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Estimation of the number of alpha-helical and beta-strand segments in proteins using circular dichroism spectroscopy

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598	Estimation of protein secondary structure from circular dichroism spectra: inclusion of denatured proteins with native proteins in the analysis. <b>2000</b> , 287, 243-51		480
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595	Unfolding pathway of the colicin E1 channel protein on a membrane surface. <b>2000</b> , 295, 679-92		56
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