

**UNITED STATES  
SECURITIES AND EXCHANGE COMMISSION  
Washington, D.C. 20549**

**FORM 8-K**

**CURRENT REPORT  
Pursuant to Section 13 or 15(d)  
of the Securities Exchange Act of 1934**

**Date of Report (Date of earliest event reported): January 10, 2022**

**MONTE ROSA THERAPEUTICS, INC.**

(Exact name of registrant as specified in its charter)

**Delaware**  
(State or other jurisdiction  
of incorporation)

**001-40522**  
(Commission  
File Number)

**84-3766197**  
(I.R.S. Employer  
Identification No.)

**645 Summer Street, Suite 102  
Boston, MA 02210**  
(Address of principal executive offices, including zip code)

**(617) 949-2643**  
(Registrant's telephone number, including area code)

**Not Applicable**  
(Former Name or Former Address, if Changed Since Last Report)

Check the appropriate box below if the Form 8-K filing is intended to simultaneously satisfy the filing obligation of the registrant under any of the following provisions:

- Written communications pursuant to Rule 425 under the Securities Act (17 CFR 230.425)
- Soliciting material pursuant to Rule 14a-12 under the Exchange Act (17 CFR 240.14a-12)
- Pre-commencement communications pursuant to Rule 14d-2(b) under the Exchange Act (17 CFR 240.14d-2(b))
- Pre-commencement communications pursuant to Rule 13e-4(c) under the Exchange Act (17 CFR 240.13e-4(c))

Securities registered pursuant to Section 12(b) of the Act:

Title of each class	Trading Symbol(s)	Name of each exchange on which registered
<b>Common Stock, \$0.0001 par value per share</b>	<b>GLUE</b>	<b>The Nasdaq Global Select Market</b>

Indicate by check mark whether the registrant is an emerging growth company as defined in Rule 405 of the Securities Act of 1933 (§ 230.405 of this chapter) or Rule 12b-2 of the Securities Exchange Act of 1934 (§ 240.12b-2 of this chapter).

Emerging growth company

If an emerging growth company, indicate by check mark if the registrant has elected not to use the extended transition period for complying with any new or revised financial accounting standards provided pursuant to Section 13(a) of the Exchange Act.

**Item 7.01. Regulation FD Disclosure.**

Monte Rosa Therapeutics, Inc. (the "Company") will be conducting meetings with investors attending the 40<sup>th</sup> Annual J.P. Morgan Healthcare Conference (the "Conference"), which is taking place virtually beginning on January 10, 2022. As part of these meetings, the Company will deliver its revised corporate presentation, furnished to this report as Exhibit 99.1 and which is incorporated herein by reference (the "Materials"). The Company will also present a portion of the Materials at the Conference.

The information in this report furnished pursuant to Item 7.01, including Exhibit 99.1, shall not be deemed "filed" for the purposes of Section 18 of the Securities Exchange Act of 1934, as amended (the "Exchange Act"), or otherwise subject to the liabilities of that section. It may only be incorporated by reference in another filing under the Exchange Act or the Securities Act of 1933, as amended, if such subsequent filing specifically references the information furnished pursuant to Item 7.01 of this report.

**Item 9.01. Financial Statements and Exhibits.**

(d) Exhibits

<u>Exhibit No.</u>	<u>Description</u>
99.1	<a href="#">Corporate presentation furnished by Monte Rosa Therapeutics, Inc. on January 10, 2022</a>
104	Cover Page Interactive Data File (embedded within the Inline XBRL document)

**Signatures**

Pursuant to the requirements of the Securities Exchange Act of 1934, the registrant has duly caused this report to be signed on its behalf by the undersigned hereunto duly authorized.

Monte Rosa Therapeutics, Inc.

Date: January 10, 2022

By: /s/ Markus Warmuth  
Markus Warmuth  
President and Chief Executive Officer

# From Serendipity to Rational Design

Taking Molecular Glue Degradors to New Heights | January 2022





## Forward-Looking Statements

These materials include express and implied “forward-looking statements,” including forward-looking statements within the meaning of the Private Securities Litigation Reform Act of 1995. Forward looking statements include all statements that are not historical facts, and in some cases, can be identified by terms such as “may,” “might,” “will,” “could,” “would,” “should,” “expect,” “intend,” “plan,” “objective,” “anticipate,” “believe,” “estimate,” “predict,” “potential,” “continue,” “ongoing,” or the negative of these terms, or other comparable terminology intended to identify statements about the future. Forward-looking statements contained in these materials include, but are not limited to, statements about our product development activities, including our expectations around the ongoing development of our QuEENTM platform and in silico tools, the advancement of our pipeline and the various products therein, including -the timing for filing our IND for our GSPT1 program and the advancement of additional programs, the expansion of our compound and debron libraries, our ability to identify additional molecular glue degraders, and our scientific predictions around clinical opportunities for our programs, including for GSPT1 program. By their nature, these statements are subject to numerous risks and uncertainties, including the impact that the current COVID-19 pandemic will have on our development activities and operations, as well as those risks and uncertainties set forth in our Quarterly Report on Form 10-Q for the third quarter ended September 30, 2021 filed with the US Securities and Exchange Commission, and any subsequent filings, that could cause actual results, performance or achievement to differ materially and adversely from those anticipated or implied in the statements. You should not rely upon forward looking statements as predictions of future events. Although our management believes that the expectations reflected in our statements are reasonable, we cannot guarantee that the future results, performance or events and circumstances described in the forward-looking statements will be achieved or occur. Recipients are cautioned not to place undue reliance on these forward-looking statements, which speak only as of the date such statements are made and should not be construed as statements of fact. We undertake no obligation to publicly update any forward-looking statements, whether as a result of new information, any future presentations or otherwise, except as required by applicable law. Certain information contained in these materials and any statements made orally during any presentation of these materials that relate to the materials or are based on studies, publications, surveys and other data obtained from third-party sources and our own internal estimates and research. While we believe these third-party studies, publications, surveys and other data to be reliable as of the date of these materials, it has not independently verified, and makes no representations as to the adequacy, fairness, accuracy or completeness of, any information obtained from third-party sources. In addition, no independent source has evaluated the reasonableness or accuracy of our internal estimates or research and no reliance should be made on any information or statements made in these materials relating to or based on such internal estimates and research.



# Monte Rosa Therapeutics Highlights

Taking molecular glue degraders (MGDs) to new heights



## **Next-generation molecular glue-based targeted protein degradation platform**

developing breakthrough small molecule drugs that selectively degrade therapeutically-relevant proteins

**Targeting the undruggable proteome** via AI-based deconvolution prediction & rational design of highly selective MGDs

**IND for GSPT1 program expected in 2022** with clinical development planned in Myc-driven tumors

**Five disclosed programs** targeting high unmet medical needs in oncology and non-oncology indications

**World-class leadership & SAB** with deep drug discovery know-how and development expertise in precision medicine

# World-Class Leadership

Deep expertise in molecular glue discovery, drug development and precision medicine



**Markus Warmuth, M.D.**  
Chief Executive Officer



**Ajim Tamboli, CFA**  
Chief Financial Officer



**Owen Wallace, Ph.D.**  
Chief Scientific Officer



**Sharon Townson, Ph.D.**  
Chief Technology Officer



**John Castle, Ph.D.**  
Chief Data Scientist



**Filip Janku, M.D., Ph.D.**  
Chief Medical Officer



**Jullian Jones, Ph.D., J.D., MBA**  
Chief Business Officer



**Silvia Buonamici, Ph.D.**  
SVP, Drug Discovery Biology



**Phil Nickson, Ph.D., J.D.**  
SVP, Head, Legal Operations



**Jennifer Champoux,**  
Vice President, Operations



# Strong Cash Position and Investor Support

Over \$455M raised since 2020 with top tier investors provides runway into late 2024



Aggregate IPO gross proceeds were approximately \$255.6 million before deducting underwriting discounts and commissions and other offering expenses and include an additional \$33.3 million in gross proceeds the company received as part of its IPO from the full exercise of the underwriters' option to purchase up to an additional 1,755,000 shares of common stock at the public offering price of \$19.00 per share.

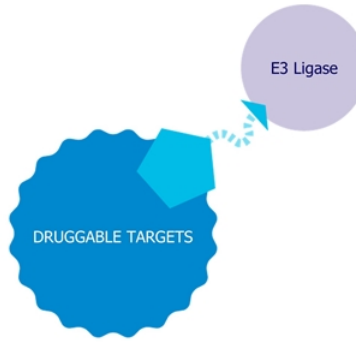
# Expanding Target Space through Molecular Glue Degraders (MGDs)

Inhibitors



Drugging  
the Druggable

PROTACs



Redrugging  
the Druggable

MGDs



Drugging  
the Undruggable



3-10% of Proteome

Uncharted Chemical and Target Space

# The Next Generation of Precision Medicine-based Small Molecule Drugs

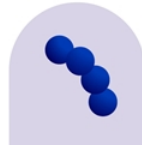
## Selectively editing the human proteome with rationally designed MGDs



Traditional small molecule inhibitors



Therapeutic Antibodies



MGDs



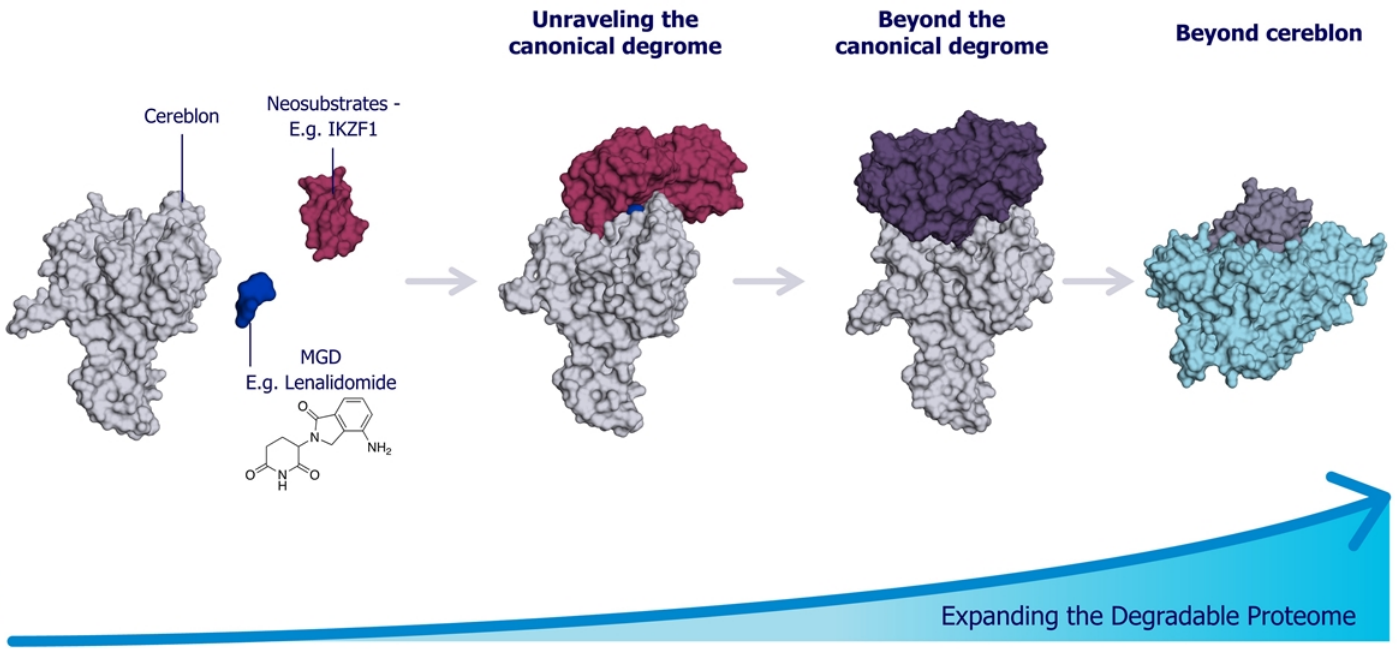
RNAi, RNA Editing



CRISPR/Gene Therapy

	Traditional small molecule inhibitors	Therapeutic Antibodies	MGDs	RNAi, RNA Editing	CRISPR/Gene Therapy
Ability to access undruggable space	X	✓	✓	✓	✓
Cellular permeability	✓	X	✓	✓	✓
Oral bioavailability	✓	X	✓	X	X
Systemic distribution	✓	✓	✓	X	X
CNS Penetration	✓	X	✓	X	X
Manufacturing scalability	✓	✓	✓	X	X

# Our Rational Approach to Unleash the Full Potential of MGDs

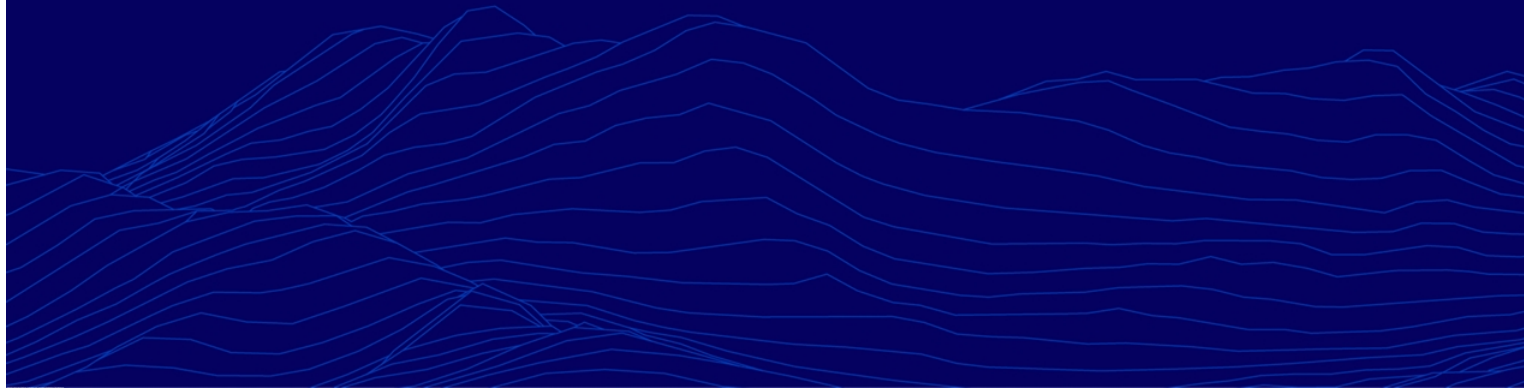






# QuEEN™ Discovery Platform

Quantitative and Engineered Elimination of Neosubstrates

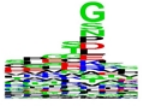




# QuEEN™ Discovery Platform: A Target-Centric Approach to MGDs

## Degron encyclopedia

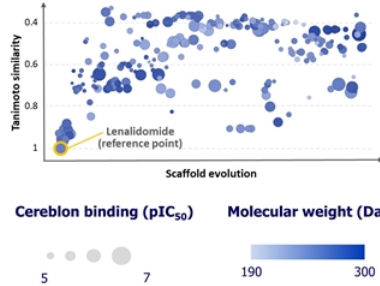
Degron discovery using proprietary AI-powered algorithm



1000s of proteins across multiple degrons

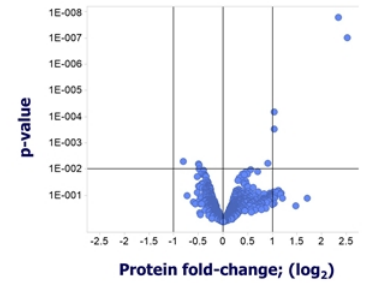
## Proprietary MGD library

Rationally designed, diverse and growing library engaging a variety of degrons



## Glueomics™ toolbox

Specialized suite of *in vitro* and *in silico* assays to discover, optimize and advance MGDs as clinical candidates



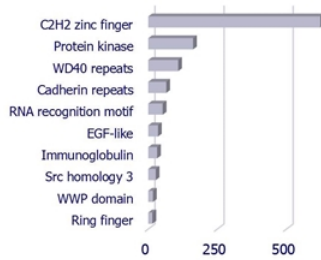
Accessing a large pool of undruggable targets with a diverse MGD library

# A Rich, Differentiated Target Space Across Protein Domains and Diseases

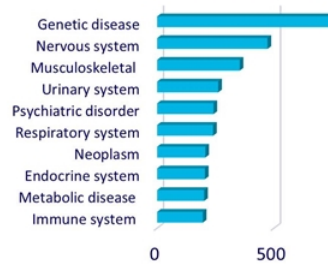
## Degron Encyclopedia

>3000 proteins containing a predicted loop degnon

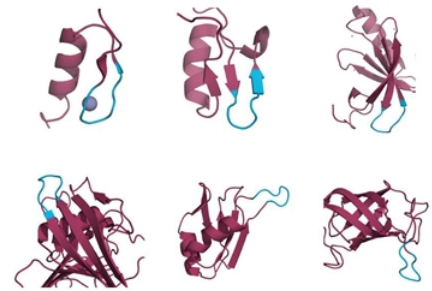
### Degron-containing domains



### Broad disease landscape



### Predicted degrons



Diverse protein domains and classes

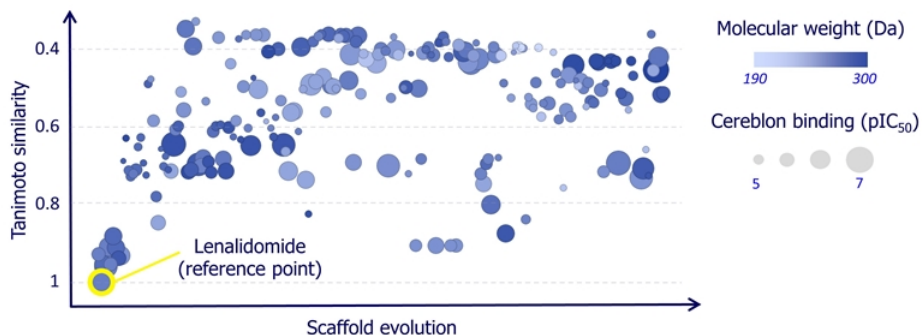
>85% degrons have unique sequence

>75% undruggable

Many highly credentialed targets

# Increasing Novelty and Structural Diversity to Match the Target Space

## Increasing MGD scaffold diversity



## Library design and expansion

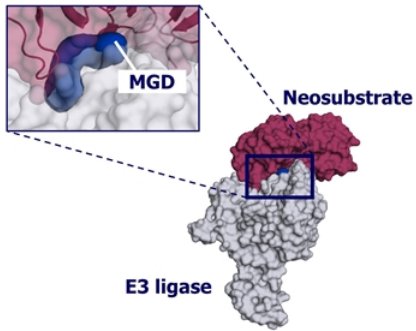
- Design focused on optimal drug-like properties
- High structural diversity and novelty
- Current library size 20K MGDs

MGD library is derived from **> 400 unique low molecular weight scaffolds** with **favorable CRBN binding** affinities

# MGDs Reprogram the Cereblon Surface

Remodeled MGD-CRBN surface enables selective engagement of neosubstrates

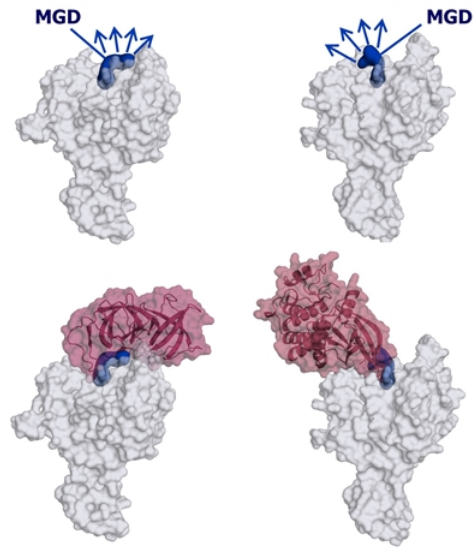
**Multiple points of contact for dialing in selectivity and potency**



Effective ternary complex generation involves

- MGD-cereblon interactions
- MGD-neosubstrate interactions
- CRBN-neosubstrate interactions

**MGDs are rationally designed to exploit key contacts to selectively engage different neosubstrates**



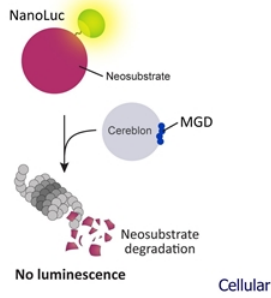
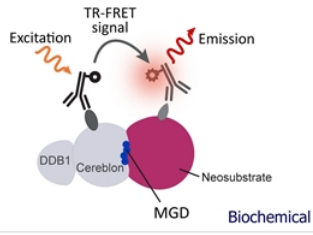
MGDs reshape the cereblon surface through different exit vector geometries

Neosubstrates are engaged selectively through unique interactions with both MGD and cereblon

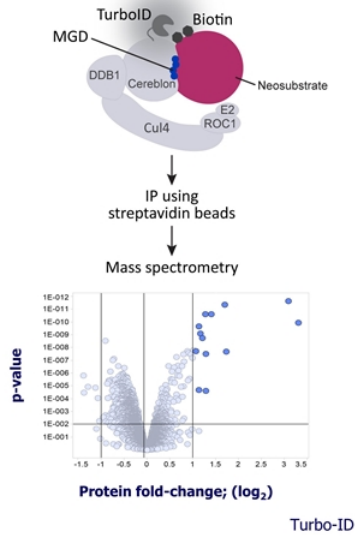
# Glueomics™ Toolbox Accelerates Identification of MGDs

Multiple assays enable rapid identification and validation of MGDs for novel targets

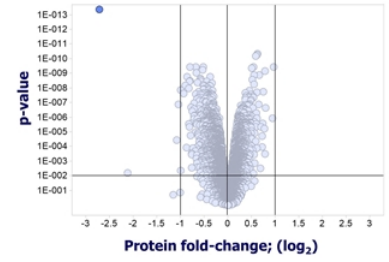
## *in vitro* screens



## Chemoproteomics - proximity



## Proteome-wide expression



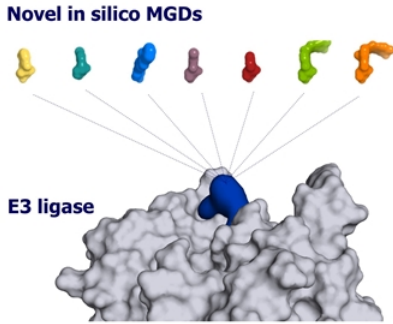
- TMT Proteomics to evaluate:
- Proteome-wide changes in protein levels
  - MGD selectivity

# Rhapsody, QuEEN's *in silico* Engine

A suite of proprietary AI-powered algorithms to design, discover and develop MGDs

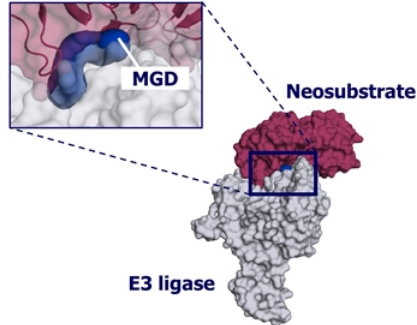
## *in silico* library generation

Creation and E3 ligase docking of novel MGDs, expanding our library to engage more targets



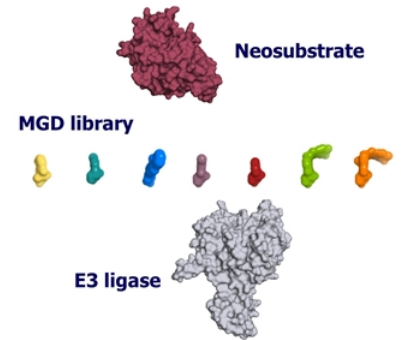
## *in silico* ternary complex models

Ternary complex models enabling MRT scientists to engineer and optimize selective MGDs



## *in silico* MGD screening

Computational screening identifying and prioritizing hits inducing binding and selective degradation



Taking MGD discovery in silico to accelerate discovery

# Leveraging a Leading Drug Discovery Platform

Purpose-built to discover and develop a wide landscape of therapeutically-relevant MGDs

## Monte Rosa's High-Value Proprietary Pipeline



### Targets

Undruggable and inadequately drugged degran-containing proteins

High level of target validation, preclinically and clinically



### Clinical Path

Programs with biomarker-based patient selection strategy and clear path to the clinic

Opportunity for rapid clinical PoC for MOA and efficacy



### Patient Benefit

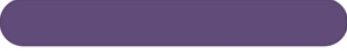


Address high unmet needs

Potential to address a wide range of therapeutically-relevant proteins in oncology and beyond

Create synergies within therapeutic areas

# Monte Rosa Pipeline

Rapidly advancing wholly owned MGD programs targeting undruggable proteins

Target / Program	Indication(s)	Discovery	IND-Enabling	Clinical	Next Anticipated Milestones	Ownership
GSPT1	NSCLC, SCLC and other MYC-driven Malignancies				IND filing mid-2022	
NEK7	Inflammatory Diseases				IND-Enabling Studies	
CDK2	Ovarian Cancer, Breast Cancer				IND-Enabling Studies	
VAV1	T and B Cell Malignancies, Autoimmune Disease				IND-Enabling Studies	
BCL11A	SCD, $\beta$ -Thalassemia				Lead Optimization	
Undisclosed	Multiple				Lead Optimization	

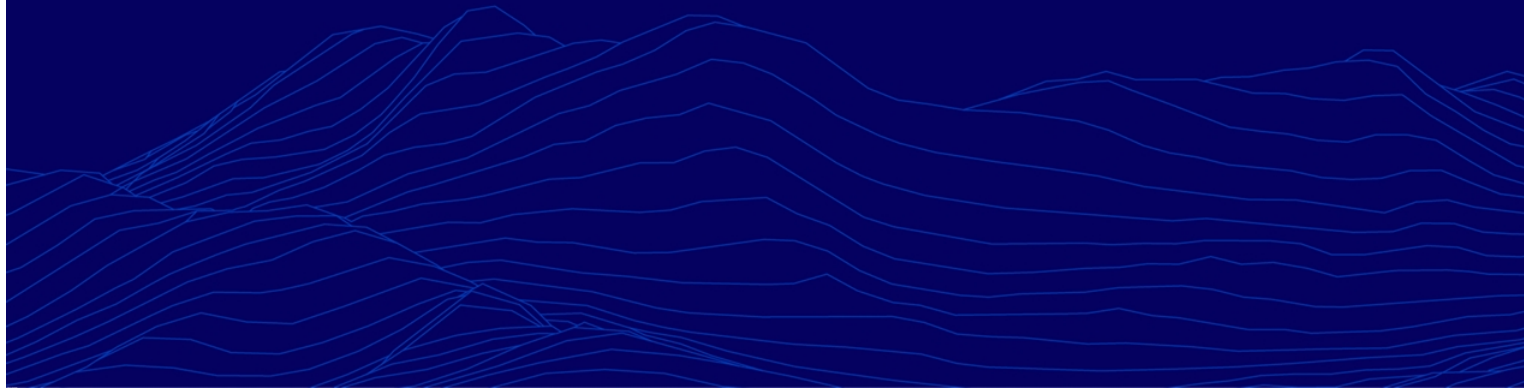
 Oncology
  Autoinflammation
  Oncology / immunology
  Genetic diseases







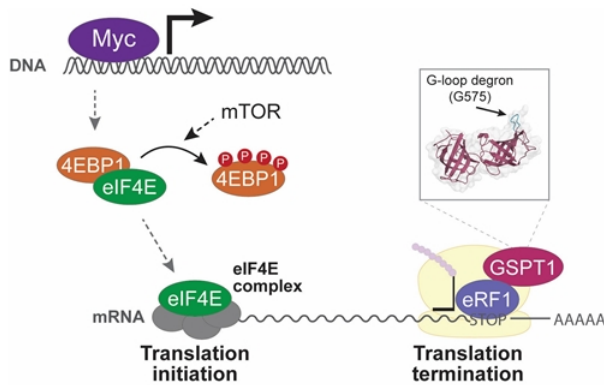
# GSPT1 Program



# Targeting Myc-driven Tumors and Their Addiction to Protein Translation

## GSPT1 is a key regulator and vulnerability of Myc-induced translational addiction

### Myc hijacks the cellular protein translation machinery creating a vulnerability to GSPT1



To sustain growth, Myc-driven tumors are **addicted to protein translation**

- Myc regulates the expression of key genes related to protein translation, including the master regulator 4EBP1 and eIF4E

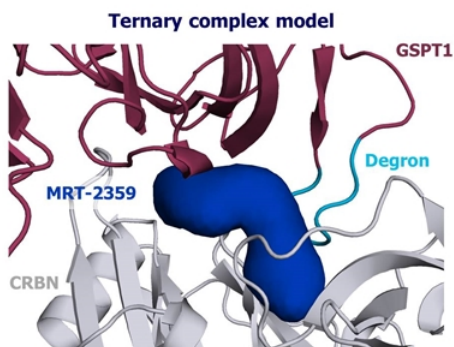
This addiction to protein translation creates a **dependency** to the translation termination factor GSPT1 a degron-containing protein

GSPT1 MGDs exploit this **vulnerability** by:

- Disrupting protein translation output
- Reducing Myc-oncogenic signaling

# MRT-2359 is a Potent and Selective GSPT1-directed MGD

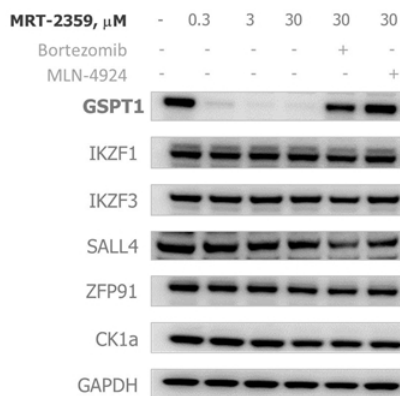
**MRT-2359 is a potent inducer of GSPT1-cereblon proximity**



***in vitro* data**

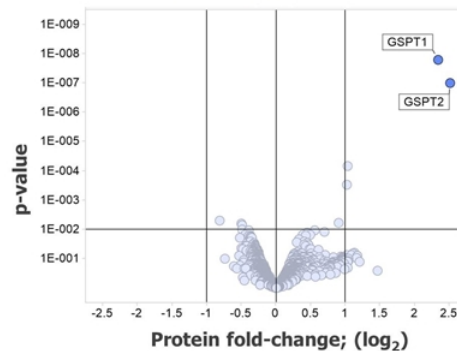
CRBN binding, $K_i$	113 nM
Ternary complex, $EC_{50}$	< 7 nM
Degradation, $DC_{50}$	80 nM

**MRT-2359 is a selective GSPT1-directed MGD**



6hr post treatment in MM1S and Kelly (SALL4)

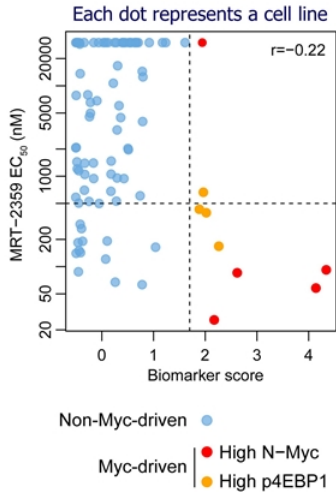
**Proximity – Turbo ID**



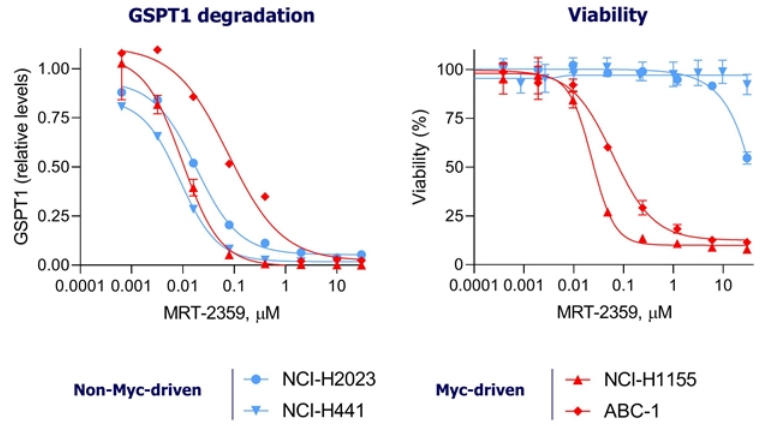
1hr post treatment

# Myc-Driven NSCLC lines are Highly Sensitive to MRT-2359

**Myc-driven NSCLC cell lines are sensitive to MRT-2359**



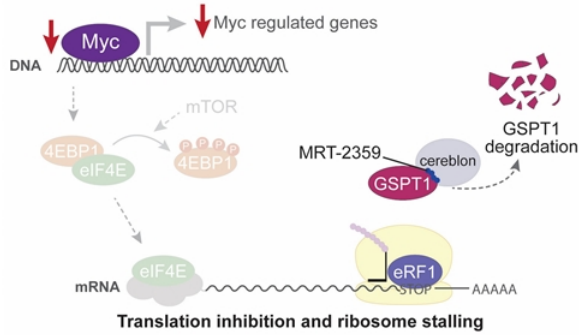
**MRT-2359 induces GSPT1 degradation in all cell models, but selective killing in high N-Myc lines only**



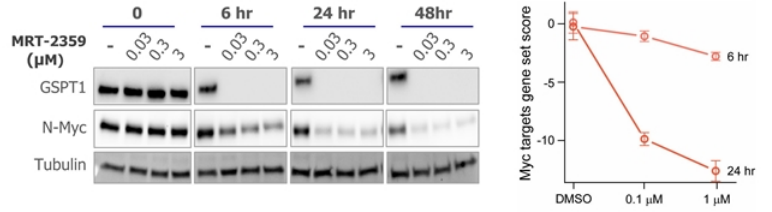
GSPT1 western blot at 6 hr (N-Myc high) and 24 hr (low). 72 hr viability assay (CTG)

# MRT-2359 Affects N-Myc Pathway only in Myc-driven Cells

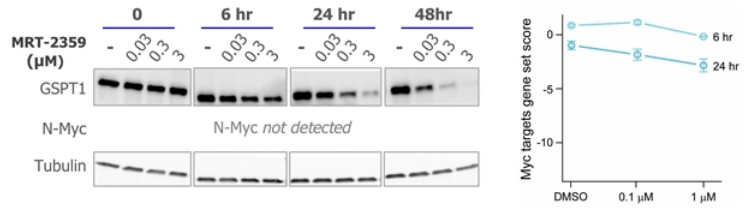
## GSPT1-directed MGD degradation affects translation, a critical vulnerability of Myc-driven cells



## Myc-driven (NCI-H1155)



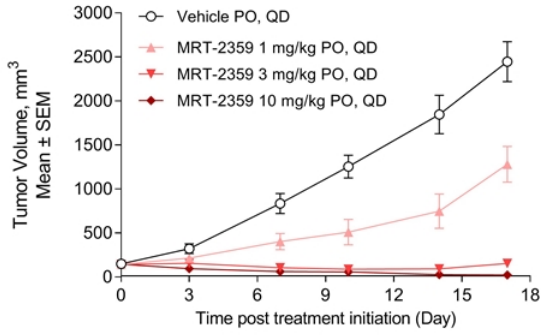
## Non-Myc-driven (NCI-H2023)



# MRT-2359 Induces Tumor Regressions in N-Myc-driven Xenograft Models

## Oral dosing of MRT-2359 shows anti-tumor activity and regressions in NCI-1155

Similar observations in other high N-Myc expression models (ABC-1, NCI-H1770)

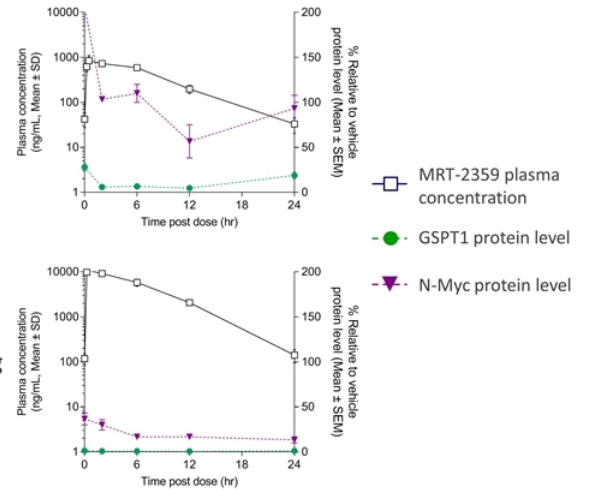


## Dose- and time-dependent degradation of GSPT1 is associated with N-Myc downregulation

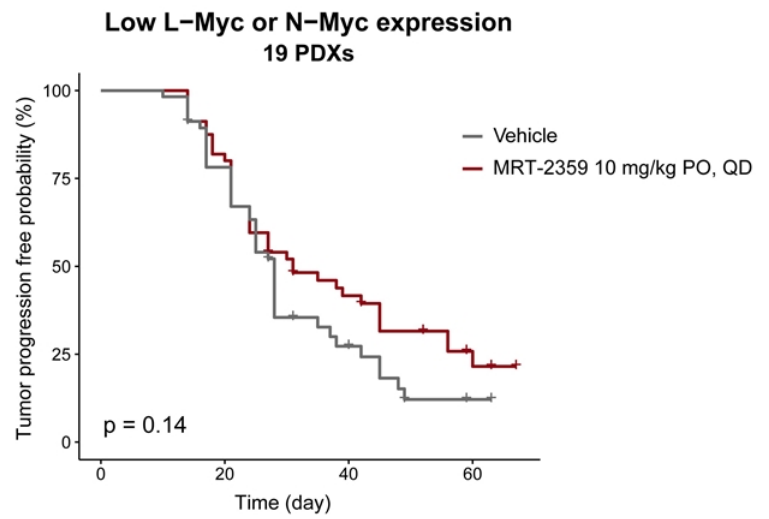
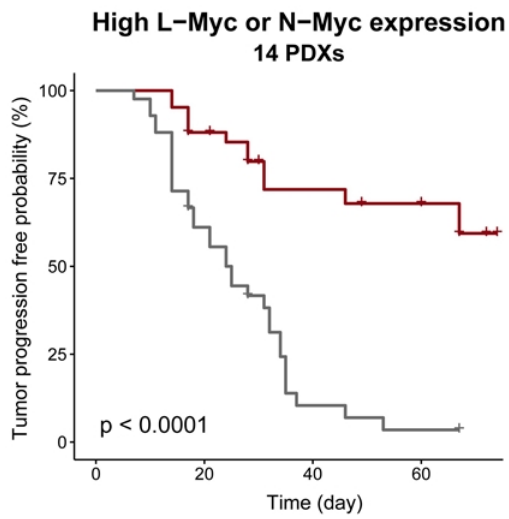
Day 5

1 mg/kg

10 mg/kg



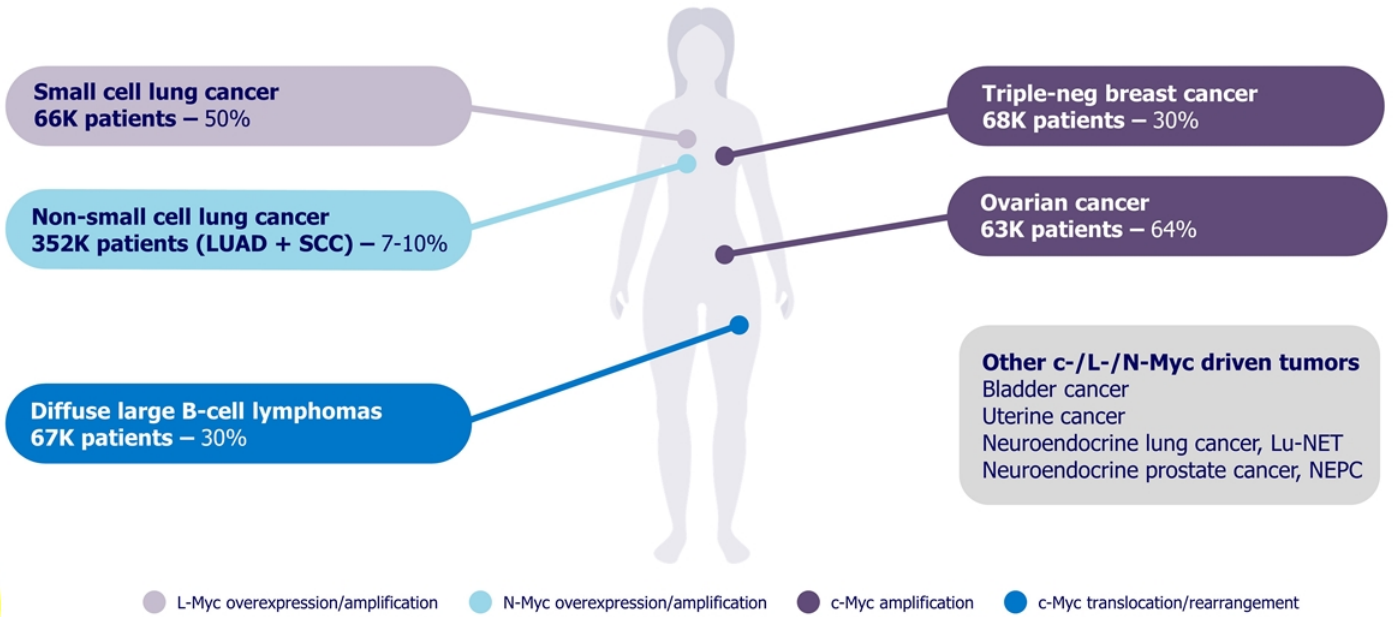
# MRT-2359 Anti-tumor Activity in L- and N-Myc-positive NSCLC PDXs



Tumor progression to  $\geq 800 \text{ mm}^3$

# Targeting Myc-positive Tumors with MRT-2359

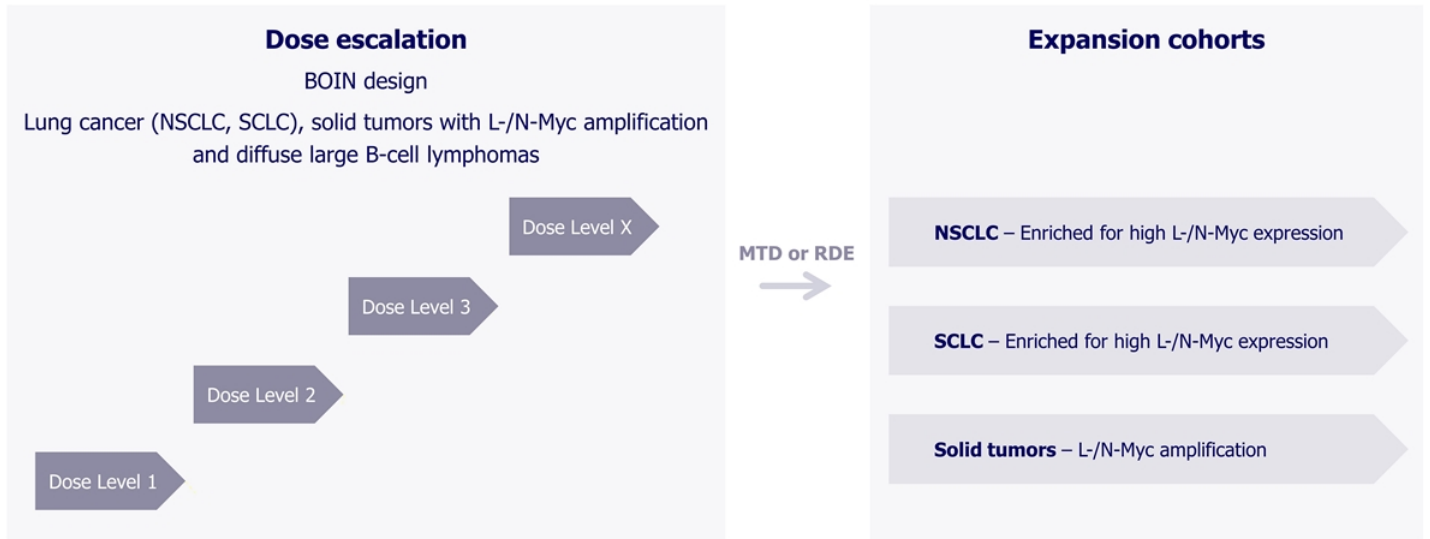
## Potential indications and patient stratification hypotheses



Patient diagnosed incidence #s, major markets (US, EU and JP): Decision Resources Group (DRG)  
Patient stratification %s: Schaub - Cell Systems 2018; Massó-Vallés – Exp. Op. therapeutic targets 2020; Sesques and Johnson - Blood 2016



# Early Phase Clinical Development



# Targeting Myc-addicted Tumors with MRT-2359

IND-enabling activities have been initiated

Rationally designed **potent and selective** GSPT1-directed MGD

Favorable **drug-like properties** and ADMET profile

**Orally bioavailable** development candidate

Robust **antitumor activity** *in vivo* in **multiple tumor models**

**IND-enabling** activities ongoing

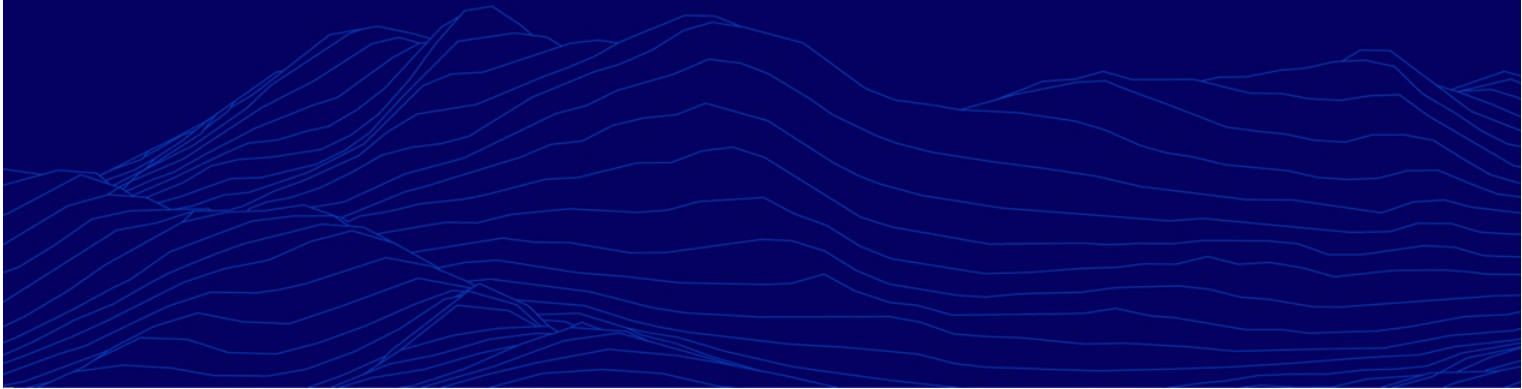
**Patient stratification hypothesis** developed and being validated

**IND filing  
expected in  
mid-2022**



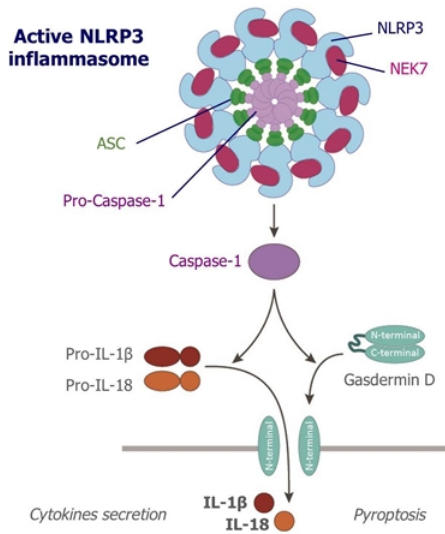


# NEK7 Program



# NEK7 (NIMA-Related Kinase 7) as a Target for Inflammatory Disease

## NEK7 is an essential regulator of the inflammasome



**Therapeutic hypothesis:** Diseases with over-activated or mutated NLRP3 inflammasome

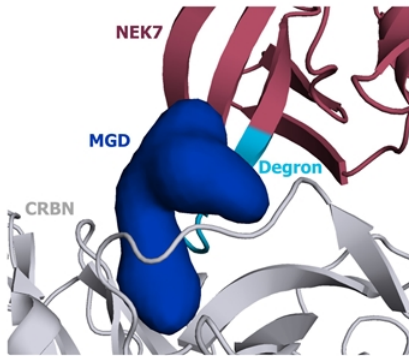
- NEK7 licenses NLRP3 assembly in a kinase independent manner
- NEK7-deficient macrophages are severely impaired in IL-1 $\beta$  and IL-18 secretion

**Clinical opportunity:** First-in-class NEK7 degraders for

- Over-activated NLRP3 inflammasome: metabolic pathologies, cardiovascular diseases, inflammatory issues and neurologic disorders
- NLRP3 activating mutations: Cryopyrin-associated periodic syndromes (CAPS)

# Rationally Designed NEK7-Directed MGDs are Selective

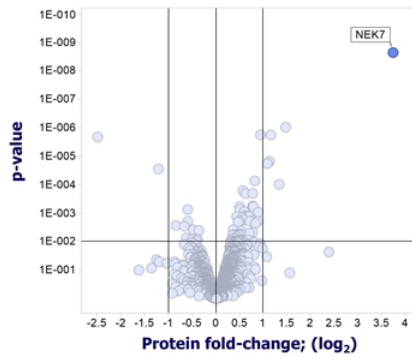
## Rhapsody model enables rapid chemistry optimization



### *in vitro* data

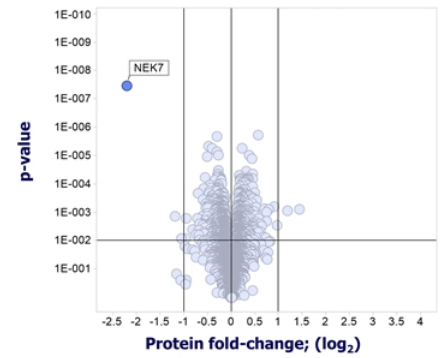
CRBN binding, $K_i$	48 nM
Ternary complex, $EC_{50}$	20 nM
Degradation, $DC_{50}$	10 nM

## Rationally designed MGDs promote selective CRBN proximity



Turbo-ID – 6hr post treatment

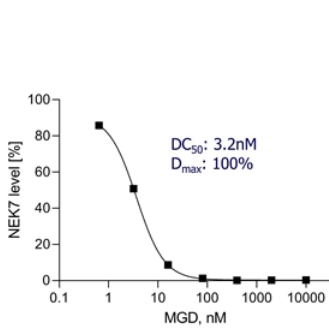
## NEK7-directed MGD promotes selective degradation of NEK7



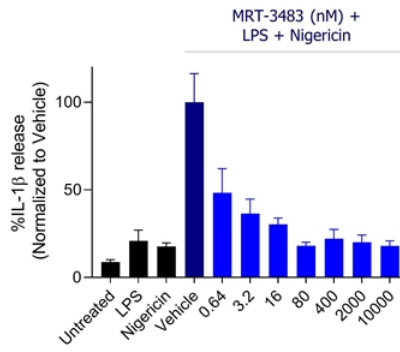
TMT-Proteomics – 24hr post treatment

# NEK7-directed MGDs Modulate NLRP3 Pathway in Human Macrophages

## MRT-3483 promotes NEK7 degradation and pathway engagement in hMDMs



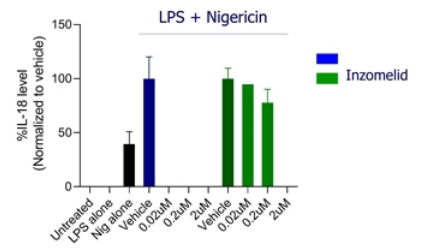
Western blot – 24 hr



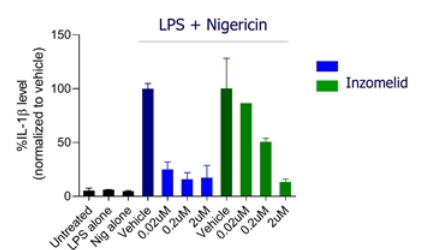
Treatment (6hr) of primed hMDMs

## NEK7-directed MGD compared to NLRP3 inhibitor

### IL-18



### IL-1β

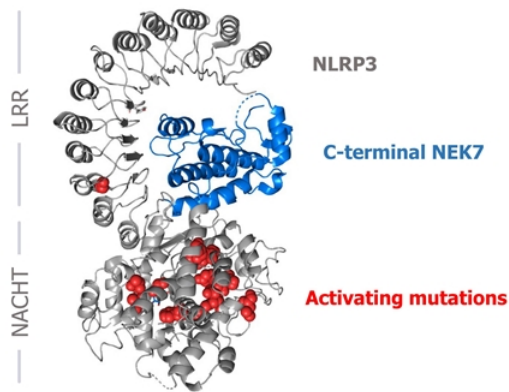


Treatment (20 hr) of primed hMDMs

CASP1 and LDH showed similar profile

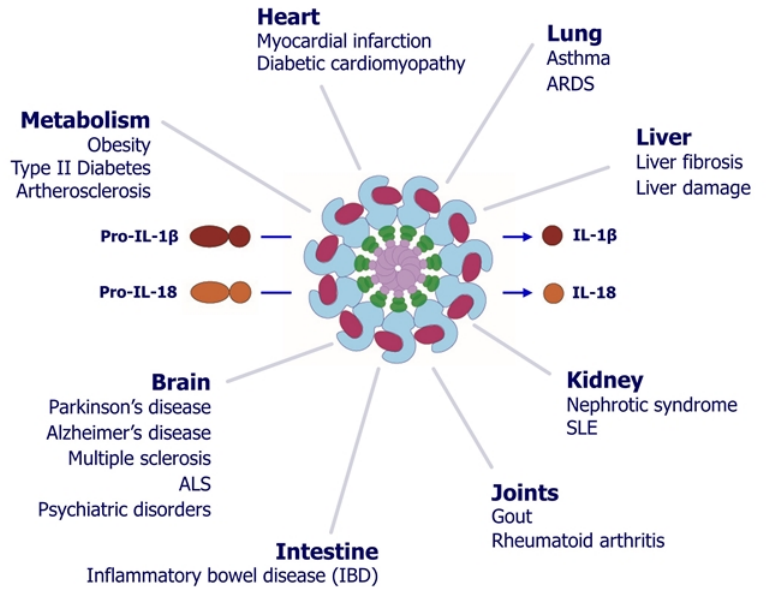
# Overactivation of the NLRP3 Inflammasome in Diseases

## NLRP3 activating mutations



NLRP3 mutations found in CAPS (Cryopyrin-associated periodic syndromes – MWS\*, FCAS\*\*, CINCA/NOMID<sup>†</sup> Syndrome) might stabilize the active form of NLRP3

## Over-activated NLRP3 inflammasome

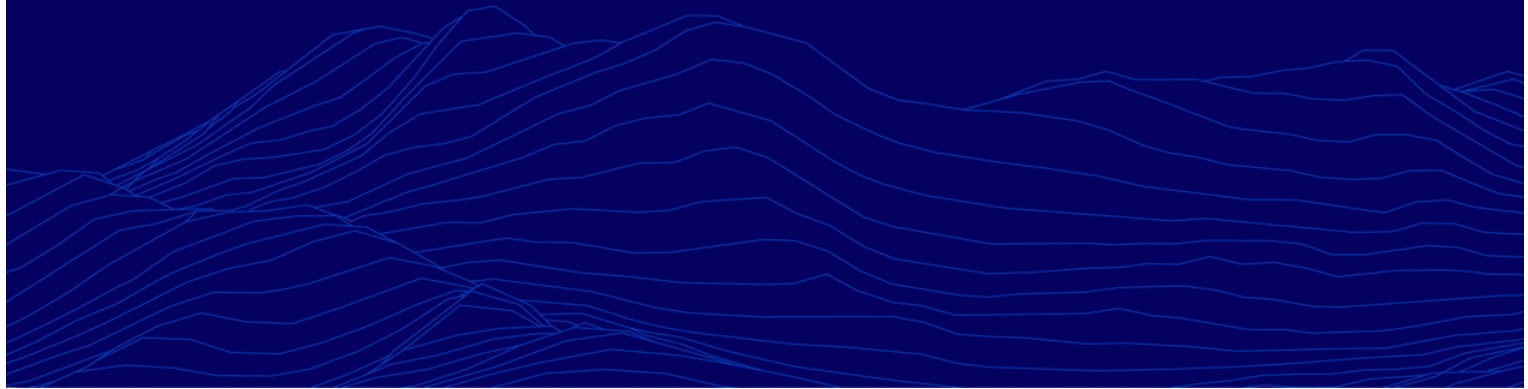


\*Muckle-Wells Syndrome

\*\*familial cold autoinflammatory syndrome, <sup>†</sup>Chronic infantile neurological cutaneous and articular/neonatal onset multisystem inflammatory disease



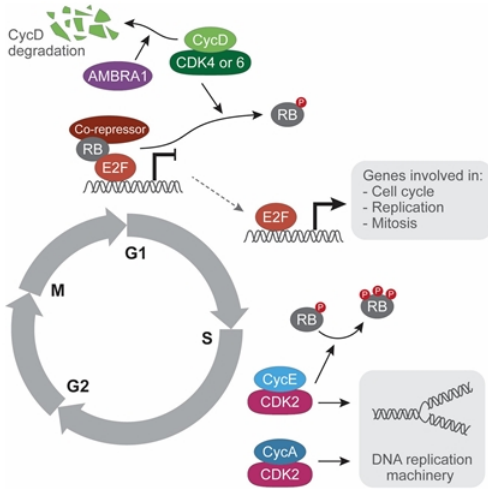
# CDK2 Program





# CDK2 as a Target for Selected Solid Tumors

CDK2 is one of the key regulators of the cell cycle



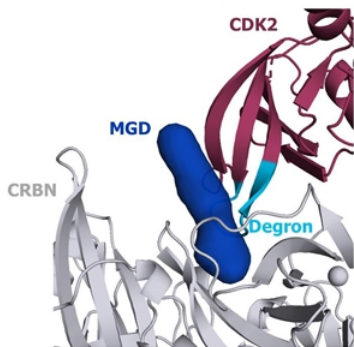
**Therapeutic hypothesis:** Tumors with CDK2 pathway activation by:

- CyclinE1/E2 amplification or loss of AMBRA1
- Loss of RB

**Clinical Opportunity:** CDK2 driven cancers: ER positive breast cancer (444K patients), ovarian cancer (63K patients), and endometrial cancer (118K patients), as well as breast cancer post treatment with CDK4/6 inhibitors

# Rationally Designed CDK2-Directed MGDs are Selective

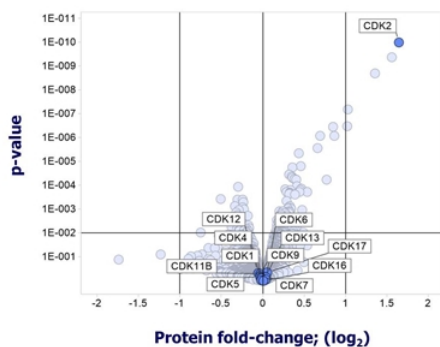
Rhapsody model enables rapid chemistry optimization



**in vitro data**

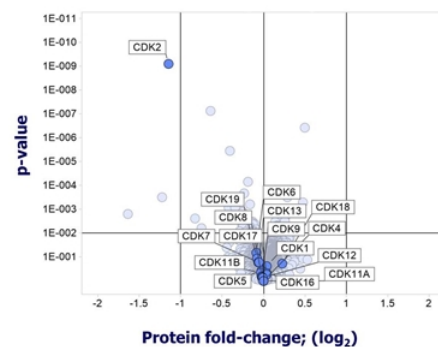
CRBN binding, $K_i$	163 nM
Ternary complex, $EC_{50}$	9 nM

Rationally designed MGD promotes selective CRBN proximity



Turbo-ID – 6hr post treatment

CDK2-directed MGD promotes selective degradation

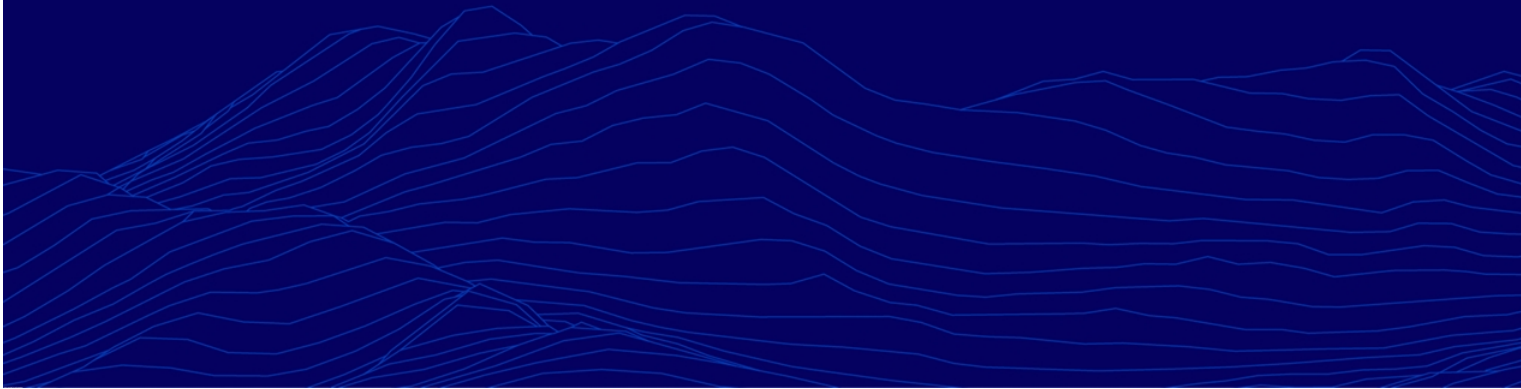


TMT-Proteomics – 24hr post treatment

Program advanced to lead optimization in Q4 2021

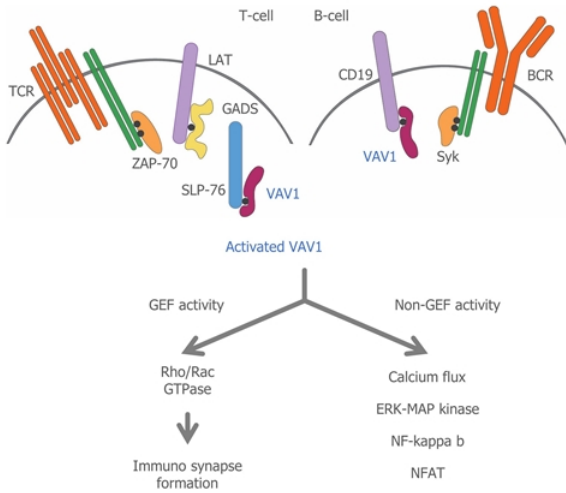


# VAV1 and BCL11a Programs



# VAV1 as a Target for Autoimmune Disease

## VAV1 plays a key role in T-cell and B-cell development and activation



**Therapeutic hypothesis:** Diseases with VAV1 activating mutations or autoimmune disorders

- VAV1 activation mutations identified in leukemia, lymphoma and lung cancer
- VAV1 KO mice improved multiple autoimmune models

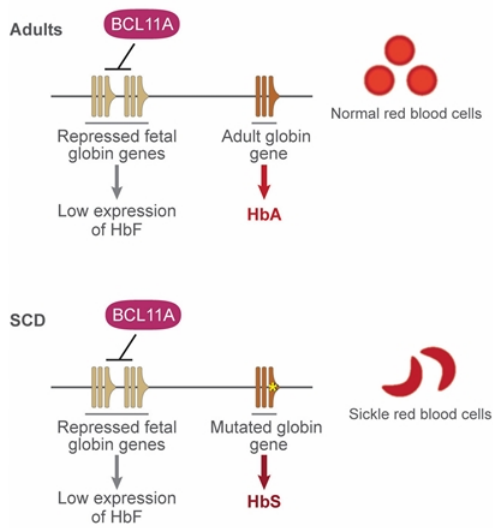
**Clinical Opportunity:** First-in-class VAV1 degraders for

- T-cell and B-cell lymphomas: DLBCL (66K patients) and Burkitt lymphoma
- Autoimmune disorders including MS (1.2M patients), myasthenia gravis (36K – 60K patients in US), and acute graft-versus-host disease (10K patients)

Patient diagnosed prevalence and incidence #s, major markets (US, EU and JP): DRG; myasthenia.org

# BCL11A as a Target for Hemoglobinopathies (SCD and $\beta$ -Thalassemia)

## BCL11A is the zinc finger transcription repressor of the fetal globin genes



**Therapeutic hypothesis:** Reactivate expression of fetal hemoglobin (HbF) to compensate for mutated adult globin

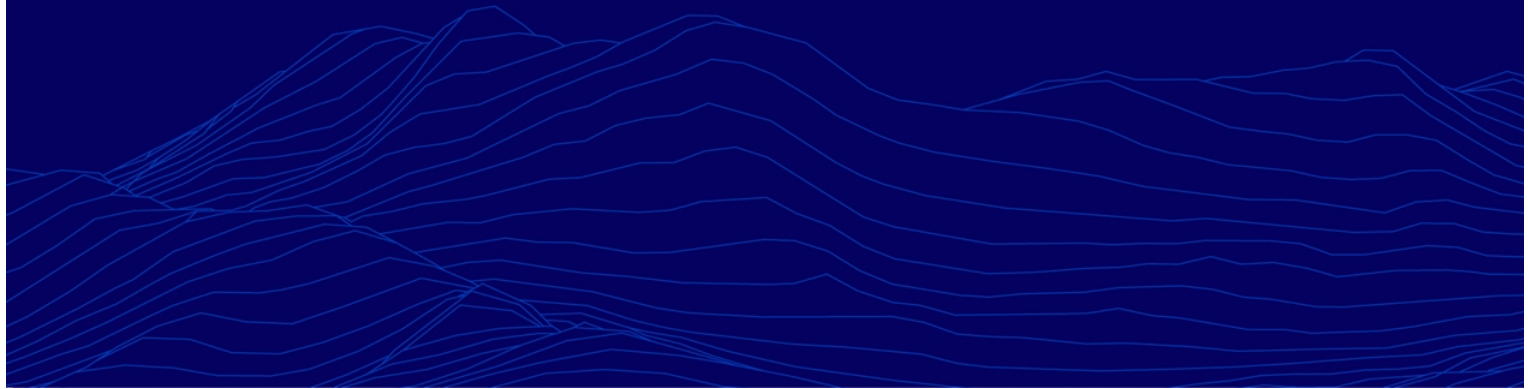
**Clinical Opportunity:** First-in-class BCL11A degraders for

- Sickle cell disease (SCD)
  - 155,000 patients (US and EU)
  - >6M patients (ROW)
- $\beta$ -thalassemia
  - 17,000 patients (US and EU)

Patient diagnosed prevalence #s: DRG; [www.notaloneinsicklecell.com](http://www.notaloneinsicklecell.com)



# Summary



# Monte Rosa Therapeutics

## From serendipity to rational design of MGDs



**Molecular glue-based targeted protein degradation platform** developing breakthrough small molecule therapeutics that selectively degrade disease-causing proteins

Proprietary, **target-centric** drug discovery platform enabling **rational design**, and anticipated rapid development, of molecular glue-based degraders targeting the **undruggable proteome in oncology** and **non-oncology disease**

Initial platform focus on **cereblon-mediated protein degradation** with **hundreds of potential targets** to address

Extensive and compelling pre-clinical *in vivo* data for GSPT1 program, demonstrating **potent anti-tumor activity** in Myc-driven tumor models

**IND filing for GSPT1 program expected in mid-2022**; additional programs at various stages of discovery

**Potential** to reprogram other E3 ligases to access more of the undruggable proteome through other degrons





Thank You

