

Morphometric characterization of fibrinogen's α C regions and their role in fibrin self-assembly and molecular organization

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Abstract

© 2017 The Royal Society of Chemistry. The flexible C-terminal parts of fibrinogen's A α chains named the α C regions have been shown to play a role in fibrin self-assembly, although many aspects of their structure and functions remain unknown. To examine the involvement of the α C regions in the early stages of fibrin formation, we used high-resolution atomic force microscopy to image fibrinogen and oligomeric fibrin. Plasma-purified full-length human fibrinogen or des- α C fibrinogen lacking most of the α C regions, untreated or treated with thrombin, was imaged. Up to 80% of the potentially existing α C regions were visualized and quantified; they were highly heterogeneous in their length and configurations. Conversion of fibrinogen to fibrin was accompanied by an increase in the incidence and length of the α C regions as well as transitions from more compact conformations, such as a globule on a string, to extended and more flexible offshoots. Concurrent dynamic turbidimetry, confocal microscopy, and scanning electron microscopy revealed that trimming of the α C regions slowed down fibrin formation, which correlated with longer protofibrils, thinner fibers, and a denser network. No structural distinctions, except for the incidence of the α C regions, were revealed in the laterally aggregated protofibrils made of the full-length or des- α C fibrinogens, suggesting a pure kinetic effect of the α C regions on the fibrin architecture. This work provides a structural molecular basis for the promoting role of the α C regions in the early stages of fibrin self-assembly and reveals this stage of fibrin formation as a potential therapeutic target to modulate the structure and mechanical properties of blood clots.

<http://dx.doi.org/10.1039/c7nr04413e>

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