

# INVENTING NEW MEDICINES

WITH TARGETED PROTEIN DEGRADATION

The Kymera logo is displayed on the left side of a wide banner. The banner's background is a composite image: on the left, a blue and purple molecular structure; on the right, a night sky with a constellation and a mountain silhouette.

**KYMER A**

# Forward-Looking Statements

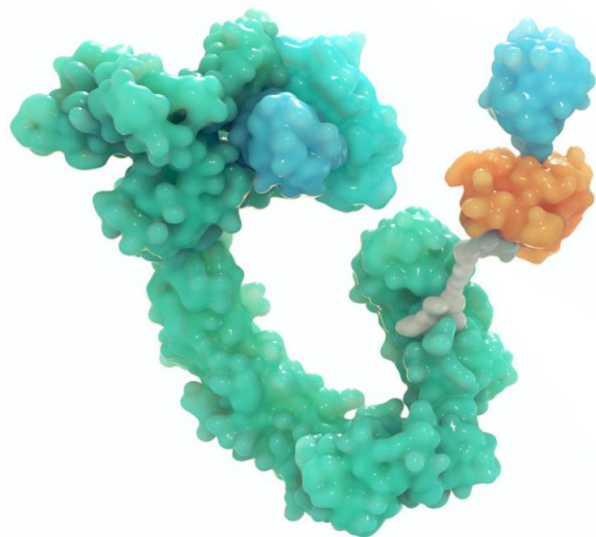
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# Kymera: A Leading TPD Company

KYMER A



## VISION

Fully integrated, **disease agnostic** protein degrader medicine company

## KEY PARTNERSHIPS



## INITIAL FOCUS

**Immune inflammation (I/I)** and **oncology**

## FIRST-IN-CLASS

**First** to show **placebo-controlled** degrader **proof-of-mechanism**

## CLINICAL PIPELINE

**2** additional **INDs** and clinical initiations expected by end of **2021**

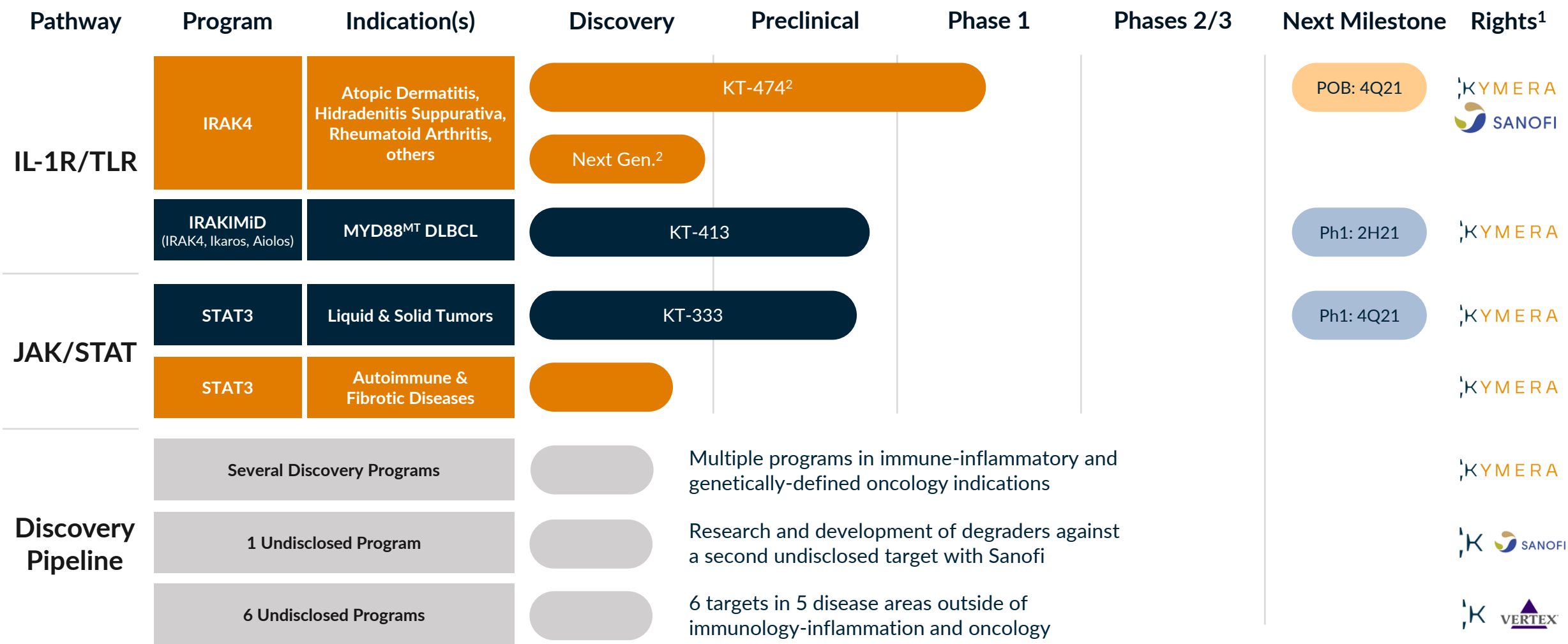
## PROOF-OF-BIOLOGY

To be established in humans in **2021**

## WELL-POSITIONED

**\$647M** cash balance\*

# Kymera's Pipeline of Novel Protein Degraders



● = Oncology ● = Immunology-Inflammation

# Near-Term Milestones Provide Significant Opportunity

Program	Compound	Indication(s)	Expected Upcoming Milestones
IRAK4	KT-474	AD, HS, RA, others	<ul style="list-style-type: none"> <li>✓ Initiated SAD portion of Phase 1 trial in healthy volunteers (Feb 2021)</li> <li>✓ Established degrader proof-of-mechanism in healthy volunteer SAD portion of Phase 1 trial (June 2021) <ul style="list-style-type: none"> <li>• Initiate enrollment in MAD portion of Phase 1 trial (July 2021)</li> <li>• Present data from atopic dermatitis cohort in non-interventional study (2H21)</li> <li>• Establish Phase 1 proof-of-biology in healthy volunteers (4Q21) and in patient cohort (1H22)</li> </ul> </li> </ul>
IRAKIMiD (IRAK4, Ikaros, Aiolos)	KT-413	MYD88 <sup>MT</sup> DLBCL	<ul style="list-style-type: none"> <li>✓ Presentation of preclinical data updates at AACR, ICML meetings (2Q21) <ul style="list-style-type: none"> <li>• Submit IND to initiate Phase 1 clinical trial in r/r B cell lymphomas (2H21)</li> <li>• Present additional KT-413 preclinical data and potential expansion strategies (2H21)</li> <li>• Establish Phase 1 proof-of-biology in patients (2022)</li> <li>• Establish Phase 1 initial clinical proof-of-concept in patients (2022)</li> </ul> </li> </ul>
STAT3	KT-333	Liquid & Solid Tumors	<ul style="list-style-type: none"> <li>✓ Nominated development candidate for liquid &amp; solid tumor indications (1Q21) <ul style="list-style-type: none"> <li>• Present additional preclinical data in liquid &amp; solid tumor indications (2H21)</li> <li>• Submit IND to initiate Phase 1 clinical trial in liquid and solid tumors (4Q21)</li> <li>• Establish Phase 1 proof-of-biology in patients (2022)</li> <li>• Establish Phase 1 initial clinical proof-of-concept in patients (2022)</li> </ul> </li> </ul>
Discovery Programs & Platform			<ul style="list-style-type: none"> <li>• Continue pipeline expansion by advancing early-stage discovery programs toward IND-enabling studies</li> <li>• Further expand Pegasus platform to generate novel degrader product candidates</li> <li>• Leverage Whole-Body Atlas to unlock new opportunities across broad therapeutic applications</li> </ul>

● = Oncology ● = Immunology-Inflammation





# Pegasus™ TPD Platform

 KYMERA

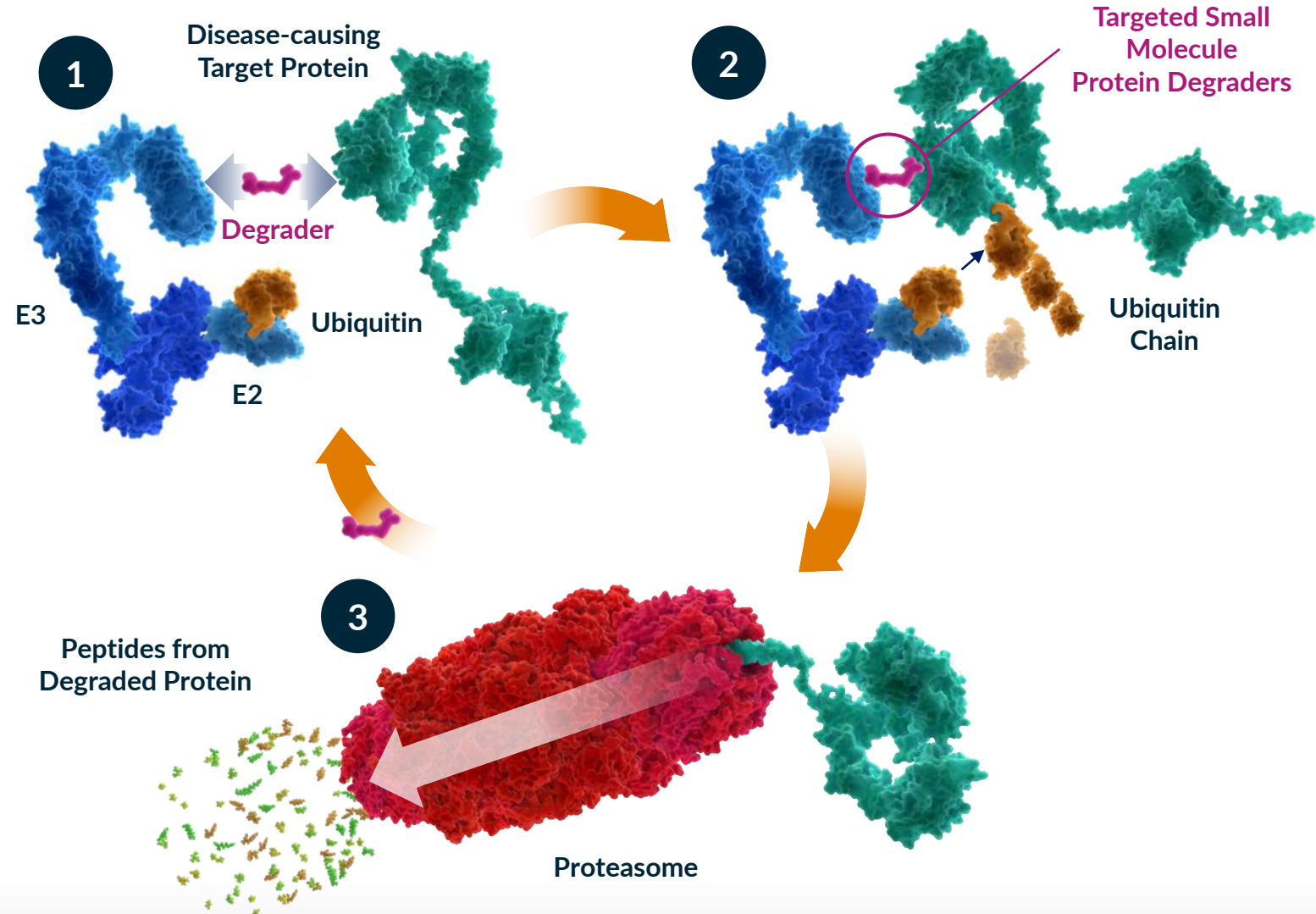


# Targeted Protein Degradation

## Biology

### Co-opting a Naturally Occurring Process to Regulate Protein Levels

- 1 E3 ligase recognizes protein
- 2 Ubiquitin chain transferred
- 3 Protein is marked for elimination



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**Broad Opportunity**  
Only Binding Site Required

**Efficient**  
Catalytic

**Prolonged Impact**  
Targeted Protein Degradation

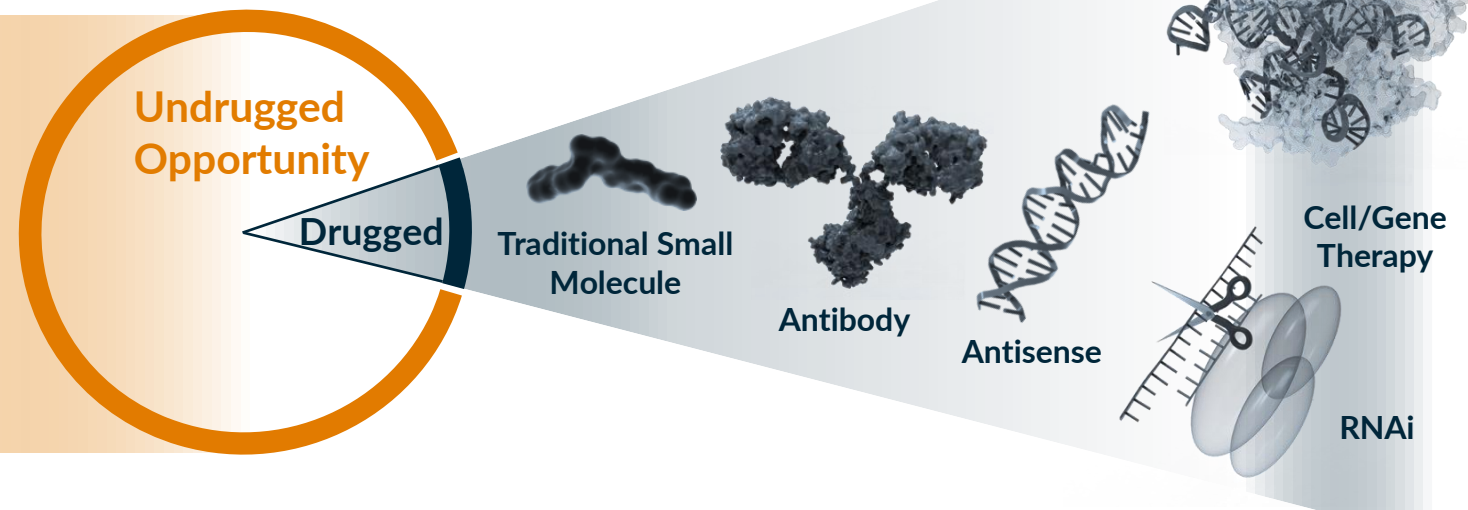
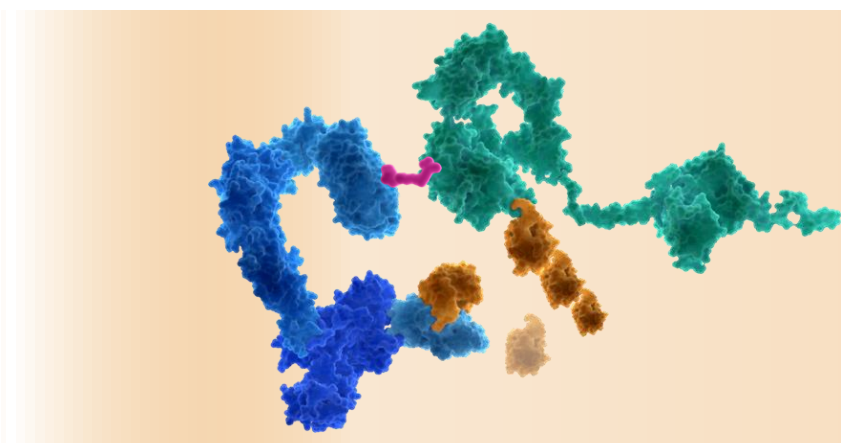
# Targeted Protein Degradation

Next Potential Breakthrough Modality to Expand Drugged Proteome

Targeted Protein Degradation

Human Proteome

Existing Modalities



✓
✓
✓
✓

**Undruggable Targets**  
Scaffold, transcript factor, multiple functions

**Efficient**  
Development / Manufacturing

**Systemic**  
Exposure

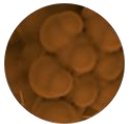
**Oral**  
Bioavailability

✓
X
X
X



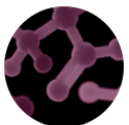
# Proprietary Pegasus™ TPD Platform

## Key capabilities



### Expanded E3 ligase toolbox

- **E3 ligase Whole-Body Atlas:** Identification of the **expression profiles of ~600 unique E3 ligases**
- Match target protein with appropriate E3 ligase based on expression, distribution, intracellular localization, and biology
- **Toolbox of proprietary ligands** leverages the E3 Ligase Whole-Body Atlas



### Understanding degradation (PK/PD) across tissue types

- **Ternary complex modeling tool optimizes the development** of highly efficient and selective degrader therapeutics
- **Quantitative System Pharmacology Model** measures and predicts diverse sets of parameters that impact protein levels
- Based on understanding of PK/PD, both *in vitro* and *in vivo*, and across different tissues and cell types



### Proprietary Chemistry

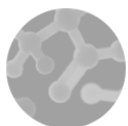
- **Comprehensive hit finding technologies toolbox:** chemoproteomics, DEL, fragment screens, *in silico*
- **Proprietary chemistry expertise** enables the design and optimization of both E3 ligases and target protein binders
- Ability to convert into degraders with optimal pharmaceutical properties tailored to specific patient populations

# Pegasus: E3 Ligase Whole-Body Atlas

Different expression profiles of E3's provide opportunity for tissue selective/restricted degradation



Expanded E3  
ligase toolbox

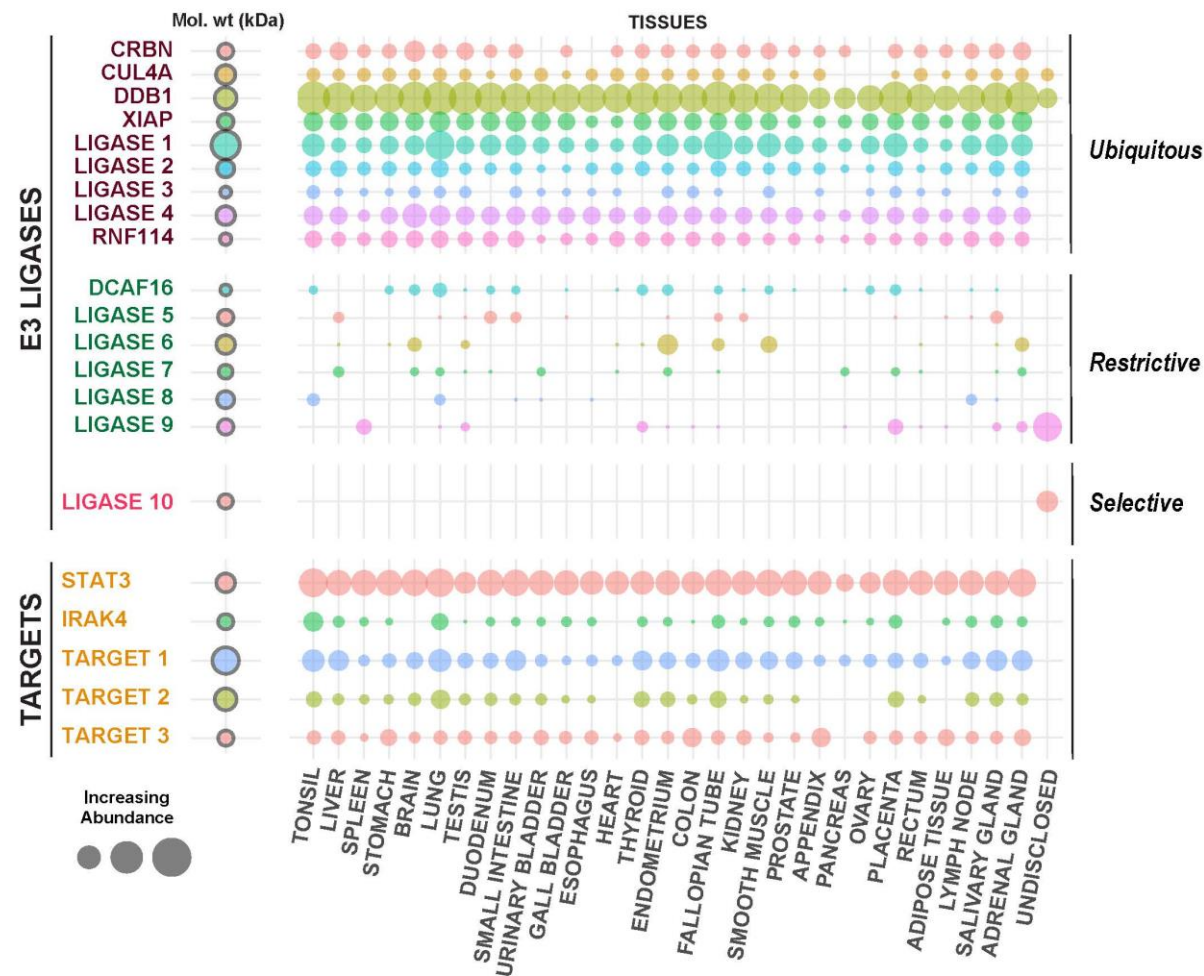


Understanding  
degradation  
(PK/PD) across  
tissue types



Proprietary  
Chemistry

- Focused on determining the expression profiles of ~600 unique E3 ligases
- Patterns mapped in both disease and healthy contexts
- Ability to match a target protein with appropriate E3 ligase based on expression and biology
- Vision to develop tissue-selective or tissue-restricted degraders to enable novel therapeutic opportunities





# IRAK4 Degradator KT-474

# IRAK4 Targeting: Clinical Validation, Human Genetics De-risking and Degradation Advantage



Unmet  
Medical  
Need



Validated  
Biology

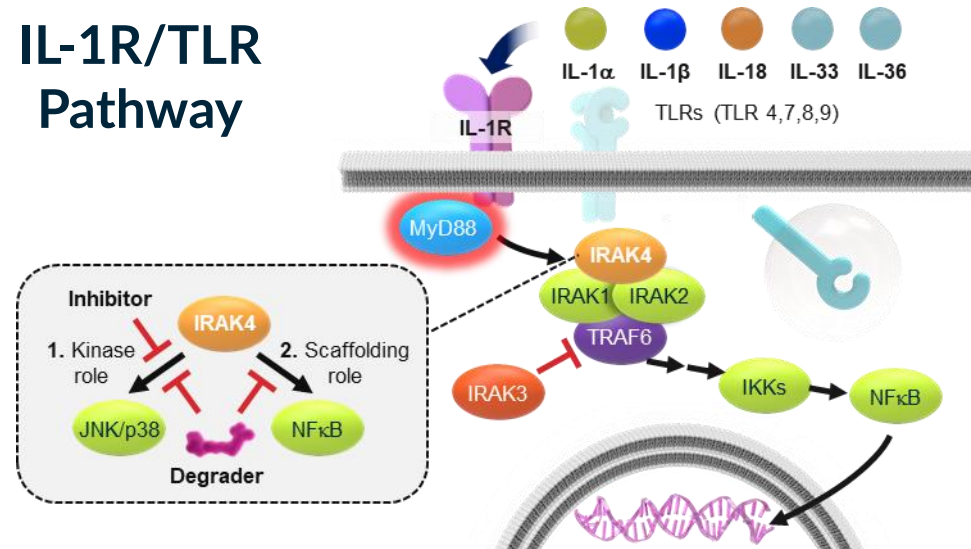


Undrugged  
Node



Precision  
Medicine  
Approach

## IL-1R/TLR Pathway



## Clinical Pathway Validation

IL1-Rα/IL-1β : Rheumatologic Diseases

IL-1α: Atopic Dermatitis

IL-1β: CANTOS Data, Atherosclerosis, Lung Cancer

IL-18: Macrophage Activation Syndrome

IL-36: Generalized Pustular Psoriasis

IRAK4 SMI: Rheumatoid Arthritis

- IRAK4 is a key component of the myddosome protein complex involved in innate immunity that mediates signals through IL-1R and TLRs
- Several commercial and clinical stage drugs have validated this pathway in multiple diseases
- Degrading IRAK4, and fully blocking IL-1R/TLR signaling, is expected to be superior to antibody-based therapies that block only single cytokines, with convenience of a daily oral therapy
- IRAK4 degradation can block pathway fully vs kinase inhibitors that partially block signaling
- Human genetics de-risk safety: adults that lack IRAK4 are healthy



# KT-474 Opportunity

Immune-inflammatory disorders collectively impacting millions of patients in the U.S.

Atopic  
Dermatitis (AD)

Total Prevalence (U.S.)

>16.0M<sup>1</sup>

Rheumatoid  
Arthritis (RA)

>1.3M<sup>2</sup>

Hidradenitis  
Suppurativa (HS)

>325K<sup>3</sup>

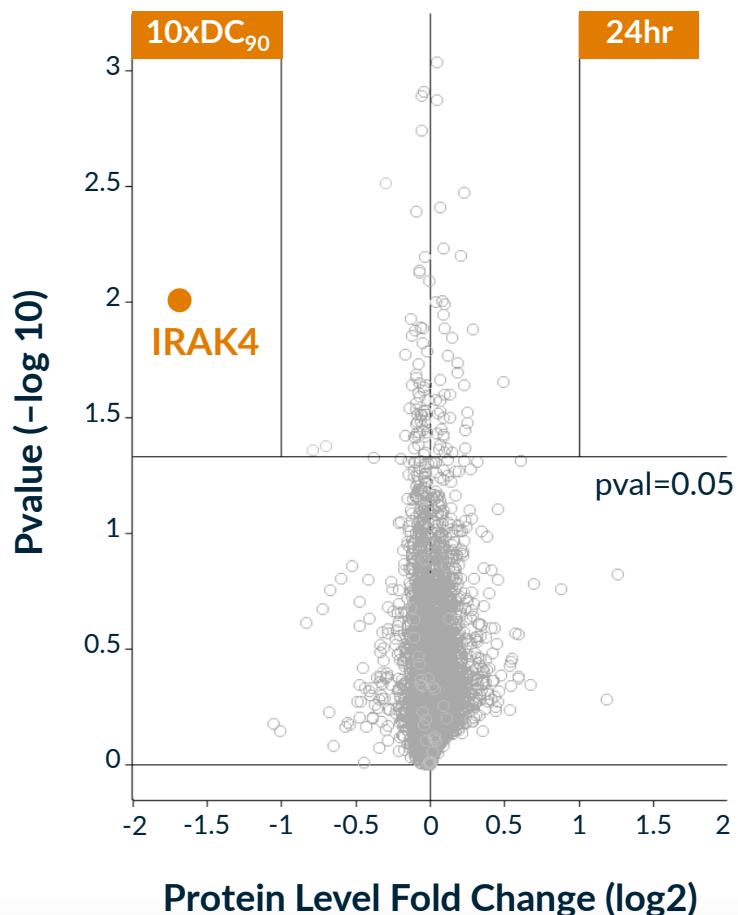
Additional  
Opportunities



- Chronic, pruritic **inflammatory skin disease**
  - Large unmet need for safe and effective oral agents for patients with AD
- 
- Chronic, systemic **autoimmune disease** that can cause irreversible joint damage
  - Multiple therapies targeting the **IL-1R/TLR pathway** are approved
- 
- Chronic and debilitating inflammatory skin disease
  - ~25% of patients with moderate-to-severe disease<sup>4</sup>
  - Adalimumab is approved, which provides some benefit to ~50% of patients with moderate-to-severe disease<sup>5</sup>
- 
- Immune-inflammatory diseases impacted by **IL-1R/TLR pathway**

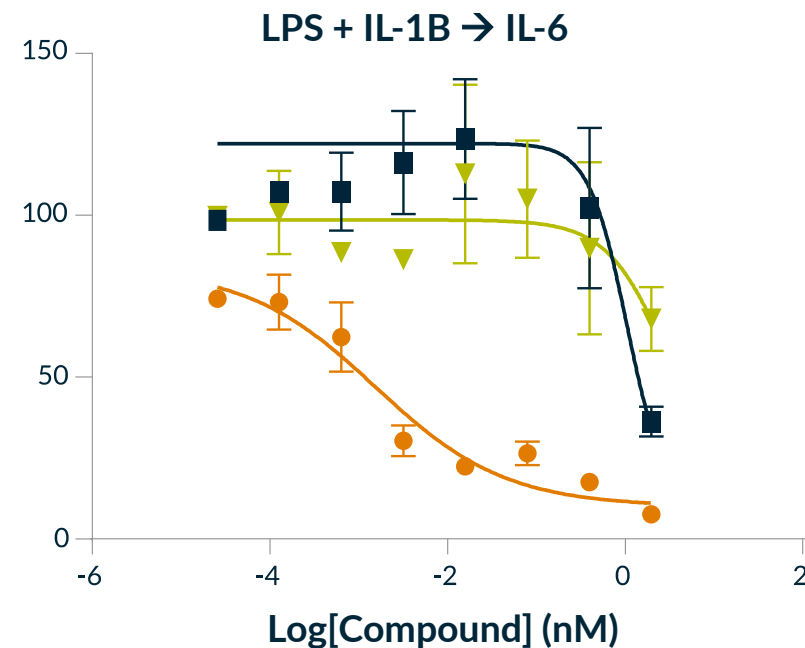
# KT-474: Potent and Specific IRAK4 Degradation Superior to Kinase Inhibition

## Degradation and Selectivity



- KT-474 DC<sub>50</sub> = 2.1 nM in human immune cells
- KT-474 only degraded IRAK4 in human immune cells at concentration 10-fold above the DC<sub>90</sub>
- KT-474 better able to inhibit IL-6 under both LPS and LPS + IL-1B than clinically active IRAK4 SM kinase inhibitor PF-06550833

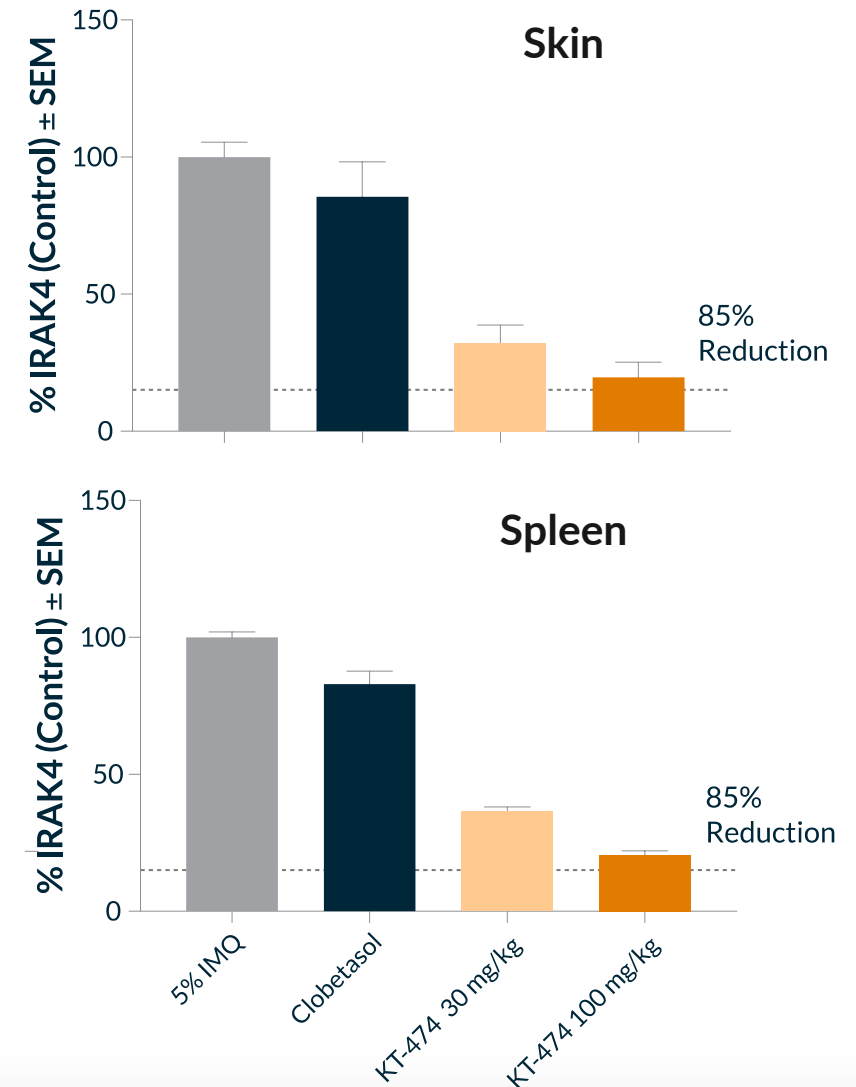
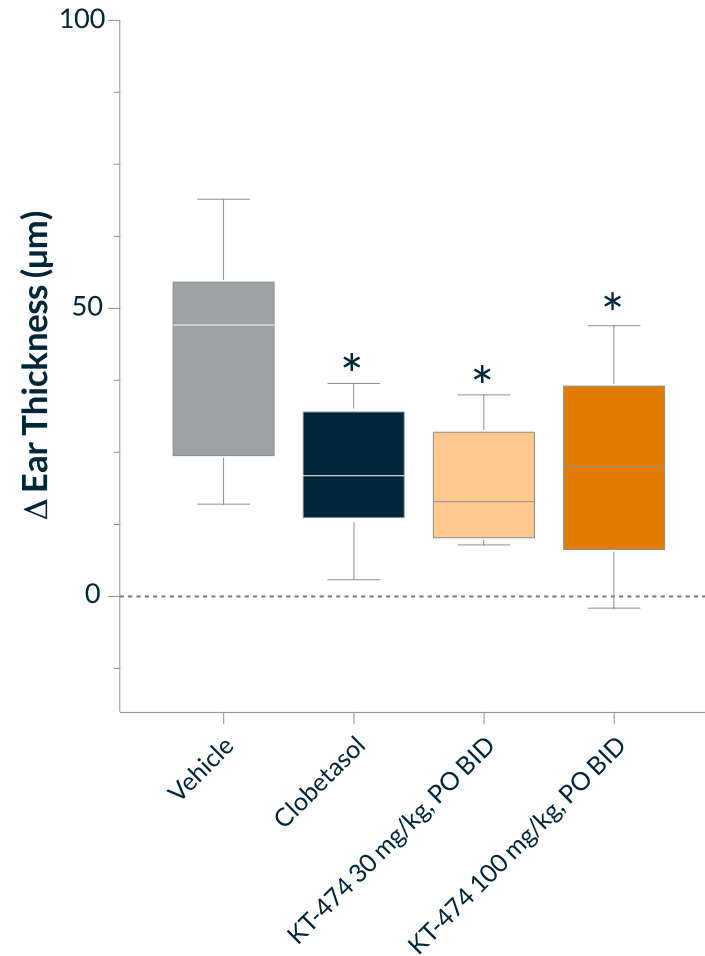
## Superiority over SM kinase Inhibitor



Legend	Compound	IL-6 IC <sub>50</sub> (nM)
●	IRAK4 Degradator	0.8
■	Negative control	450
▼	IRAK4 SMI (PF-06550833)	N/A

# 85% IRAK4 Degradation Sufficient for Maximal *In Vivo* Efficacy in Preclinical Models

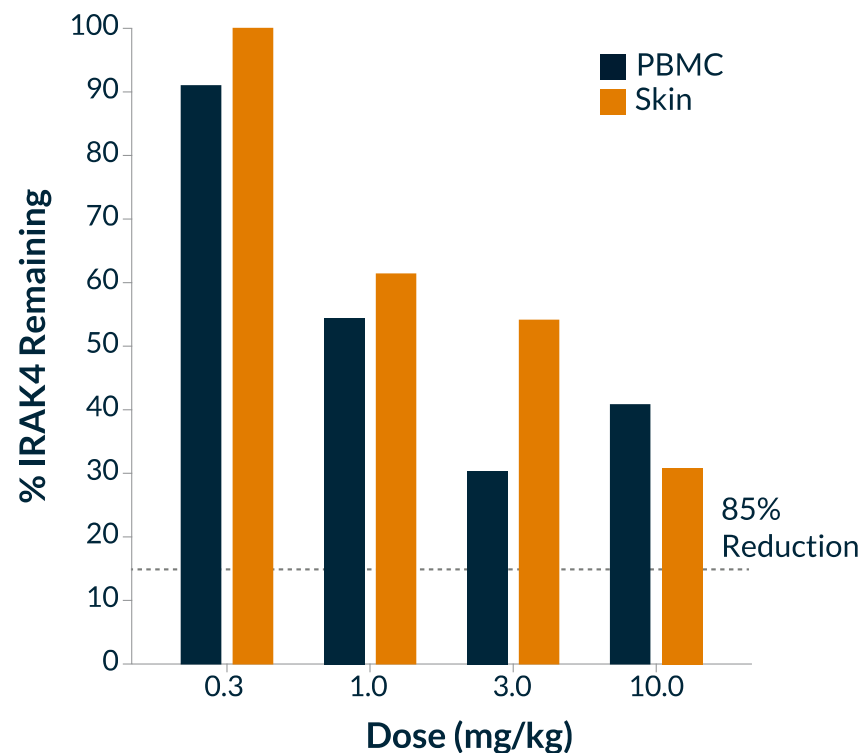
- Ability to inhibit topical skin thickening induced by imiquimod was measured in a mouse model of psoriasis
- Orally dosed KT-474 inhibited thickening, a reflection of local and systemic inflammation, comparable to a topic corticosteroid after 2 or 4 days of dosing
- Full efficacy at doses achieving at 65-80% IRAK4 reduction in skin and spleen. In other models KT-474 has demonstrated full efficacy with 85% degradation



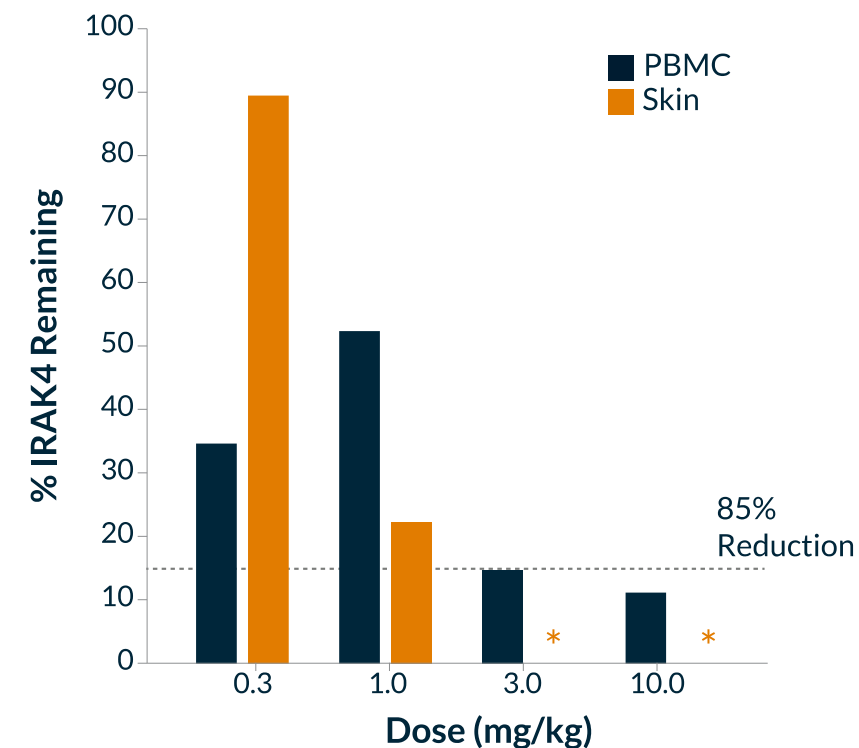
# KT-474 in Dog: Multi-dosing Required to Achieve Target Degradation

- Orally-administered KT-474 achieves >85% knockdown of IRAK4 at Day 7 with repeat dosing in MAD study
- Multiple doses (MAD) lead to optimal degradation profile vs SAD upon reaching steady-state
- Consistency of IRAK4 knockdown observed across peripheral blood mononuclear cells (PBMC) and skin tissue

**Dog Single Ascending Dose (SAD)**  
*IRAK4 Knockdown at Day 1*



**Dog Multiple Ascending Dose (MAD)**  
*IRAK4 Knockdown at Day 7*



\* = Below Limit of Quantitation



# KT-474 Interim Phase 1 SAD Results

# KT-474 Phase 1 Trial Design

Double-blind, Placebo-controlled, Single Ascending Dose (SAD) and Multiple Ascending Dose (MAD) trial

## Three-part Phase 1 Design



- **7 cohorts**  
(up to 56 adult healthy subjects)
- **8 per cohort**  
(6:2 randomization)
- **Single dosing**  
(starting dose 25 mg)
- **5 cohorts**  
(up to 60 adult healthy subjects)
- **12 per cohort**  
(9:3 randomization)
- **14x daily doses**  
(starting dose 25 mg)
- **1 cohort**  
(up to 20 AD and HS patients)
- **Open-label**
- **14x daily doses**

## Endpoints

### Primary

- Safety & tolerability

### Secondary/ Exploratory

SAD & MAD

- Pharmacokinetic measures  
(half-life, bioavailability)
- IRAK4 knockdown in PBMC

### Exploratory

MAD only

- IRAK4 knockdown in skin biopsies
- Proinflammatory cytokine and chemokine levels in skin biopsies
- C-reactive protein and cytokine levels in plasma
- Ex vivo response of whole blood to TLR agonists and IL-1 $\beta$

# KT-474 Phase 1 Trial Goals

Establishing proof-of-mechanism and proof-of-biology

## De-risking Milestones

1

### SAD Portion

Healthy Volunteers

#### Oral Bioavailability and Proof-of-Mechanism

- Efficacious plasma exposures that are safe and well-tolerated
- Proof-of-mechanism with IRAK4 knockdown following single KT-474 dose
- Predictable PK/PD supporting oral daily dosing regimen

2

### MAD Portion

Healthy Volunteers

#### Optimal IRAK4 Reduction and Proof-of-Biology

- $\geq 85\%$  IRAK4 knockdown in skin and blood with daily dosing x 14 days that is safe and well-tolerated
- Proof-of-biology with systemic anti-inflammatory effect: reduction in plasma hsCRP and inhibition of whole blood ex vivo response to TLR agonists and IL-1 $\beta$
- Establishment of maximum effective dose

3

### MAD Portion

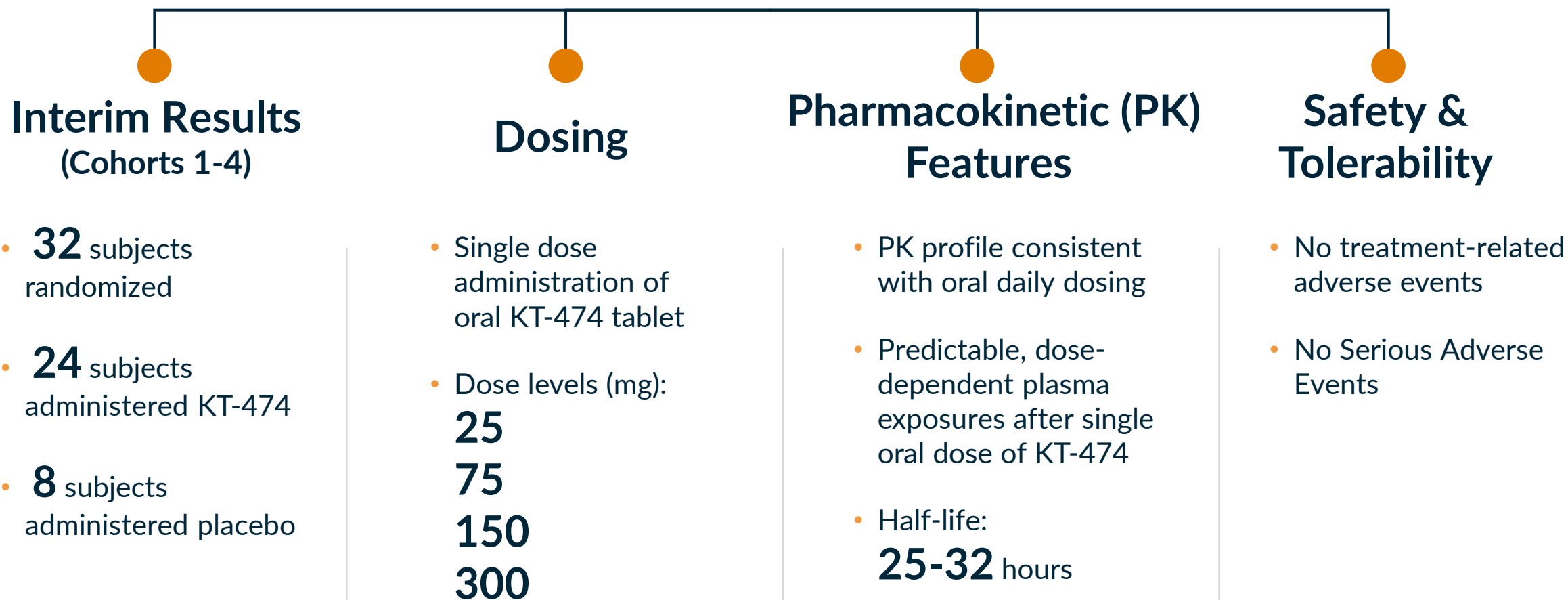
Patient Cohort

#### Establish Proof-of-Biology in Patients

- $\geq 85\%$  IRAK4 degradation in diseased skin and blood
- Anti-inflammatory effect in diseased skin and reduction of plasma cytokines and hsCRP
- Confirmation of dose for subsequent Phase 2 studies



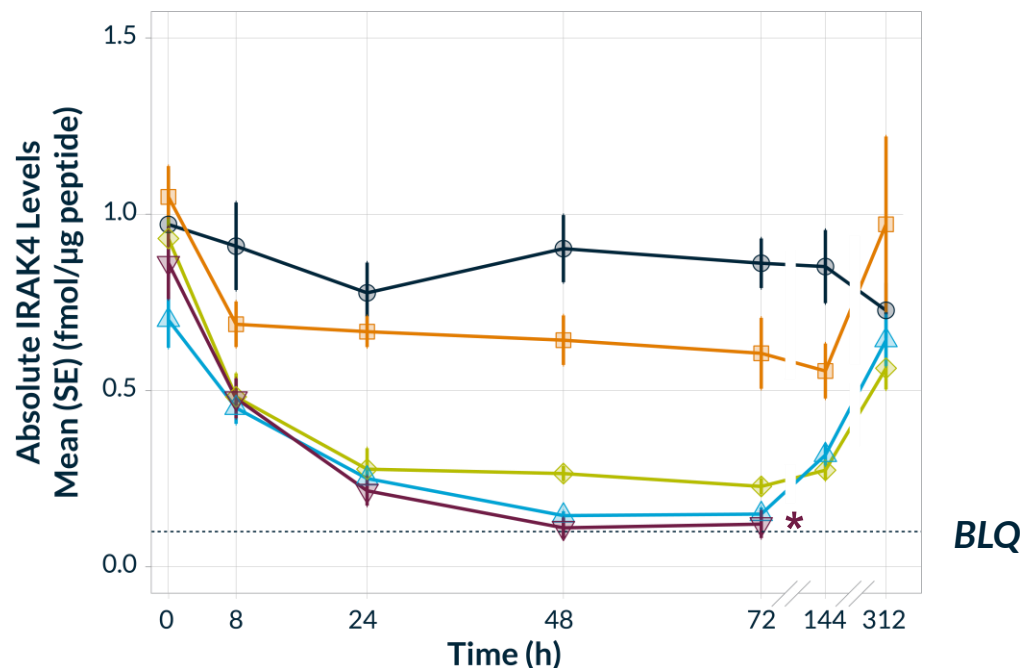
# KT-474 Interim Phase 1 Healthy Volunteer SAD Overview



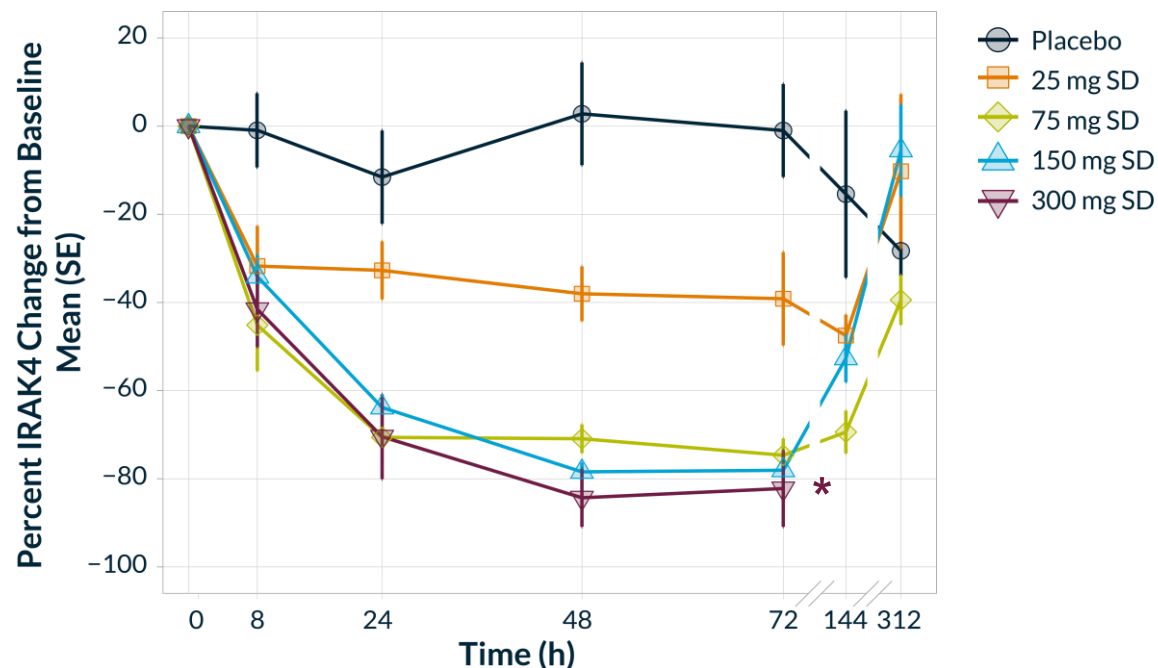


# KT-474 Achieved Profound IRAK4 Degradation after Single Oral Dose that Lasted for at Least 6 Days

## Absolute IRAK4 Levels



## Mean % Reduction of IRAK4

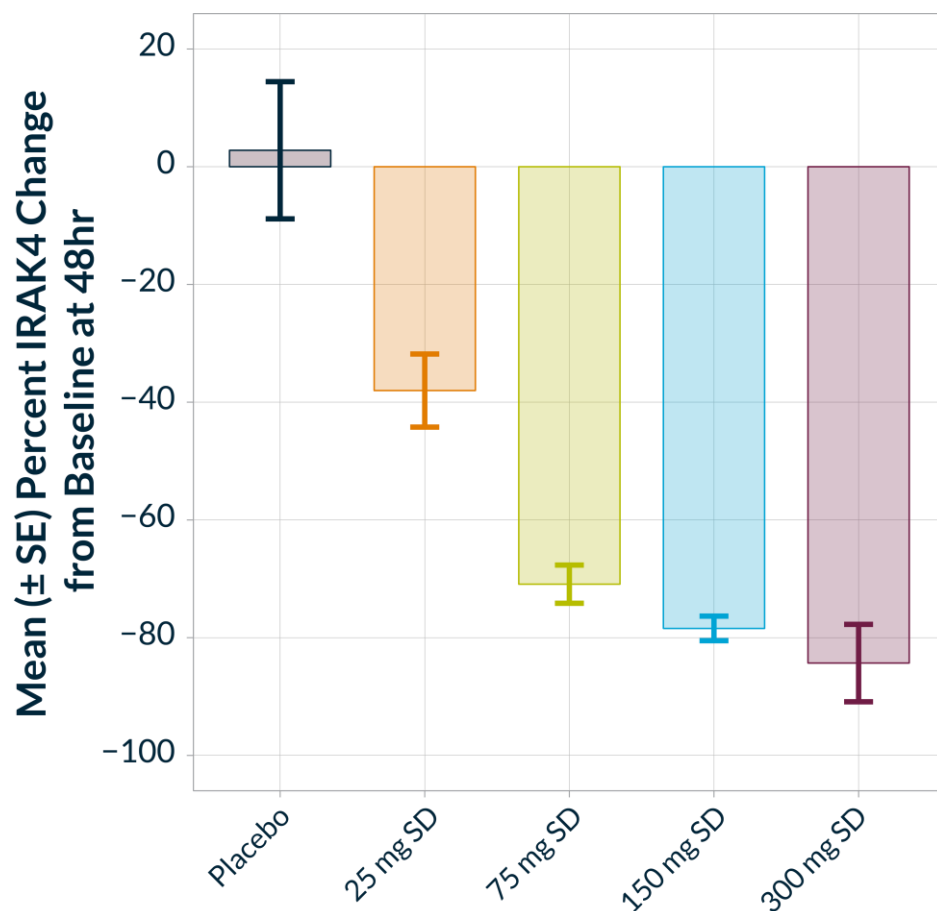


BLQ = Below Limit of Quantitation

\* SAD4 144/312 h PD timepoints pending

- Measured by mass spectrometry in circulating PBMC
- IRAK4 levels nadired at 48-72 hours
- IRAK4 reduction lasted for at least 144h (6 days post-dose) in all dose groups

# IRAK4 Degradation >85% Achieved Following Single KT-474 Dose



## Percent IRAK4 Reduction in PBMC at 48 Hours Post-Dose using Mass Spectrometry

	Placebo (n=8)	Cohort 1 (n=6)	Cohort 2 (n=6)	Cohort 3 (n=6)	Cohort 4 (n=6)
KT-474 dose	-	25 mg	75 mg	150 mg	300 mg
Mean IRAK4 Change	+3%	-38%	-71%	-78%	-84%
Median IRAK4 Change	+16%	-41%	-71%	-78%	-90%
<i>p value*</i>		0.0057	<0.0001	<0.0001	<0.0001

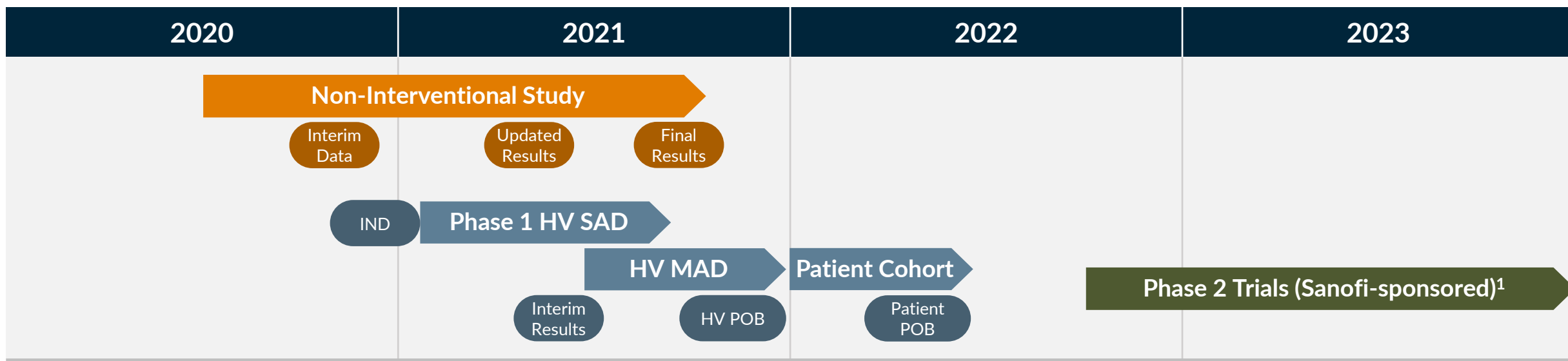
\* *p-values relative to placebo*

# Interim Results from Phase 1 Healthy Volunteer SAD

## Summary and Next Steps

- **KT-474 interim Phase 1 results demonstrate degrader proof-of-mechanism, first time for TPD in a placebo-controlled study**
  - Median IRAK4 reduction of 90% ( $p < 0.0001$  vs placebo) and maximum reduction of 94% at 48 hours following single dose of 300 mg, with sustained degradation that lasted for at least 6 days at all dose levels
  - Based on potency, PK and PD profiles with deep and sustained multiday degradation, expect to achieve biologically relevant (85%) level of degradation with repeat dosing at lower doses; selected MAD starting dose of 25 mg
  - Demonstrated predictable, dose-dependent and biologically active plasma exposures, and half-life that supports oral daily dosing
  - No treatment-related adverse events or serious adverse events observed to date
  - Demonstrating Phase 1 target degradation of >85% de-risks KT-474 ability to reach clinically relevant biological effects in future development as a potential best-in-class anti-inflammatory oral drug
- **FDA lifted partial clinical hold following review of interim healthy volunteer SAD results**
  - Dose escalation in SAD portion of Phase 1 to continue, including assessment of food-effect
  - In July, plan to initiate MAD portion of Phase 1 in healthy volunteers assessing daily dosing of KT-474 for 14 days
- **Expect to present updated results from healthy volunteer SAD/MAD portions in Q4'21**
  - Data to include IRAK4 degradation in skin and PBMC and effects on inflammatory biomarkers after repeat dosing
  - Optimal dose from MAD healthy volunteer portion to be evaluated in an open label cohort of patients with atopic dermatitis and hidradenitis suppurativa

# KT-474 Development Plan



## Non-Interventional

- 40 patients (HS n=30; AD n=10)
- Biomarker endpoints in blood and skin: IRAK4, cytokines, acute phase reactants
- Data updates:
  - Interim: **Oct 2020**
  - Updated HS: **May 2021**
  - Final AD: **2H21**

## Phase 1

- SAD dosing initiated **1Q21**
- SAD/MAD studies: healthy volunteers (HV) and AD/HS patients
- Endpoints: primary - **Safety**; secondary - **Proof-of-Biology**
- Data updates:
  - Interim SAD proof-of-mechanism: **June 2021**
  - HV proof-of-biology: **4Q21**
  - Patient proof-of-biology: **1H22**

## Phase 2

- Randomized, placebo-controlled trials in patients in potential indications such as AD, HS, RA, others





IRAKIMiD

 KYMERA

# IRAKIMiD

A combo in a single molecule

- MYD88 mutation drives differentiation and proliferation in ~25% of diffuse large B cell lymphoma (DLBCL)
- IMiDs downregulate IRF4, increasing IFN signaling and further suppressing NFκB activation and show activity in lymphoma
- Inhibiting both MYD88 and IRF4-dependent NFκB and activating IFN signaling drive cell death in MYD88-mutant lymphomas and leads to full and durable responses *in vivo*
- Combining two therapeutically relevant pathways in a single molecule has the potential to be first single agent targeted therapy agent in lymphoma (MYD88-mut)

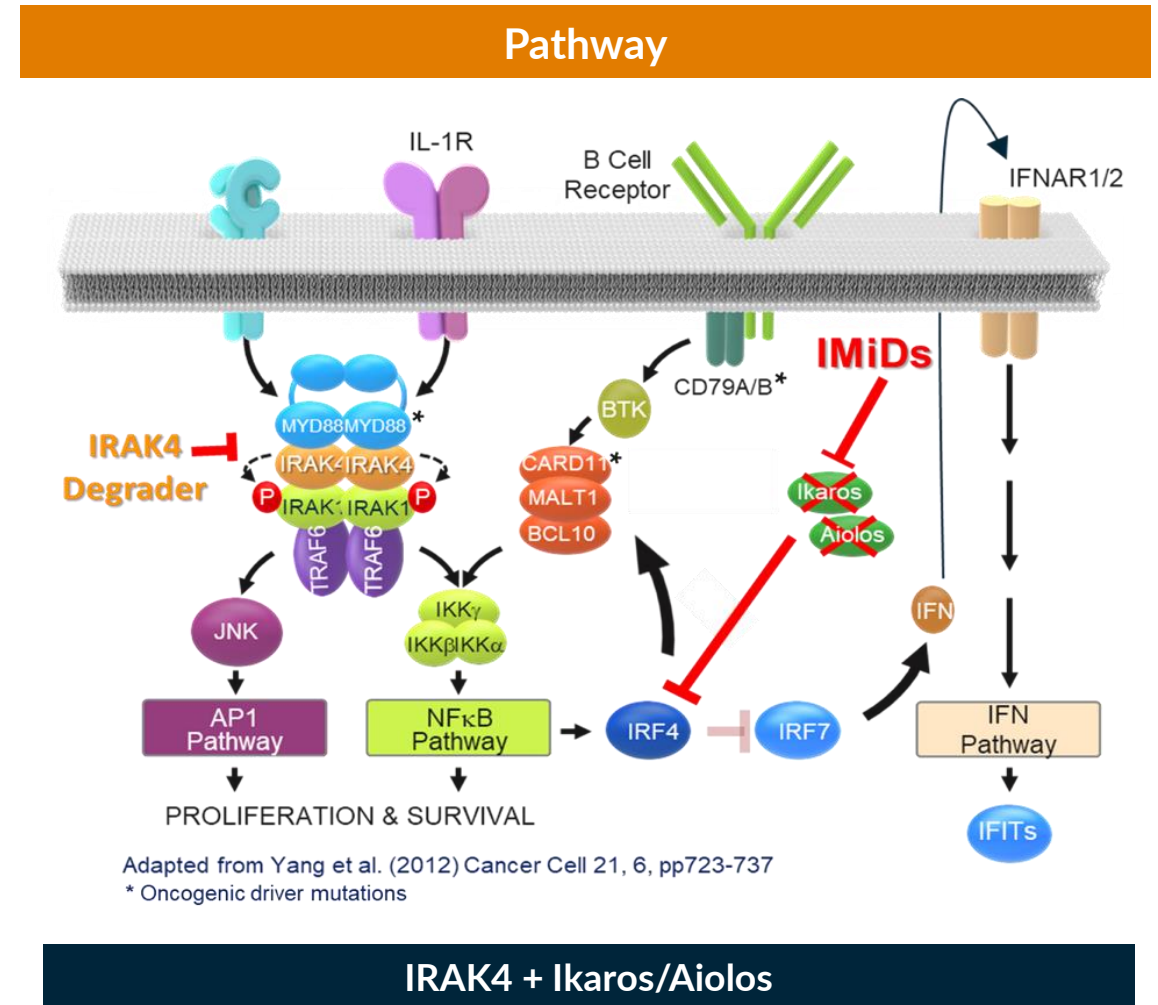
## Indications/Expected Timeline

### MYD88-mutant DLBCL

Current: KT-413 in IND-enabling activities

IND/Phase 1 initiation: 2H 2021

Phase 1 proof-of-biology in patients: 2022





# KT-413 Opportunity

Potential to be first precision medicine in DLBCL to target a genetically defined population (MYD88-mut)

MYD88-mutant  
DLBCL

Patient Impact (U.S.)

~7.0k<sup>1</sup>  
per year

Other  
MYD88-mutant  
B cell Lymphomas

>1.0k<sup>2</sup>  
per year

Additional  
Cancers

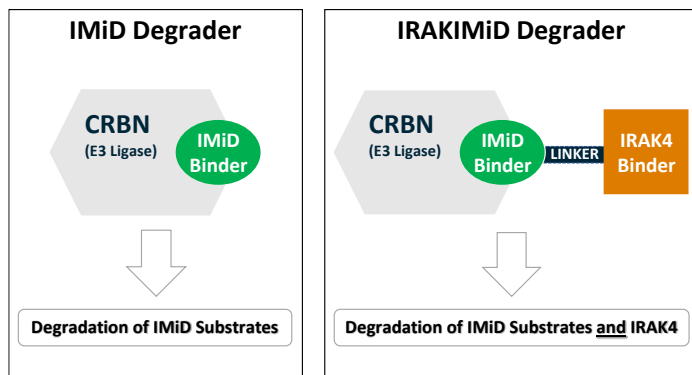


- MYD88 is mutated in at least 25% of DLBCL patients, the most common subtype of non-Hodgkin's lymphoma<sup>1</sup>
- Front-line treatment includes **R-CHOP** (chemo/rituximab)
- DLBCL 5-year survival rate is ~64%, and MYD88 mutations in DLBCL are associated with poorer survival following frontline R-CHOP chemotherapy<sup>3</sup>

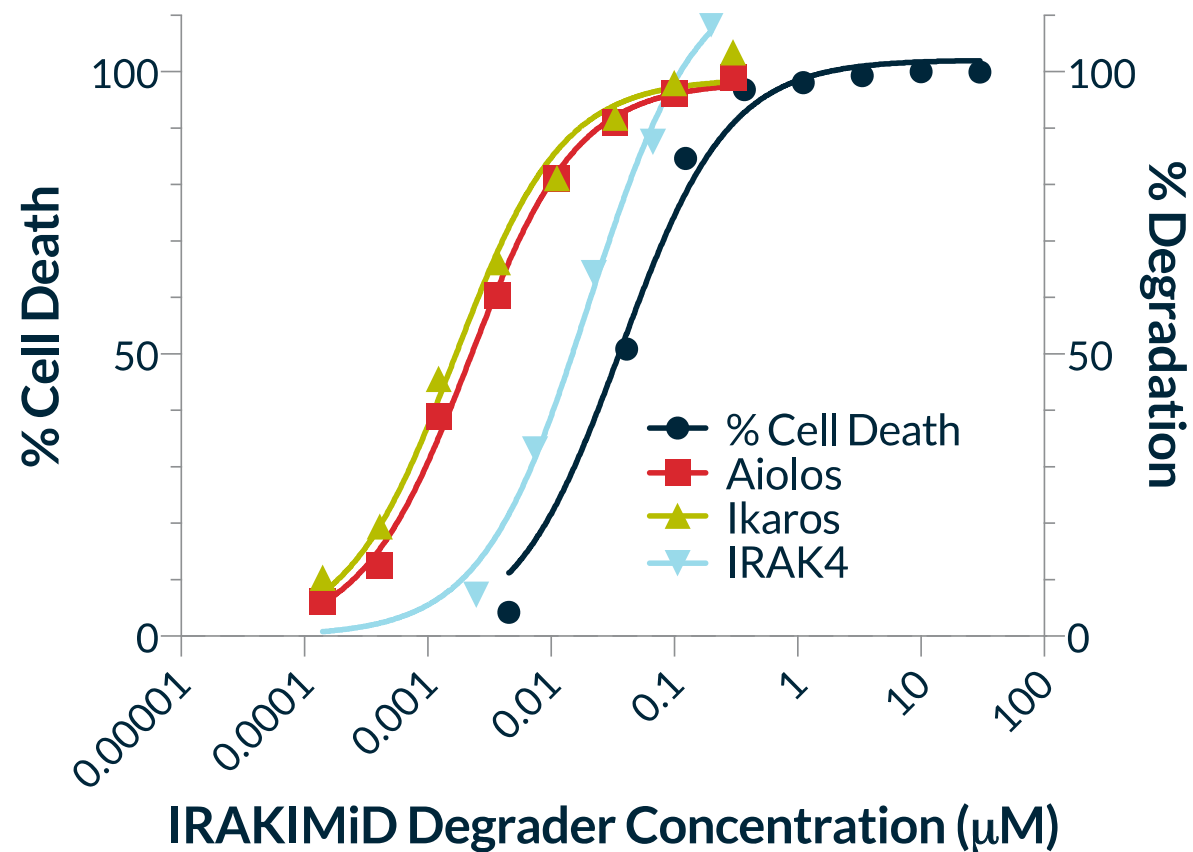
- MYD88 is mutated in approximately 90% of **Waldenström's macroglobulinemia** cases and 70% of primary central nervous system lymphoma<sup>4,5</sup>

- **IL1R/TLR/NFκB**-driven cancers, AML & MDS subsets with long isoform of IRAK4 (IRAK4L)

# Degradation of IRAK4, Ikaros and Aiolos Correlates to Cell Killing



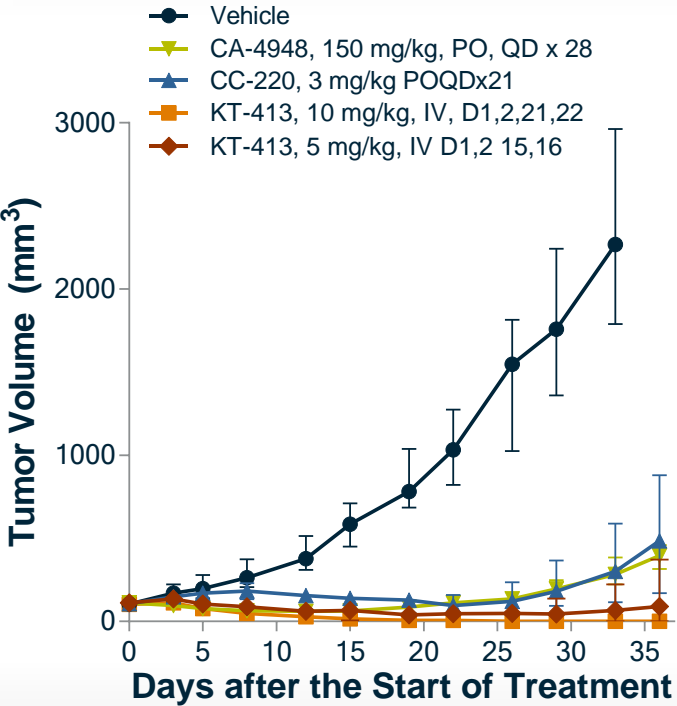
- IRAK4, Ikaros and Aiolos degradation measured in MYD-88-mutated OCI-Ly10 cells after 24 h of drug exposure
  - $IRAK4\ DC_{50} = 4\text{ nM}$
  - $Ikaros/Aiolos\ DC_{50} = 2/2\text{ nM}$
- Degradation correlates with cell killing effects
  - $IC_{50} = 31\text{ nM}$





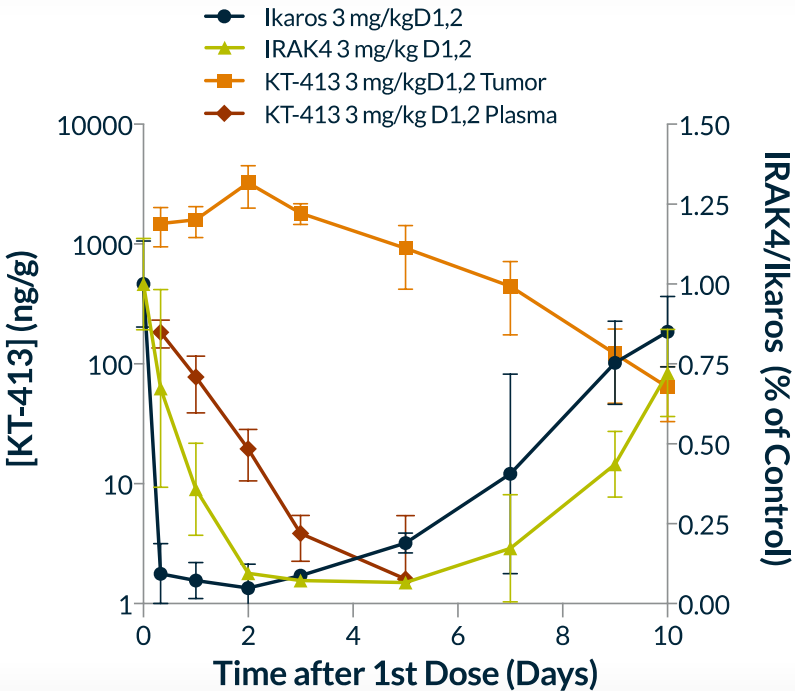
# KT-413 is Highly Active on Intermittent Dosing Regimens and Superior to Clinically Active CA-4948 and CC-220

- In the OCI-Ly10 MYD88<sup>MT</sup> xenograft model, intermittent dosing of KT-413 induced strong antitumor activity, including complete or partial regressions
  - Superior activity compared to the clinically active IRAK4-inhibitor CA-4948 or the latest generation IMiD CC-220 alone
- Minimally active dose of 3 mg/kg D1,2 showed extended tumor exposure and strong degradation of both IRAK4 and IMiD substrates that was maintained for at least 72h



Drug	CR	PR	SD	PD
CA-4948	0	0	3	4
CC-220	0	1	4	2
KT-413 (5 mpk)	2	2	3	-
KT-413 (10 mpk)	5	2	-	-

CR: <10mm<sup>3</sup> tumor on D26  
PR: >50% regression from baseline  
SD: <50% regression to 20% increase in tumor volume  
PD: >20% tumor growth on D26



# KT-413 Shows Regressions in MYD88<sup>MT</sup> Patient-Derived Xenograft Models

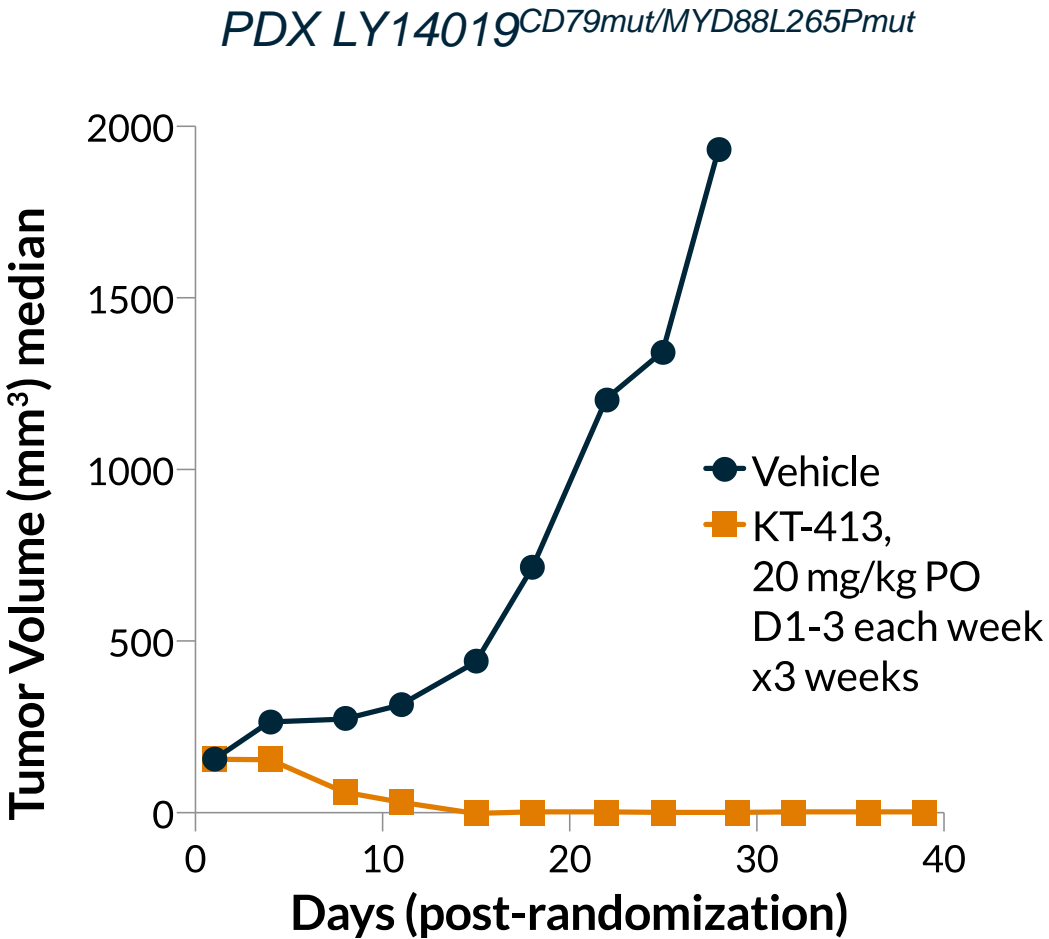
Model	MYD88	CD79B	TNFAIP3	Other	KT-413 (%TGI)
LY14019	L265P	MT	MT		100
LY2264	L265P	MT		IRF4	100
LY2298	L265P	MT		BCL2/BCL6	90
LY12699	L265P	MT			87
LY2345	WT		MT		70
LY2301	WT				30
LY0257	L265P			BCL2/BCL6/IKZF3	0

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

- Activity is observed regardless of co-mutations that activate NFkB and IRF4 pathways
- The non-responsive MYD88<sup>MT</sup> 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

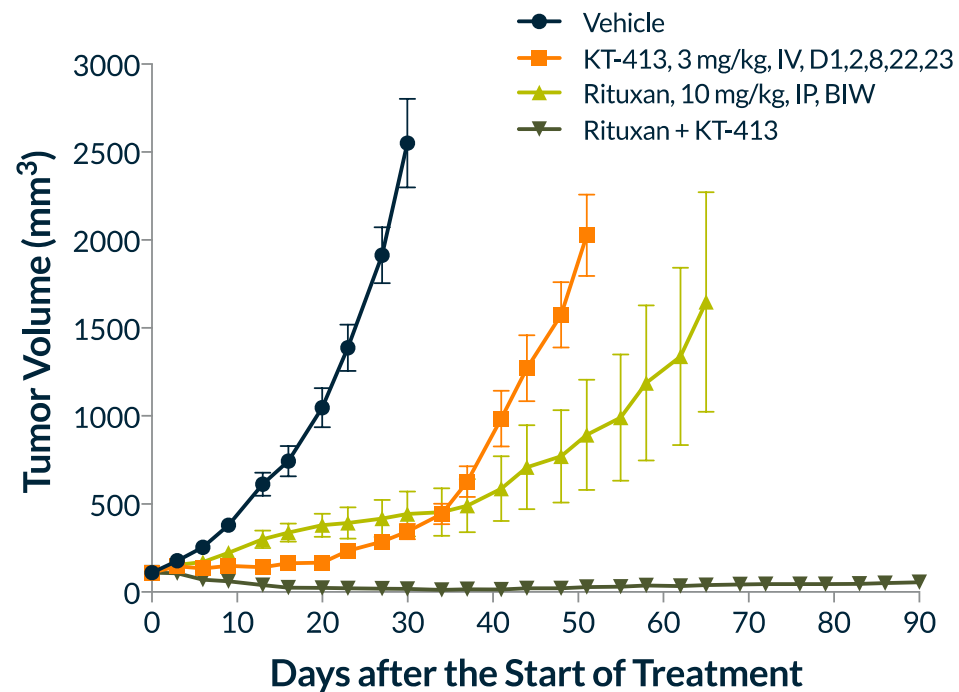
**Some level of tumor growth inhibition observed in MYD88-WT PDX**

- May be consistent with IMiD activity of KT-413

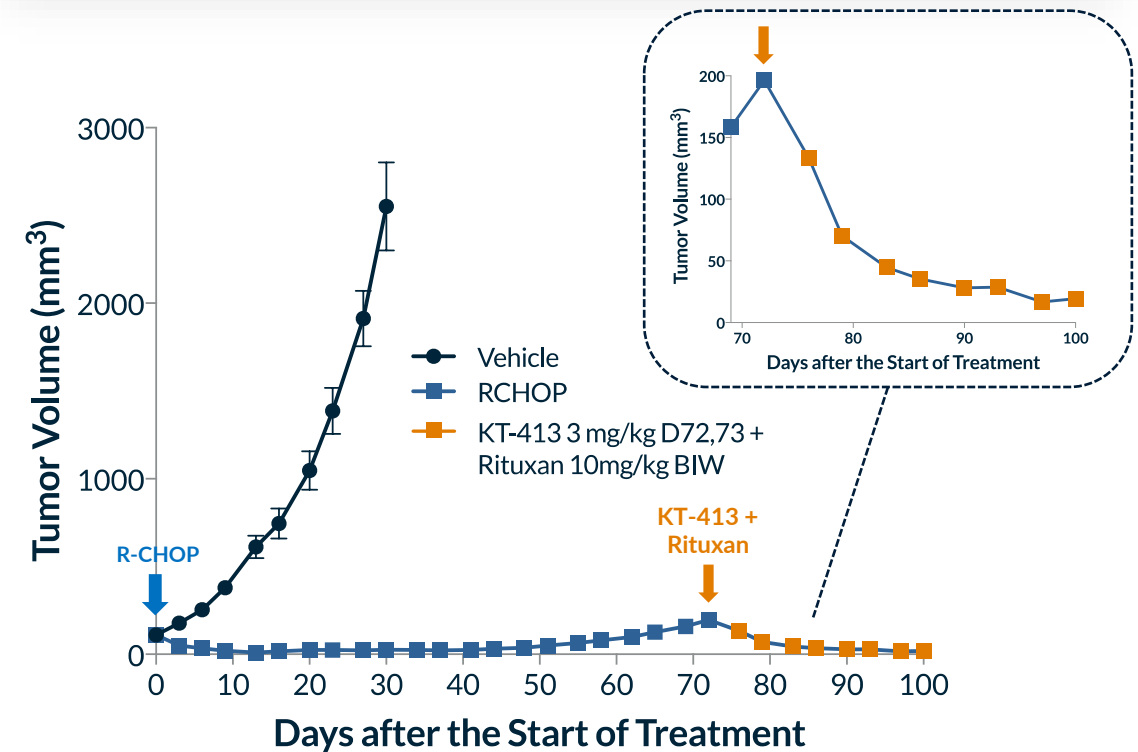


# KT-413 has Synergistic Activity in Driving Deep Tumor Regressions in Combination with Other Therapies in Preclinical Models

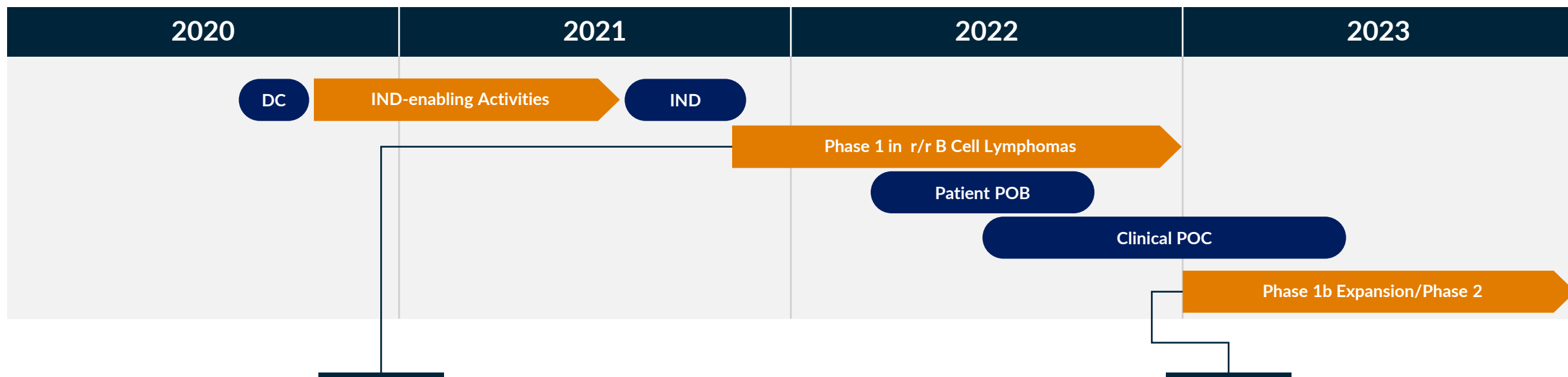
- KT-413 administered on intermittent schedules demonstrated deep and durable regressions in combination with rituximab in MYD88MT OCI-Ly10 xenografts



- KT-413 + rituximab showed strong tumor regressions in tumors that relapsed following initial R-CHOP treatment



# KT-413 Development Plan



- **Multi-center Phase 1 dose escalation study (US sites) start in 2H21**
- Relapsed/refractory B cell lymphomas, including MYD88-mutant DLBCL
- Objectives include safety, tolerability, PK and PD (proof-of-biology) and preliminary clinical activity
- Clinical and biomarker endpoints
- **POB to be presented in 2022**

- Phase 1b expansion cohorts in DLBCL (MYD88-mut and -wt) and other MYD88-mut lymphomas, including Waldenstrom's macroglobulinemia and primary central nervous system lymphoma
- Objectives include safety, tolerability and clinical activity of monotherapy and select combinations
- Clinical and biomarker endpoints
- Potential expansion in other indications





**STAT3**



# STAT3 Biology and Degradar Rationale

- STAT3 is a traditionally largely undrugged transcription factor activated through cytokine and growth factor receptors via JAKs and non-JAKs mediated mechanisms
- High degree of validation of JAK-STAT pathway in oncology and immuno-oncology supported also by numerous publications
- STAT3 plays a role in tumor biology, evasion of immune surveillance and inflammation/fibrosis
- No known drugs specifically affect STAT3 broadly across all relevant cell types

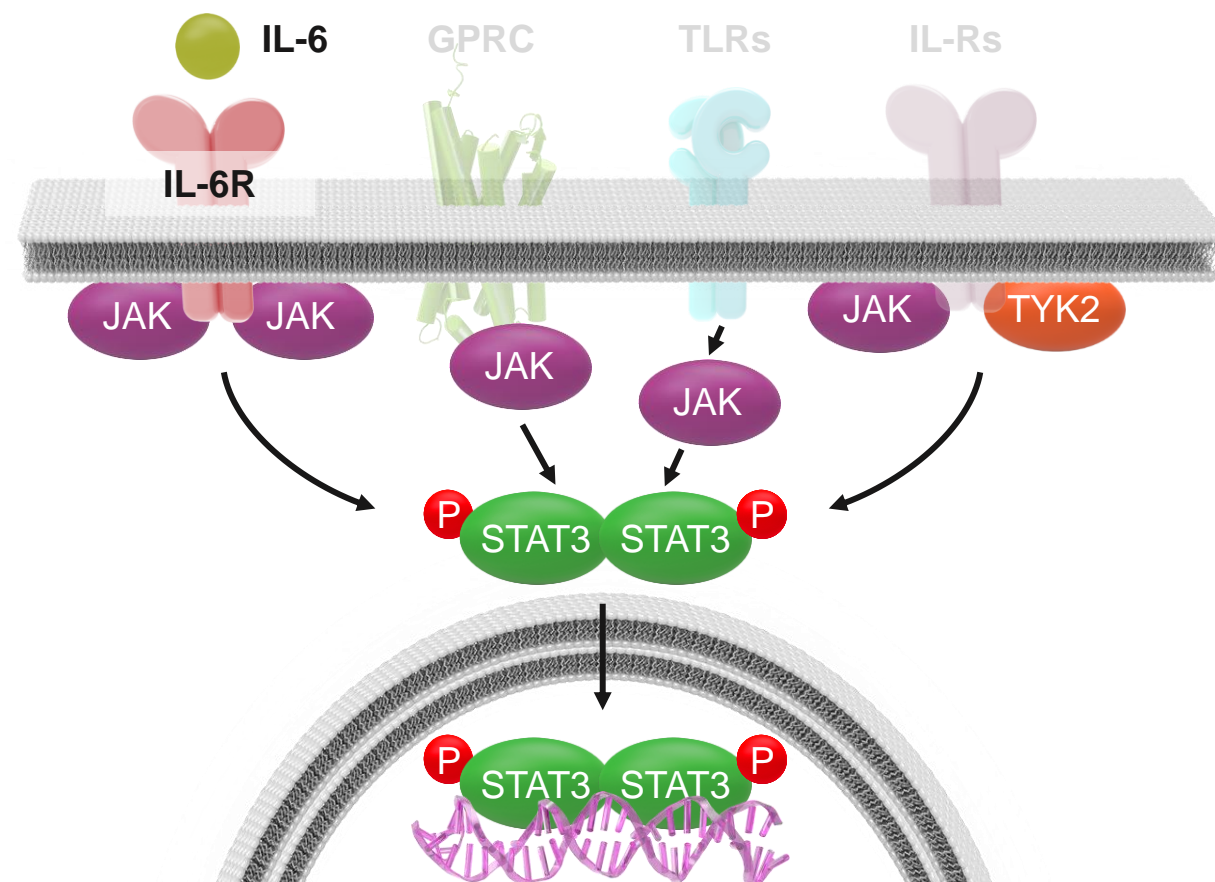
## Indications/Expected Timeline

### Hematological Malignancies/Solid Tumors and Autoimmune/Fibrosis

Nomination of development candidate: 1Q 2021 ✓

IND/Phase 1 initiation: 4Q 2021

Phase 1 proof-of-biology in patients: 2022



# STAT3 Opportunity in Oncology & Autoimmunity

First-in-class opportunity to address STAT3-driven pathology across large and diverse indications

## Patient Impact (U.S.)

### Cancer

**~5.0k** per year<sup>1</sup>  
Peripheral T-cell Lymphoma

**~2.0k** per year<sup>2</sup>  
Cutaneous T-cell Lymphoma

**~200.0k** per year<sup>3</sup>  
NSCLC

Liquid Tumors

**Genetically-defined STAT3 mutation and/or hyperactivation**

*PTCL, CTCL, T-LGL leukemia*

**STAT3 activation and dependency**

*DLBCL, AML, multiple myeloma*

Solid Tumors

**Cell Intrinsic: STAT3 role in EMT/TKI resistance**

*Combinations in TKI / chemotherapy resistant settings*

**Cell Extrinsic: STAT3 role in IO**

*T-cell infiltrated tumors. Combinations with immune-modulators*

### I/I Fibrosis

**>40.0k**<sup>4</sup>  
Systemic Sclerosis

**>16.0M**<sup>5</sup>  
Atopic Dermatitis

**>40.0k**<sup>6</sup>  
Idiopathic Pulmonary Fibrosis

Autoimmune

**STAT3 GOF syndrome**

*Genetically-defined population characterized by enteropathy, arthritis, dermatitis, lung disease*

**Immune-inflammatory**

*Systemic sclerosis, atopic dermatitis, rheumatoid arthritis, Crohn's disease/ulcerative colitis*

Fibrosis

**Chronic inflammation / fibrosis**

*Idiopathic pulmonary fibrosis, CKD/renal fibrosis*

# KT-333 Highly Specific Degradation of STAT3

CANCER

Liquid Tumors

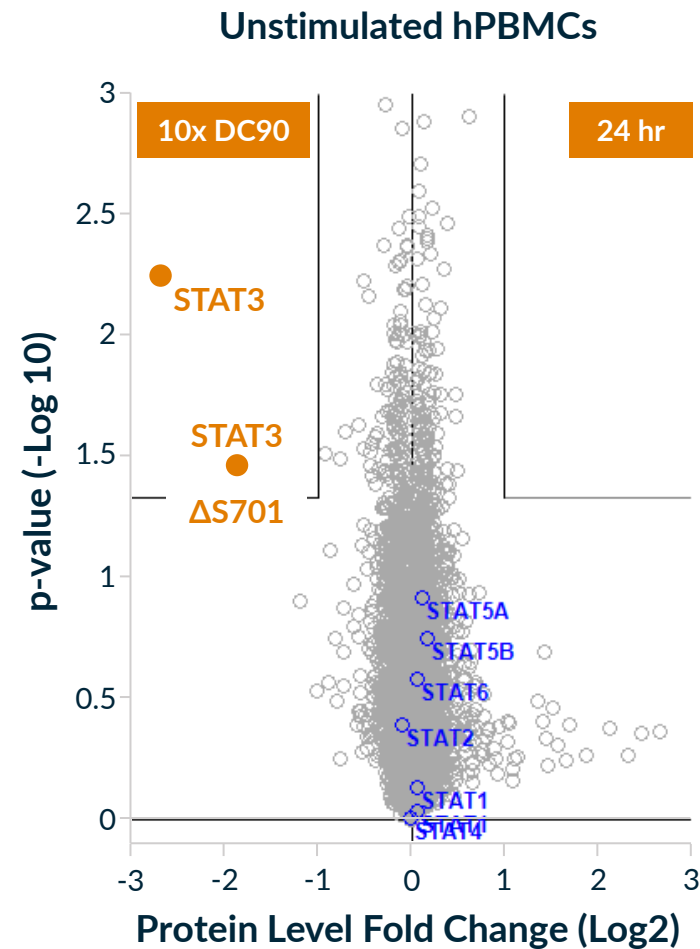
Solid Tumors

I/I  
FIBROSIS

Autoimmune

Fibrosis

- Deep mass spectrometry-based proteomics to assess STAT3 specificity performed
- hPBMC and tumor cells (SU-DHL-1) treated with KT-333 degrader
- STAT3 was the only protein to be degraded with statistical significance
- Data demonstrate highly selective degradation profile



● STAT Family Members: STAT1, STAT2, STAT3, STAT4, STAT5A, STAT5B, STAT6



# Full and Durable Regressions Across Multiple *in vivo* Preclinical Tumor Models

CANCER

Liquid Tumors

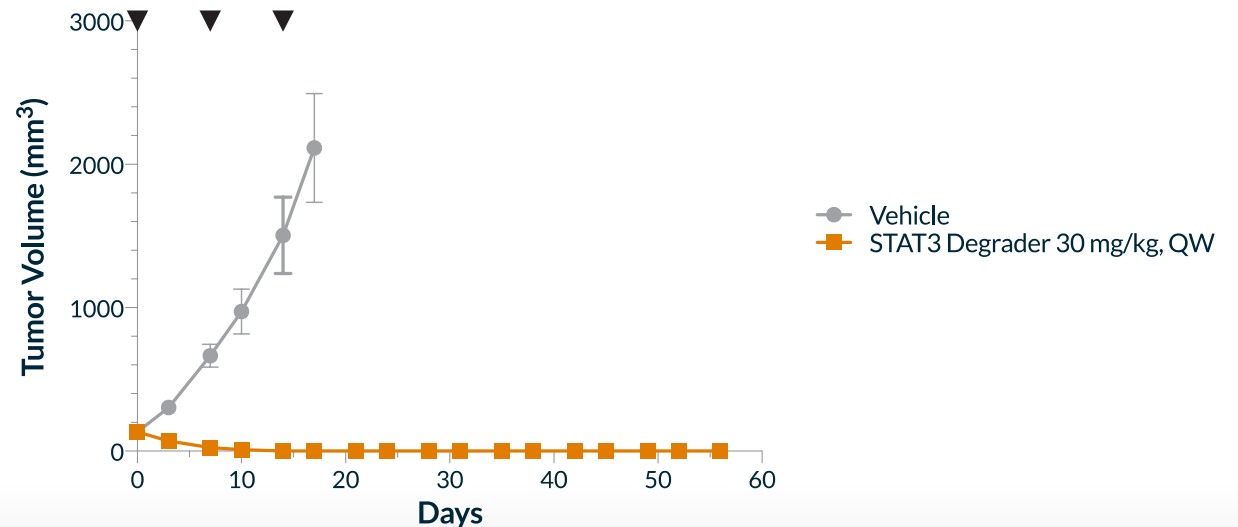
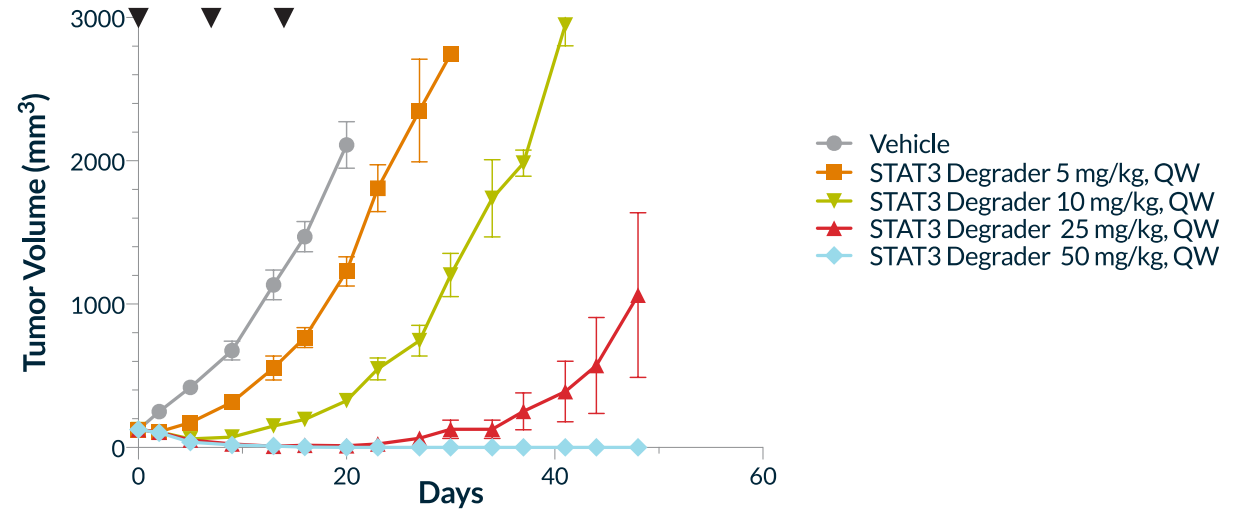
Solid Tumors

- Mice bearing STAT3-dependent ALK+ ALCL SU-DHL-1 (above) and STAT3-driven ALK+ ALCL xenograft model SUP-M2 (below) tumors dosed with STAT3 degrader
- Dose and degradation dependent tumor growth inhibition observed with once-a-week IV dosing
- 30 mg/kg sufficient to drive full tumor regression that was durable for multiple weeks after the last dose

I/I  
FIBROSIS

Autoimmune

Fibrosis



# STAT3 Degradar *In Vivo* Active in Preclinical PD-1/L-1 Refractory Solid Tumor Model

CANCER

Liquid Tumors

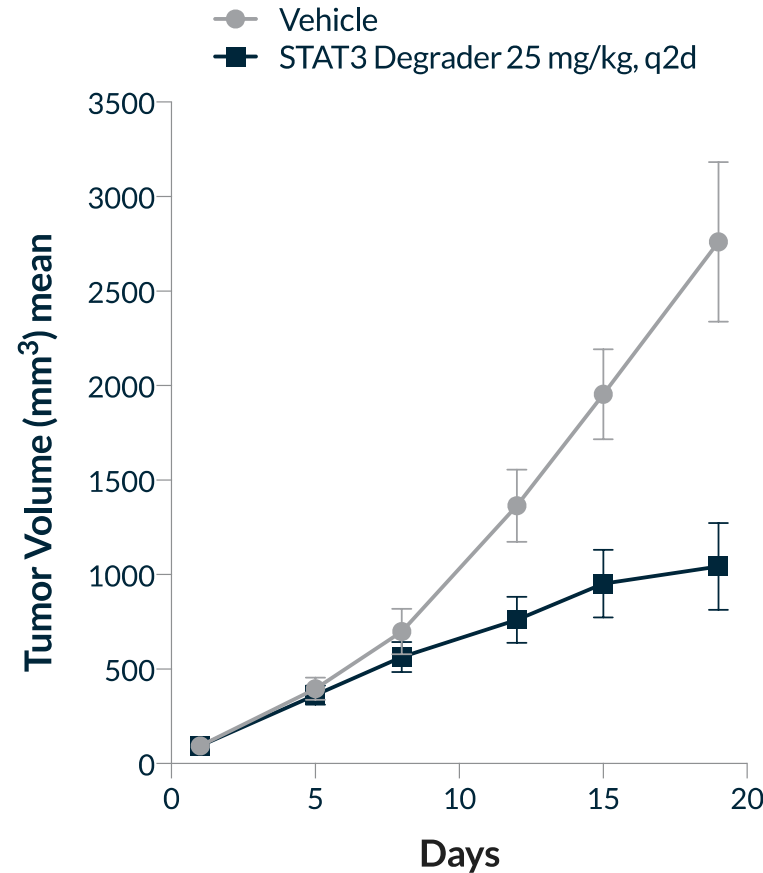
Solid Tumors

I/I  
FIBROSIS

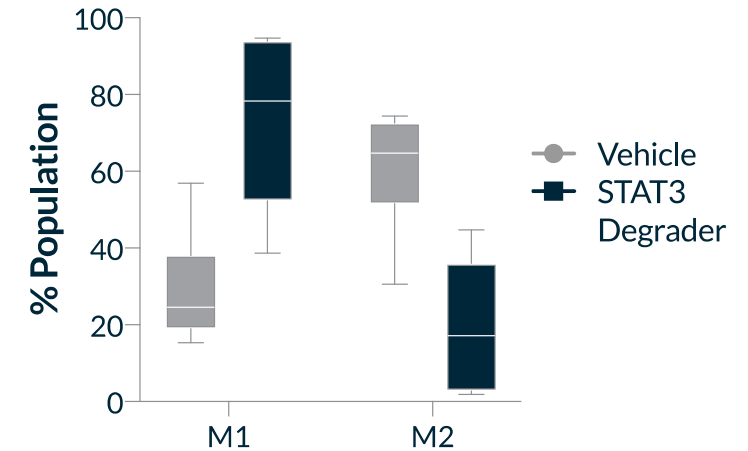
Autoimmune

Fibrosis

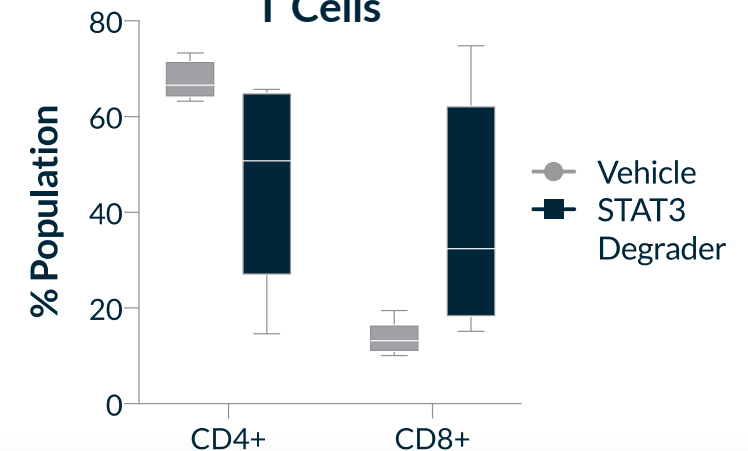
- Kymera's STAT3 degrader assessed in colorectal cancers (CT-26) known to be refractory to approved immunotherapies
- STAT3 degrader significantly reduced tumor growth when administered every two days
- Analysis of tumors showed synergistic modulation of immune cells (M2/M1 and T cells) within the tumor microenvironment to favor an anti-tumor response



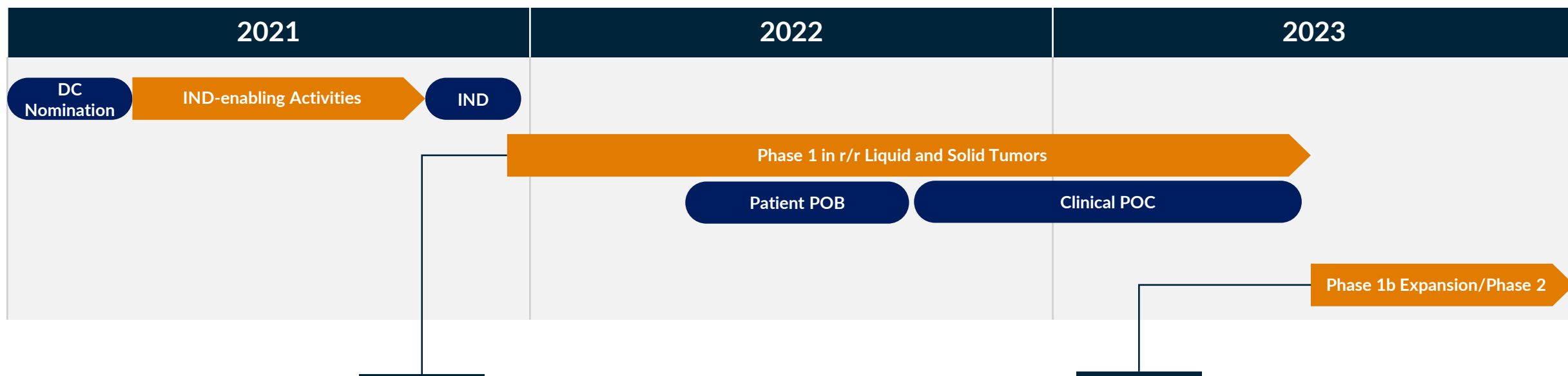
## Macrophages (M1/M2)



## T Cells



# STAT3 Degradar Development Plan in Liquid & Solid Tumors



- **Multi-center Phase 1 dose escalation study start in 4Q21**
- Safety, tolerability, PK and PD (proof-of-biology) and preliminary clinical activity
- Clinical and biomarker endpoints
- **POB to be presented in 2022**

- Phase 1b expansion cohorts in STAT3-dependent liquid tumors
- Objectives include safety, tolerability and clinical activity of monotherapy and select combinations
- Separate Phase 2 in solid tumors

# THANK YOU



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The bottom section of the slide features a large, horizontal banner. On the left side of the banner, the Kymera logo is displayed, with the 'K' and 'Y' in orange and 'YMER A' in white. The background of the banner is a composite image. The left half shows abstract, glowing blue and purple lines and shapes. The right half shows a night sky with a starry background and a constellation of stars connected by thin white lines. Below the sky, there is a dark silhouette of a mountain range and a forest of trees.