Molecular Glue Degraders (MGDs) – From the Bench to the Clinic

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Forward-Looking Statements

This communication includes 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 herein include, but are not limited to, statements about our ability to grow our product pipeline, statements around the Company's OuEENTM discovery engine and its potential to identify novel degradable protein targets and rationally designed MGDs with unprecedented selectivity, statements around the potential of the Company's MGDs against a broad spectrum of targets, statements about the advancement and timeline of our preclinical and clinical programs and our pipeline and the various products therein, statements about the ongoing clinical development of our GSPT1 degrader referred to as MRT-2359, including our expectations for the nature, significance, and timing for our disclosure of any updated data from our Phase 1/2 clinical trial of MRT-2359 in MYC-driven solid tumors in the second half of 2024, statements about Phase 2 development of MRT-2359, including timing of initiation and enrollment of any Phase 2 expansion cohorts in the second half of 2024, statements about the ongoing Phase 1 SAD/MAD study for the VAV1-directed degrader, referred to as MRT-6160 and our expected timing of disclosure of initial clinical data therefrom, our expectations of indications for the potential future clinical development for a VAV1-directed degrader, including MRT-6160, including the relevance of preclinical data for such indications, and our expectations regarding the potential clinical benefit for a VAV1-directed degrader, including MRT-6160, statements concerning our expectations regarding our ability to nominate and the timing of our nominations of additional targets, product candidates, and development candidates, statements regarding regulatory filings for our development programs, including the planned timing of such regulatory filings, such as IND applications, and potential review by regulatory authorities, our use of capital, expenses and other financial results in the future, availability of funding for existing and future programs, statements related to our strategic agreements, goals of such agreements and any related milestone, royalty or other payments related thereto, statements about the transaction with Novartis, including the closing thereof, as well as our expectations of success for our programs, strength of collaboration relationships and the strength of our financial position, among others. By their nature, these statements are subject to numerous risks and uncertainties, including those risks and uncertainties set forth in our most recent Annual Report on Form 10-K for the year ended December 31, 2023, filed with the U.S. Securities and Exchange Commission on March 14, 2024, 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. 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Our Molecular Glue Degraders (MGDs) Edit the Proteome



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Monte Rosa's rationally designed MGDs have potential applications in Oncology, Immunology, Neuroscience and other therapeutic areas

Key Advantages of Our Rationally Designed MGDs

Unique Target Space

Unprecedented Selectivity

Catalytic Mechanism of Action







Unique insights into anatomy of protein-protein-MGD interaction allows unprecedented MGD selectivity



Long lasting, catalytic protein degradation effect creates differentiated target product profiles

Monte Rosa Pipeline and Upcoming Milestones

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Target	Compound	Indication(s)	Discovery	IND-Enabling	Clinical	Next Anticipated Milestone	Ownership
GSPT1	MRT-2359	NSCLC, SCLC and other MYC-driven Malignancies				RP2D and Phase 1 data in H2 2024	
VAV1	MRT-6160	Autoimmune Disease – Systemic and CNS				Phase 1 data in Q1 2025	<mark>₺</mark> NOVARTIS*
NEK7	MRT-8102	IL-1β/NLRP3 driven				IND submission in H1 2025	
	LO (2 nd generation)	Inflammatory Diseases				Development candidate	
CDK2	LO	Breast Cancer				Development candidate in 2024	
CCNE1 (Cyclin E	E1) LO	CCNE1 amplified tumors				Development candidate	
Discovery Targe	its -	Multiple				Lead optimization	
Discovery Targe	its -	Oncology and Neurological Diseases				Undisclosed	Roche
		Oncology	Immunology	Inflammation	Various		

* Monte Rosa has signed an exclusive global license agreement with Novartis for this asset. This transaction is subject to customary closing conditions, including regulatory clearance.



GSPT1/MRT-2359

Targeting MYC-driven Tumors and Their Addiction to Protein Translation Through GSPT1 Degradation





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



in vitro data

113 nM

< 7 nM

1 - 20 nM

CRBN binding, K_i

Ternary complex, EC₅₀

Degradation, DC₅₀

(in disease relevant cell lines)

MRT-2359 is a potent GSPT1-directed MGD

MRT-2359 induces selective GSPT1 degradation and shows favorable ADME/DMPK profile



Protein fold-change (log₂)

No degradation of	other known	cereblon	neosul	bstrat	es
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ADMET profil	e
CYP DDIs	> 30 µM
hERG inhibition patch clamp	EC ₅₀ > 30 µM
Oral bioavailability all species	~50%

MRT-2359 Has Optimized Depth of Degradation To Achieve Preferential Activity in MYC High Cancer Cells



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MRT-2359 displays preferential activity in MYC driven NSCLC cells



Non-optimal GSPT1 MGD (MRT-2136) shows limited preferential activity



Pharmacogenomic Profiling Identifies Multiple MYC-Driven Tumor Types With Sensitivity to MRT-2359 – Non-small Cell Lung Cancer

Pharmacogenomic profiling of cancer cell lines identifies N-Myc high / NDRG1 low NSCLC as potential indication





Preclinical Validation of Activity of MRT-2359 in Lung Cancer PDX Models



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PD modulation

Targeted mass spectrometry in 7 representative models



Pharmacogenomic Profiling Identifies Multiple MYC-Driven Tumor Types With Sensitivity to MRT-2359 – Prostate Cancer

Pharmacogenomic profiling of cancer cell lines identifies AR positive prostate cancer as potential indication – MRT-2359 reduces expression of GSPT1, MYC, CCND1 and AR Signature





MRT-2359 Leads to Tumor Regressions in Preclinical Models of Castration Resistant Prostate Cancer and ARV7-driven Prostate Cancer



MRT-2359-001 Phase 1/2 Clinical Study Design

Phase 1: Dose Escalation

Lung cancer, high-grade neuroendocrine tumors and solid tumors with N-/L-MYC amplification **Phase 2: Expansion Cohorts**

Retrospective stratification per N-/L-MYC expression



**

21/7 = 21 days on drug, 7 days off drug.

MRT-2359 Induces Optimal GSPT1 Degradation in Tissue Biopsies*



- GSPT1 degradation assessed from pretreatment screening biopsies and biopsies taken at day 19
- Matched biopsies obtained from 11 patients across the 3 cohorts analyzed
- GSPT1 expression assessed using targeted mass spectrometry
- PD modulation seen in tissue biopsies in line with PD modulation seen preclinically at efficacious dose levels using same assay (targeted mass spectrometry)



⁶ Based on optimal PD modulation in preclinical studies as presented on 10/17/23

MRT-2359-001 – Preliminary Efficacy Data*

- As of September 7th, 2023, of 15 evaluable patients treated across 3 cohorts, tumors from 6 patients were identified as biomarker positive
- Of these 6 biomarker positive patients, 2 have experienced a PR (1 confirmed, 1 unconfirmed) and 1 patient has SD
 - PR (-59%) HG NE bladder carcinoma
 - uPR (-41%) NSCLC with SCLC/NE transformation
 - SD (0%) SCLC (remains on therapy for > 4 months)
- In addition, one patient with NSCLC and unclear biomarker status remains on therapy for > 7 months with stable disease
- No clinical activity seen in biomarker negative patients





VAV1/MRT-6160

VAV1 is a Key Regulator of T- and B-cell Receptor Activity



VAV1-directed MGDs have the potential to modulate

VAV1 signaling increases cytokine production, proliferation, and differentiation

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TCR = T-cell receptor. BCR = B-cell receptor. IL-2, IL-17 and IL-6 are cell signaling molecules (cytokines) that promote immune response. sIgG is the most common circulating antibody.

Therapeutic hypothesis:

- VAV1 is a pivotal scaffolding protein and signaling molecule downstream of both the T-cell and B-cell receptors – confirmed by multiple CRISPR screens & VAV1 knockout (KO) mice
- VAV1 degradation is predicted to impact both T- & B-cell function and has the potential to treat a broad set of autoimmune diseases

Clinical Opportunity:

Autoimmune/inflammatory disorders including inflammatory bowel disease (4.1M patients), rheumatoid arthritis (6.2M patients), multiple sclerosis (1.3M patients), and myasthenia gravis (~300K patients)

Patient diagnosed prevalence #s, major markets (US, EU and JP): Decision Resources Group (DRG)

Unique Mechanism of VAV1 MGD May Achieve Broad Clinical Response





- Patients are currently treated by various drugs focused on single cytokine or cell-type specific therapies which best address specific endotypes
- Instead, deep modulation of multiple cytokine pathways with a first-in-class VAV1 oral MGD may benefit patients presenting a broad range of pathophysiology

MRT-6160 is a Potent and Highly Selective VAV1-directed MGD

MRT-6160 is a potent VAV1-directed MGD

MRT-6160 induces highly selective VAV1 degradation and has a favorable ADME/DMPK profile

No degradation of other known cereblon neosubstrates

ADMET profile				
CYP DDIs	IC ₅₀ > 30 μM			
hERG inhibition patch clamp	EC ₅₀ > 30 μM			
Oral bioavailability all species	> 50%			

<i>in vitro</i> data	<i>in vitro</i> data			
CRBN binding, IC ₅₀	670 nM			
Ternary complex, EC ₅₀	11 nM			
Degradation, DC ₅₀ /D _{max}	7 nM / 97 %			
(Jurkat)				

MRT-6160 Blocks T-cell-Mediated B-cell Activity in BioMAP® Profile

BioMAP[®] Diversity Plus Platform (Eurofins). Shark tooth plots show relative expression levels of indicated proteins in Drug treated vs. DMSO controls. 3C/4H, Venular endothelial cells; LPS/SAg, Venular endothelial cells + PBMC; BT, PBMC + B cells; BF4T, Bronchial epithelial cells + dermal fibroblasts; BE3C. Bronchial epithelial cells; CASM3C, Coronary artery smooth muscle cells; HDF5CGF, Dermal fibroblasts; KF3CT, keratinocytes + dermal fibroblasts; MyoF, lung fibroblasts; IMphg, macrophages + venular epithelial cells

MRT-6160 Ameliorates T Cell Transfer-Induced Colitis Better than Standard of Care

11.

Non-pathogenic CD45RB^{low} or pathogenic CD45RB^{high} cells were transferred into SCID mice to induce colitis. Mice were treated with vehicle, MRT-6160 (PO QD), or anti-TNF (IP Q3D) from Day 0 to Day 42 and assessed for disease every 3 days (*left*) or with vehicle, MRT-6160, or S1PR antagonist (etrasimod; PO QD), or JAKi (upadacitinib; PO BID) from Day 17 to Day 45 and assessed for disease every 3 days (*left*) or with vehicle, MRT-6160, or S1PR antagonist (etrasimod; PO QD), or JAKi (upadacitinib; PO BID) from Day 17 to Day 45 and assessed for disease every 3 days (*left*) or with vehicle, MRT-6160, or S1PR antagonist (etrasimod; PO QD), or JAKi (upadacitinib; PO BID) from Day 17 to Day 45 and assessed for disease every 3 days (*left*) or with vehicle, MRT-6160, or S1PR antagonist (etrasimod; PO QD), or JAKi (upadacitinib; PO BID) from Day 17 to Day 45 and assessed for disease every 3 days (*left*) or with vehicle, MRT-6160, or S1PR antagonist (etrasimod; PO QD), or JAKi (upadacitinib; PO BID) from Day 17 to Day 45 and assessed for disease every 3 days (*left*) or with vehicle, MRT-6160, or S1PR antagonist (etrasimod; PO QD), or JAKi (upadacitinib; PO BID) from Day 17 to Day 45 and assessed for disease every 3 days (*left*) or with vehicle, MRT-6160, or S1PR antagonist (etrasimod; PO QD), or JAKi (upadacitinib; PO BID) from Day 17 to Day 45 and assessed for disease every 3 days (*left*) or with vehicle, MRT-6160, or S1PR antagonist (etrasimod; PO QD), or JAKi (upadacitinib; PO BID) from Day 17 to Day 45 and assessed for disease every 3 days (*left*) or with vehicle, MRT-6160, or S1PR antagonist (etrasimod; PO QD), or JAKi (upadacitinib; PO BID) from Day 17 to Day 45 and assessed for disease every 3 days (*left*) or with vehicle, MRT-6160, or S1PR antagonist (*left*) or with vehicle, MRT-6160,

MRT-6160 Reduces Inflammation-Mediated Damage of the Colon and Cytokine Production in a T-Cell Transfer Model of Ulcerative Colitis

MRT-6160 reduces inflammation-mediated damage and swelling of the colon

MRT-6160 reduces cytokine production in the mesenteric lymph node and colon

Hematoxylin and eosin-stained histopathology sections from colon at end of study

Flow cytometric (*upper row*) and cytokine bead array (*lower row*) analysis of mesenteric lymph node CD4+ T cells and colon tissue respectively

Phase 1 Biomarker Strategy to Demonstrate MRT-6160 Pharmacodynamic Effects

Phase 1 SAD/MAD in Healthy Volunteers

Provide early insights into safety, PK/PD, and effects on key immunomodulatory signaling pathways

VAV1 protein degradation

- Flow cytometry on T and B cells: whole blood (WB)
- Targeted Mass Spec: PBMCs
- Potential: Mature B cell typing in MAD

Key downstream PD

- Flow cytometry for CD69 protein on T & B cells: WB
- Immunoassay for IL-2, IL-6, IL-17, BAFF, CCL3/4
- hs C-reactive protein

Phase 1 SAD/MAD study ongoing, clinical data anticipated Q1 2025

VAV1 is an Upstream Targeting Node Associated with Clinically Validated Pathways

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New positions opening soon!

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- Computational Chemistry

Thank You

