

# PDL1-targeted synthetic receptor engineering for T cell immunotherapy in solid tumor

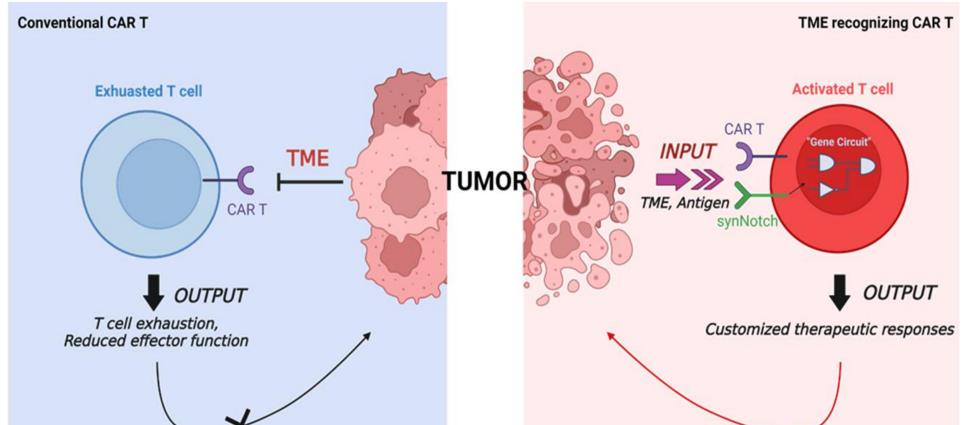
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## Introduction

## Figure 1 | Schematic of overcoming solid TME by Synnotch signaling.

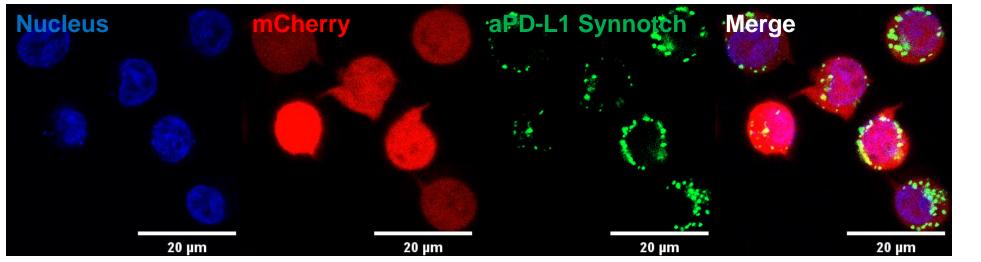


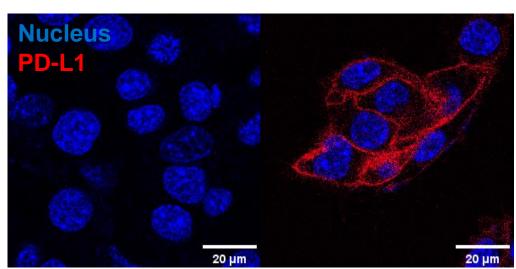
CAR T cell therapy has demonstrated significant potential, particularly in hematological malignancies, but its efficacy in solid tumors is hindered by the immunosuppressive tumor microenvironment (TME). Within the TME, immune checkpoints such as PD-1 induce T cell exhaustion, limiting their ability to sustain an antitumor response. This study introduces a synNotchbased CAR T platform designed to **overcome** these challenges by recognizing TME-specific antigens and incorporating a genetic circuit for IL-10 cytokine secretion. This system enhances **immune activation**, supports effector molecule production, and inhibits PD-1 signaling, thereby preventing T cell exhaustion and enabling a more durable anti-tumor response, even in the presence

## Results

Figure 5 | Effector cells and Target cells express each component after lentiviral transduction





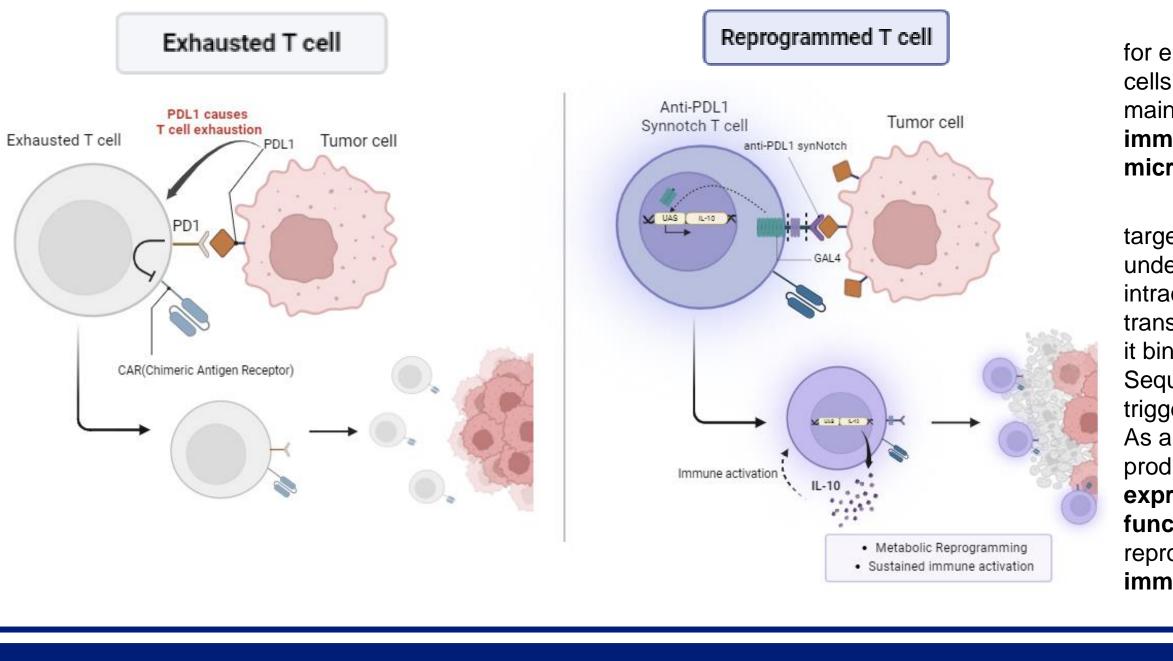




### Effective tumor cell killing

of immunosuppressive signals.

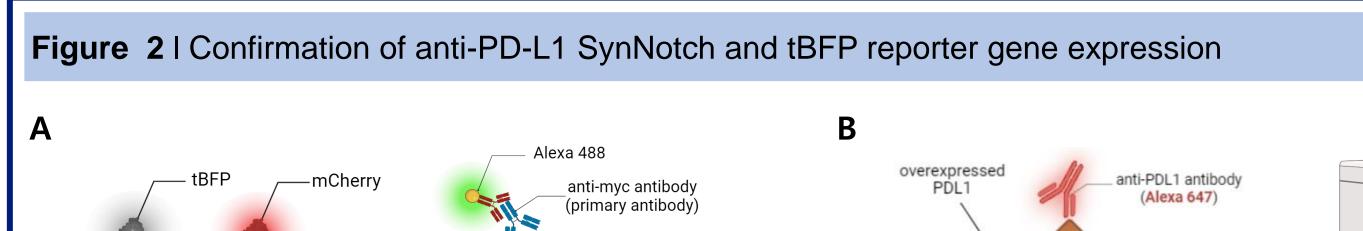
## Mechanism

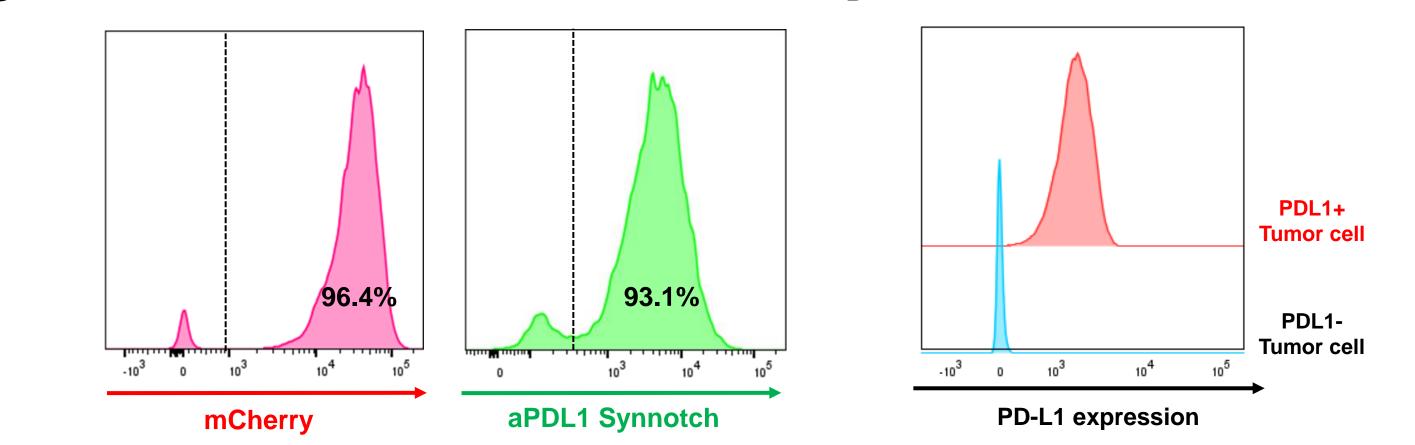


The proposed mechanisms for engineering SynNotch CAR T cells that resist exhaustion and maintain effector functions in **immunosuppressive tumor microenvironments (TME).** 

When SynNotch binds its target antigen (PD-L1), it undergoes cleavage, releasing an intracellular transcription factor that translocates to the nucleus. There, it binds to the Upstream Activation Sequence (UAS) promoter region, triggering IL-10 gene expression. As a result, engineered T cells produce and secrete IL-10. IL-10 expression activates the functions of metabolic reprogramming and sustained immune activation.

# Experimental Methods





#### Figure 5 I

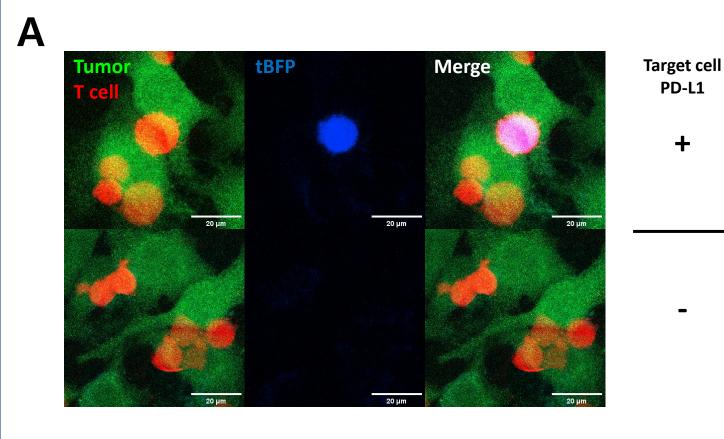
A) Confirmation of anti-PD-L1 synNotch expression on the membrane of Jurkat T cells by confocal microscopy. Cells were stained with DAPI (Nucleus) and antimyc (primary antibody) and anti-mouse Alexa fluor 488 (secondary antibody).

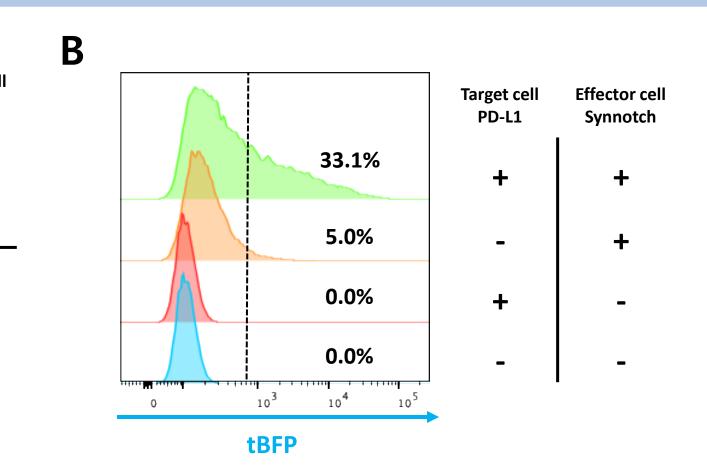
**B)** Confirmation of PD-L1 expression on the membrane of B16F10 tumor cells by confocal microscopy. Cells were stained with DAPI (Nucleus) and anti-PD-L1 APC conjugated antibody (primary antibody).

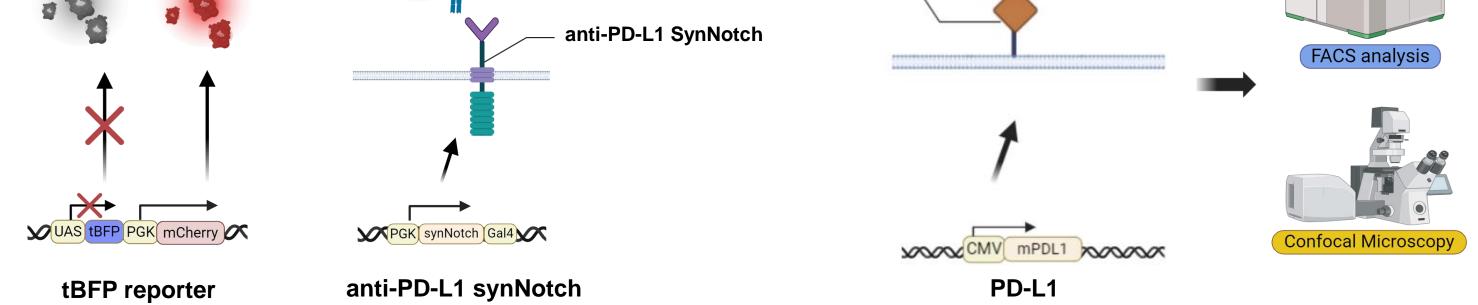
## **C)** Confirmation of anti-PD-L1 synNotch expression on the membrane of Jurkat T cell by Flow cytometry.

**D)** Confirmation of PD-L1 expression on B16F10 Tumor cells by Flow cytometry. FACS result shows that expresses each fluorescence component accordingly. (PD-L1and PD-L1+ tumor cells).

## **Figure 6 I** Jurkat T-cell expresses tBFP selectively when co-cultured with PDL1 overexpressing B16F10 tumor cells.





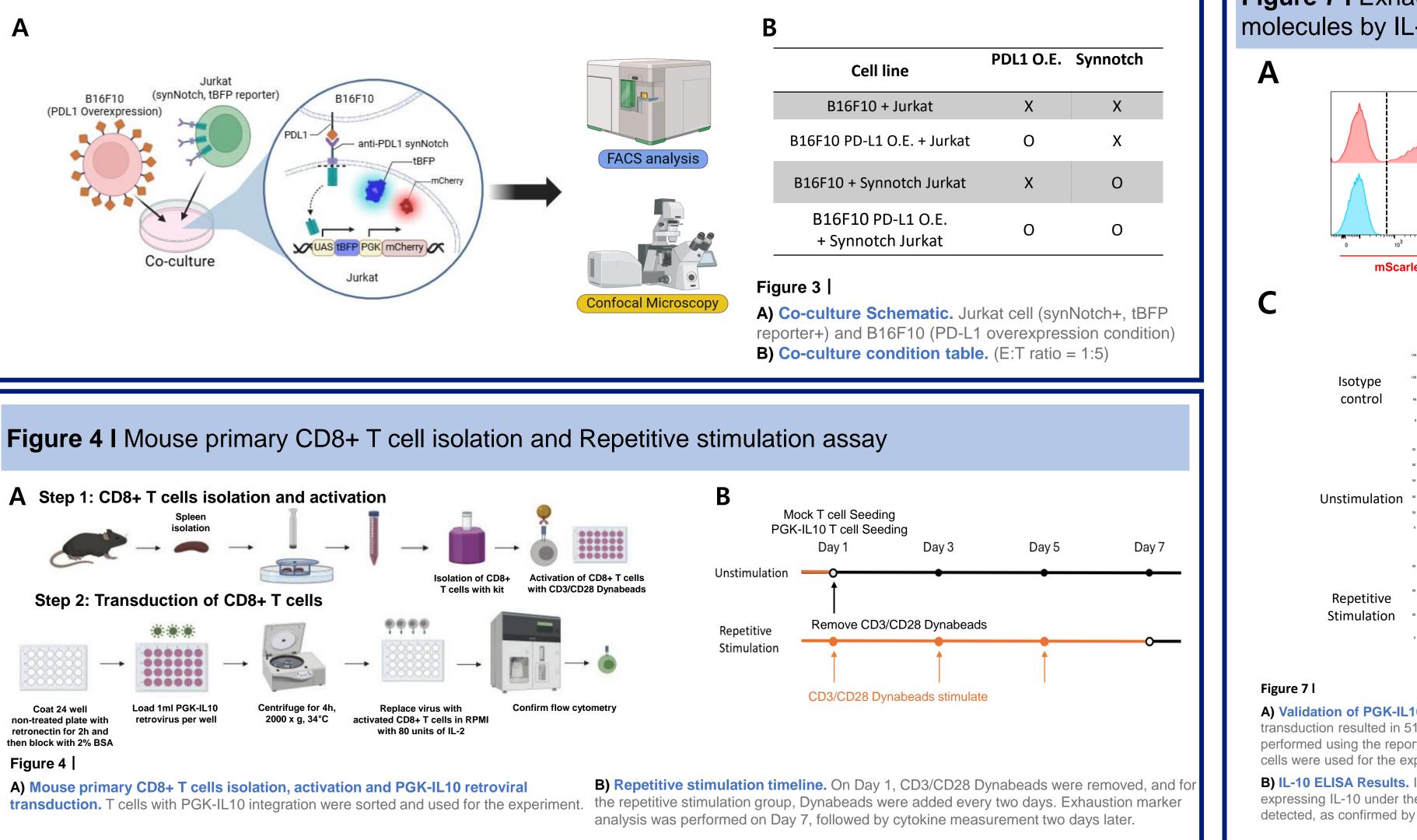


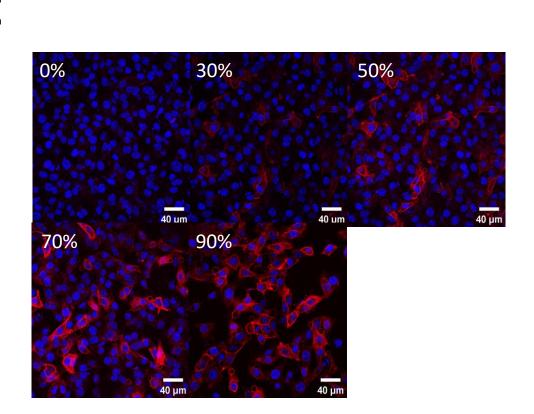
#### Figure 2 I

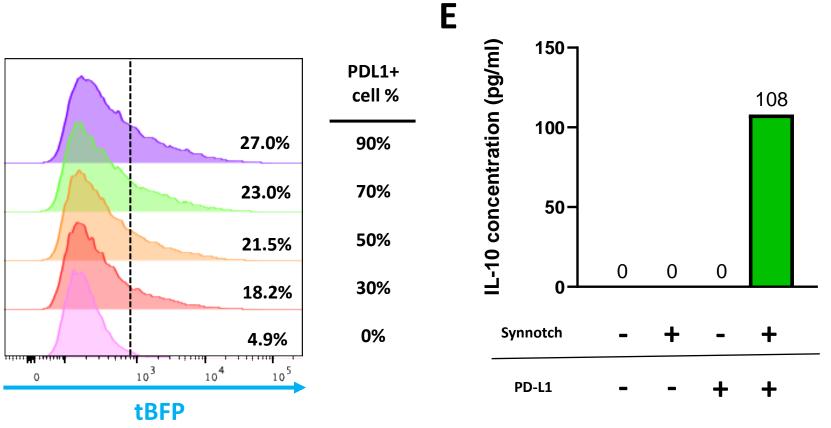
**A)** Confirmation of tBFP reporter and synNotch transduction via antibody staining. Two plasmids were synthesized using a lentivector plasmid, one tBFP reporter (UAS-tBFP-PGK-mCherry) and synNotch (PGK-anti-PDL1-synNotch-Gal4) plasmid. Lenti-X HEK293T was used as host. Lentivirus was harvested and transduced to Jurkat T cell line. ICC was conducted with anti-myc (primary antibody) and anti-mouse-488 (secondary antibody). Fluorescence data was obtained through Confocal Microscopy and FACS analysis.

**B)** Confirmation of PD-L1 transduction via antibody staining. A plasmid for PDL1(pHR-CMV-mPDL1) was synthesized using Lenti-X HEK293T as a host. After harvest, the lentivirus was transduced to the B16F10 cell line. ICC was conducted with anti-PDL1 antibody expressing Alexa 647 signal. Fluorescence data were obtained through Confocal Microscopy and FACS analysis.

**Figure 3 I** Co-culture for confirming selective binding of anti-PDL1 synNotch T cells to PDL1 overexpressing tumor cells.



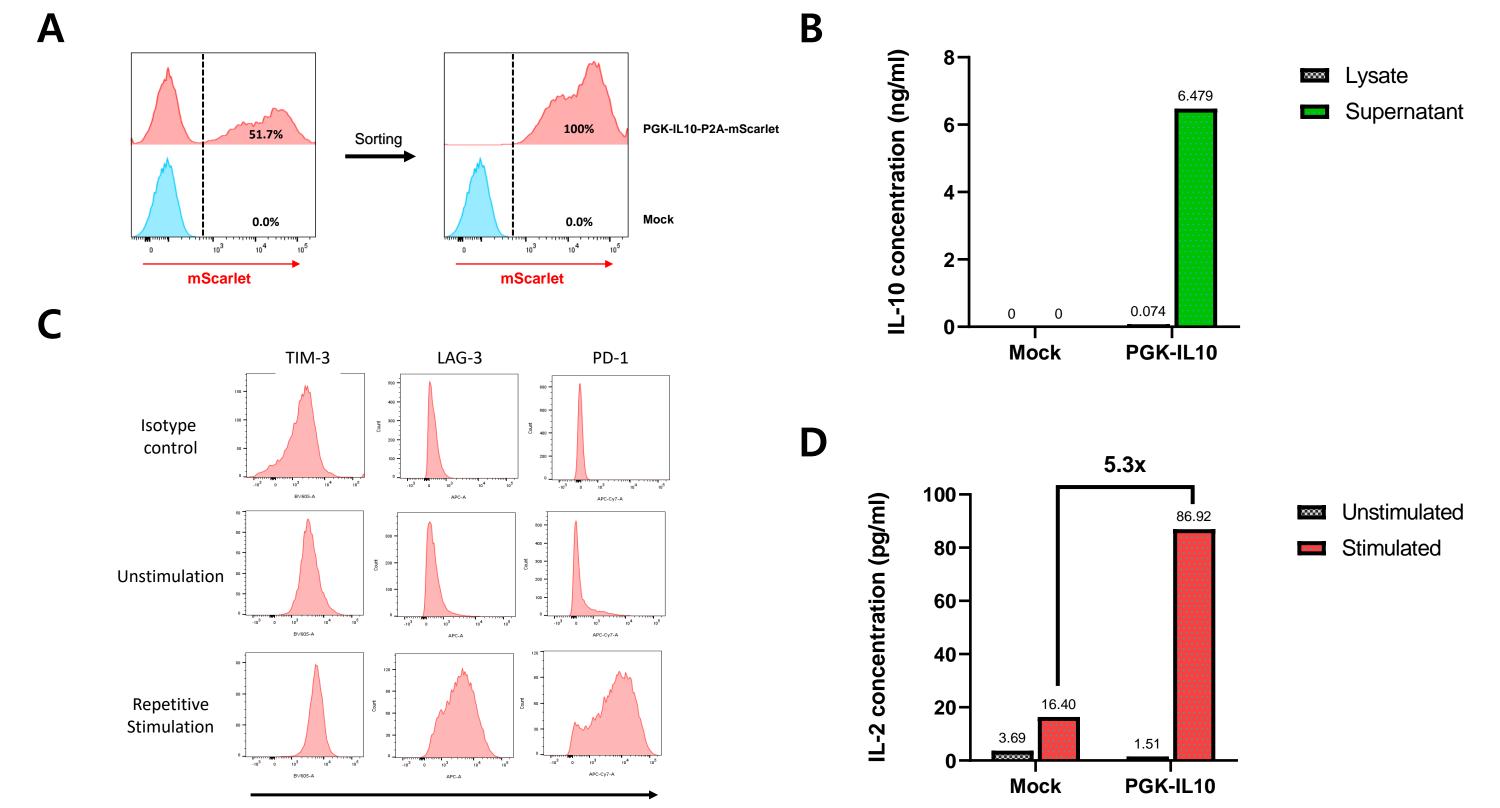




#### Figure 6 I

A,B) Comparison of relative tBFP signals. ICC imaging data shows that T cell SynNotch activation occurs only when PD-L1 is expressed in tumor cells. Flow cytometry data shows that T cell SynNotch activation occurs only when PD-L1 is expressed in tumor cells. As the percentage of PD-L1<sup>+</sup> cells increases, SynNotch activation in cocultured T cells also increases.

## **Figure 7 I** Exhaustion marker analysis after primary T cell stimulation and relative increase in effector molecules by IL-10.



#### Ire 7 I Exhausted

**A)** Validation of PGK-IL10 expression in primary mouse T cells. Following isolation, transduction resulted in 51.7% expression of PGK-IL10. Subsequently, sorting was performed using the reporter fluorescent protein mScarlet, and only PGK-IL10-expressing cells were used for the experiment.

**B) IL-10 ELISA Results.** IL-10 was detected only in the supernatant of primary T cells expressing IL-10 under the PGK promoter. In contrast, IL-10 in the cytosol was not detected, as confirmed by lysate measurements.



**D)** IL-2 ELISA Results. Cytokine secretion was measured after sorting PD-1<sup>+</sup> LAG-3<sup>+</sup> cells from the repetitive stimulation group. IL-2 expression was higher in the stimulation group compared to the unstimulated group, and the PGK-IL10-expressing group showed a 5.3-fold increase in IL-2 levels.

## Conclusion & Further Study

Our study demonstrates that SynNotch-based engineered **T cells can selectively recognize PD-L1-expressing tumor cells and sustain immune activation** through **IL-10 secretion.** This system effectively enhances effector molecule production and reduces exhaustion marker expression, thereby overcoming immunosuppressive signals in the tumor microenvironment. However, some limitations remain, including slight **leakage from the UAS promoter** and the potential for further **optimization of activation efficiency**. Addressing these issues will improve the precision and efficacy of the system. For future studies, we aim to evaluate this approach in **in vivo tumor models**. Specifically, we will assess whether **SynNotch-driven IL-10 secretion significantly enhances anti-tumor cytotoxicity** while maintaining T cell functionality within the immunosuppressive tumor microenvironment. These findings will provide valuable insights for the clinical translation of SynNotch-based T cell therapies.