

Development of a Novel Interleukin-27 (IL-27) Targeted Gene Therapy for Therapeutic Applications

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INTRODUCTION

SARS-CoV-2 relies on the angiotensin-converting enzyme 2 (ACE2) transmembrane peptidase as its cellular receptor and is responsible for **7,000,000 global deaths** since 2019¹. Current therapeutics (e.g., plasmapheresis, hydroxychloroquine, etc.) prove ineffective in many patient populations and support the need for more targeted therapies².

Interleukin-27 (IL-27) is a pleiotropic cytokine with anti-inflammatory and antiviral properties³. We designed a novel plasmid encoding IL-27 with an ACE2-targeting peptide (IL-27^{ACE2pep}) and proposed an IL-27 delivery via adipose stromal mesenchymal cells (ASC) to models of lung inflammation, exploring *in vitro* potential of ASC in delivering therapeutics to target cells. Elucidating the effects of IL-27^{ACE2pep} hASC CM on A549-ACE2 establishes a foundation for developing life-saving IL-27 targeted therapies for COVID-19 and a multitude of other conditions, such as ACE2-overexpressing tumors and autoimmune conditions.

HYPOTHESIS

We hypothesize that human ASC conditioned media (hASC CM) containing IL-27^{ACE2pep} will **reduce both ACE2 expression and viral entry** in A549-ACE2 cells, inhibiting inflammation.

RATIONALE

Our *in silico* analysis demonstrated the remarkable ability for IL-27 to induce antiviral genes and to restore cytokine balance, positioning this as a potentially life-saving translational therapy. When previously tested on HEK293-ACE2, IL-27^{ACE2pep} hASC CM caused a STAT1-mediated upregulation of ACE2, increasing SARS-CoV-2-Spike pseudotyped lentivirus entry. Thus, we sought to determine whether A549-ACE2, a lung epithelial cell line, would elicit the same response.

The applications of this therapy extend to a plethora of immune reactions associated with cytokine release syndrome, including autoimmune diseases, viral infections, and immunotherapy treatments for cancer (such as CAR T-cell therapy). Additionally, this therapy can be applied as anti-cancer agents for tumors overexpressing ACE2, such as papillary renal cell carcinoma (94% overexpress ACE2)⁴. Therefore, the preclinical experimentation on models of lung inflammation are promising for a wide variety of therapeutic applications and reinforce the importance of this research.

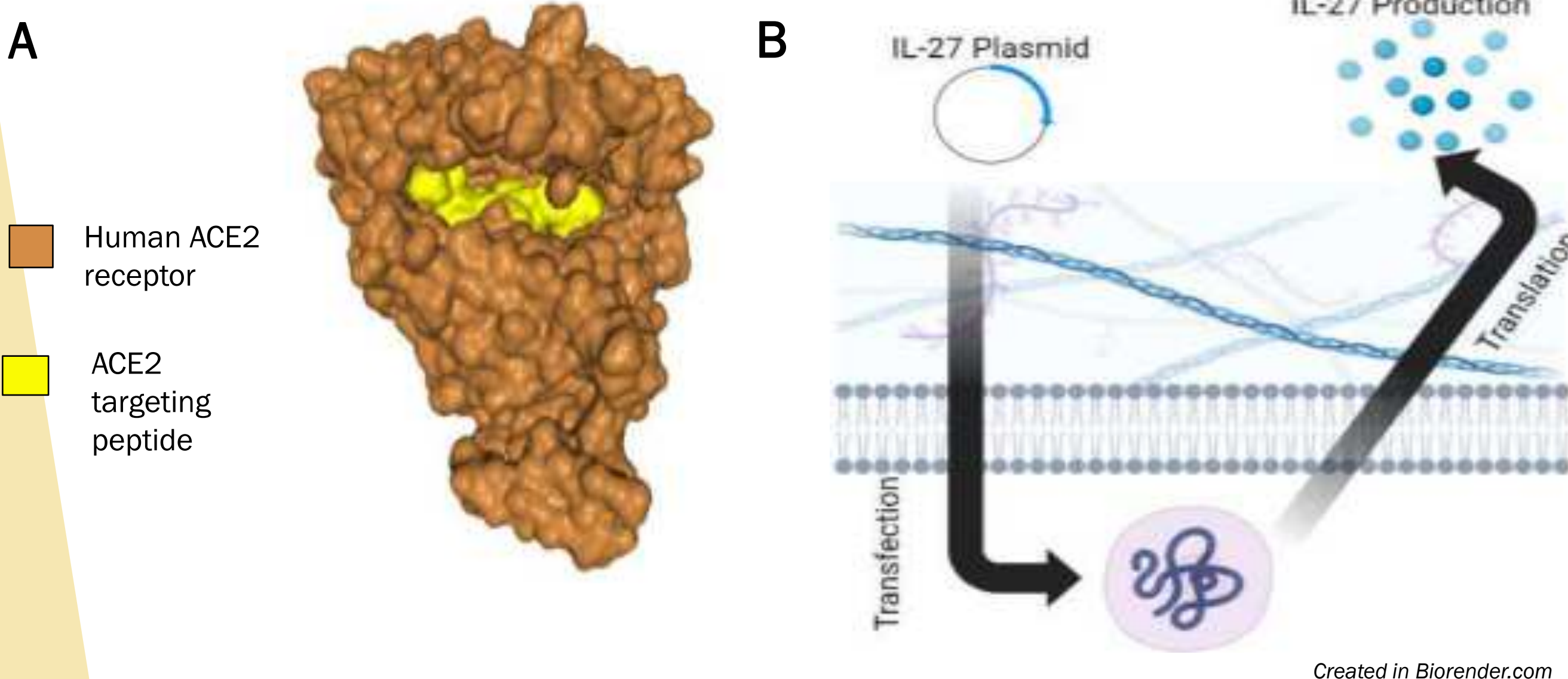


Fig 1. Overview of Novel IL-27^{ACE2pep} Gene Delivery

(A) Predicted binding of the ACE2-targeting peptide to Human ACE2 receptor using HPEPDOCK.

(B) Transfection of the IL-27 plasmid initiates translation and consequential production of IL-27. Human adipose stromal/stem cells (hASC) were used to produce IL-27, and the conditioned media was collected and stored for future experiments.

METHODOLOGY AND RESULTS

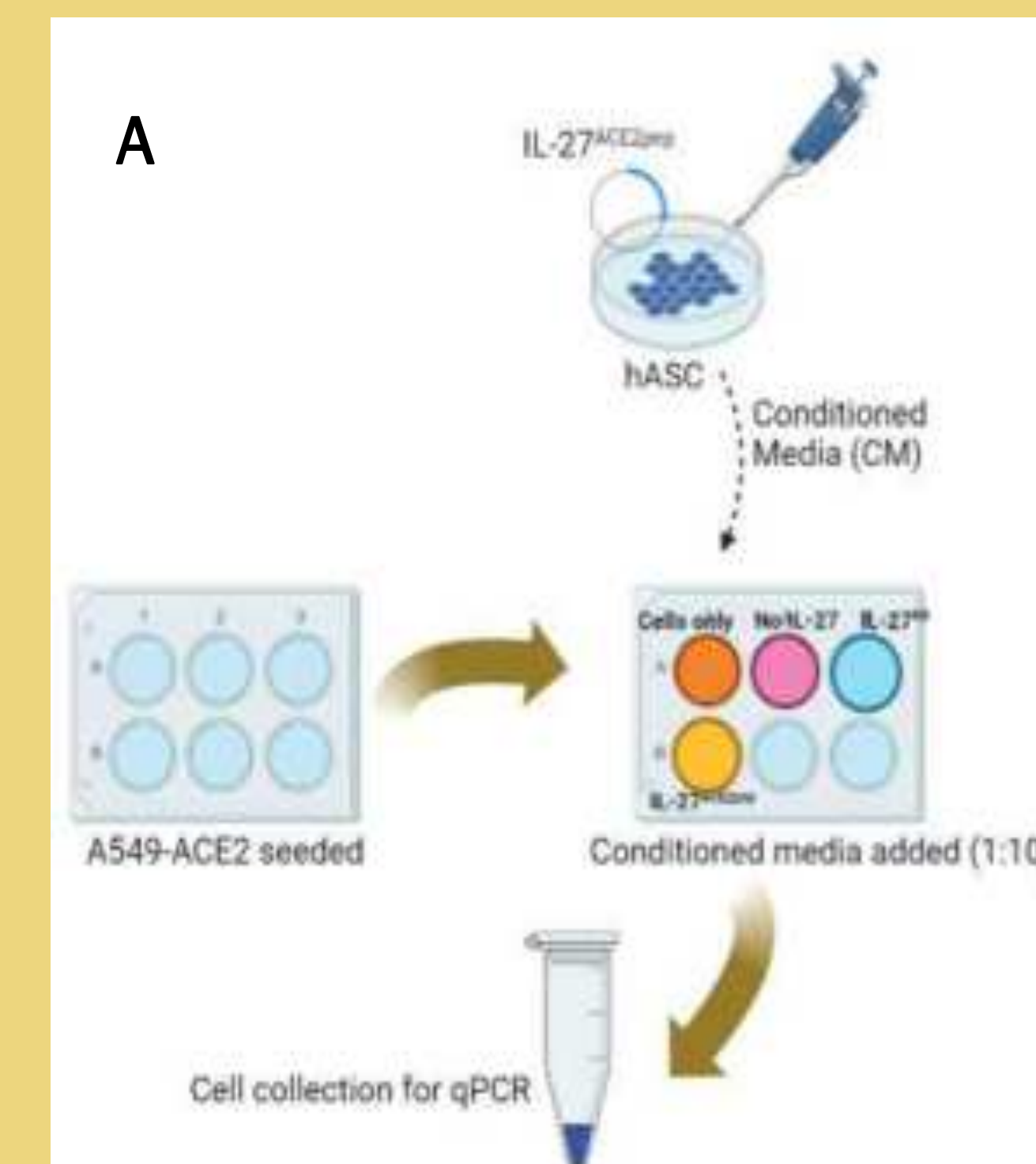


Fig 2. Determining gene expression in A549-ACE2 cells following treatment of IL-27 hASC CM

(A) Methodology overview to determine gene expression in A549-ACE2 cells following treatment of IL-27 hASC CM. 150,000 A549-ACE2 cells/well were seeded in a six-well plate. The cells were then treated with conditioned media with the following treatments: cells only, empty vector (No IL-27), non-targeting IL-27 (IL-27^{NS}), and ACE2 targeting IL-27 (IL-27^{ACE2}). The cells were collected 48 hours after treatment and analyzed via qPCR.

(B) Gene expression fold change in A549-ACE2 cells treated with IL-27^{NS} and IL-27^{ACE2}. Target genes analyzed were ACE2, dACE2, STAT1, and STAT3, normalized to GAPDH. Data is represented as mean ± S.D. (**p<.001).

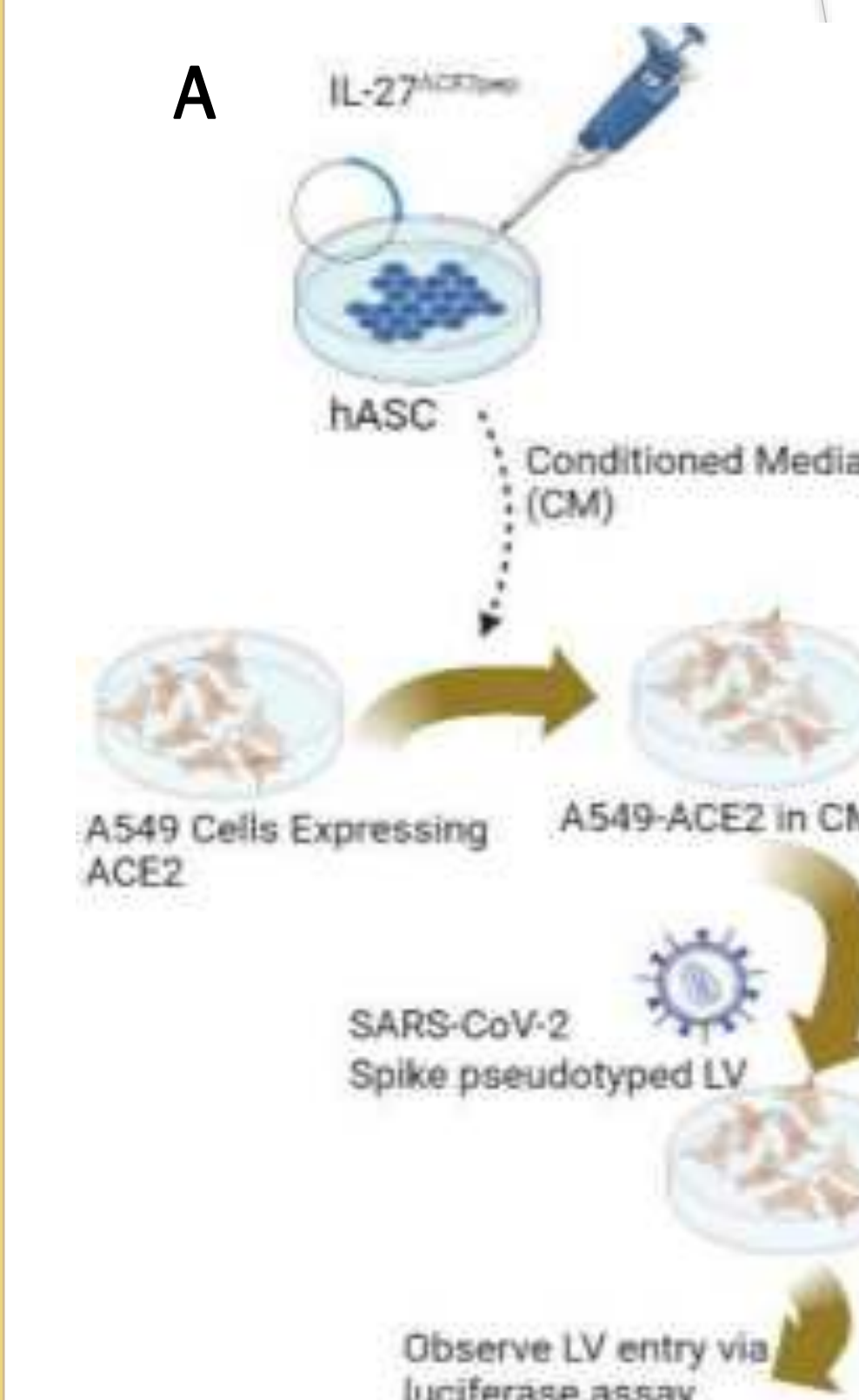
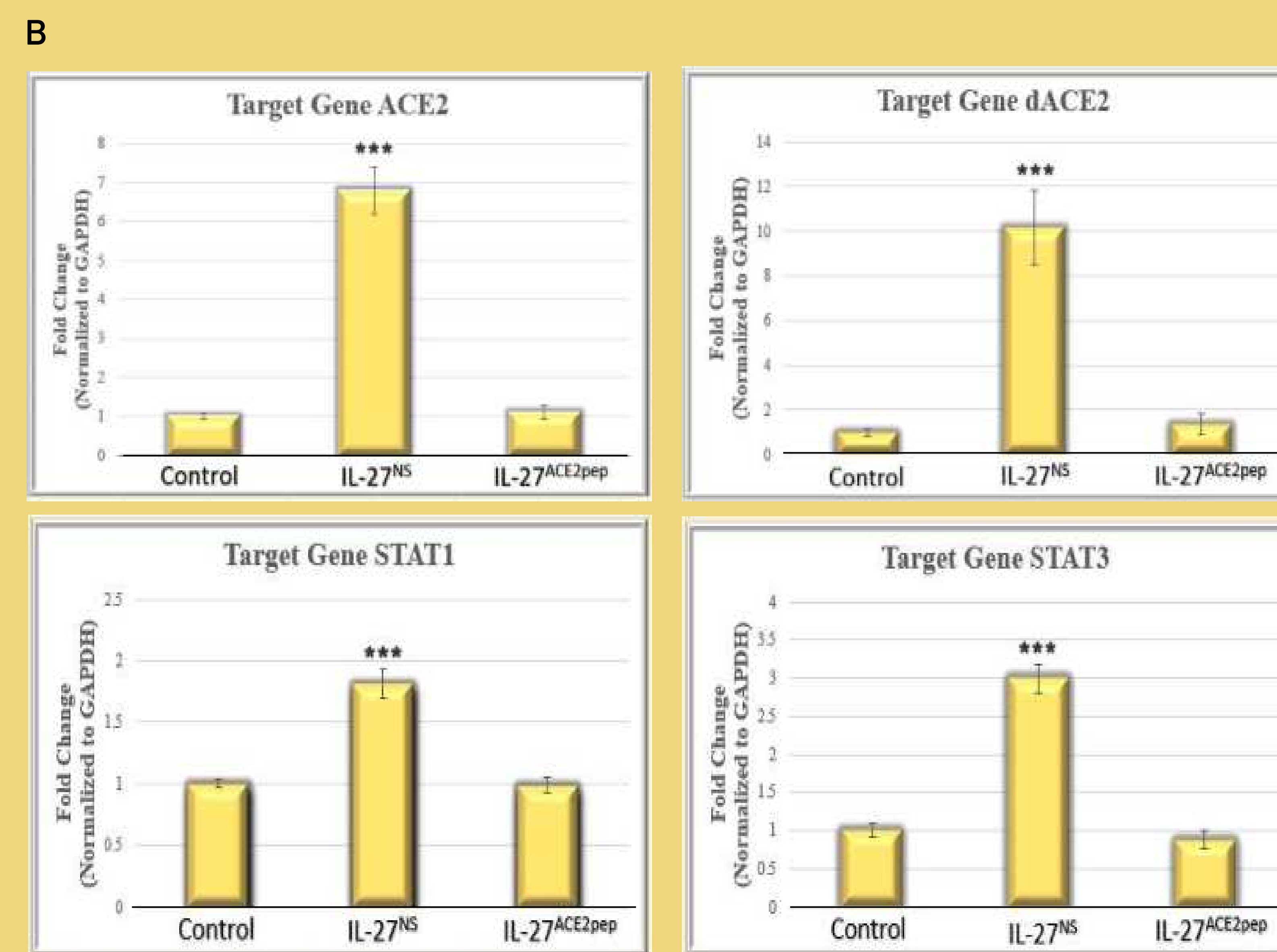
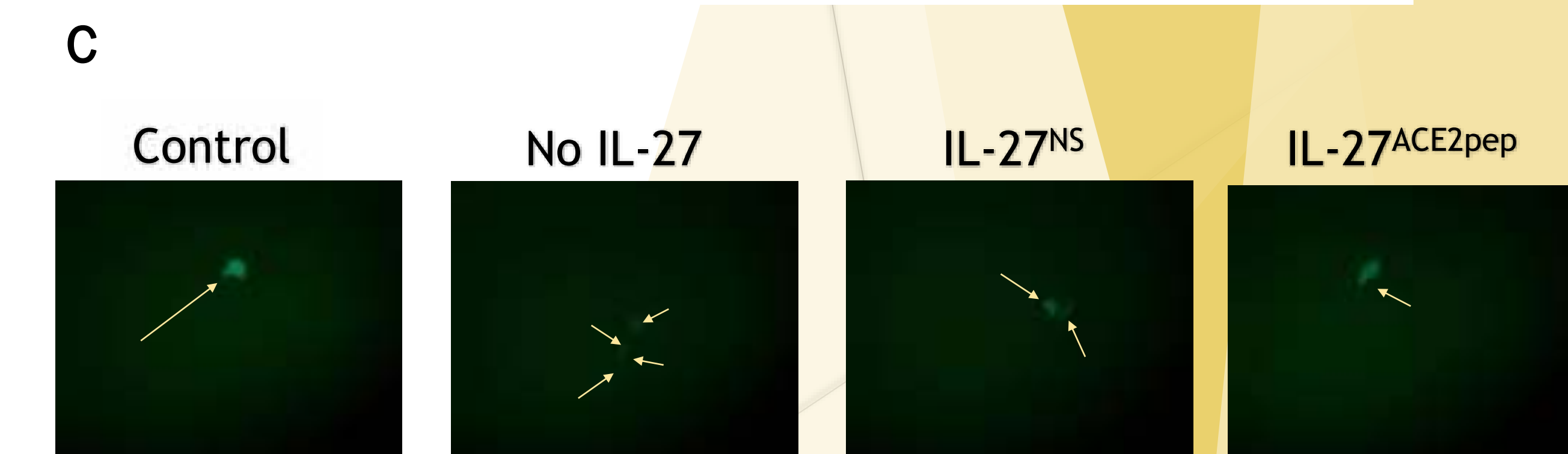
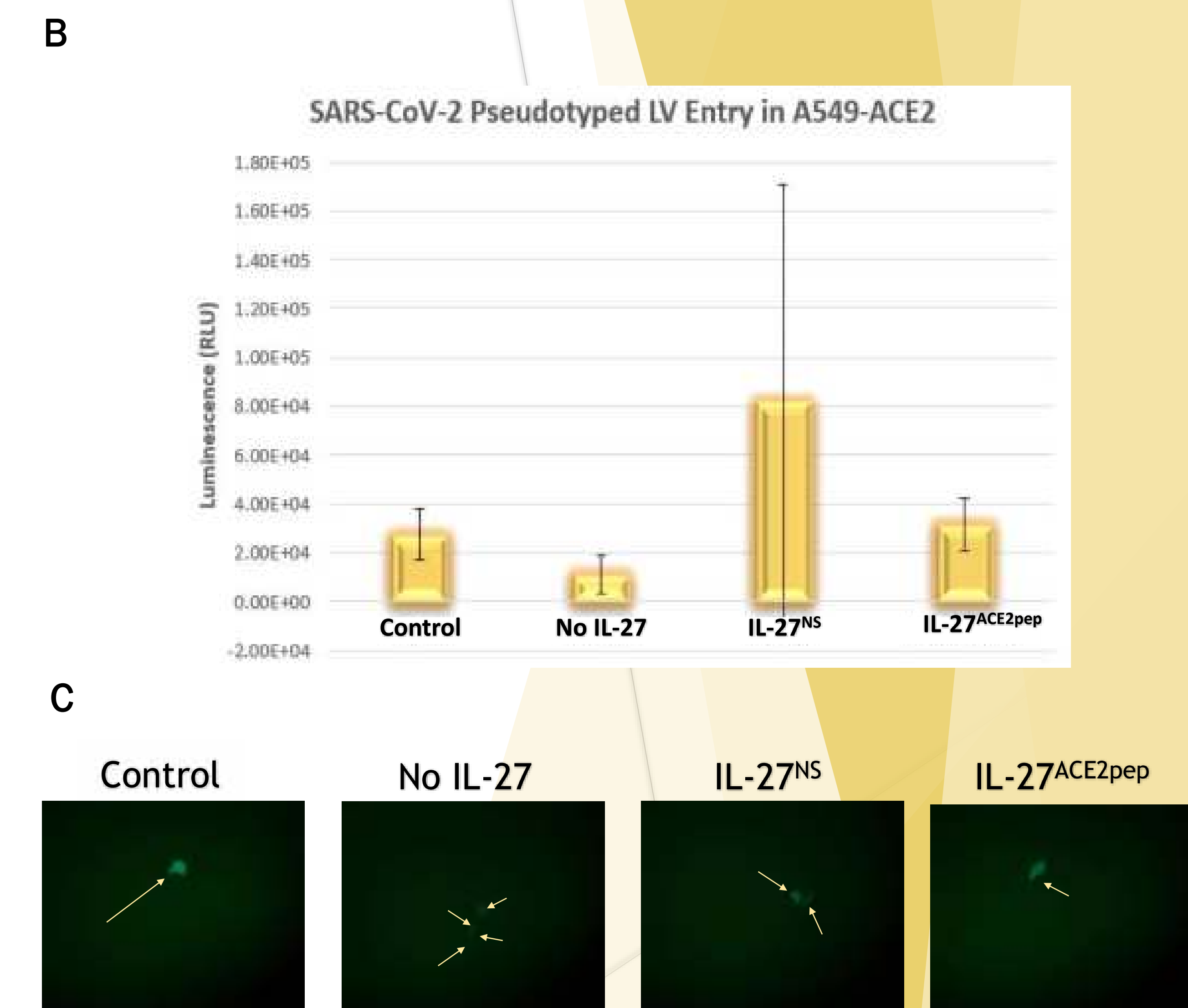


Fig 3. Determining SARS-CoV-2-Spike pseudotyped lentivirus entry in A549-ACE2 following treatment of IL-27 hASC CM

(A) Methodology overview to determine SARS-CoV-2-Spike pseudotyped lentivirus entry in A549-ACE2. Cells were seeded in 96-well plate at 5,000 cells/well. The cells were then treated with/without conditioned media (same as Fig 3) for 4 hours before media is replaced, and pseudotyped LV was added. Luciferase assay was conducted at 48h post LV transduction.

(B) SARS-CoV-2-Spike pseudotyped LV entry in A549-ACE2 measured in luminescence (RLU). SARS-CoV-2 pseudotyped LV expresses synthetic firefly luciferase (Luc2). Data is represented as mean ± S.D. and paired t-test was used to determine significance.

(C) Representative images of A549-ACE2 taken at 48 hours post LV transduction. SARS-CoV-2-Spike pseudotyped LV expresses green fluorescence protein (ZsGreen1). Images were taken using an inverted fluorescent microscope (Olympus IX71) at 10X magnification.



DISCUSSION & FUTURE DIRECTIONS

Our *in vitro* analysis demonstrated a **reduced trend of LV entry** following treatment with IL-27^{ACE2pep} compared to IL-27^{NS}. This is likely due to the **differences in ACE2 expression**, in which **non-specific IL-27 (IL-27^{NS}) hASC CM significantly upregulated ACE2 expression**, whereas **IL-27^{ACE2pep} CM did not**. Analyzed together, the results suggest existing differences in responses between HEK293-ACE2 and A549-ACE2 cells to our therapy and show promise in mitigating the frequently life-threatening effects of COVID-19 and other conditions. Additionally, the demonstrated ability for IL-27^{ACE2pep} to downregulate ACE2 expression in A549-ACE2 cells is significant for the field of oncology and represents substantial potential in translational therapy for tumors overexpressing ACE2.

Future directions include further **optimization** of the protocol for SARS-CoV-2-Spike pseudotyped LV entry assay in order to more accurately determine viral entry and then analysis of **STAT1 activity** following treatment of A549-ACE2 cells.

AUTHOR CONTRIBUTION STATEMENT

JD: Design of the poster and graphical illustrations in Figures 1B, 2A, 3A; contributed and generated data in Figures 2B and 3B
GM: Design of the experiments; generated the graphical illustration in Figure 1A and data in Figure 3C
WS: Design and cloning of the IL-27 plasmids
MLF: Oversaw the project design and execution; contributed funding

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