

A Novel Top-Down Proteomics Method Empowered by Photocleavable Surfactant for **Comprehensive Analysis of Phospholamban Proteoforms** <u>Austin Carr¹, Kyle Brown¹, Andrew Alpert⁴, Song Jin¹, Ying Ge^{1,2,3}</u>

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Overview

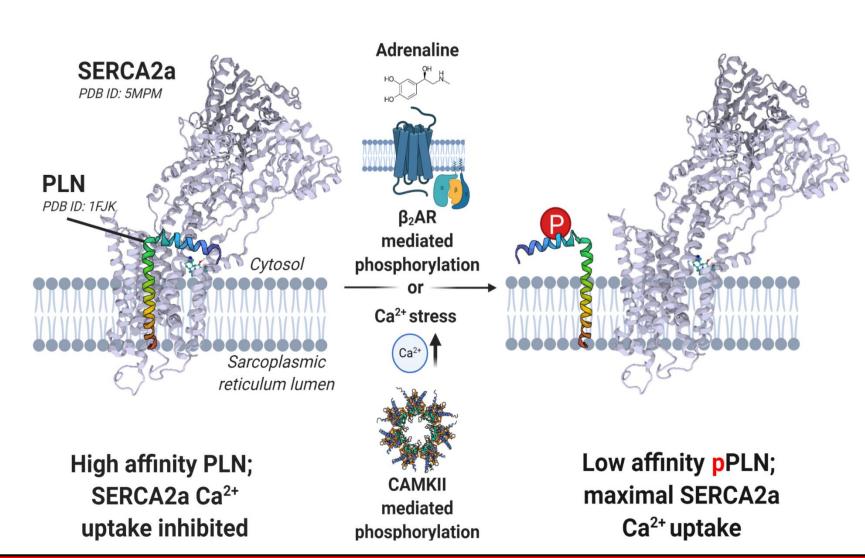
- (PLN), > Phospholamban critical, membrane protein, cardiac was extracted from swine heart using Azo, a spectrometry (MS)-compatible, mass photocleavable surfactant [1].
- >Hydrophilic interaction chromatography (HILIC) methods were developed for PLN analysis.
- >HILIC sample preparation methods reproducibly detect PLN and limit the amount of TFA adducts.

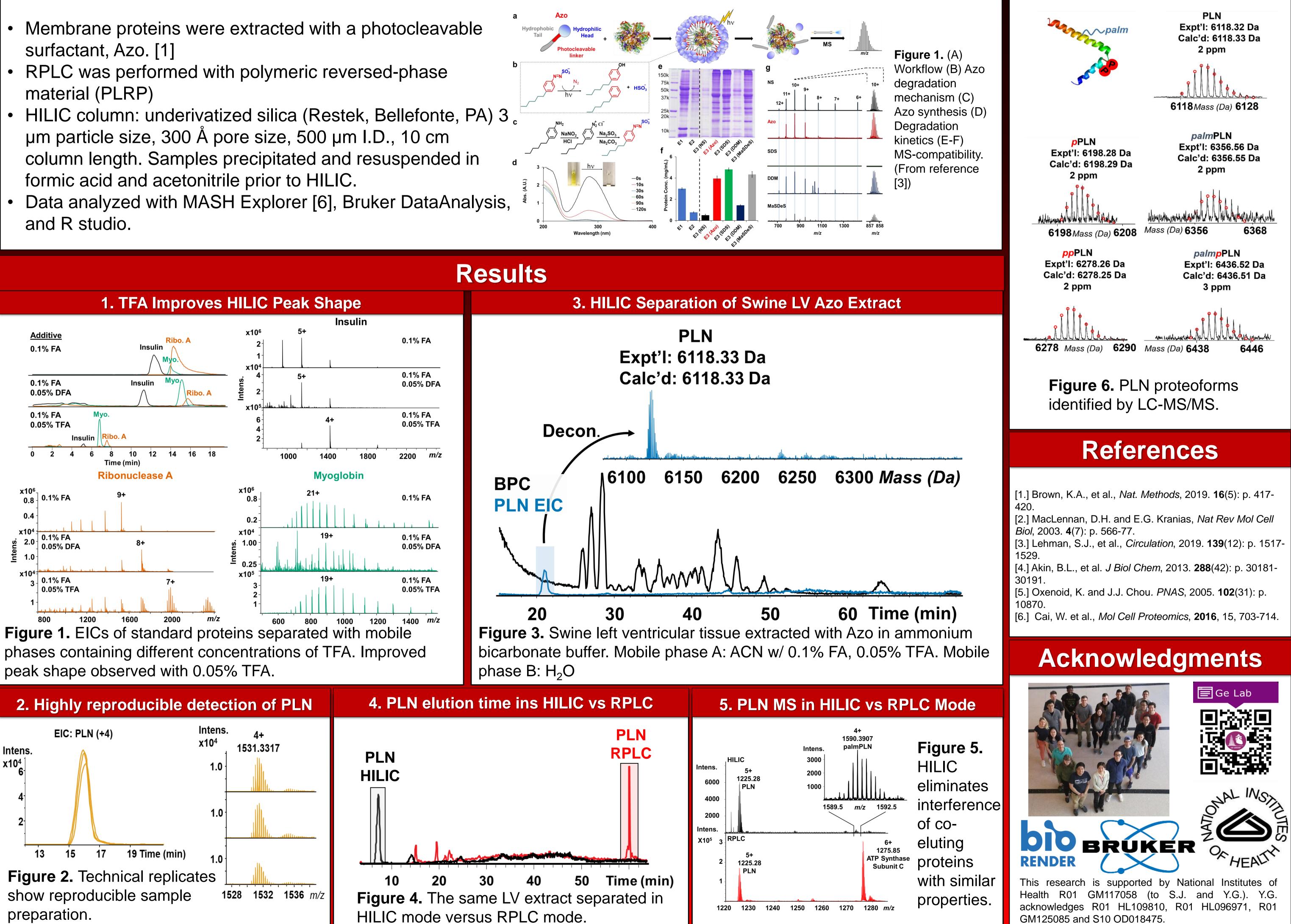
- Membrane proteins were extracted with a photocleavable surfactant, Azo. [1]
- material (PLRP)
- Data analyzed with MASH Explorer [6], Bruker DataAnalysis, and R studio.

Introduction

Phospholamban (PLN) is an important Ca²⁺ handling regulating protein sarco(endo)plasmic reticulum Ca²⁺-ATPase 2a (SERCA2a), responsible for 70% of cytosolic Ca²⁺ handling [2]. As some heart disease is currently hypothesized to be a result of aberrant Ca²⁺ handling [2], methods for studying the major Ca²⁺-handling proteins are critical. The regulatory role of PLN in heart disease is not fully understood, due to its hydrophobicity and lack of antibodies against rare PLN proteoforms. Herein, we have developed an unbiased, top-down MS method for characterization of PLN proteoforms in cardiac disease model enabled by Azo and HILIC.

Phosopholamban phosphorylation pathways





Methods



lm	PLN Expt'l: 6118.32 Da Calc'd: 6118.33 Da 2 ppm
	6118 Mass (Da) 6128
Da Da	<i>palm</i> PLN Expt'l: 6356.56 Da Calc'd: 6356.55 Da 2 ppm
208	Mass (Da) 6356 6368
)a)a	<i>palmp</i> PLN Expt'l: 6436.52 Da Calc'd: 6436.51 Da 3 ppm
290	Mass (Da) 6438 6446