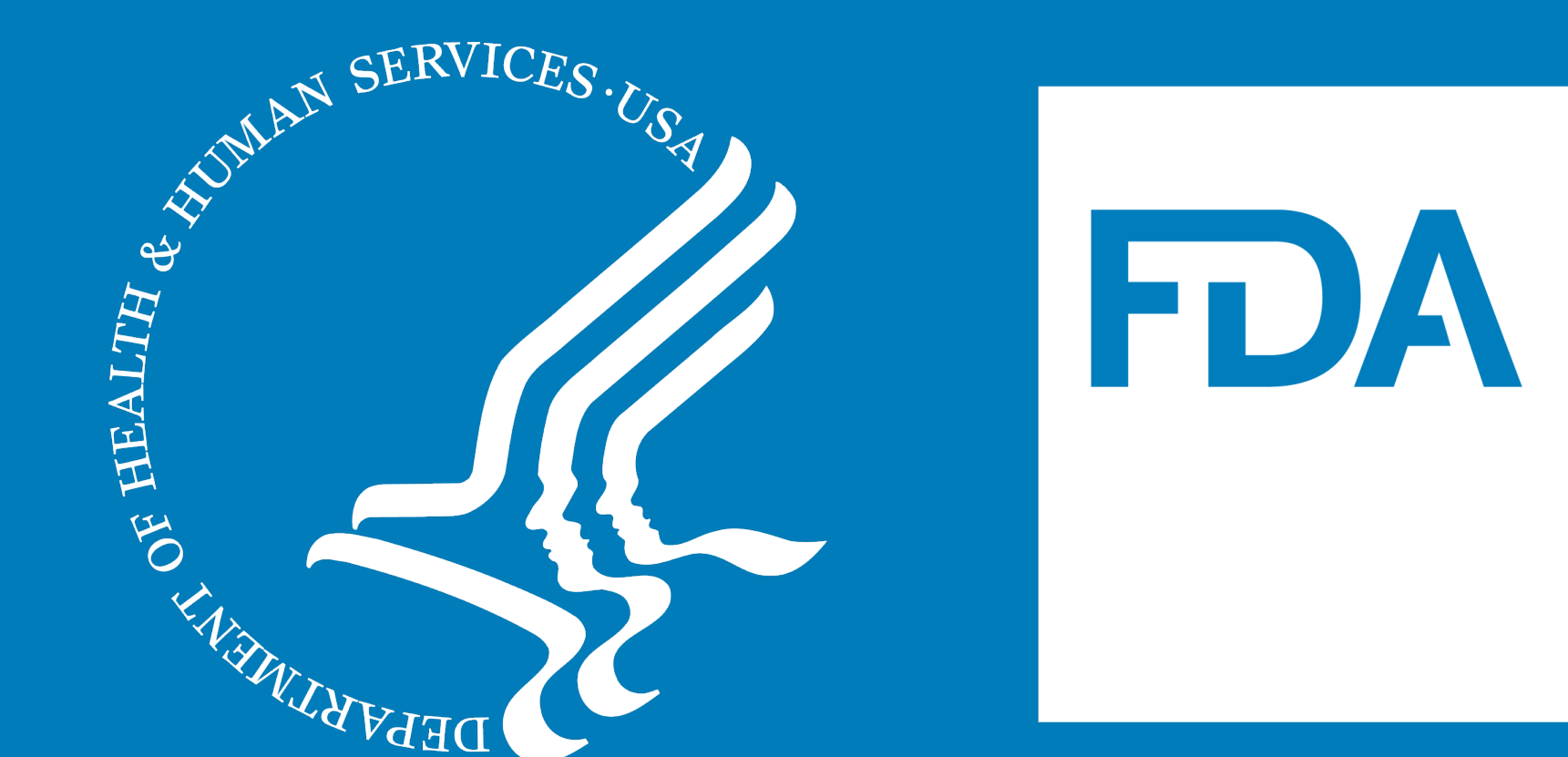


The Transcriptome Landscape of 3D-cultured Placental Trophoblasts Reveals Activation of TLR2 and TLR3/7 in Response to *Trypanosoma cruzi*

Erica Silberstein¹, Charles C. Chung² and Alain Debrabant¹
¹ OBRR/DETTD/LEP; ² CBER/HIVE



Introduction

Chagas Disease and Congenital Transmission

- Chagas disease (CD) is caused by the bloodborne protozoan parasite *Trypanosoma cruzi* (*T. cruzi*).
- CD is transmitted by triatomine bugs, from **mother-to-baby** and by blood transfusion and organ transplantation.
- Congenital infections account for over 20% of new cases worldwide.
- The risk of vertical transmission is 1-5%.
- 40,000 infected women of childbearing age live in the United States, where an estimated 63-315 babies acquire *T. cruzi* infection from their mothers per year.
- The placental syncytiotrophoblasts (SYNs) form an effective physical and immunological barrier against pathogens.
- T. cruzi* parasites circulating in maternal blood can be transmitted to the fetus through the SYNs.

Previous Findings

- We have found that 3D-grown cultures of SYNs exhibit morphological and secretory characteristics comparable to placental trophoblasts (form syncytia and express hormones) and are highly resistant to *T. cruzi* infection. Silberstein et al. *Fron. Microbiol.* 2021; doi:10.3389/fmicb.2021.626370.

The Human Placenta

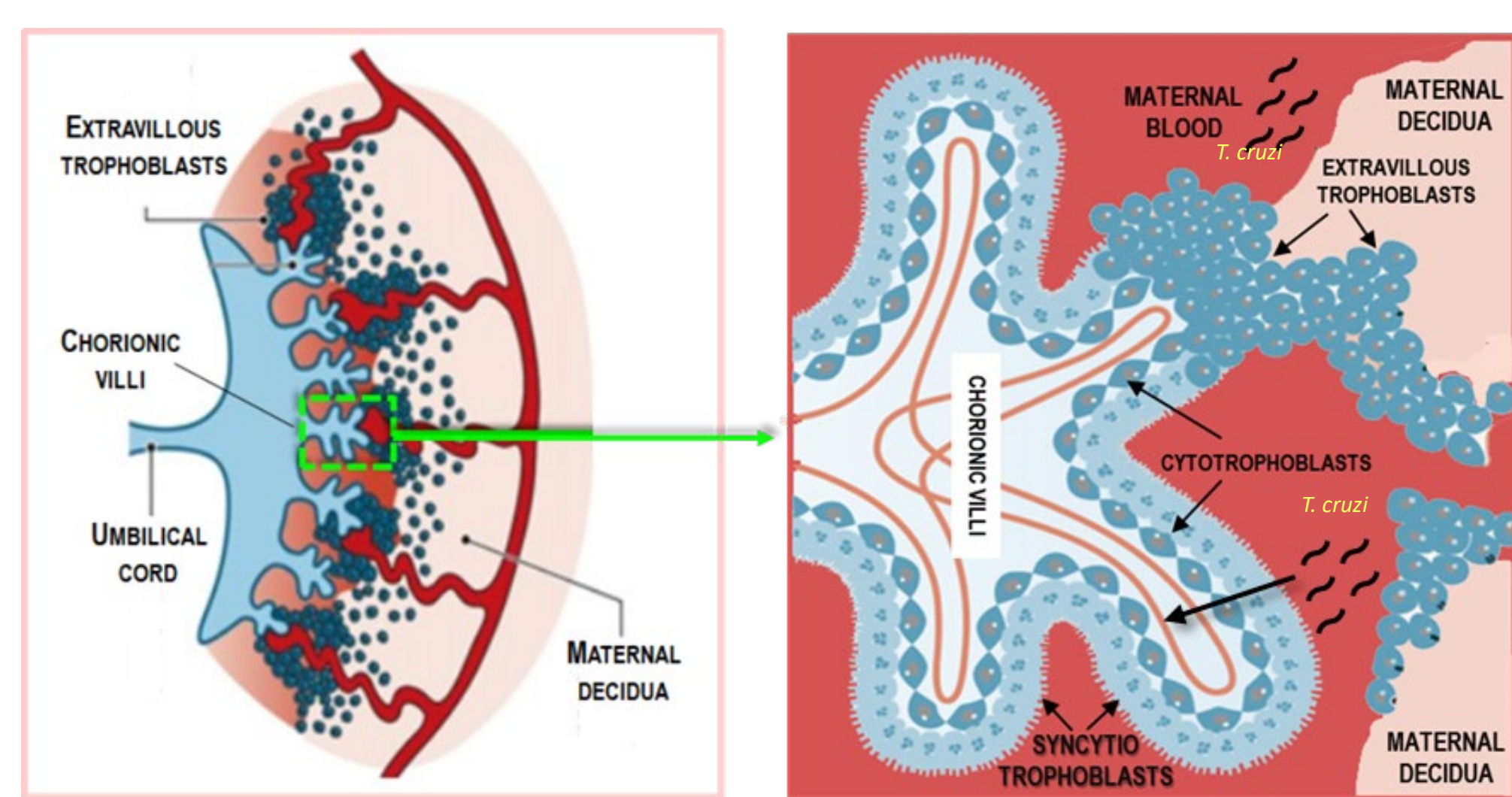


Figure 1. Structure of the human placenta. Adapted from “Immune responses at the maternal-fetal interface”; *Sci Immunol.* 2019; doi: 10.1126/sciimmunol.aat6114.

Objectives

- Identify the defense mechanisms that restrict *T. cruzi* placental infection using a three-dimensional (3D) cell culture system of human syncytiotrophoblasts (SYNs).
- Explore the role of Toll-like receptors in the control of *T. cruzi* infection of the placenta.

Methodology

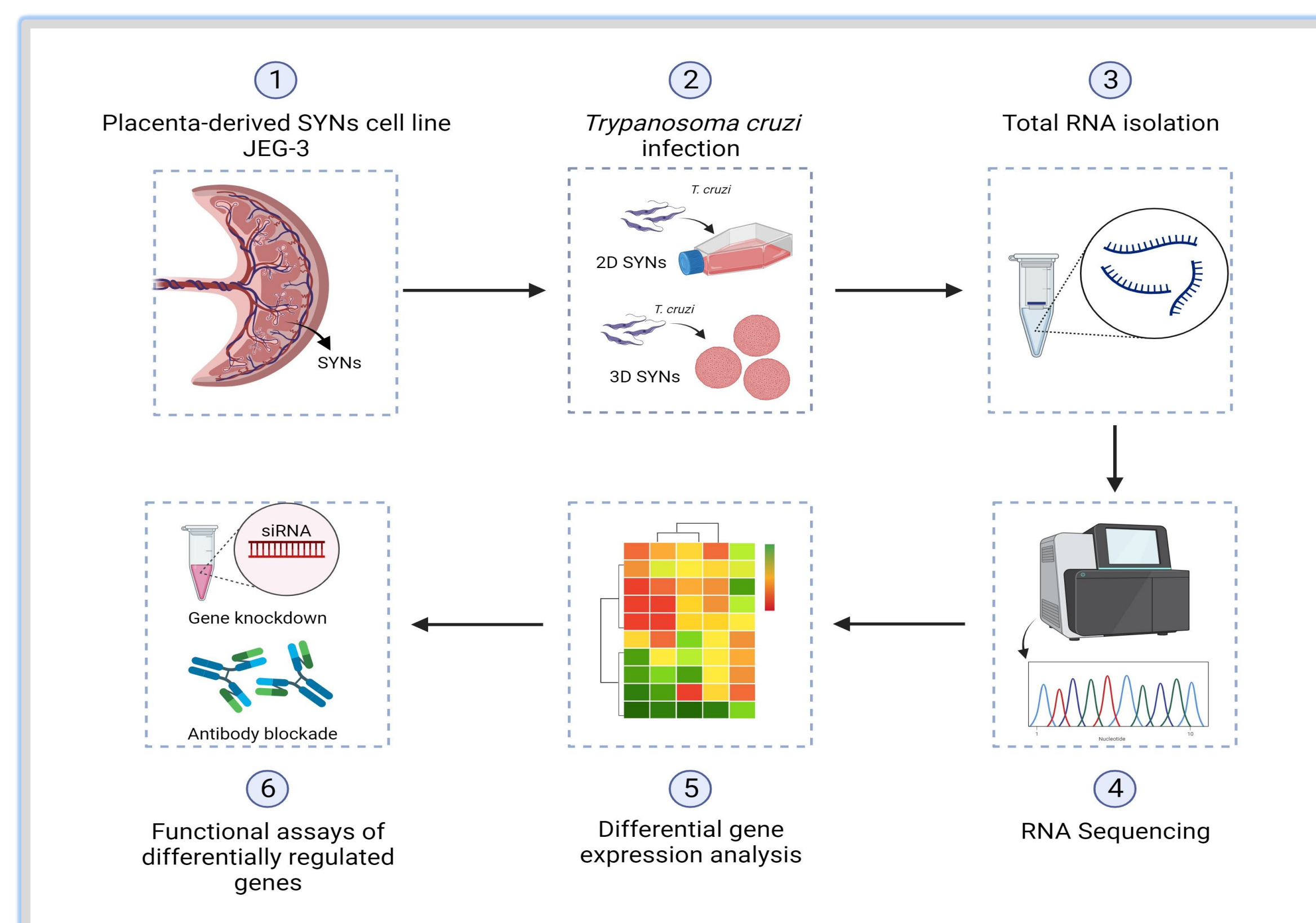


Figure 2. Experimental workflow for the whole transcriptome analysis of 2D and 3D-grown SYNs cultures unexposed or exposed to *T. cruzi*.

Results

Alterations in global gene expression profiles of 3D and 2D-grown SYNs

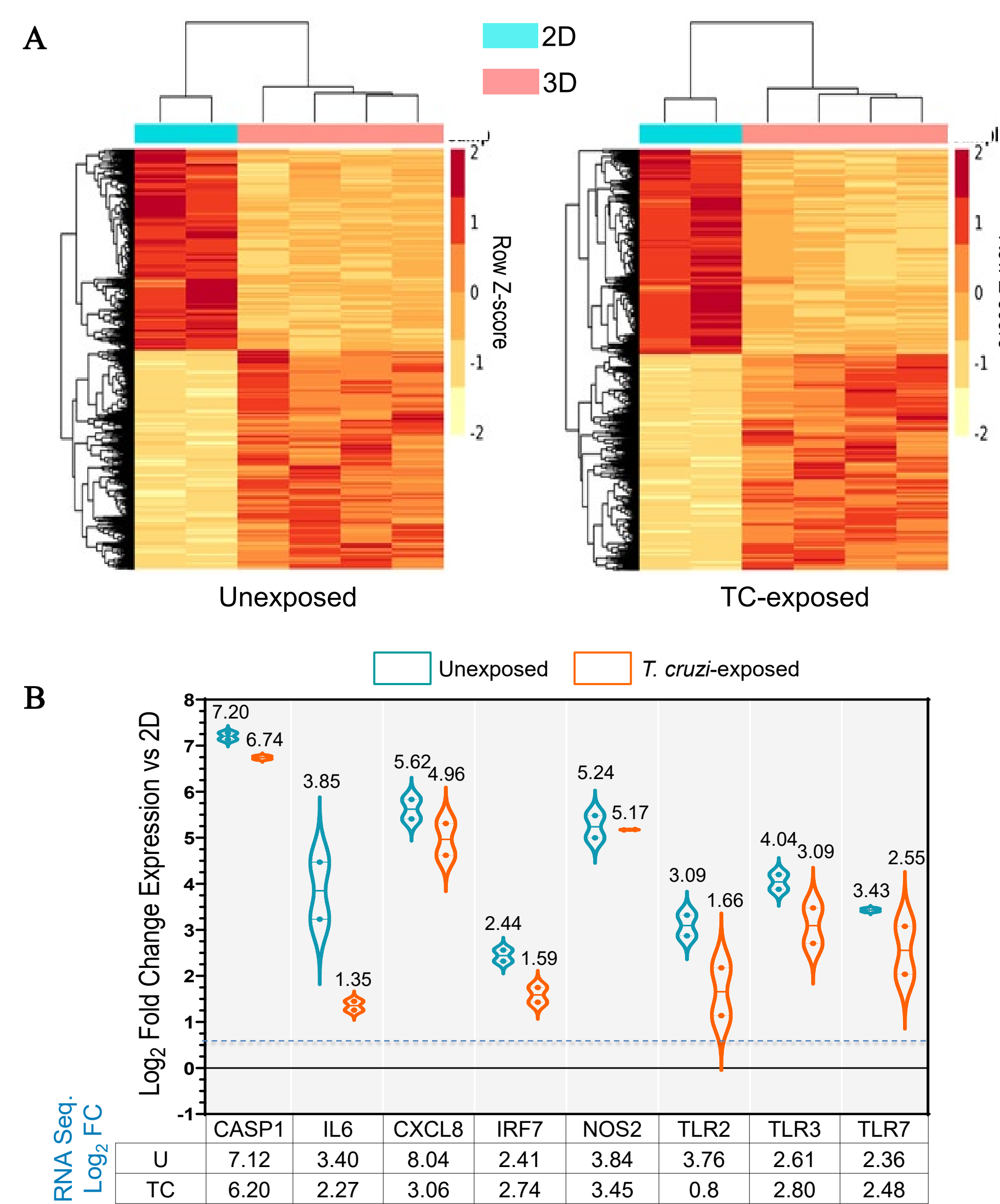


Figure 3. (A) Hierarchical cluster analysis of normalized read counts of differentially expressed genes between 3D and 2D SYNs unexposed or exposed to *T. cruzi*. (B) Validation of RNA-Seq data by qRT-PCR. mRNA quantification of selected genes carried out by qRT-PCR in unexposed or *T. cruzi*-exposed 3D SYNs.

Results

Pathway analysis reveals activation of processes associated with cellular immune defenses, pathogen-influenced signaling and cellular stress/injury in 3D-cultured SYNs

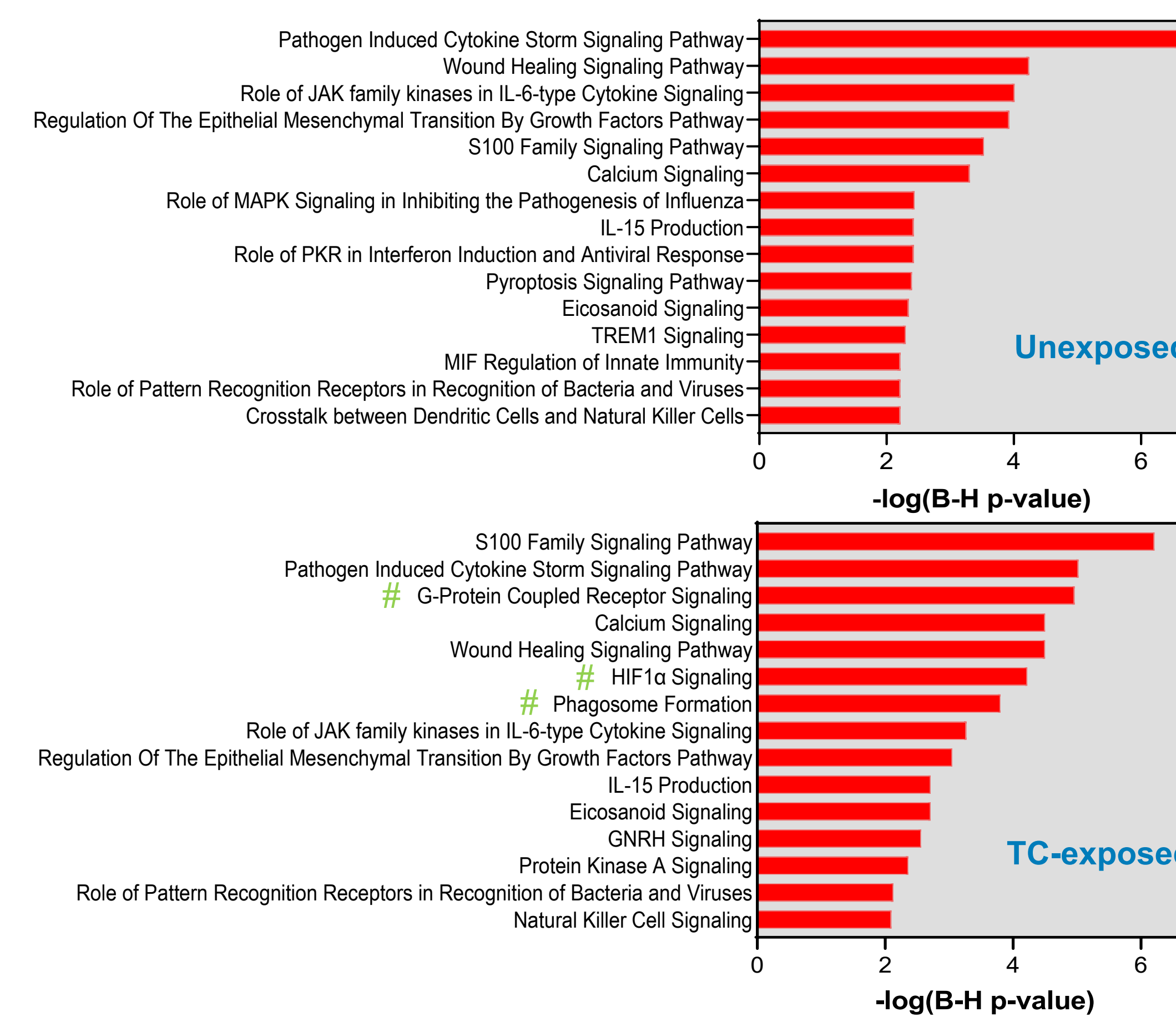


Figure 4. Top significant canonical pathways identified by IPA in 3D SYNs compared to 2D SYNs, in unexposed and *T. cruzi*-exposed cultures. The red-colored bars represent the $-\log(B-H p\text{-value})$ calculated for each pathway. Only pathways with $z\text{-score} \geq 2$ and $B-H p\text{-value} \geq 0.05$ were considered significant.

Toll-like receptors and cytokines are common genes in the top activated canonical pathways

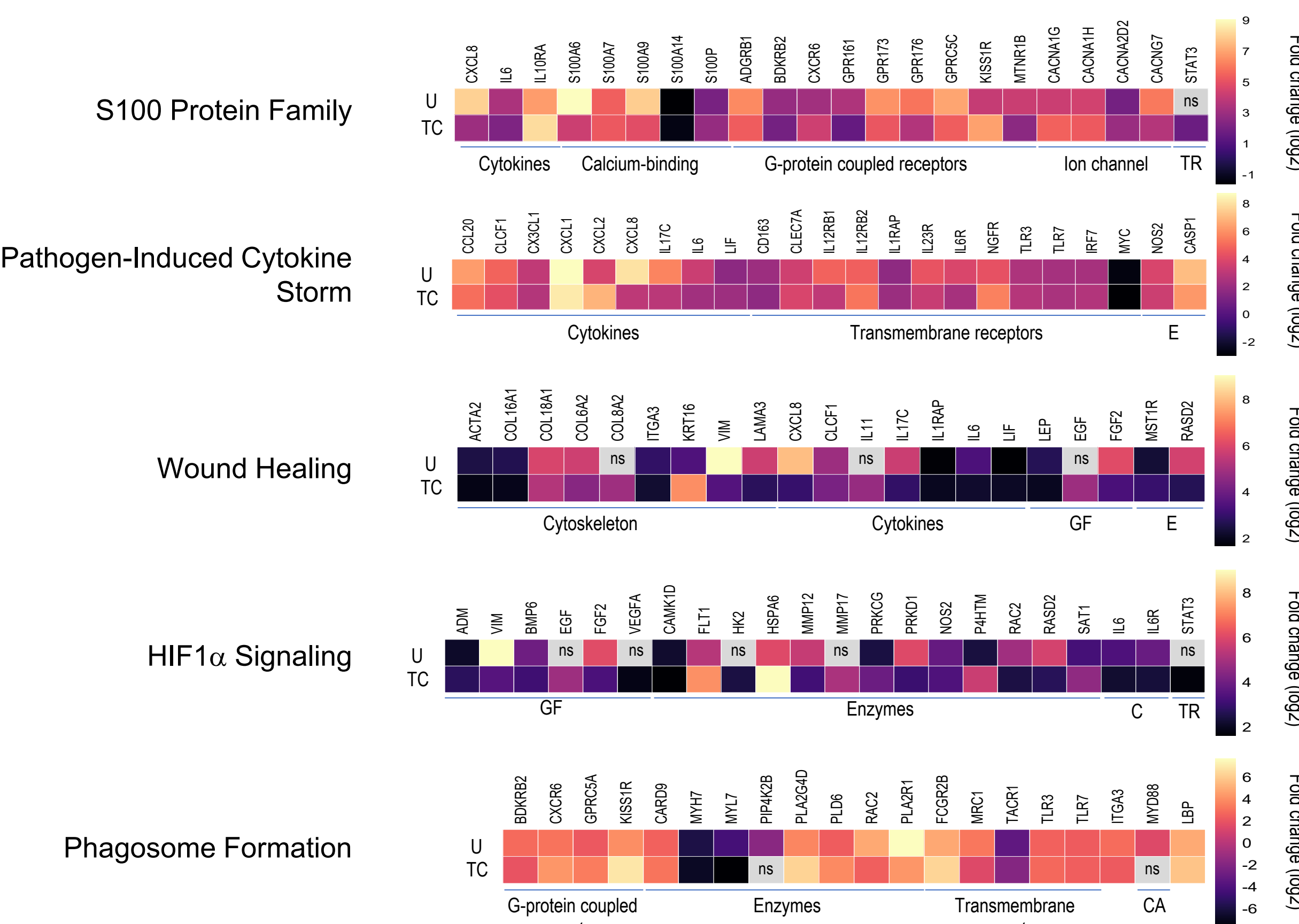


Figure 5. Heatmaps of top differentially expressed genes. Only genes with \log_2 fold change $\geq \pm 1.5$ and $p\text{-value} \geq 0.05$ were considered significant. ns: no significant \log_2 fold change or B-H $p\text{-values}$. U: unexposed; TC: *T. cruzi*-exposed. CA: cytosolic adapter protein; C: cytokines and cytokine receptors; E: enzymes; GF: growth factors; TR: transcription regulators.

Inhibition of TLRs significantly decreases expression of cytokines and interferons promoting *T. cruzi* growth

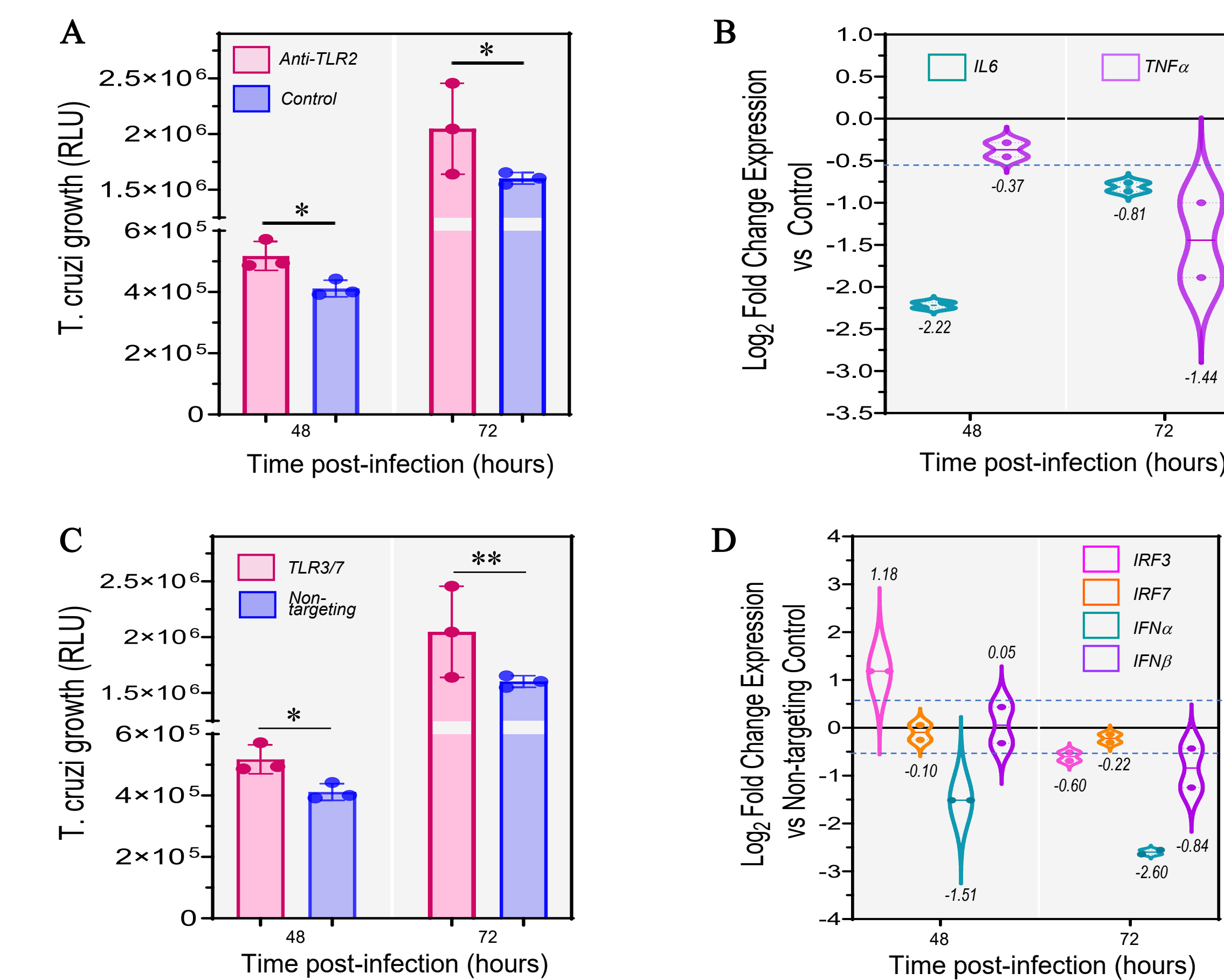
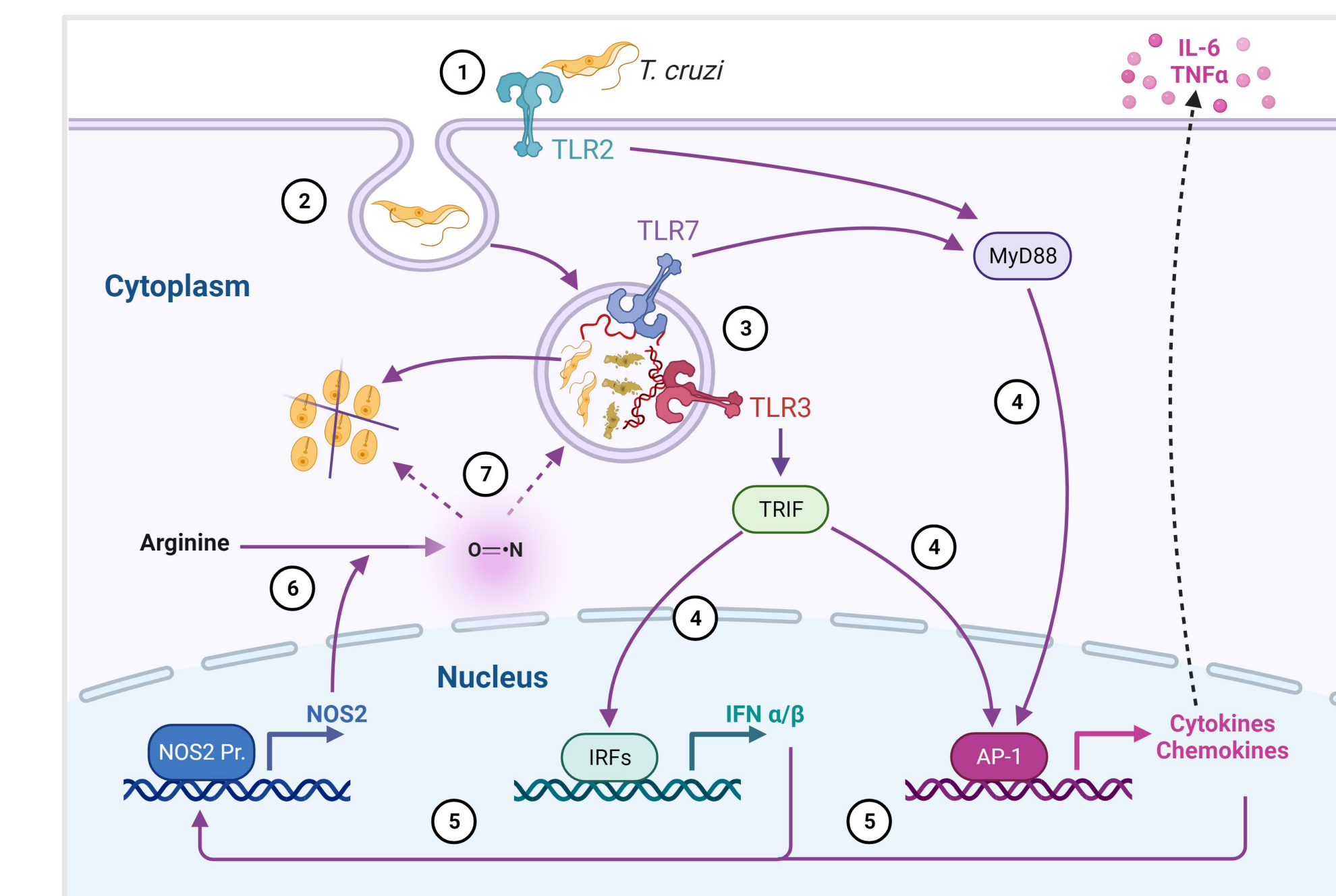


Figure 6. 2D SYNs were treated with anti-hTLR2 (TLR2 blockade) or transfected with TLR3/TLR7 siRNA mix, followed by *T. cruzi* infection. (A) *T. cruzi* growth over time after TLR2 blockade. (B) qRT-PCR mRNA quantification after TLR2 blockade. (C) *T. cruzi* growth over time after TLR3/TLR7 siRNA transfection. (D) qRT-PCR mRNA quantification after TLR3/TLR7 siRNA transfection. * $p \leq 0.05$; ** $p \leq 0.01$.

Model of immune response mechanisms against *T. cruzi* mediated by Toll-like receptors in 3D-grown SYNs



Conclusions

SYNs resist infection by constitutively expressing pro-inflammatory molecules and modulating multiple defense mechanisms that interfere with the parasite's intracellular life cycle, contributing to parasite killing and infection control