Targeting STAT3 with Selective Protein Degraders for the Treatment of PTCL

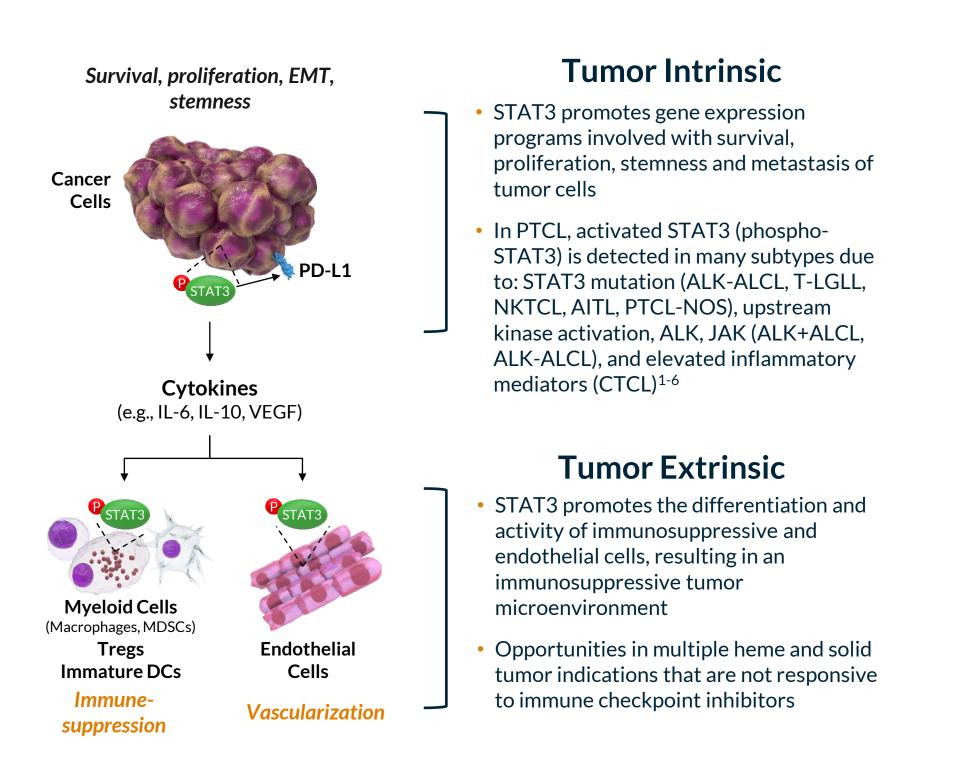
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INTRODUCTION

of Signal Transducer and Activator of Transcription 3 (STAT3) is dysregulated in many cancers including aggressive hematological malignancies with high unmet medical need. Aberrant activation of STAT3 can promote the establishment and progression of malignant cells through regulation of cell survival and proliferation pathways and suppression of anti-tumor immunity, also known as tumor intrinsic and tumor extrinsic mechanisms, respectively

Selective targeting of STAT3 has been challenging, but targeted protein degradation mediated by heterobifunctional small molecule degraders is a novel therapeutic modality to target difficult-to-drug oncogenic proteins. These molecules bind to both the target protein and an E3 ligase, enabling the formation of a ternary complex which leads to ubiquitination and proteasomal degradation of the target protein.

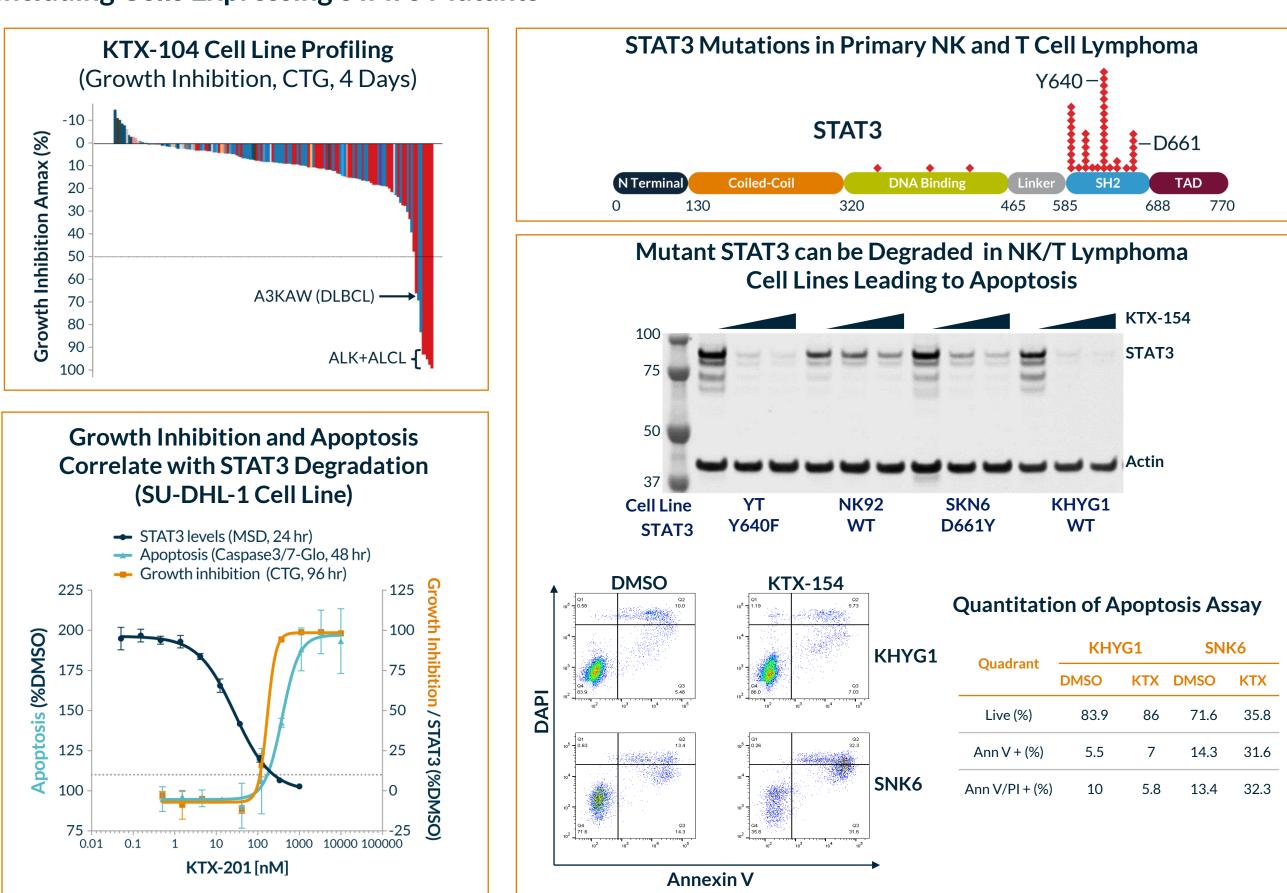


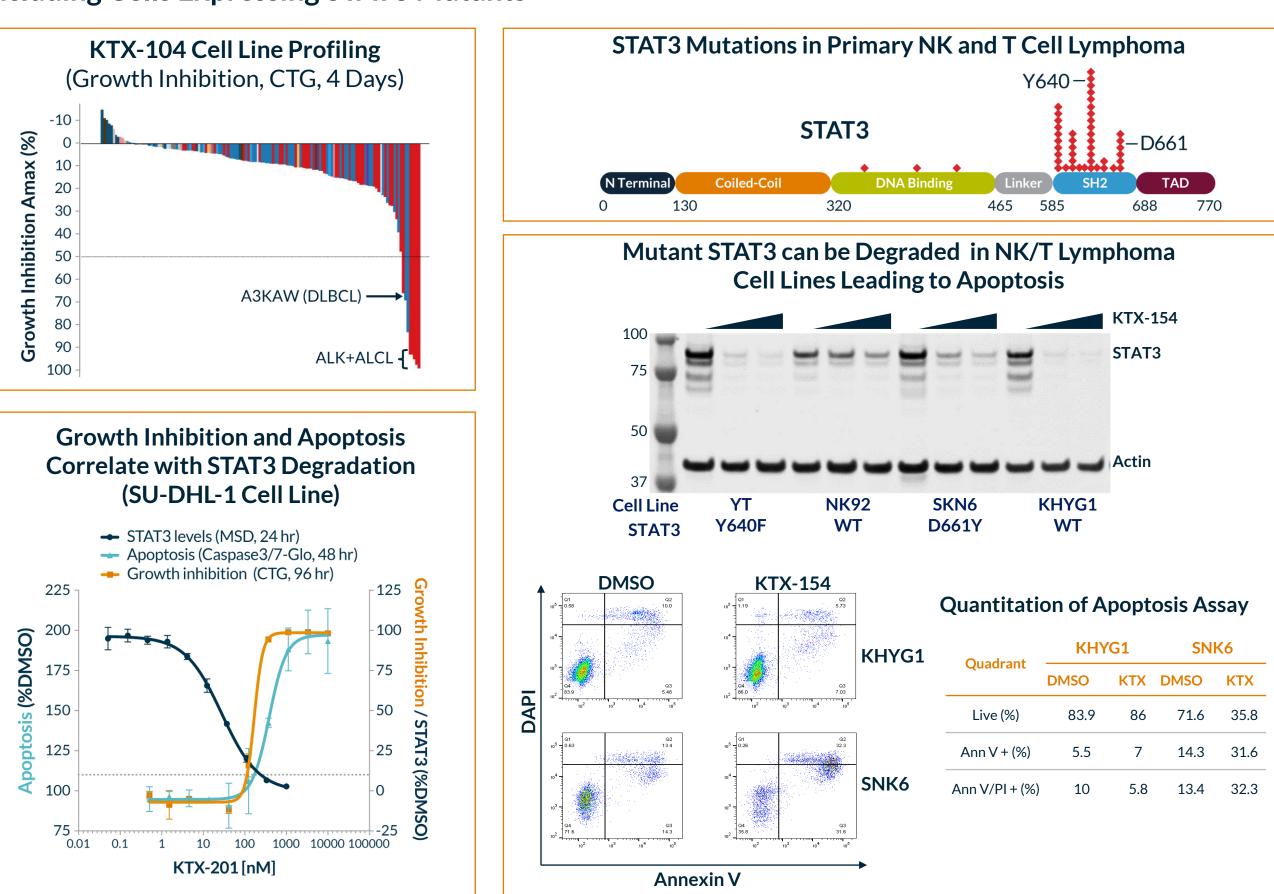
AIM

We have discovered a series of heterobifunctional STAT3 degraders that potently and selectively degrade STAT3. We evaluated the in vitro and in vivo activity and molecular mechanisms of degrading STAT3 in models representing different subtypes of mature T and NK cell lymphoma.

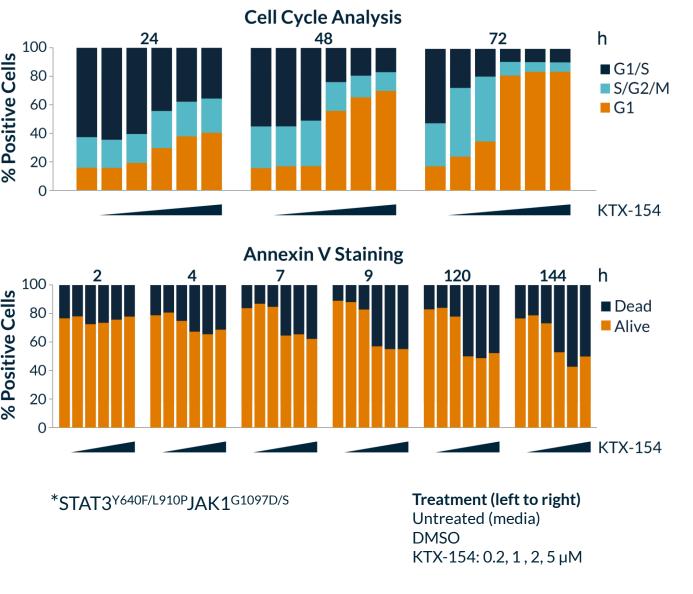
RESULTS

Figure 1: STAT3 Degraders are Highly Active against NK and T Cell Lymphoma Cell Lines Including Cells Expressing STAT3 Mutants





Time and Dose-dependent Cell Cycle Arrest and Cell Death of the BELLI (STAT3^{mt}JAK1^{mt*)} ALK-primary ALCL Model



METHODS

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In vitro Assays: All cell lines were cultured according to recommended procedures unless otherwise noted. For degradation assays, cells were treated with compounds for 24 h, and total STAT3 protein assessed by western blotting or MSD assay.

Viability was assessed using cells treated with degraders for 4 days and assayed by CTG (SU-DHL-1) or MTT (KHYG1, SNK6) assay. For KHYG1 and SNK6 assays, cells were cultured in low IL-2 (50 ng/mL). Cell cycle and cell death analysis was performed by flow cytometry using cells stained with propidium iodide (PI) or PI and AnnexinV (BELLI) or DAPI/Annexin V(KHYG1 SNK6).

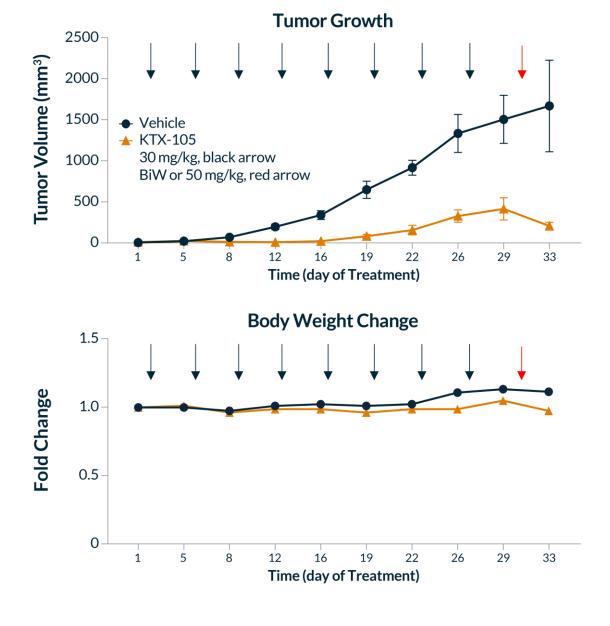
In vivo Experiments: Subcutaneous xenografted tumors were established in immune compromised mice. STAT3 degraders were formulated in buffered PBS and administered by iv infusion. For PK/PD study, a single administration of STAT3 degrader was administered on day 0. Plasma and tumor levels of drug were measured by MS, and tumor STAT3 levels measured by western blotting.

Transcriptomic and Proteomic Analysis: RNAseq was performed on cells treated with STAT3 degraders to identify differentially expressed genes. Pathway analysis was performed using Gene Set Enrichment Analysis (GSEA).

Proteomics was conducted using Tandem Mass Tag. Protein association was evaluated by STRING analysis with Gene Ontology (GO) classifiers.

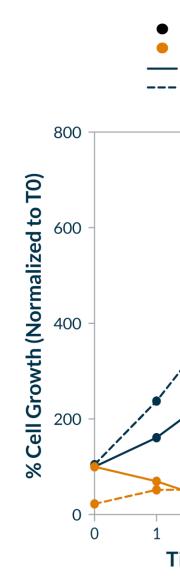
Figure 2: Activity Against STAT3-mutant ALK-negative ALCL Patient Derived Cells

Intermittent Dosing of KTX-105 Significantly Inhibits Growth of BELLI Xenograft Tumors and is Tolerated



Intermittent (QW) Dosing

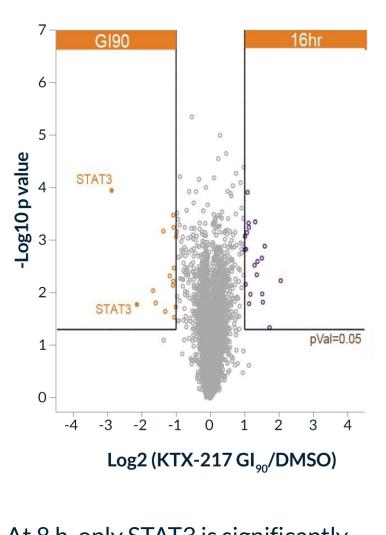




• Strong anti-tumor activity is observed when 90% or greater degradation of STAT3 is maintained for at least 48 h in vitro and in vivo in ALK+ALCL

Rapid and Profound Down-regulation of STAT3 and STAT3-Regulated Targets with Figure 4 **Tight Correlation between mRNA and Protein Changes**





- down-regulated
- down-regulated with STAT3 showing greatest decrease in abundance

CONCLUSIONS

- Kymera has discovered a series of potent and selective STAT3 degraders.
- STAT3 degraders show activity against both wild-type and clinically-relevant mutant forms of STAT3 resulting in growth arrest and increased cell death of ALK+ALCL as well as STAT3-mutant NK lymphoma and ALK-ALCL cell lines in vitro and in vivo.
- Transcriptomic and proteomic analyses of STAT3 degrader-treated cells reveal tightly correlated changes in proximal STAT3-dependent genes and proteins in ALK+ALCL cell lines.

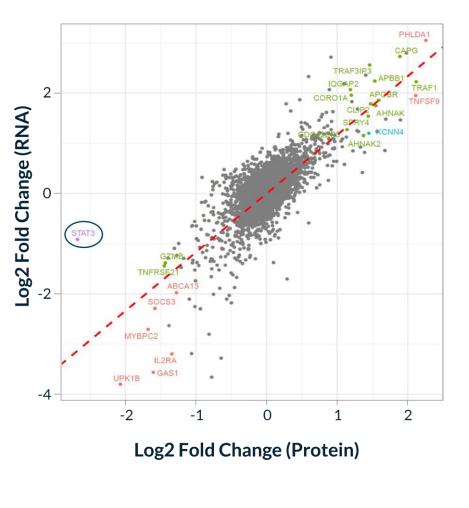
Content in SU-DHL-1 Model Can be Achieved with **Figure 5:** Time-Dependent Changes in Transcription in SU-DHL-1 Cells Upon STAT3 Degradation STAT3-dependent Transcriptional Changes Increase Over Time Dose-dependent TGI in SU-DHL-2 Wash-out of KTX-201 **PK-PD Relationship** Xenograft Model with KTX-201 from SU-DHL-1 cells 8 h KTX-201 Conc. vs. Time STAT3 Degrader 5 mg/kg, QW Plasma PK Tumor PK • DMSO KTX-201, 1 μM STAT3 Degrader 10 mg/kg, QW ★ STAT3 Degrader 25 mg/kg, QW — Cell Growth STAT3 vs. Control vs. Time --- STAT3 levels STAT3 Degrader 50 mg/kg, QW - Tumor PD 3000 🗸 🗸 🗸 Representative Genes Associated with Cell Cycle Regulation are Decreased at 48 h 2000 CCNB1 (Cyclin B1) 0.1 0.01 0.001 **GSEA** Reveals Decrease in Cell Cycle Signatures and Increase in Interferon Signaling Time (davs) E2F Targets nterferon- α Response

Deep Proteomics Shows Strong Reduction of STAT3 at 16 h in SU-DHL-1 Cells

• At 8 h, only STAT3 is significantly

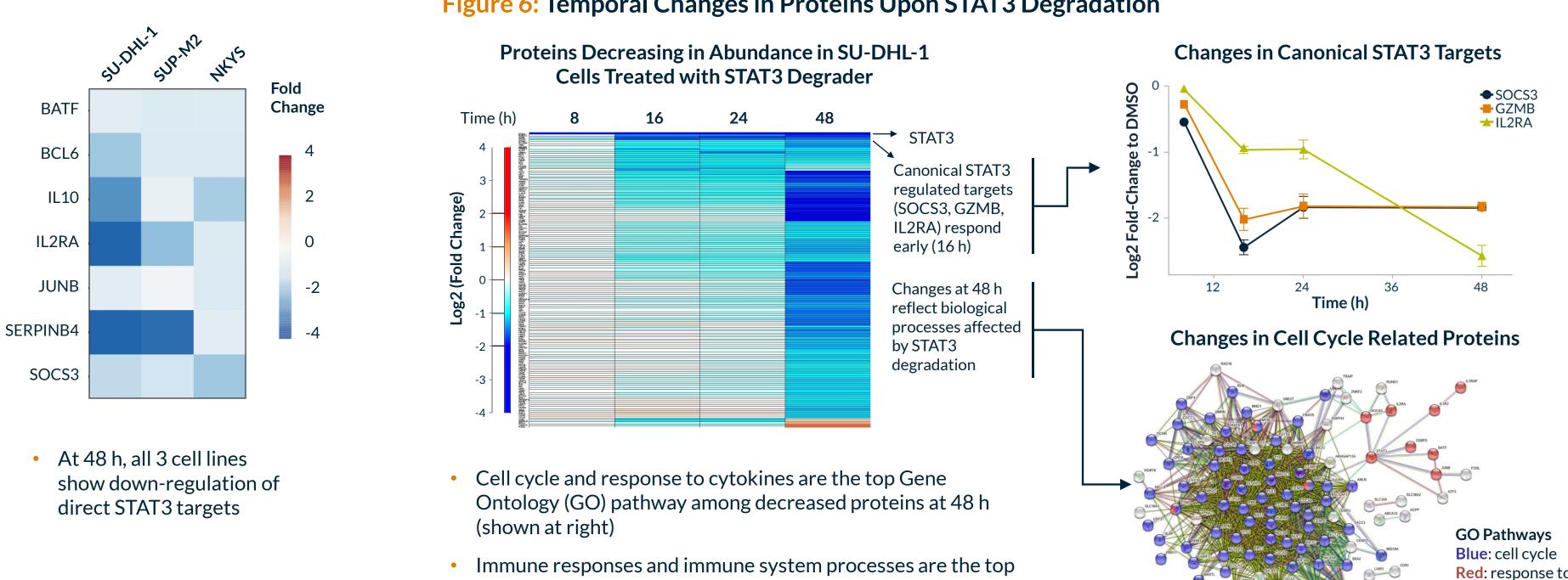
• At 16 h, 16 proteins are significantly

Changes in Protein Levels are **Tightly Correlated with Changes in** Transcript Levels at 24 h



STAT3 (circled) shows significant decrease in protein levels in SU-DHL 1 cells but minimal change in RNA consistent with degrader mechanism while most other genes/proteins have highly correlated changes

A Subset of STAT3 Target Genes is Down-regulated Across Three **PTCL Cell Lines**

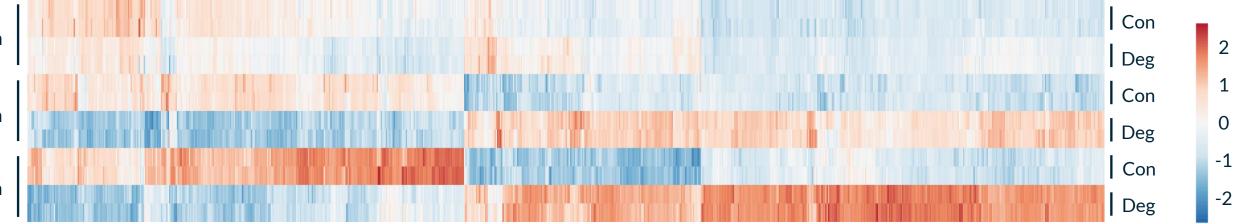


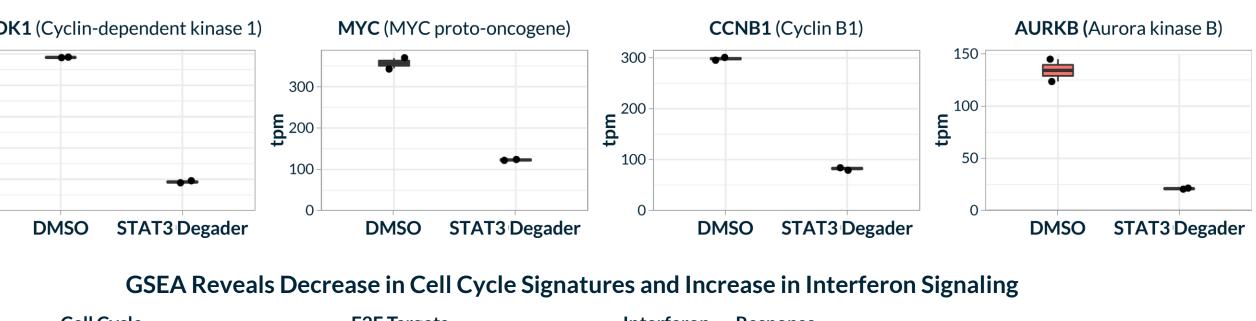
GO pathways among upregulated proteins at 48 h (not shown)

 Subset of transcriptional responses to STAT3 degrader is conserved between ALK+ ALCL and STAT3-mutant NK and T cell lymphoma cell lines.

- Pathway analyses confirm down-regulation of STAT3-regulated processes including cytokine responses at 16-24 h and consistent down-regulation of cell cycle signatures and up-regulation of immune pathways at 48 h suggesting modulation of tumor cell-intrinsic processes and the potential to regulate cell-cell interactions in the tumor microenvironment.
- Additional in vivo models of PTCL and related heme malignancies are being investigated, but these data illustrate the therapeutic potential of STAT3 degraders for the treatment of cancer with aberrant STAT3 activation.

KYMERA





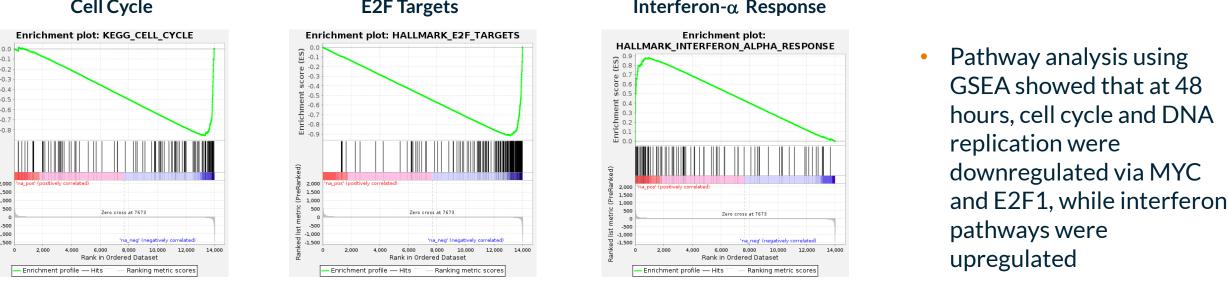


Figure 6: Temporal Changes in Proteins Upon STAT3 Degradation

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