# Preclinical Development of LYL119, a ROR1-Targeted CAR T-Cell Product Candidate Incorporating Four Novel T-Cell Reprogramming Technologies to Overcome Barriers to Effective Cell Therapy for Solid Tumors

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#### Background

- T-cell exhaustion and lack of durable stemness, defined as the ability of cells to proliferate, persist, and self-renew, are key barriers to effective T-cell therapy in solid tumors<sup>1–2</sup>
- Ongoing efforts to overcome these barriers include genetic and epigenetic T-cell reprogramming: - Genetically reprogramming T cells through c-Jun overexpression delays exhaustion and results
- in increased proliferation, sustained cytokine production, and durable antitumor activity<sup>1,3–4</sup>
- NR4A transcription factors may contribute to exhaustion and reduced T-cell function by limiting the activity of c-Jun.<sup>5–6</sup> NR4A3 gene KO synergizes with c-Jun overexpression to further enhance resistance to exhaustion, resulting in improved antitumor activity<sup>7</sup>
- Epigenetic reprogramming with Lyell's Epi-R<sup>™</sup> manufacturing protocols preserves stem-like qualities by controlling T-cell proliferation and differentiation with optimized proprietary cell culture media and other manufacturing steps<sup>1,8–10</sup>
- Epigenetic reprogramming with Lyell's Stim-R<sup>™</sup> technology, a synthetic biomimetic designed to present precise and more physiological T-cell activation signals during manufacturing, further improves T-cell polyfunctionality, persistence, and antitumor activity<sup>11</sup>
- These four T-cell reprogramming technologies are combined to create LYL119, an investigational ROR1-targeted CAR T-cell product enhanced with c-Jun overexpression, NR4A3 KO, and Epi-R and Stim-R technologies to overcome barriers to successful T-cell therapy in solid tumors (Figure 1)

Figure 1: LYL119, an investigational ROR1-targeted CAR T-cell product enhanced with Lyell's T-cell reprogramming technologies



#### Methods

- Healthy donor T cells were transduced with a ROR1 CAR tri-cistronic lentiviral vector with (c-Jun–2A–ROR1 scFv-BBz–2A–EGFRt) or without (CD19t–2A–ROR1 scFv-BBz–2A–EGFRt) c-Jun overexpression
- NR4A3 was knocked out using CRISPR/Cas9 ribonucleoprotein delivery via electroporation. Control ROR1 CAR T-cells were edited at a control locus (human CD19) that is not expressed in T cells
- T cells were stimulated with an optimized Stim-R formulation or a standard benchmark activation reagent ("Reagent"
- T-cell products for all conditions were produced using the Epi-R manufacturing protocol
- CAR T-cell cytotoxicity and cytokine production were evaluated in vitro after primary and serial antigen-stimulation assays designed to promote exhaustion
- CAR T cells were evaluated in vivo using ROR1-expressing A549 or H1975 human lung cancer xenograft models in NSG HLA dKO mice

## Abbreviations

4-1BB, tumor necrosis factor ligand superfamily member 9; AUC, area under the curve; CAR, chimeric antigen receptor; Cas9, CRISPR associated protein 9; CCL5, chemokine ligand 5; CD, cluster of differentiation; CRISPR, clustered regularly interspaced short palindromic repeats: dKO. double knockout; EGFR, epidermal growth factor receptor; EGFRopt, optimized variant of epidermal growth factor receptor; GNLY, granulysin; gRNA, guide RNA; HLA, human leukocyte antigen; HSD, honest significant difference; IFNy, interferon gamma; IL-2, interleukin 2; KO, knockout: NR4A, nuclear receptor subfamily 4A; ns, not significant; RFU, relative fluorescent units; RNA, ribonucleic acid; ROR1, receptor tyrosine kinase-like orphan receptor; scFv, single-chain variable fragment; SEM, standard error of the mean; TCF7, transcription factor 7; TIGIT, T-cell immune receptor with immunoglobin and ITIM domains; UMAP, uniform manifold approximation and projection.

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#### Results

**NR4A3 KO and c-Jun overexpression synergize to reduce exhaustion and maintain ROR1 CAR T-cell function in vitro and in vivo** 

- NR4A3 KO + c-Jun ROR1 CAR T cells demonstrated improved tumor cell killing and produced high levels of cytokines (data not shown), and reduced surface expression of inhibitory receptors after repetitive antigen stimulation compared with ROR1 CAR T cells with c-Jun overexpression alone (**Figure 2**)
- *NR4A3* KO + c-Jun ROR1 CAR T cells displayed transcriptional profiles consistent with reduced terminal exhaustion and increased retention of memory T cells following serial stimulation (Figure 3)
- NR4A3 KO + c-Jun ROR1 CAR T cells had robust antitumor efficacy in vivo with activity observed at a 7-fold reduction in CAR T-cell dose and a 10-fold greater expansion in blood compared with control CAR T cells enhanced with c-Jun overexpression alone (Figure 4)





(B) NR4A3 KO + c-Jun ROR1 CAR T cells had superior cytotoxicity with significant differences at the end of the seventh sequential stimulation with H1975 target cells. (C) NR4A3 KO + c-Jun ROR1 CAR T cells have reduced expression of TIGIT and increased CD127 expression after four stimulations. Shapes correspond to CAR T cells derived from each donor. Asterisks indicate significant differences compared to NR4A3 KO + c-Jun ROR1 CAR T cells. Error bars represent mean ± SEM. \*p<0.05; \*\*p<0.005; \*\*\*p<0.001; \*\*\*\*p<0.0001 by unpaired t-test (**A**, **B**) or paired t-test (**C**).

Figure 3: Transcriptomic analysis revealed that *NR4A3* KO reduces terminal exhaustion and enhances memory-like cell proportions following serial restimulation with antigen



(A) Single-cell RNA-seq clustering analysis identified a terminally exhausted cluster (C6, brown) and a CD127-hi cluster (C1, green) of CD8+ NR4A3 KO and controledited ROR1 CAR T cells with c-Jun overexpression following 7 days of in vitro antigen restimulation using A549 target cells. (B) Cluster C6 was enriched for expression of genes identified in a previously published T-cell terminal exhaustion gene set,<sup>6</sup> whereas cluster C1 was enriched for expression of CD127. (C) The proportion of cells in the terminally exhausted cluster C6 was reduced, and the proportion in the CD127-hi cluster C1 was increased in NR4A3 KO + c-Jun ROR1 CAR T cells compared to control-edited + c-Jun ROR1 CAR T cells. Closed shapes represent CAR T cells derived from one of three independent donors.



![](_page_0_Figure_43.jpeg)

H1975 (**A-C**) and A549 (**D-E**) xenograft in vivo models, NR4A3 KO + c-Jun ROR1 CAR T cells had potent antitumor activity (A, D), increased expansion of peripheral blood CD3 CAR T cells derived at Day 14 (B) of Day 13 (E) post T-cell injection, and improved survival of tumor-bearing mice (**C**). Data were averaged for CAR cells derived from 3 independen donors with n = 10 mice/group/dono(A-C) or 1 donor with n = 5 mice/group (D-E). Asterisks indicate significant differences compared to NR4A3 KO + c-Jun ROR1 CAR T-cell-treated animals. Error bars represent mean ± SEM. \*p<0.05; \*\*p<0.005; \*\*\*p<0.001; \*\*\*\*p<0.0001 by Tukey's one-way ANOVA (A, D), unpaired t-test (B, E), or log-rank Mantel-Cox test (C).

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Stim-R mock

0

0 10 20 30 40 50

Davs after T-cell injection

Reagent

![](_page_0_Figure_46.jpeg)

ROR1 CAR T cells produced with Stim-R technology showed improved antitumor activity (A) and overall survival (B) compared to mock untransduced Stim-R T cells (Stim-R mock) and Reagent CAR T cells. (C) ROR1 CAR T cells produced with Stim-R technology showed higher peak CAR T-cell numbers. prolonged persistence. and higher total CAR T cells detected over the course of the study as assessed by calculating the AUC compared to ROR1 CAR T cells produced with Reagent. Data in (A) represent individual tumor growth curves of n = 10 mice per condition. Data in (C) represent mean  $\pm$  SEM of n = 10 mice per condition. \*\*\*\*p<0.0001 by Mantel-Cox test (B) or one-way ANOVA, followed by Tukey's HSD post hoc test for each metric (C); only Reagent and Stim-R-produced CAR T-cell conditions shown.

Days after T-cell injection

![](_page_0_Picture_49.jpeg)

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![](_page_0_Picture_50.jpeg)

LYL119 ROR1 CAR T cells, incorporating NR4A3 KO + c-Jun + Epi-R + Stim-R technologies, demonstrate superior in vitro and in vivo activity

- LYL119 demonstrates prolonged cytotoxicity and enhanced cytokine production in vitro compared with control cells (**Figure 7**)
- LYL119 has potent antitumor efficacy and superior CAR T-cell expansion in vivo compared with control-edited ROR1 CAR T cells (with only c-Jun and Epi-R and Stim-R technologies; Figure 8)

#### Figure 7: LYL119 demonstrates prolonged cytotoxicity and enhanced cytokine production in vitro

![](_page_0_Figure_55.jpeg)

Stim-R + Epi-R + c-Jun Stim-R + Epi-R + c-Jun + NR4A3 KO (LYL119)

(A) LYL119 CAR T cells showed superior target clearance compared to controls. (B) LYL119 CAR T cells showed a trend of higher IFNv and IL-2 production after repeated stimulation. Representative data from 1 in 3 donors is shown (A, B). \*\*p=0.0017 compared to "Reagent + Epi-R + c-Jun + NR4A3 KO". Statistical comparisons were performed using the AUC of the final round of restimulation in three donors using a one-way ANOVA analysis.

![](_page_0_Figure_58.jpeg)

LYL119 has potent antitumor activity (A) and enhanced expansion of peripheral blood CD3+ CAR T cells (B) in the H1975 xenograft model. Data are CAR T cells derived from 1 independent donor with n = 5 mice/group. Asterisks indicate significant differences compared to LYL119-treated animals. Error bars represent mean ± SEM. \*p<0.05; \*\*p<0.005; \*\*\*p<0.001; \*\*\*\*p<0.0001 by Tukey's one-way ANOVA (A) or unpaired t-test (B)

#### Conclusion

- NR4A3 KO and c-Jun overexpression synergize to reduce CAR T-cell exhaustion and enhance ROR1 CAR T-cell function
- Stim-R technology generates ROR1 CAR T cells with enhanced proliferative capacity and sustained antitumor activity
- Preliminary data suggest that ROR1 CAR T cells genetically modified to knock out NR4A3 expression and overexpress c-Jun and epigenetically reprogrammed using Epi-R and Stim-R technologies demonstrate additive functional enhancements to further improve CAR T-cell antitumor efficacy in vitro and in vivo
- To test whether this strategy can enhance the efficacy of T-cell therapy in patients with ROR1-expressing solid tumors, Lyell is developing LYL119, an investigational ROR1-targeted CAR T-cell product candidate that incorporates these four reprogramming technologies

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