

## RESEARCH BASED TEMPLATE

Submissions must not exceed 300 words (excluding title & authors). The document **must not** be password protected or saved as read only as this may result in your abstract failing to upload successfully. Use Arial 12 point type only. Please structure your submission using the subheadings below. If the abstract does not fit the headings, please put full abstract beneath introduction and we will remove the headings once submitted.

# Optimization of Lipid Nanoparticle-based CRISPR-activation to Improve HIV Latency Reversal

## Authors:

Moso MA<sup>1,2</sup>, Cevaal PM<sup>1</sup>, Tan A<sup>1</sup>, Fisher BM<sup>1</sup>, Johnston A<sup>4</sup>, Teo S<sup>4</sup>, Chen M<sup>4</sup>, Fareh M<sup>5,7</sup>, Quadeer A<sup>3,7</sup>, McKay M<sup>3,7</sup>, Lewin SR<sup>1,2,8</sup>, Roche M<sup>1</sup>

<sup>1</sup>Department of Infectious Diseases, The University of Melbourne at the Peter Doherty Institute for Infection and Immunity, <sup>2</sup>Victorian Infectious Diseases Service, The Royal Melbourne Hospital at the Peter Doherty Institute for Infection and Immunity, <sup>3</sup>Department of Microbiology and Immunology, The University of Melbourne at The Peter Doherty Institute for Infection and Immunity, <sup>4</sup>Monash Institute of Pharmaceutical Sciences, Monash University; <sup>5</sup>Cancer Immunology Program, Peter MacCallum Cancer Centre, <sup>6</sup>Sir Peter MacCallum Department of Oncology, The University of Melbourne, <sup>7</sup>Department of Electrical and Electronic Engineering, The University of Melbourne, <sup>8</sup>Department of Infectious Diseases, Alfred Hospital and Monash University.

## Background:

Current latency reversal agents (LRAs) lack specificity and potency. We previously harnessed the inherent sequence specificity of CRISPR activation (CRISPRa) to induce HIV transcription *ex vivo* but observed incomplete proviral reactivation, which may be related to poor expression of the CRISPRa construct in primary CD4+ T cells. CRISPRa may also be limited by the vast sequence diversity of HIV. Here, we optimised CRISPRa in a lipid nanoparticle (LNP)-based delivery system to improve its activity and coverage for diverse HIV subtypes.

## Methods:

CRISPRa expression was quantified by intracellular staining (ICS) for dCas9 protein. mRNA expression was optimised by altering its 5' and 3' Untranslated Regions (UTRs). LNPs were modified by conjugating T cells specific targeting antibodies (CD7) to its surface. CRISPRa activity in primary T cells was evaluated using CD25-targeting gRNA and measurement of surface CD25 expression by flow cytometry. Cross-subtype gRNAs were designed by generating variants of HIV-targeting gRNAs matched to non-B sequences from the Los Alamos database. CRISPRa HIV latency reversal was assessed in the J-Lat A2 cell line via GFP expression.

## Results:

High expression of dCas9 was observed in JLatA2 (90.6% dCas9+) but not in activated (3.2%) or resting CD4+ T cells (0.9%). Altered dCas9 mRNA enhanced expression in J-Lat A2 (9.7-fold increase in dCas9 expression) but not in resting CD4+ T cells. CD7-targeted LNPs improved CRISPRa activity (6.1-fold increase in CD25 expression). gRNA pools of up to 9 gRNAs improved coverage to greater than 75% for subtype A1, B, C, D, AE sequences. The CRISPRa pan-genotypic LNPs

## RESEARCH BASED TEMPLATE

Submissions must not exceed 300 words (excluding title & authors). The document **must not** be password protected or saved as read only as this may result in your abstract failing to upload successfully. Use Arial 12 point type only. Please structure your submission using the subheadings below. If the abstract does not fit the headings, please put full abstract beneath introduction and we will remove the headings once submitted.

retained activity in the J-Lat A2 cell line, with slight reduction in potency (70.6% activity Cf. subtype B specific gRNAs).

### **Conclusion:**

CRISPRa activity was improved through alteration in mRNA and use of CD7-targeted LNPs. Sequence coverage was improved by combination of gRNA variants with a minor loss of potency.

**Disclosure of Interest Statement:** No conflicts of interest to disclose.