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1 10000 000	this identifier to cite or link to this item: http://hdl.handle.net/123456789/212				
Title:	Structural and dynamical insights into the molten-globule form of ovalbumin.				
Authors:	Bhattacharya, M. (/jspui/browse?type=author&value=Bhattacharya%2C+M.) Mukhopadhyay, S. (/jspui/browse?type=author&value=Mukhopadhyay%2C+S.)				
Keywords:	Biphasic Conformational change				
Issue Date:	2012				
Publisher:	American Chemical Society.				
Citation:	Journal of Physical Chemistry B, 116 (1), pp. 520-531.				
Abstract:	Ovalbumin is a 45 kDa egg-white glycoprotein which belongs to the class of serpin superfamily. We have studied the structural properties of both native and partially unfolded molten-globule forms of ovalbumin using a diverse array of spectroscopic tools. Time-resolved fluorescence measurements provided important structural and dynamical insights into the native and molten-globule states. Fluorescence anisotropy decay analysis indicated that there is a conformational swelling from the native to the molten-globule form of ovalbumin. We have also carried out rededge excitation shift measurements to probe the dipolar relaxation dynamics around the intrinsic tryptophan residues. Additionally, stopped-flow fluorescence experiments revealed that the conformational transition from the native to the molten-globule form proceeds in a stepwise manner involving a burst-phase with a submillisecond conformational change followed by biphasic slower conformational reorganizations on the millisecond time scale leading to the final molten-globule state.				
URI:	http://pubs.acs.org/doi/abs/10.1021/jp208416d (http://pubs.acs.org/doi/abs/10.1021/jp208416d)				
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