# Development of a mode of action-based testing suite for the screening of regenerative macrophage cell therapy candidates for end-stage liver disease and beyond

# Introduction

Patients with end-stage liver disease (ESLD) have extensive fibrosis and inflammation, which cause loss of organ function and portal hypertension, resulting in decompensation events and limited life expectancy (median ~2years after first decompensation; median ~9 months after second decompensation event)<sup>1</sup> Currently, there are no licensed therapies to treat ESLD<sup>1</sup>. Macrophages with a proregenerative phenotype control inflammation and promote fibrosis remodeling, thereby coordinating liver regeneration and offering a potential therapeutic avenue in these patients<sup>2,3,5</sup>.

Autologous, non-engineered, pro-regenerative macrophages have been tested in patients with advanced cirrhosis in the academic MATCH clinical studies, which show that the therapy is well tolerated and improves transplant-free survival<sup>4,6,8</sup>. Further enhancement of pro-regenerative macrophage potency via engineering is needed to ensure durable clinical responses in a more severe patient population with ESLD.

To ensure that the best candidate was progressed to clinical trial, Resolution Therapeutics proceeded with an extensive screening process to select the most effective payloads. The screening process consisted of establishing a mode of action (MoA) based screening platform and validating on a clinically relevant cell type, candidate screening and refinement, and finally, candidate selection.

Payloads were considered effective if they maintained or enhanced phenotypic attributes as well as enhanced anti-inflammatory and anti-fibrotic functions when compared to the clinically-tested non-engineered macrophages.

Method

RTX001 was envisioned as a monocyte-derived, autologous macrophage cell therapy that has undergone *engineering* to enhance macrophage function. Monocytes are derived from leukapheresis and cultured as per protocol<sup>4,7</sup> to derive macrophages which are then engineered *ex vivo* using mRNA before being cryopreserved.

Resolution established a modular in vitro platform based on RMT MoA (Fig 1) from which it could compare potential candidates against non-engineered macrophages (MATCH-like, clinically-tested regenerative macrophages). An extensive screening program with candidate genes selected from literature5 and previous preclinical experiments was then conducted (Fig 2).

Finally, we combined the selected candidate payloads, and we tested the combination of IL10 and MMP9 against IL10 alone and non-engineered (MATCH-like) macrophages (Fig 3), comparing their functional phenotype and secretion profile.

## References

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Establish comprehensive MoA understanding and its relationship with efficacy

Develop modular *in vitro* testing suite

Verify modular testing suite with clinically relevant cells

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### Results



### Figure 3: Candidate Refinement and Selection



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**HLA-DR Fold Change** 

Fig 3B Phenotypic characterisation (Fold Change): CD86 (Left) and HLA-DR (Right) expression in RTX001/IL10 engineered macrophages compared to a non-engineered (NTrx) control. Data displayed as an MFI fold change from a Naïve macrophage



Fig 3E Human MMP9 Secretion Measured secretion of human MMP9 in the conditioned media of RTX001/IL10 engineered macrophages compared to non engineered controls (NTrx)



Fig 3F MMP Activity Measured activity of human MMPs in the conditioned media of RTX001/IL10 engineered macrophages compared to non engineered controls (NTrx)

Monocyte Migration

Fig 3H Monocyte recruitment capacity PBMCs cultured in conditioned media from macrophages or a Non-engineered (NTrx)



Fig 3I Phagocytic capacity. Percentage of RTX001/IL10 engineered and Nonengineered (NTrx) macrophages phagocytosing E.coli-coated beads.

# **Contact Information**