

Supplementary Materials

Transforming Growth Factor β 1 Modulates Sex Differences in Cardiac Myofibroblast Activation on Hydrogel Biomaterials

Authors:

Mason N. Faust^{1,2†}, Ashley K. Nguyen^{1,2†}, Rayyan M. Gorashi^{1,2}, Nicole E. Félix Vélez^{1,2}, Meaghan C. Loud^{1,2}, and Brian A. Aguado^{1,2,3*}

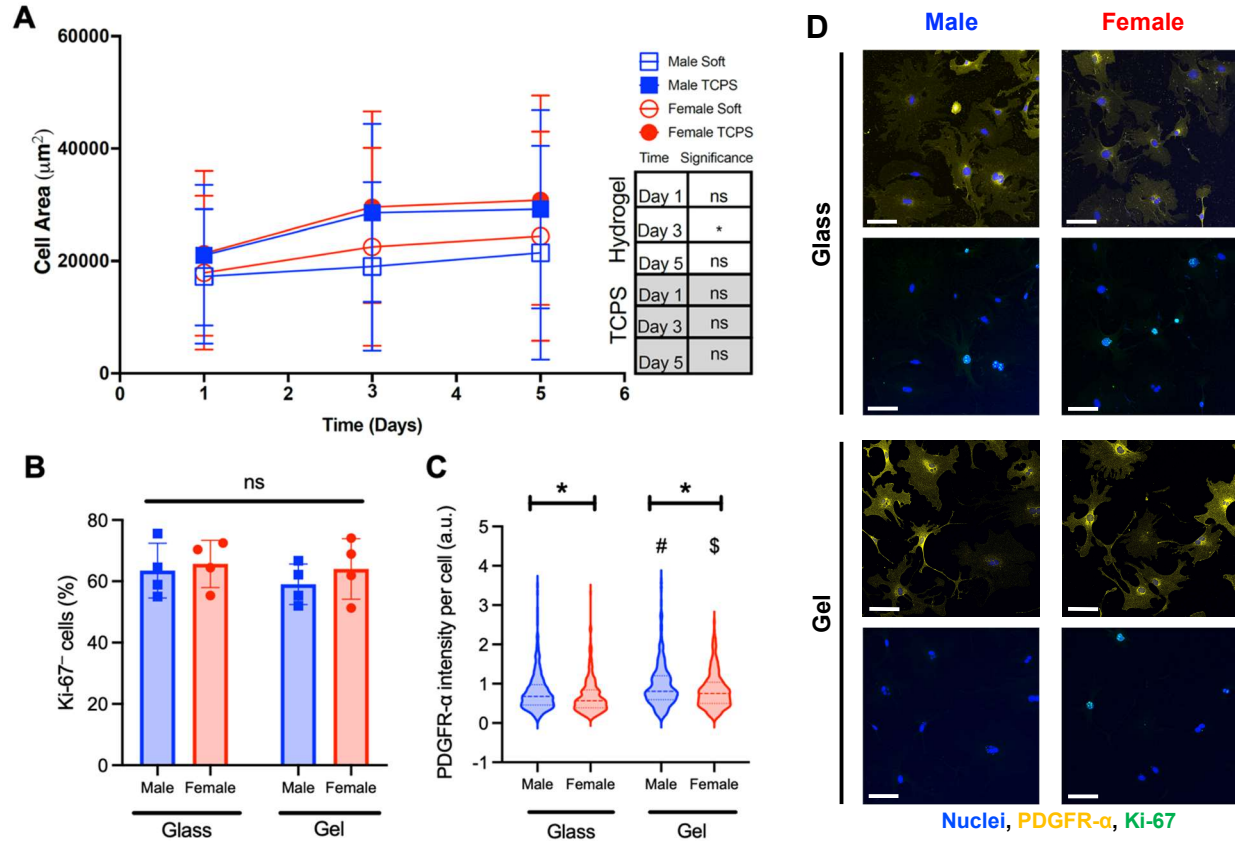
[†]These authors contributed equally.

Affiliations:

1. Shu Chien-Gene Lay Department of Bioengineering, University of California San Diego, La Jolla, CA 92093, USA.
2. Sanford Consortium for Regenerative Medicine, La Jolla, CA 92037, USA.
3. Program in Materials Science and Engineering, University of California San Diego, La Jolla, CA 92093, USA

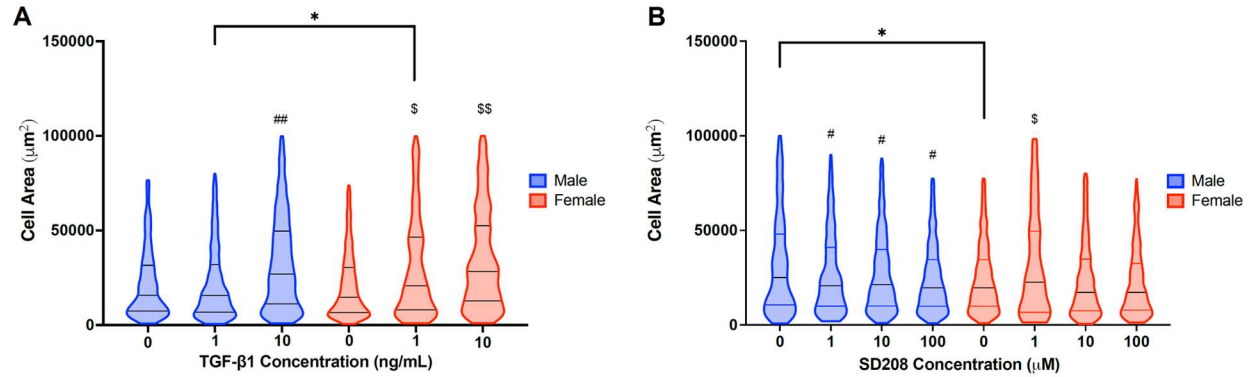
*Address correspondence to:

Brian A. Aguado, Ph.D.
2880 Torrey Pines Scenic Drive
University of California, San Diego
La Jolla, CA 92037
Email: baguado@ucsd.edu



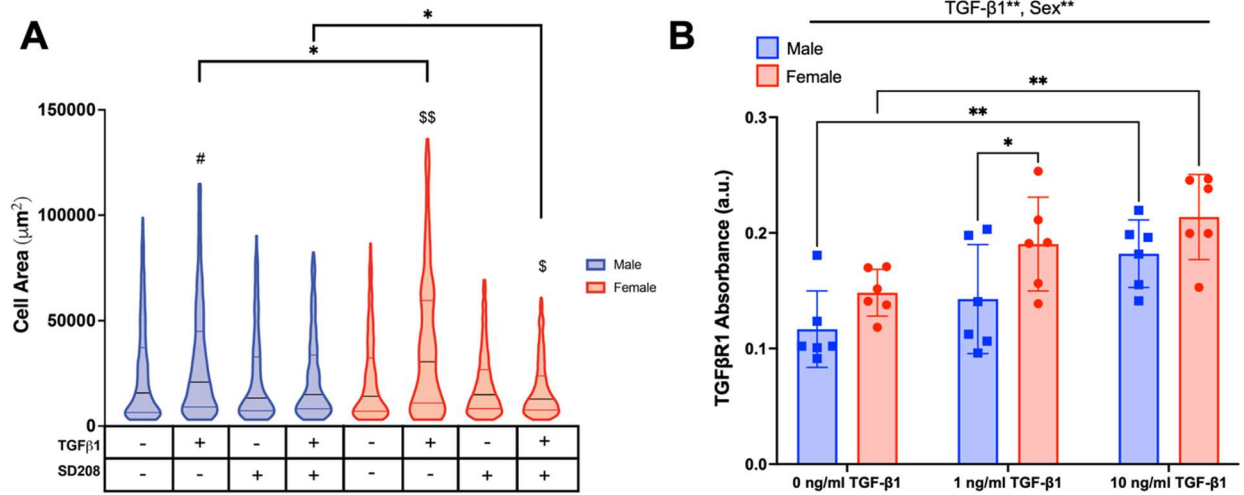
Supplementary Figure 1: Phenotype of cardiac fibroblasts cultured on hydrogels or TCPS.

(A) Quantification of cell area in male and female CFs cultured on hydrogels and TCPS at various time points. N=3 hydrogels, mean $\pm$ S.D. shown. Significance was determined using two-way nonparametric ANOVA ($P < 0.05$) and a Cohen's d test (*d > 0.2), denoting effect size between sex in the table provided. (B) Quantification of Ki-67⁺ cells (%) for male and female cardiac fibroblasts cultured on glass and hydrogels. N=4 gels, n>196 cells per gel, mean $\pm$ S.D. shown. Significance was determined using two-way ANOVA with Tukey's post-hoc test. (C) Quantification of PDGFR- α mean intensity per cell for male and female cardiac fibroblasts cultured on glass and hydrogels. N=4 hydrogels, n>196 cells per gel. Statistical significance was determined using two-way ANOVA ($P < 0.05$) with Tukey's post-hoc test and effect size determined using Cohen's d test (*d > 0.2) denoting effect size between sex. Hashtag indicates effect size in male group relative to glass control (#d > 0.2); dollar sign indicates effect size in female group relative to glass control (\$d > 0.2). (D) Representative immunofluorescence images of PDGFR- α (yellow) and Ki-67 (green) in male and female cardiac fibroblasts cultured on glass and hydrogels. Scale is 100 μm .

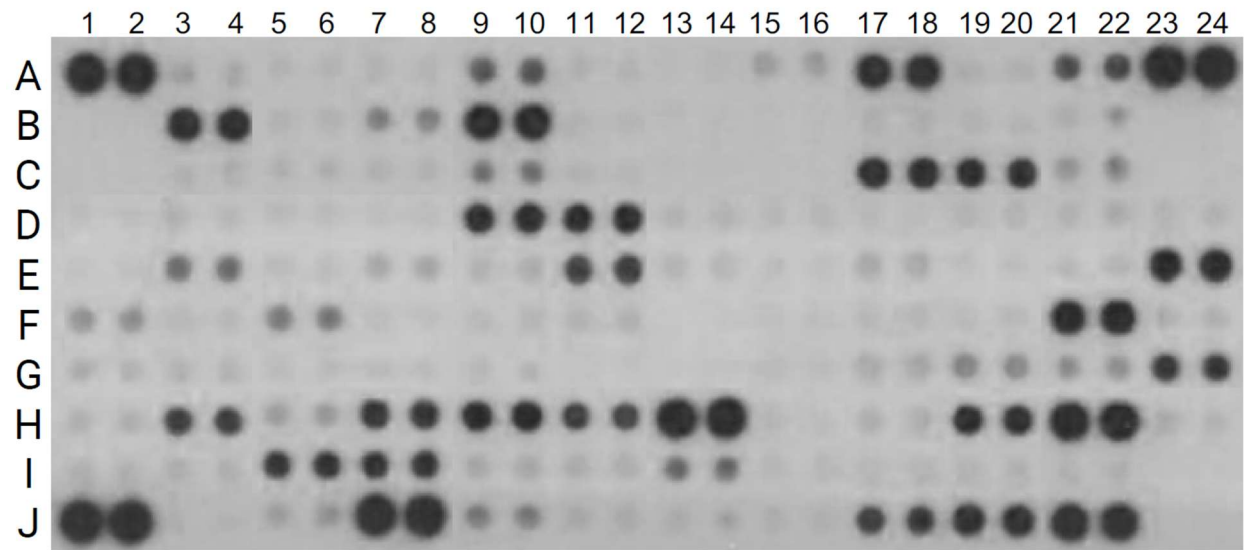


Supplementary Figure 2: Cell area for cardiac fibroblasts treated with TGF β -1 or SD208.

Quantification of cell area in male and female CFs cultured on hydrogels and TCPS when treated with either (A) 0, 1, and 10 ng/mL TGF β -1 or (B) with 0, 1, 10, and 100 μM of SD208. N=3 hydrogels, n>432 cells, mean $\pm$ S.D. shown. Significance was determined using two-way nonparametric ANOVA ($P < 0.05$) and a Cohen's d test (*d > 0.2, **d > 0.5, ***d > 0.8, and ****d > 1.4) denoting effect size between sex; asterisk indicates effect size in male group (#d > 0.2, ##d > 0.5, ###d > 0.8, and ####d > 1.4) denoting effect size relative to male control; dollar sign indicates effect size in female group (\$d > 0.2, \$\$d > 0.5, \$\$\$d > 0.8, and \$\$\$\$d > 1.4) denoting effect size relative to female control.



Supplementary Figure 3: Cell area for cardiac fibroblasts treated with combined media. (A) Quantification of cell area for male and female cardiac fibroblasts cultured on hydrogels TGF-β1 and SD208. N=3 hydrogels, mean ± S.D. shown. Significance was determined using two-way nonparametric ANOVA ($P < 0.05$) and a Cohen's d test (* $d > 0.2$, ** $d > 0.5$, *** $d > 0.8$, and **** $d > 1.4$) denoting effect size between sex; asterisk indicates effect size in male group (# $d > 0.2$, ## $d > 0.5$, ### $d > 0.8$, and #### $d > 1.4$) denoting effect size relative to male control; dollar sign indicates effect size in female group (\$ $d > 0.2$, \$\$ $d > 0.5$, \$\$\$ $d > 0.8$, and \$\$\$\$ $d > 1.4$) denoting effect size relative to female control. **(B)** Quantification of TGFβ1 protein expression in male and female cardiac fibroblasts were treated with 0, 1, or 10 ng/mL TGF-β1 via in-cell ELISA. n=6 per group, mean ± S.D. shown. Significance was determined using two-way ANOVA ($P < 0.05$) with Tukey's post-hoc test. Main effects of TGF-β1 treatment ($P = 0.0011$) and sex ($P = 0.0092$) were significant.



Supplementary Figure 4: Sample Cytokine Array Blot. Row/column IDs for each cytokine spot is provided in Supplementary Table 2.

Supplementary Table 1: RT-qPCR Primer Sequences

Gene Name	Primer Sequence	Direction
<i>B2M</i>	GGTCTTTCTGGTGCTTGTCT	Forward
	ACGTAGCAGTTCAGTATGTTTCG	Reverse
<i>COL1A1</i>	TGGTGAAGCAGGCAAGC	Forward
	GAAACCTCTCTCGCCTCTTG	Reverse
<i>FN</i>	GAGCTATCCATTTACCTTCAGA	Forward
	TTGTTCGTAGACACTGGAGAC	Reverse
<i>ACTA2</i>	TCAGCGTTCAGCCTCC	Forward
	CCAGAGCCATTGTCGC	Reverse

Supplementary Table 2: Cytokine Array Protein Targets

Spot(s)	Protein Name	Spot(s)	Protein Name	Spot(s)	Protein Name
A1, A2	Reference Spots	D13, D14	DKK-1	G17, G18	IL-27 p28
A3, A4	Adiponectin/Acrp30	D15, D16	DPPIV/CD26	G19, G20	IL-28A/B
A5, A6	Amphiregulin	D17, D18	EGF	G21, G22	IL-33
A7, A8	Angiopoietin-1	D19, D20	Endoglin/CD105	G23, G24	LDL R
A9, A10	Angiopoietin-2	D21, D22	Endostatin	H1, H2	Leptin
A11, A12	Angiopoietin-like 3	D23, D24	Fetuin A/AHSG	H3, H4	LIF
A13, A14	BAFF/BLyS/TNFSF13B	E1, E2	FGF acidic	H5, H6	Lipocalin-2/NGAL
A15, A16	C1q R1/CD93	E3, E4	FGF-21	H7, H8	LIX
A17, A18	CCL2/JE/MCP-1	E5, E6	Flt-3 Ligand	H9, H10	M-CSF
A19, A20	CCL3/CCL4/MIP-1 α/β	E7, E8	Gas 6	H11, H12	MMP-2
A21, A22	CCL5/RANTES	E9, E10	G-CSF	H13, H14	MMP-3
A23, A24	Reference Spots	E11, E12	GDF-15	H15, H16	MMP-9
B3, B4	CCL6/C10	E13, E14	GM-CSF	H17, H18	Myeloperoxidase
B5, B6	CCL11/Eotaxin	E15, E16	HGF	H19, H20	Osteopontin (OPN)
B7, B8	CCL12/MCP-5	E17, E18	ICAM-1/CD54	H21, H22	Osteoprotegerin/TNFRSF11B
B9, B10	CCL17/TARC	E19, E20	IFN- γ	H23, H24	PD-ECGF/Thymidine phosphorylase
B11, B12	CCL19/MIP-3 β	E21, E22	IGFBP-1	I1, I2	PDGF-BB
B13, B14	CCL20/MIP-3 α	E23, E24	IGFBP-2	I3, I4	Pentraxin 2/SAP
B15, B16	CCL21/6Ckine	F1, F2	IGFBP-3	I5, I6	Pentraxin 3/TSG-14
B17, B18	CCL22/MDC	F3, F4	IGFBP-5	I7, I8	Periostin/OSF-2
B19, B20	CD14	F5, F6	IGFBP-6	I9, I10	Pref-1/DLK-1/FA1
B21, B22	CD40/TNFRSF5	F7, F8	IL-1 α /IL-1F1	I11, I12	Proliferin
C3, C4	CD160	F9, F10	IL-1 β /IL-1F2	I13, I14	Proprotein Convertase 9/PCSK9
C5, C6	Chemerin	F11, F12	IL-1ra/IL-1F3	I15, I16	RAGE
C7, C8	Chitinase 3-like 1	F13, F14	IL-2	I17, I18	RBP4

C9, C10	Coagulation Factor III/Tissue Factor	F15, F16	IL-3	I19, I20	Reg3G
C11, C12	Complement Component C5/C5a	F17, F18	IL-4	I21, I22	Resistin
C13, C14	Complement Factor D	F19, F20	IL-5	J1, J2	Reference Spots
C15, C16	C-Reactive Protein/CRP	F21, F22	IL-6	J3, J4	E-Selectin/CD62E
C17, C18	CX3CL1/Fractalkine	F23, F24	IL-7	J5, J6	P-Selectin/CD62P
C19, C20	CXCL1/KC	G1, G2	IL-10	J7, J8	Serpin E1/PAI-1
C21, C22	CXCL2/MIP-2	G3, G4	IL-11	J9, J10	Serpin F1/PEDF
D1, D2	CXCL9/MIG	G5, G6	IL-12 p40	J11, J12	Thrombopoietin
D3, D4	CXCL10/IP-10	G7, G8	IL-13	J13, J14	TIM-1/KIM-1/HAVCR
D5, D6	CXCL11/I-TAC	G9, G10	IL-15	J15, J16	TNF- α
D7, D8	CXCL13/BLC/BCA-1	G11, G12	IL-17A	J17, J18	VCAM-1/CD106
D9, D10	CXCL16	G13, G14	IL-22	J19, J20	VEGF
D11, D12	Cystatin C	G15, G16	IL-23	J21, J22	WISP-1/CCN4