Letter to the Editor

Can Pre-analytical Mistake Bearing Irisin Concentrations Be an Indicator of Coronary Artery Disease?

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Dear Editor,

It was with great interest that I read the article, entitled “Serum Irisin Level Can Predict the Severity of Coronary Artery Disease in Patients with Stable Angina,” published in the Korean Circulation Journal, a leading journal in cardiology and cardiac surgery. I would like to congratulate Efe et al. on their contributions to our knowledge of the circulatory physiology and biochemistry of the heart, as well as the mechanisms underlying stable angina. However, considering a number of pre-analytical errors and the incorrect citation of some sources, which I specify below, I think it would be useful to revisit and reinterpret this article.

Irisin is a peptide molecule containing 112 amino acids. Peptide/protein molecules are destroyed very rapidly by proteases. The human body produces more than 700 proteases and their concentration varies from one disease to another. In order to correctly measure the concentrations of peptide and protein molecules, biological samples taken from patients must be put into tubes containing a protease inhibitor like aprotinin. Otherwise, because the degree of destruction by proteases remains unclear, accurate measurement of peptide/protein concentrations is not possible. In the study, researchers put 10-mL blood samples into tubes containing ethylenediaminetetraacetic acid (EDTA) and stored the tubes at −80°C until analysis; they did not add any protease inhibitor to the tubes. EDTA in this context is a calcium-binding anticoagulant molecule. When samples taken from the patients are placed into tubes containing anticoagulants, the resulting biological sample will be plasma, not serum. Consequently, Efe et al. studied the irisin molecule not in serum (as indicated in their title), but in plasma. When the reported irisin concentrations were evaluated, the standard deviation value was very high. For instance, the standard deviation of irisin value (299.54±123.20 ng/mL) in the control group is almost half of the main value (299.54 ng/mL). In the study, researchers put 10-mL blood samples into tubes containing ethylenediaminetetraacetic acid (EDTA) and stored the tubes at −80°C until analysis; they did not add any protease inhibitor to the tubes. EDTA in this context is a calcium-binding anticoagulant molecule.
to 60 times higher than normal. This was likely due to different ELISA kits being used for the measurement of irisin concentration (AdipoGen ELISA kit; per the manufacturer, normal irisin measurement ranges of 0.2–2.0 µg/mL can be calculated 200–2,000 ng/mL). Cross-reactions with non-specific serum proteins can also influence the accuracy of ELISA kits.\(^5\) However, the researchers did not address the reasons for these high values. Efe et al.\(^1\) noted that the irisin decrease they found in stable coronary artery disease in their study resulted from the need for adenosine triphosphate (ATP) synthesis, as irisin overproduction would cause more heat and block ATP synthesis. Thus, they stated that they agreed with Boström et al.\(^2\) and Park et al.,\(^9\) who claimed that decreases in irisin served as a compensatory mechanism for increased ATP demand. However, the mechanism they explain was suggested by Aydin and colleagues,\(^7\) not by Boström et al.\(^2\) or Park et al.\(^9\). Likewise, that the heart muscle produces more irisin than the skeletal muscle was reported first by Aydin et al.,\(^7\) not by Boström et al.\(^2\).

In consideration of the pre-analytical mistakes indicated above, irisin concentrations well outside of previously reported ranges, and failure to correctly cite sources, we think it will be useful to revisit and reinterpret the article.

REFERENCES

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