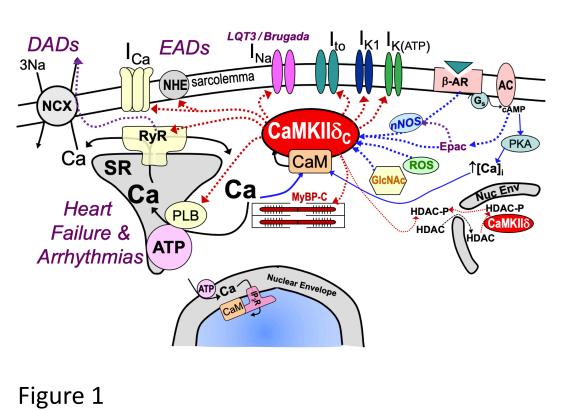
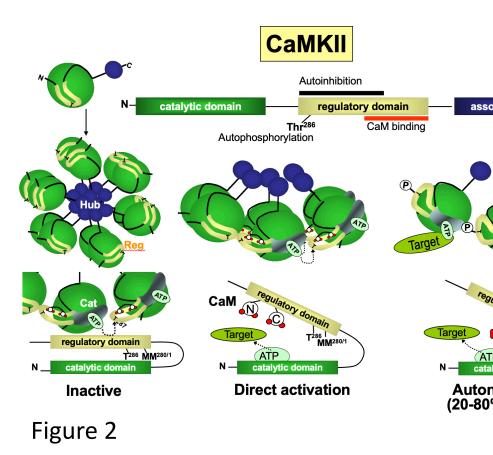


The role of calmodulin-dependent protein kinase II nitrosylation (CaMKII) in ischemia-reperfusion injury Brandon Weiss¹, Kenneth S. Ginsburg², Adam Wilder², Juliana Mira Hernandez², Julie Bossuyt², Donald M. Bers²

Introduction

- CaMKIId is an important regulator of cardiac function and dysfunction in pathological states. It critically modulates ion channels, Ca handling proteins (phospholamban/SERCA, RyR), myofilaments, nuclear transcription and energetics/metabolism. (Fig1) [1]
- CaMKII holoenzyme becomes activated by binding of Ca-CaM, resulting in autophosphorylation of neighboring subunits at T286 and an autonomously active kinase. Other CaMKII targets include T17 on phospholamban and S2814 on RyR (Fig2) [1]
- in addition to autophosphorylation, other post-translational modifications were recently identified that also promote autonomous CaMKII activation: oxidation (M281/282), O-GlcNAcylation (S280) and S-nitrosylation (C290). Interestingly CaMKIId has a second regulatory nitrosylation site (C273) that inhibits instead of activates the kinase. [2]
- PTMs that promote autonomous activity are implicated in cardiac pathology but little is known about the interplay between these PTMs. Oxidized CaMKII contributes to apoptosis post-MI and atrial fibrillation and O-GlcNAcylation contributes to hyperglycemia-induces SR Ca leak and arrhythmia. The role of CaMKII Snitrosylation has yet to be investigated.
- Previous studies have suggested a key role for CaMKII and nitrosylation in IR injury [3]
- This study will look at the role of nitrosylation at both regulatory sites on CaMKII: C290 and C273S.





Hypothesis

Nitrosylation at the C290 site is detrimental to recovery from acute IR injury, while nitrosylation at the C273 site improves recovery from IR injury.

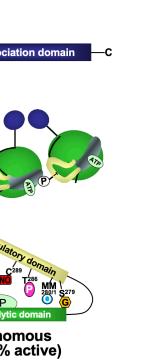
Materials & Methods

Phase I: Data Collection

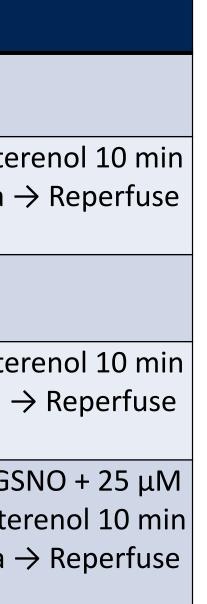
- Utilized both wild type and novel C290A knock-in mice resistant to nitrosylation at the C290 site
- Hearts excised from mice, aorta cannulated for Langendorff perfusion. Each heart was subjected to one of the 6 following experimental protocols:

Group	Experimental Protocol
WT Control	Hearts hung on rig and perfused for 90 mins NO ISCHEMIC EVENT
WT IR	Hearts hung & stabilized on rig for 20 mins \rightarrow Add 10 nM isoproterenol 10 min prior to global ischemia \rightarrow initiate 20 minutes of global ischemia \rightarrow Reperfuse for 40 minutes
C290A Control	Hearts hung on rig and perfused for 90 mins NO ISCHEMIC EVENT
C290A IR	Hearts hung & stabilized on rig for 20 mins \rightarrow Add 10 nM isoproterenol 10 min prior to global ischemia \rightarrow Initiate 20 minutes of global ischemia \rightarrow Reperfuse for 40 minutes
WT GSNO + EGTA	Hearts hung & stabilized on rig for 20 mins \rightarrow Add 100-150 μ M GSNO + 25 μ M EGTA 17.5 minutes prior to global ischemia \rightarrow Add 10 nM isoproterenol 10 min prior to global ischemia \rightarrow Initiate 20 minutes of global ischemia \rightarrow Reperfuse for 40 minutes
WT L-NAME	Hearts hung & stabilized on rig for 20 mins → Add 1 nM L-NAME 17.5 minutes prior to global ischemia → Add 10 nM isoproterenol 10 min prior to global ischemia →Initiate 20 minutes of global ischemia → Reperfuse for 40 minutes

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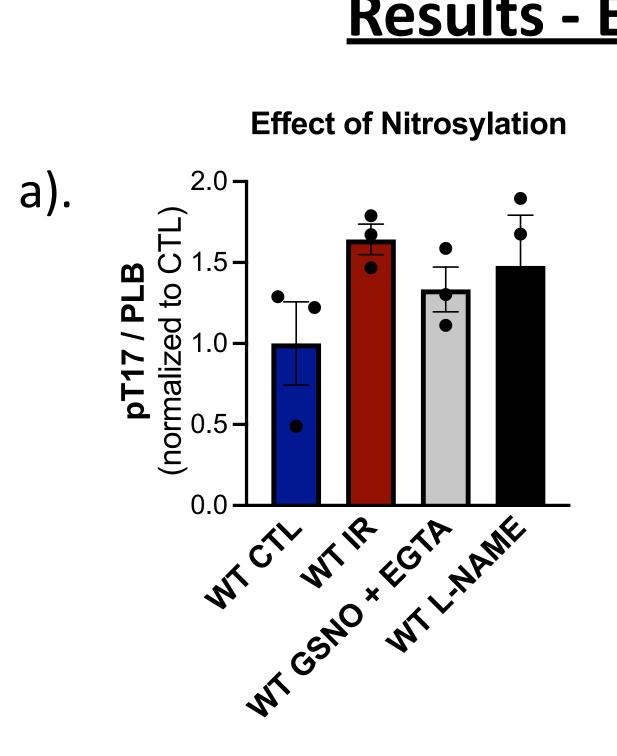


b).

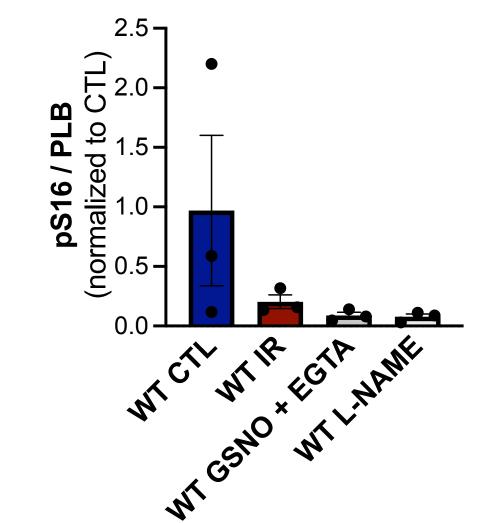


Phase II: Data Analysis

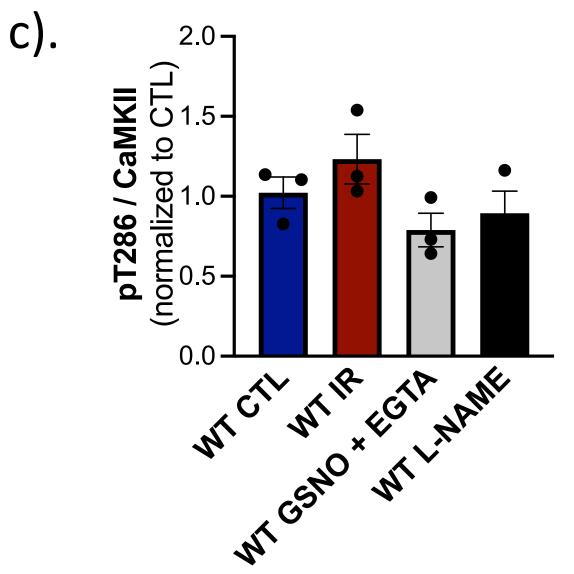
- Following perfusion protocol, hearts were flash-frozen and stored at -80C
- Hearts were homogenized as previously described (ref https://pubmed.ncbi.nlm.nih.gov/33926209/) • After protein quantification with BCA assay, samples were run on criterion TGX gels 4-20% before transfer to 0.2 μm nitrocellulose
- Blots were probed with one of the following antibody solutions
- Anti-pT17 and anti-PLB (Badrilla), Anti-pS16 and anti-PLB (Badrilla), Anti-CaMKII delta (custom antibody) • Followed by anti-rabbit IRDye800 and anti-mouse IR Dye680LT before scanning with the Sapphire Biomolecular Imager
- (Azure Biosystems).
- Blots were subsequently analyzed with Image J



Effect of Nitrosylation



Effect of Nitrosylation



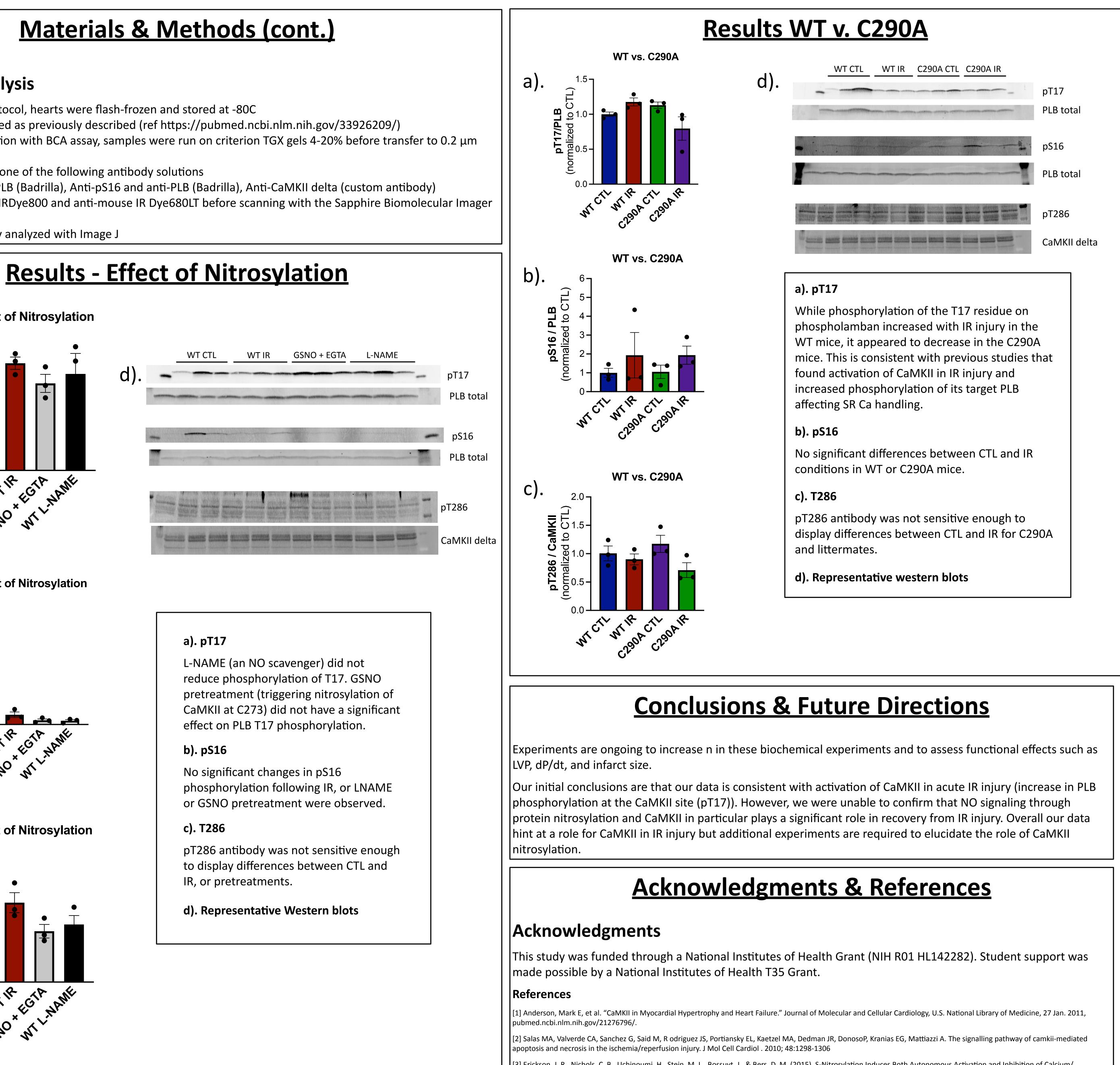
d).

a). pT17

b). pS16

c). T286

IR, or pretreatments.



[3] Erickson, J. R., Nichols, C. B., Uchinoumi, H., Stein, M. L., Bossuyt, J., & Bers, D. M. (2015). S-Nitrosylation Induces Both Autonomous Activation and Inhibition of Calcium/ Calmodulin-dependent Protein Kinase II δ. The Journal of biological chemistry, 290(42), 25646–25656. https://doi.org/10.1074/jbc.M115.650234

