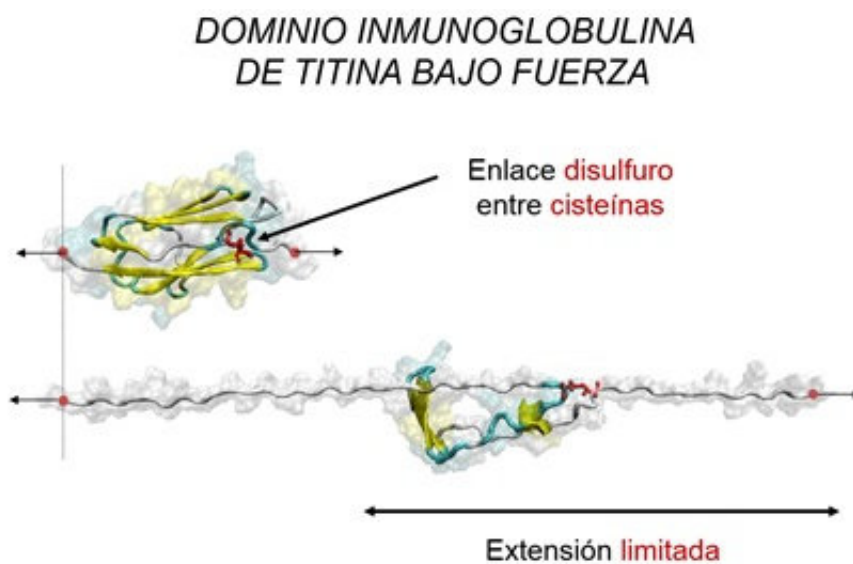


Scientists identify a key mechanism regulating a protein required for muscle and heart function

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Three-dimensional structure of one of the more than 100 immunoglobulin domains in titin. A disulfide bond is highlighted in red between two cysteines brought into proximity by the protein folding. When the immunoglobulin domain is subject to mechanical force, as during muscle stretching or in the diastolic phase of heart contraction, the domain opens up but is prevented from completely unfolding by the presence of the disulfide bond. Credit: CNIC

Scientists at the Centro Nacional de Investigaciones Cardiovasculares Carlos III (CNIC) and Columbia University in New York have discovered an important mechanism in the regulation of a protein that

plays an essential role in the function of skeletal muscle and the heart. The study, published today in *Nature Communications* and coordinated by CNIC researcher Jorge Alegre-Cebollada, describes a new mechanism in the regulation of the elasticity of the giant protein titin. Titin, explained Alegre-Cebollada, is a key protein in the functioning of striated muscles throughout the body, particularly in the heart: "the proof of this is that mutations in the titin gene are a common cause of diseases affecting the muscles of the body and the heart."

Titin is the largest [protein](#) in the human body and as such has a multitude of functions. According to Jorge Alegre-Cebollada, "in simple terms we can think of titin as a 'molecular spring' that allows muscle cells to contract in synchrony." However, it is not a simple spring, and the many mechanisms that determine titin elasticity include the unfolding of specific regions in its structure called immunoglobulin domains. In all, titin elasticity is determined by the concerted action of more than 100 immunoglobulin domains within the protein.

Using bioinformatic and structural biology approaches, the research team found that the immunoglobulin domains have a high cysteine content. This amino acid confers special properties. Jorge Alegre-Cebollada explained that "when 2 cyteines in a protein come close to one another, they can form a chemical link between different parts of the polypeptide chain called a disulfide bond." The researchers observed that many of the immunoglobulin domains in titin form disulfide links and that the cysteines participating in them can change dynamically, a process called isomerization. "The most interesting finding was that the formation and isomerization of [disulfide bonds](#) causes major changes in the elastic properties of titin."

The formation of disulfide bonds is an example of a broader class of biochemical transformations known as reduction-oxidation (redox). It has long been known that many disease processes affecting the [heart](#),

including myocardial infarction, involve sudden and drastic changes in the redox state of the heart muscle.

Dr. Alegre-Cebollada's group is currently investigating how our cells modify the [titin](#) redox state as a mechanism to modulate skeletal and heart [muscle](#) activity and how different diseases can interfere with the mechanical action of the protein, resulting in loss of functionality. Alegre-Cebollada concludes, "Our mechanical findings were made possible by reconstituting contractile systems in vitro. While we have learned a lot through this approach, the challenge now is to understand how these basic principles operate in a living organism. This is the focus of our current research, using a multidisciplinary approach that combines advanced techniques in physiology, biology, and biochemistry."

The study is the result of a collaboration between Dr. Alegre-Cebollada and Professor Julio Fernández at Columbia University. Professor Fernández is a pioneer in the development of single-molecule biophysical techniques for investigating the mechanical properties of proteins, and his collaboration with Dr. Alegre-Cebollada has made it possible to describe the biochemical regulation of protein mechanics.

More information: David Giganti et al, Disulfide isomerization reactions in titin immunoglobulin domains enable a mode of protein elasticity, *Nature Communications* (2018). [DOI: 10.1038/s41467-017-02528-7](https://doi.org/10.1038/s41467-017-02528-7)

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