

INTRODUCTION

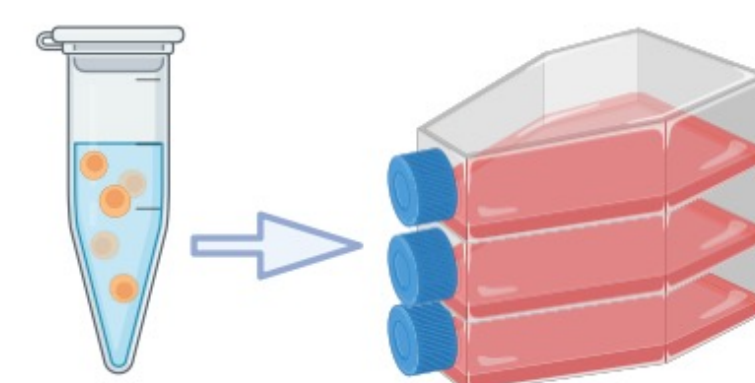
- Beta-blockers are some of the most prescribed drugs today, most notably among 22.3% of the geriatric population in the United States.¹
- These drugs are used to inhibit beta-adrenergic signaling that consequently lower heart rate and contractility, culminating in lower blood pressure.²
- Dopamine has long been shown to be involved in cardiovascular disease, and its receptors in cardiac tissue have been pharmacological targets for congestive heart failure.³
- Our lab has previously highlighted the importance of dopamine receptor 3 (D3R) and its role in fibroblast proliferation during wound healing.⁴
- This project seeks to understand whether there is cross-talk between $\beta 2$ and D1 receptors, as well as any effect on the expression of matrix metalloproteinases (MMPs) and tissue inhibitor metalloproteinases.⁵
- Understanding the interaction between these two signaling pathways can provide deeper insights into the cardiac remodeling process following physiological stress.

MATERIALS & METHODS

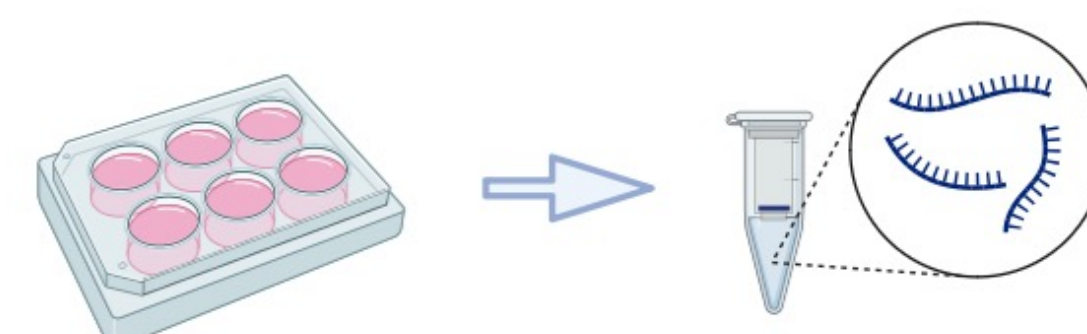
Cell Culture: Cryopreserved human cardiac fibroblasts (hCFB) (C12375) from PromoCell were centrifuged, mixed and suspended in media and serum, incubated at 37°C and 5% CO₂ in cell culture flasks. Cells were cultured from passages 3-7. Treatments were designed using dopamine (DA), D1R antagonist SCH 39166 hydrobromide (SCH), D3R agonist pramipexole dihydrochloride (PPX), and $\beta 2$ agonist isoproterenol.

Real Time quantitative PCR: RNA was extracted and isolated from control and treatment groups for subsequent analysis with RT-qPCR. Control groups with dopamine were used as the standard of comparison for statistical analysis.

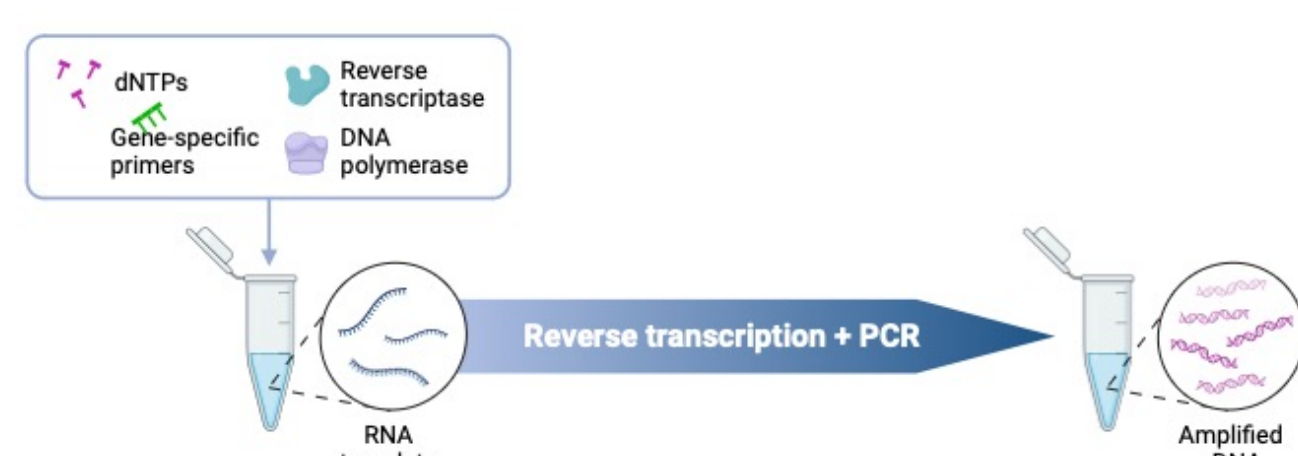
Phase 1: hCFb Culture



Phase 2: RNA Extraction



Phase 3: cDNA Synthesis



Phase 4: Gene Expression Quantification via RT-qPCR

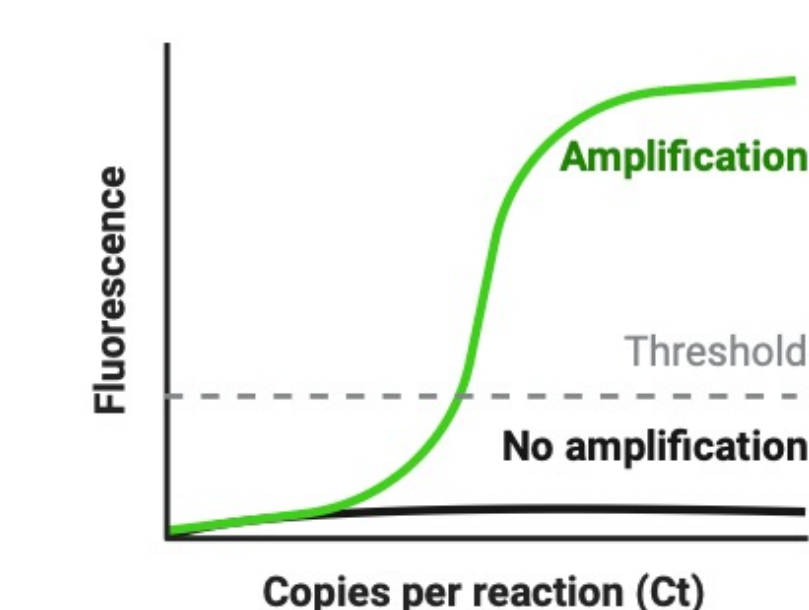


Figure 1. Schematic of experimental design.

RESULTS

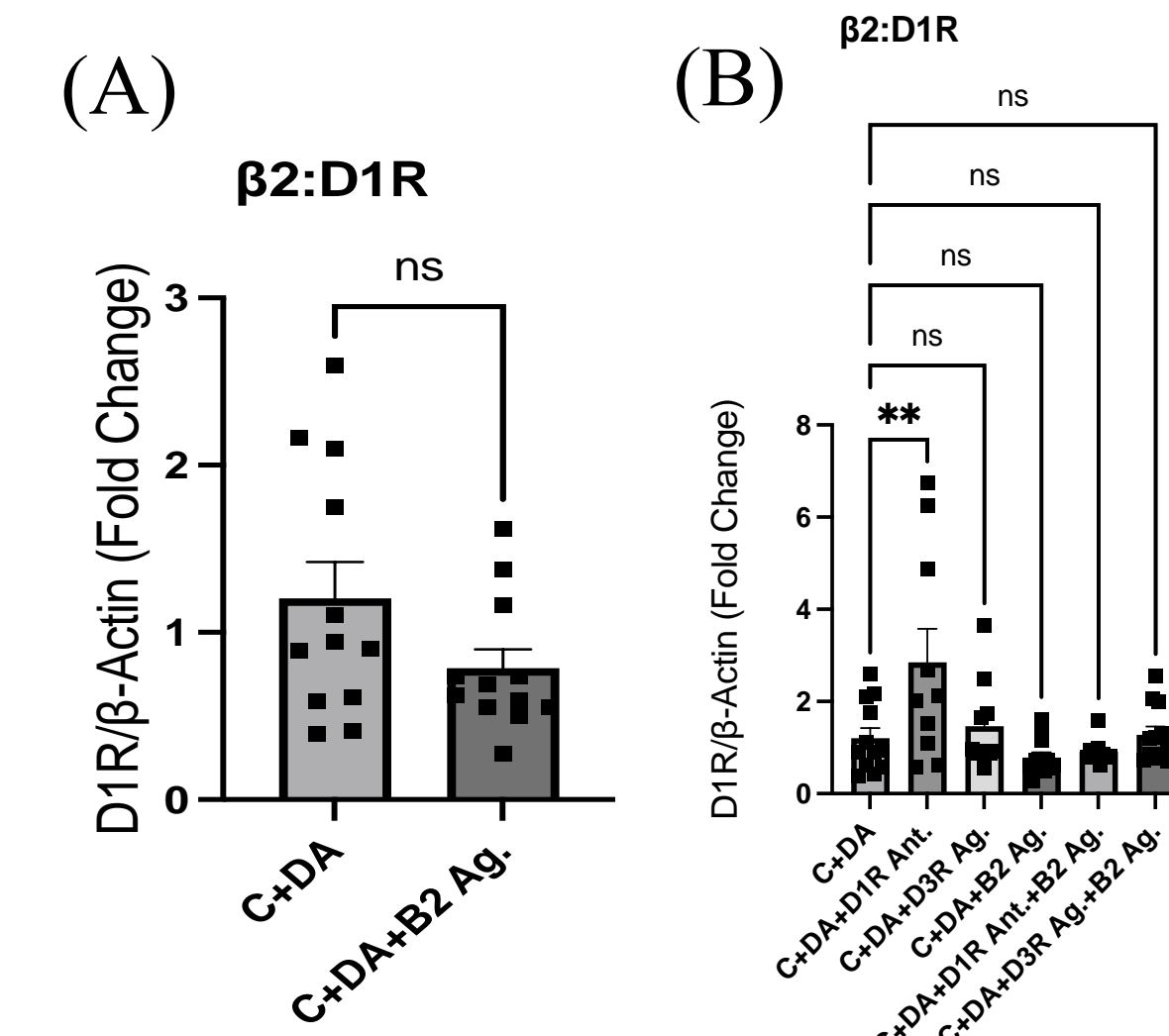


Figure 2: $\beta 2$ stimulation on D1R gene expression ($\beta 2$:D1R). (A) Comparing the gene expression of D1R between control+dopamine and $\beta 2$ agonist. $\beta 2$ agonist had no significant alterations in D1R gene expression. (B) SCH treatment increased expression of D1R (* $p < 0.01$). Other treatment groups showed no significant differences as compared to the control.

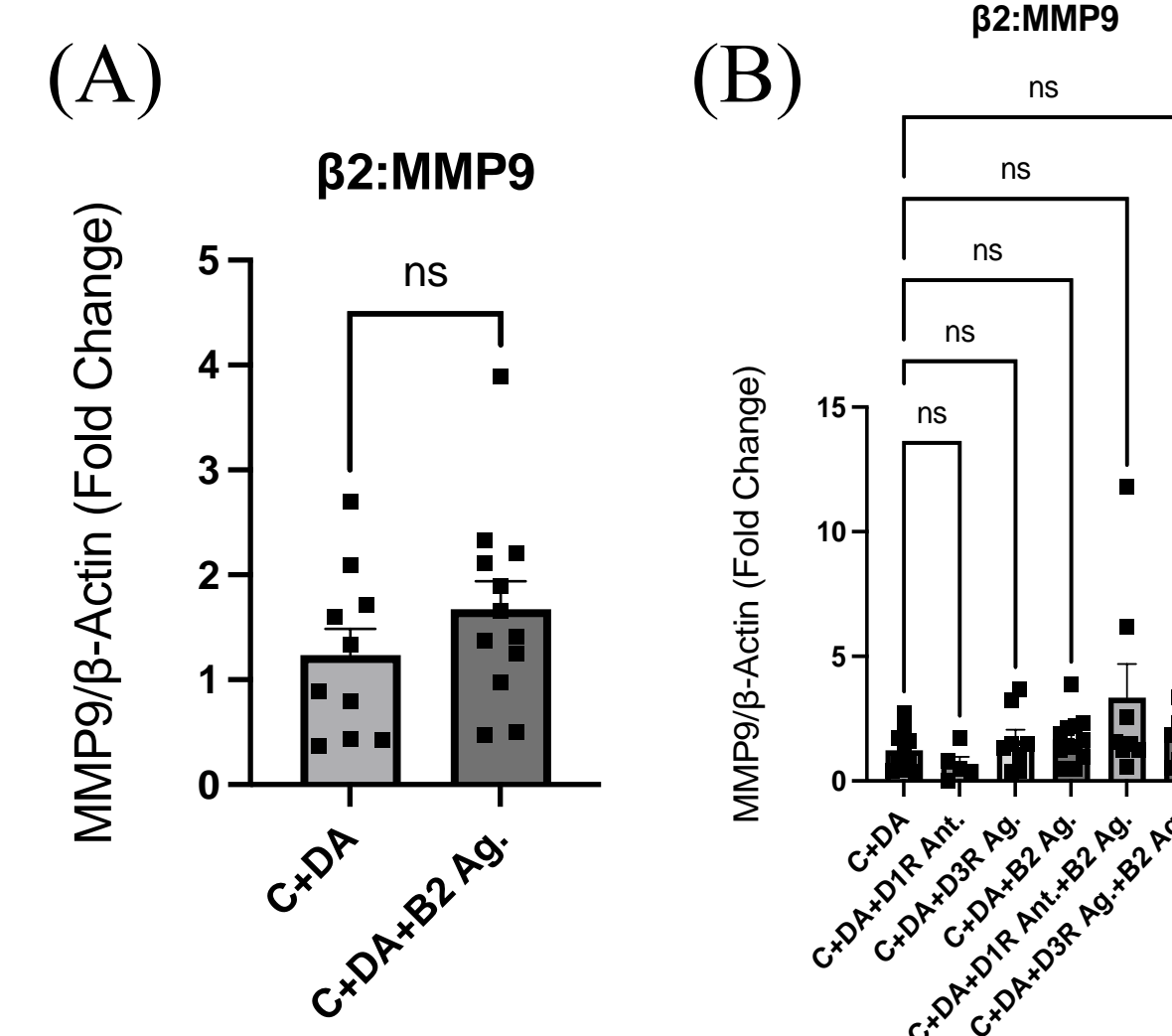


Figure 4: $\beta 2$ stimulation on MMP9 gene expression ($\beta 2$:MMP9). (A) Comparing the gene expression of MMP9 between control+dopamine and $\beta 2$ agonist. $\beta 2$ agonist had no significant decrease in MMP9 gene expression. (B) Treatment groups showed no significant differences as compared to the control.

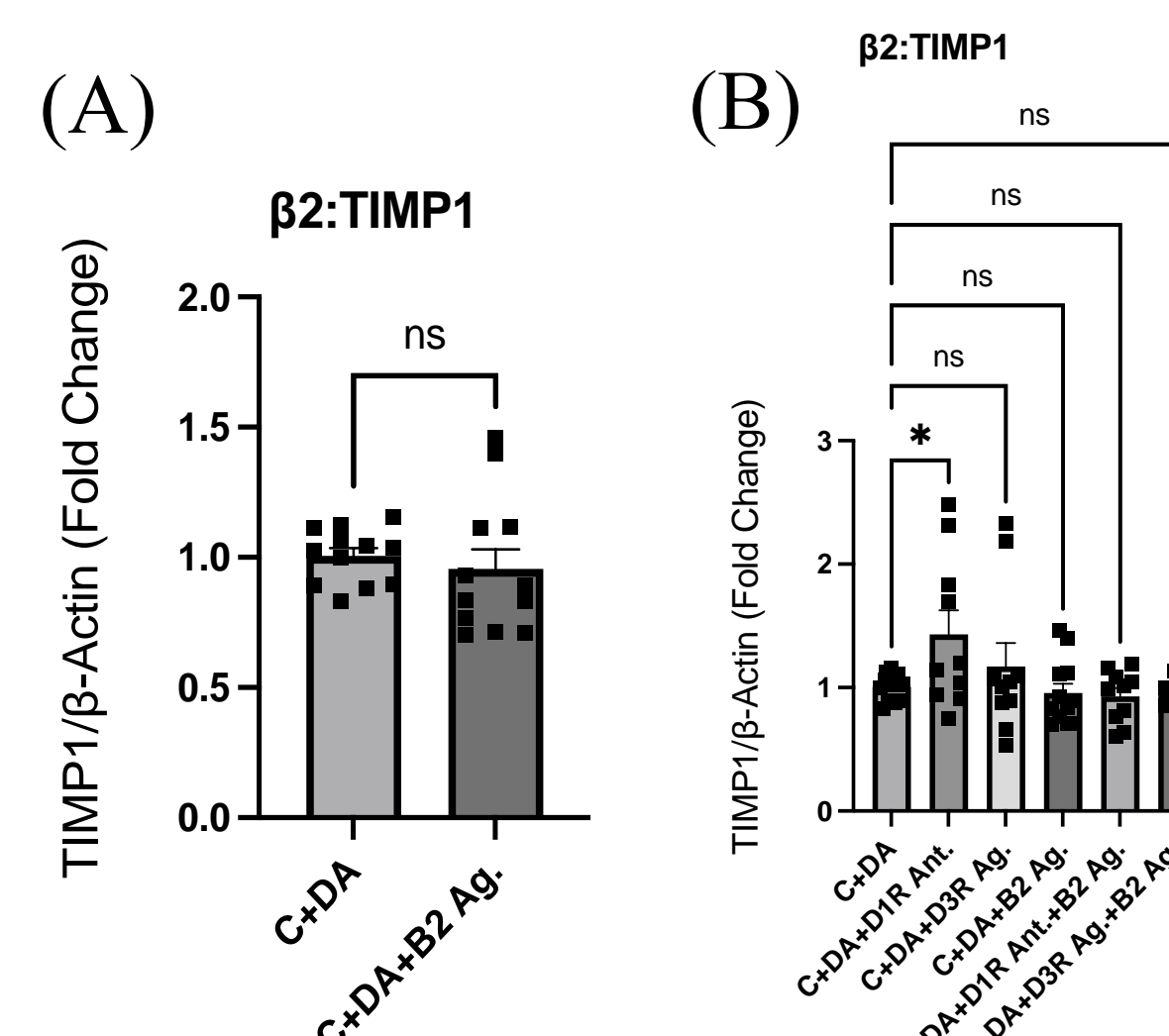


Figure 6: $\beta 2$ stimulation on TIMP1 gene expression ($\beta 2$:TIMP1). (A) Comparing the gene expression of TIMP1 between control+dopamine and $\beta 2$ agonist. $\beta 2$ agonist had no significant decrease in TIMP1 gene expression. (B) SCH treatment showed increased expression of TIMP1 (* $p < 0.05$). Other treatment groups showed no significant differences as compared to the control.

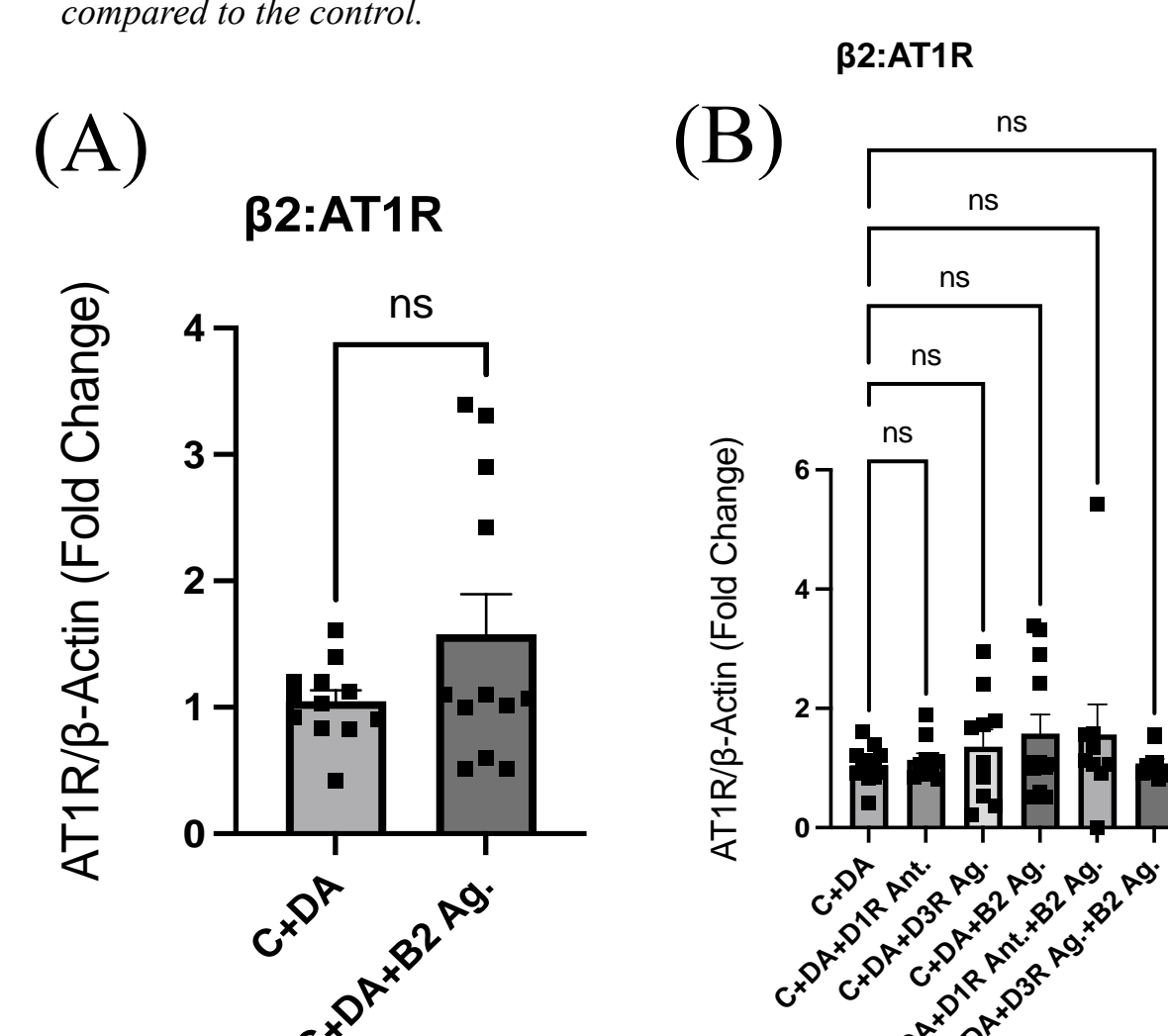


Figure 8: $\beta 2$ stimulation on AT1R gene expression ($\beta 2$:AT1R). (A) Comparing the gene expression of AT1R between control+dopamine and $\beta 2$ agonist. $\beta 2$ agonist had no significant decrease in AT1R gene expression. (B) Treatment groups showed no significant differences as compared to the control.

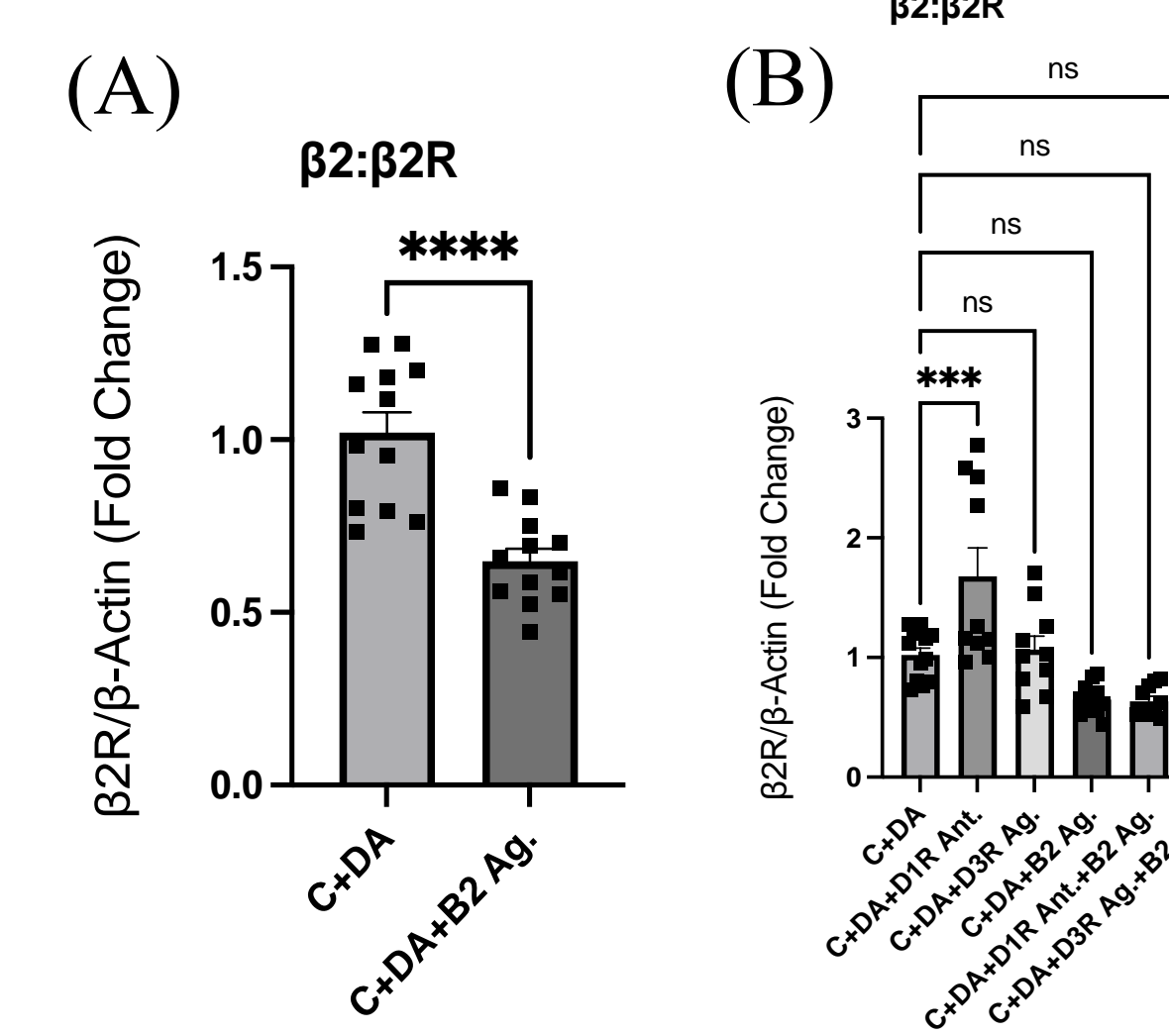


Figure 3: $\beta 2$ stimulation on $\beta 2R$ gene expression ($\beta 2$: $\beta 2R$). (A) Comparing the gene expression of $\beta 2R$ between control+dopamine and $\beta 2$ agonist. $\beta 2$ agonist had significant decrease in $\beta 2R$ gene expression (* $p < 0.0001$). (B) SCH treatment increased expression of $\beta 2R$ (* $p < 0.001$). Other treatment groups showed no significant differences as compared to the control.

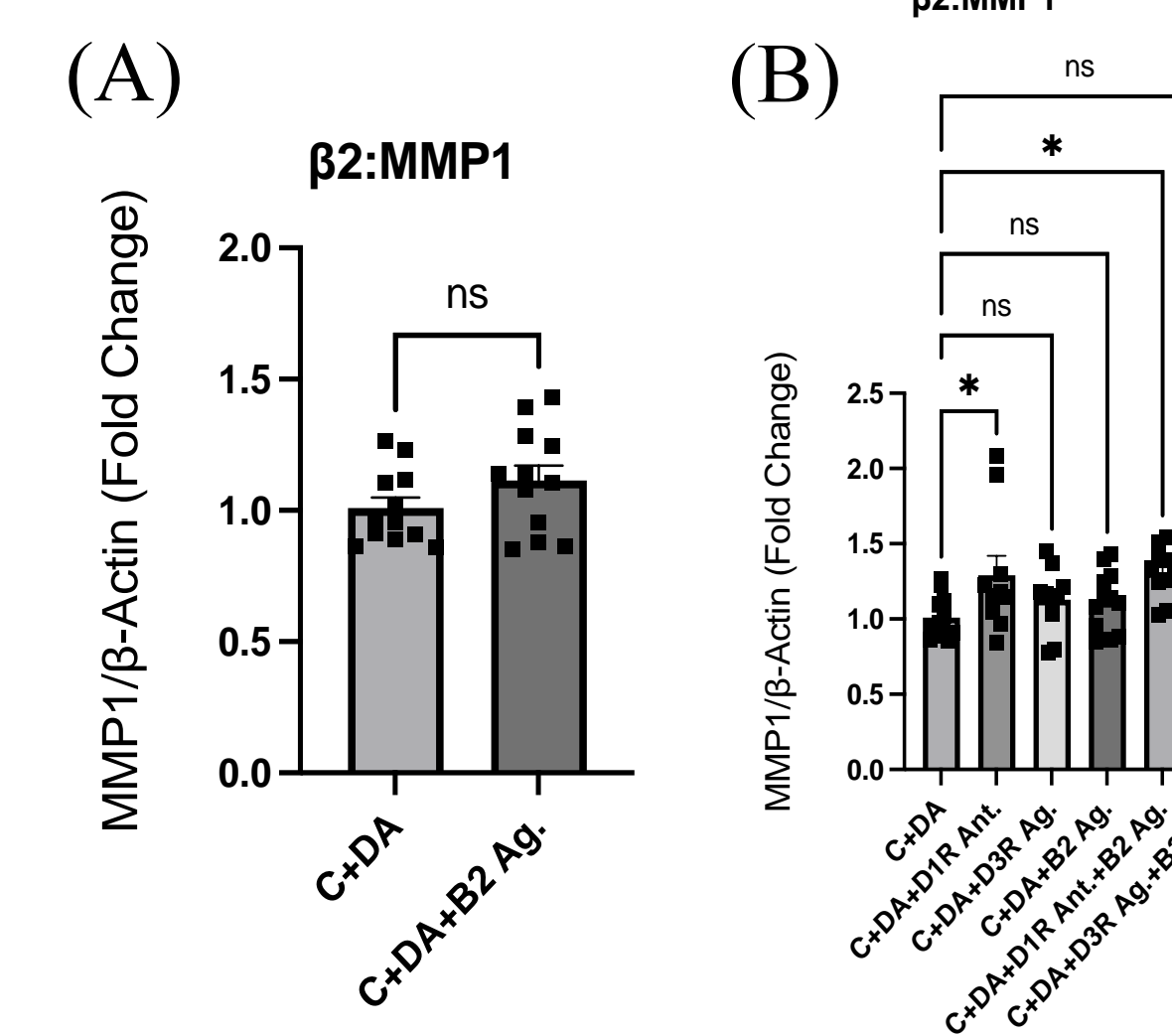


Figure 5: $\beta 2$ stimulation on MMP1 gene expression ($\beta 2$:MMP1). (A) Comparing the gene expression of MMP1 between control+dopamine and $\beta 2$ agonist. $\beta 2$ agonist had no significant decrease in MMP1 gene expression. (B) SCH treatment and SCH+ $\beta 2$ agonist both increased expression of MMP1 (* $p < 0.05$). Other treatment groups showed no significant differences as compared to the control.

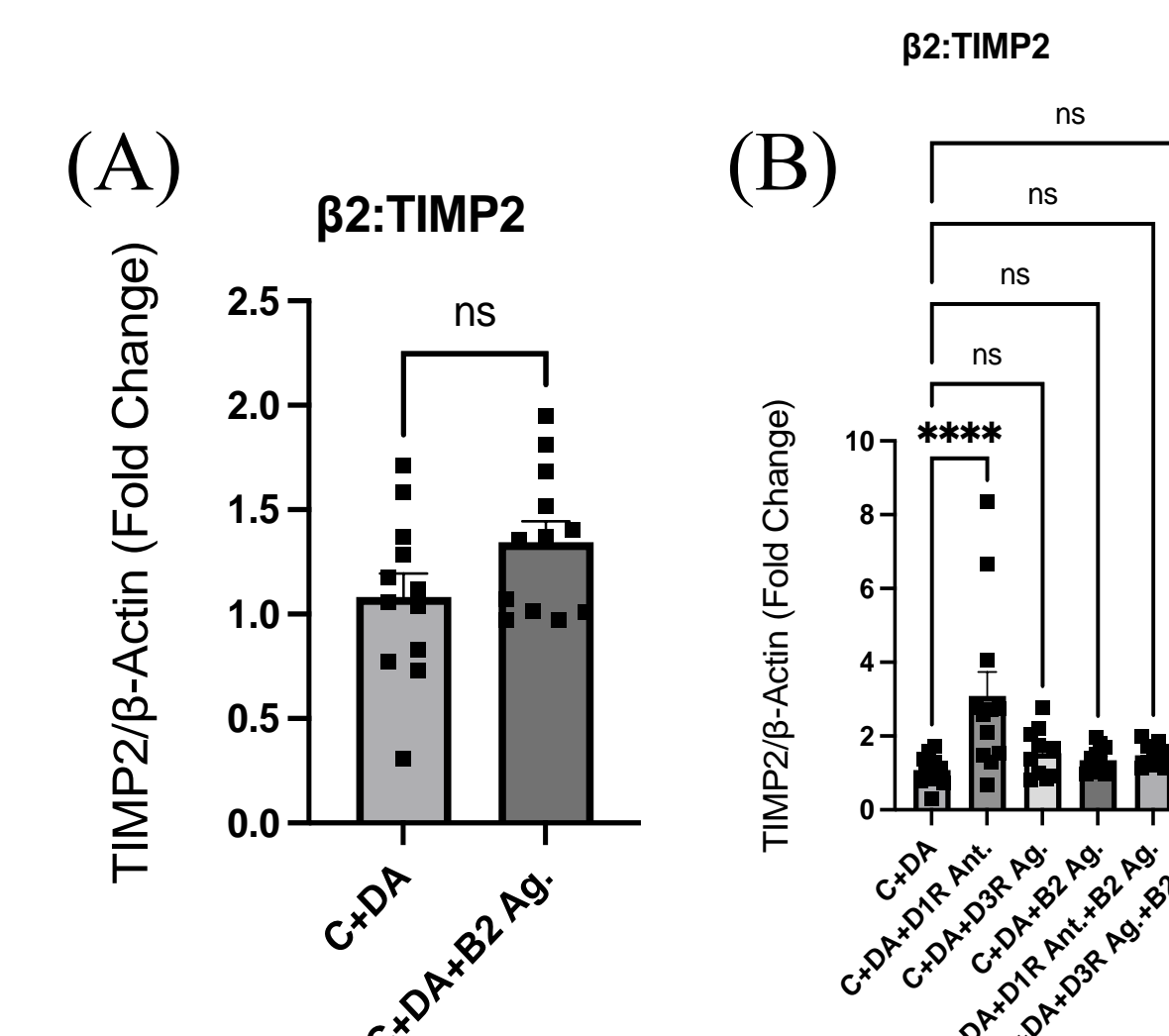


Figure 7: $\beta 2$ stimulation on TIMP2 gene expression ($\beta 2$:TIMP2). (A) Comparing the gene expression of TIMP2 between control+dopamine and $\beta 2$ agonist. $\beta 2$ agonist had no significant decrease in TIMP2 gene expression. (B) SCH treatment showed increased expression of TIMP2 (* $p < 0.0001$). Other treatment groups showed no significant differences as compared to the control.

Summary of Results:

- Presence of dopamine lead to $\beta 2$ agonist decreasing expression of $\beta 2R$.
- $\beta 2$ agonist showed marginal increase in MMP1 expression when in combination with D1R antagonist, as compared to D1R antagonist alone.
- D1R antagonist increased expression of D1R, $\beta 2R$, MMP1, TIMP1, and TIMP2 genes.

DISCUSSION

- $\beta 2$ levels decreased under a negative feedback system when exposed to isoproterenol, potentially serving as a mechanism preventing tachycardia.
- D1R antagonist has been shown to behave similarly to angiotensin II and $\beta 2$ regarding hypertensive activity. This experiment showed D1R antagonist increases levels of $\beta 2$ to maintain contractility properties, to which its antagonistic properties might help prevent ventricular arrhythmia. This suggests some cross-talk between the two receptors.
- Tissue inhibitor metalloproteinases (TIMPs) were upregulated by D1R antagonist. D1R antagonist's influence on MMP1 upregulation was insignificantly increased by the presence of isoproterenol.
- D1R antagonist is a profound stimulator of multiple genes involved in the structure, function, and maintenance of the cardiac extracellular matrix.**

FUTURE DIRECTIONS

- Repeat RT-qPCR experiments to confirm results and resolve discrepancies in data.
- Conduct enzyme-linked immunosorbent assays (ELISA) to determine protein expression of MMP1 and MMP9.
- Further postulate signaling pathways of SCH involving upregulation of MMPs and TIMPs, as well as potential cross-talk with other adrenergic receptors.

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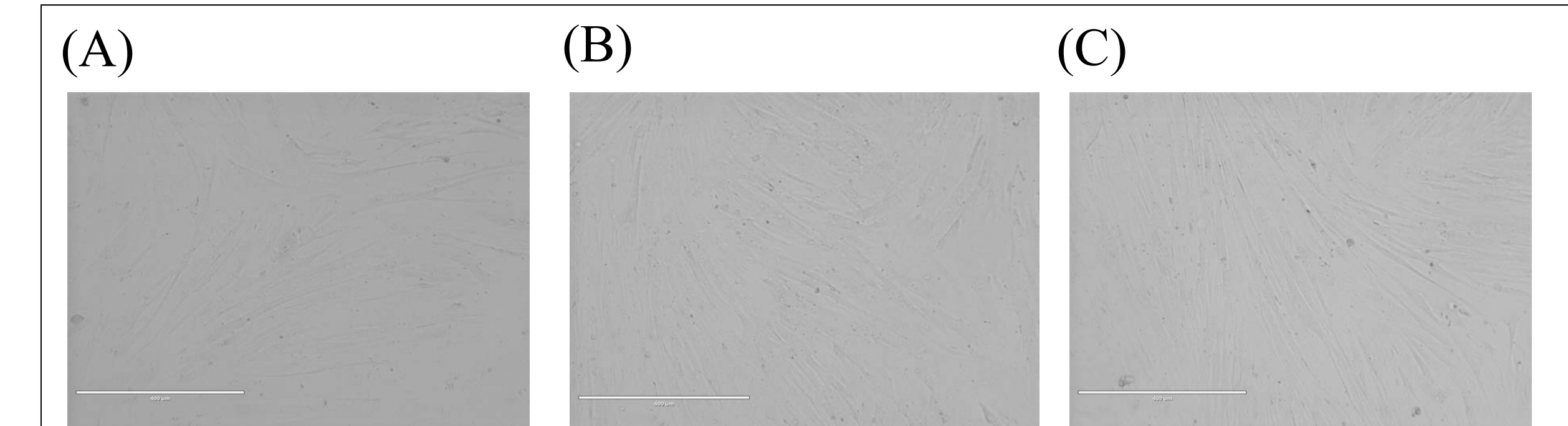


Figure 9. Human cardiac fibroblasts, passage 7.