg)	19	< 12
Rat PPB (Fu)	0.0024	0.0013
Rat IV CL (mL/min/kg)	22	21
Rat PO PK (10 mg/kg) %F AUC (μM*hr)	4 0.35	14 1.2

IRAK4 Degrader vs IRAKIMiD Degrader: Degradation

Parameter	KTX-545	KT-413 IRAKIMiD Degrader	
	IRAK4 Degrader		
OCI-Ly10 Cell IRAK4 DC ₅₀ (nM)	1.0	6.0	
OCI-Ly10 Cell Ikaros DC ₅₀ (nM)	> 1,000	2.0	
OCI-Ly10 Cell Aiolos DC ₅₀ (nM)	> 1,000	2.0	

- IRAK4 degrader KTX-545 exhibits no effect on levels of Ikaros or Aiolos
- IRAKIMiD degrader KT-413 induces strong degradation of IRAK4, Ikaros, and Aiolos
- IRAKIMiD degrader KT-413 degrades IRAK4 more slowly than Ikaros and Aiolos



IRAK4 and IMiD Degraders vs IRAKIMiD Degrader: Cytotoxicity

- DLBCL cell line OCI-Ly10 expresses most prevalent MYD88 mutation (L265P)
- IRAK4 degrader KTX-545 exhibits no OCI-Ly10 cytotoxicity
- IRAKIMiD degrader KT-413 shows robust OCI-Ly10 cytotoxicity
- IRAKIMiD degraders can show strong and broad activity across several MYD88 mutant cell lines

Parameter	KTX-545	KT-413	
	IRAK4 Degrader	IRAKIMiD Degrader	
OCI-Ly10 Cell IRAK4 DC ₅₀ (nM)	1.0	6.0	
OCI-Ly10 Cell Ikaros DC ₅₀ (nM)	> 1,000	2.0	
OCI-Ly10 Cell Aiolos DC ₅₀ (nM)	> 1,000	2.0	
OCI-Ly10 Cell CTG IC ₅₀ (nM)	> 10,000	9.0	



KT-413 Highly Active on Intermittent Dosing Regimens

- Minimally active dose of 3.0 mg/kg D1,2 showed extended tumor exposure and strong degradation of both IRAK4 and IMiD substrates that was maintained for at least 72 h
- In the OCI-Ly10 cell MYD88^{MT} xenograft model, intermittent dosing of KT-413 induced strong antitumor activity, including complete or partial regressions



KT-413 Shows Regressions in MYD88^{MT} Patient-Derived Xenograft (PDX) Models in Mouse

Model	MYD88	CD79B	TNFAIP3	Other	KT-413 (%TGI)	ian		PDX Ly14019 ^{CD79MT/MYD88MT}	
LY14019	L265P	MT	MT		100		-000	- Vehicle	
LY2264	L265P	MT		IRF4	100	15 <mark>(</mark> و 15	500	- K1-413, 20 mg/Kg PO D1-3 QW	
LY2298	L265P	MT		BCL2/BCL6	90	<u>ب</u> 10	000 -	per la comparte de la	
LY12699	L265P	MT			87	Inm		×	
LY2345	WT		MT		70	r Vo	500 -		
LY2301	WT				30	own	0		
LY0257	L265P			BCL2/BCL6/IKZF3	0	F	0	Days (post-radomization)	40

KT-413 shows strong tumor growth inhibition (> 85% TGI) in 4/5 MYD88-Mutated DLBCL PDX Models in mouse

- Activity is observed regardless of co-mutations that activate NFkB and IRF4 pathways
- The non-responsive MYD88^{MT} model LY0257 harbors a mutation in Aiolos and is reported to be insensitive to lenalidomide. The functional consequence of Aiolos mutations in IRAKIMiD and IMiD response is being investigated

Lessons Learned in the Optimization of IRAKIMiDs

- Challenges:
 - Managing independent SAR for degradation of IRAK4 and IMiD substrates
 - Optimizing physicochemical and pharmacokinetic properties in bRo5 chemical space
 - Developing appropriate in vitro assays for evaluating pharmacological properties
- Opportunities:
 - Embracing the linker: it is not just a bystander
 - Leveraging subtle structure modifications to POI ligand or linker for profound impact on degradation potency
 - Exploiting minor structure modifications to linker for large impact on selectivity, oral bioavailability and pharmacokinetics
 - Using ternary complex modeling to inspire design
 - Understanding parameters (e.g. permeability, efflux ratio, etc.) that have impact on oral bioavailability and developing models to predict those parameters



Modulating Pharmacokinetics and Safety: CRBN ligand, linker, IRAK4 ligand

Acknowledgments

