

Virtual symposium celebrating the 50th anniversary of the Protein Data Bank



May 4-5, 2021

Day 1 Abstracts







Exploration of a novel mechanism of cardiac troponin release: in silico evidence of non-classical secretion

Jose M. Gonzalez-Rayas¹, Jose A. Hernandez-Hernandez¹, Rosa D. Lopez-Sanchez¹, Ana L. Rayas-Gomez², Jose M. Gonzalez-Yanez², Carlos Jerjes-Sanchez¹

¹Tecnologico de Monterrey, School of Medicine and Health Science, ²Hospital San Jose de Queretaro

Cardiac troponin is one of the most important biomarkers in cardiology. Traditionally, it has represented a biochemical sign of necrosis, a characteristic of acute myocardial infarction (AMI). However, with the arrival of high-sensitivity detection assays, cardiac troponin is now detected in healthy patients and in a number of uncommon scenarios unrelated to AMI (such as strenuous exercise, rapid atrial pacing and COVID-19). This suggests that troponin could be released as a consequence of alternative processes other than the well-stablished necrotic cell death. To try to explain this phenomenon, multiple mechanisms of troponin release have been postulated, although there is still no consensus on this matter. Furthermore, the term myocardial injury (a troponin determination above the upper reference limit, not necessarily associated with a specific diagnosis) was included in the Fourth Universal Definition of Myocardial Infarction. Nevertheless, the histological or pathobiological process responsible for myocardial injury has not been determined. Here, we studied the possibility of troponin release through non-classical secretion, a pathway specially triggered by cellular stress and inflammation. The protein sequences of the eight different isoforms of troponin (I, T and C of fast-twitch skeletal, slow-twitch skeletal and cardiac muscle) were retrieved from UniProtKB and the Protein Data Bank. SecretomeP 2.0, a neural networks-based program, was used to predict if any of these proteins undergoes non-classical release. Signal P 5.0 and TMHMM 2.0 were also applied to rule out the possibility of classical (endoplasmic reticulum/Golgi dependent) and Type IV (transmembrane) non-classical secretion. After analyzing the sequences, the cardiac T isoform was predicted to be non-classically secreted, as well as the I, T and C subunits of the fast-twitch skeletal muscle and the I subunit of the slow-twitch skeletal muscle. None of the troponin isoforms was found to be subject to classical or Type IV secretion. We believe that troponin T release through a non-classical pathway may contribute to the pathobiological definition of myocardial injury. There are already some experimental studies showing the release of troponin through extracellular microvesicles (a type of non-classical secretion) in various cell lines. Nonetheless, more evidence is needed to fully understand this intriguing phenomenon in cardiomyocytes.