

Characterization of transient intermediates of a calmodulin- peptide complex

Yoshinobu IZUMI^{*1}, Yuji JINBO¹, Shigeo KUWAMOTO¹, Nobuyuki Miho¹, Hidenori YOSHINO²,
Yuzuru HIRAGI³, Hiroshi KIHARA⁴

¹Graduate School of Science and Engineering, Yamagata University, Yonezawa 992-8510, Japan

²Department of Chemistry, Sapporo Medical University, Sapporo, 060-8556 Japan

³Institute for Chemical Research, Kyoto University, Uji, 611-0011 Japan

⁴Department of Physics, Kansai Medical University, Uyama, Hirakata, 573-1136, Japan

Introduction

In the present work we have measured the dissociation kinetics of a Ca^{2+} -saturated calmodulin-peptide complex. The peptide studied is the Ca^{2+} -calmodulin dependent protein kinase IV calmodulin target-site. The effects of Ca^{2+} removal with a chelator were monitored using a SR-SAXS stopped-flow measurement. The present result allows the characterization of transient intermediates for the dissociation of 4Ca^{2+} -calmodulin (4Ca^{2+} -CaM)-peptide complex. The result obtained is compared with other studies on the dissociation of calmodulin-peptide complexes. We confirm that the rate of the slowest step is determined by the contribution of a kinetic relaxation mechanism involving the intermediate species 2Ca^{2+} -CaM-peptide, with two Ca^{2+} ions bound in the C-terminal domain, which has been previously suggested [1].

Materials and Methods

A 19-residue peptide having the sequence (RRKLKAAVKAVVASSRLGS; CaMKIVp) and bovine brain CaM were used. Stopped-flow experiments were performed using an instrument for SAXS with a stopped-flow apparatus (Unisoku Co Ltd).

Results and Discussion

The changes in SR-SAXS profiles on mixing of CaMKIVp with 4Ca^{2+} -CaM (at 25°C) were almost complete within the instrument deadtime, indicating a bimolecular association was complete within a few msec. We therefore studied the dissociation process of 4Ca^{2+} -CaM-peptide complex (C^+). The results in Fig. 1 indicate that the molecular weight of the CaM-peptide complex does not change during the dissociation process, suggesting that the peptide binds to CaM even in the absence of Ca^{2+} , which is supported by a recent result[2]. Furthermore, the dissociation pathway is characterized by biphasic kinetics as shown in Fig. 2. The first event is the loss of two Ca^{2+} ions from the N-terminal lobe, followed the loss of two Ca^{2+} ions from the C-terminal lobe. The intermediate species (I^+) with two Ca^{2+} ions in the C-domain is observed within the first 50-100 msec of the

dissociation pathway. A subsequent activated state (A^*) is observed at about 250 msec of the dissociation pathway. Final event is a conformational change in CaM-peptide complex (A^+).

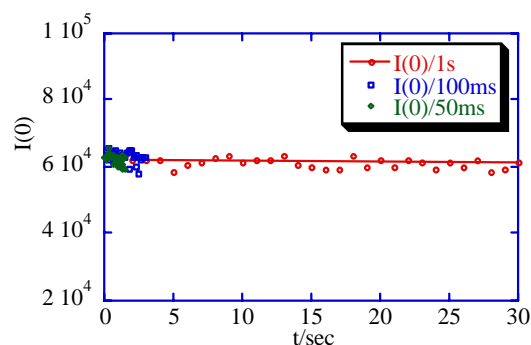


Fig. 1 Time course of the forward scattering amplitude $I(0)$ for the dissociation of 4Ca^{2+} -CaM-peptide complex.

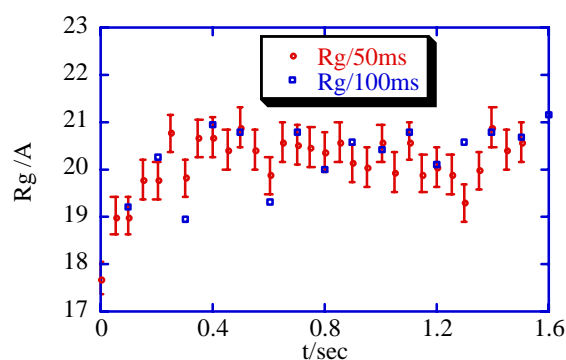


Fig. 2 Time course of the radius of gyration of gyration R_g for the dissociation of 4Ca^{2+} -CaM-peptide complex.

References

- [1] S.E.Brown et al., J.Biol.Chem. 276 3389(1998).
- [2] Y.Izumi et al. FEBS Lett. 495 126(2001).

* yizumi@dip.yz.yamagata-u.ac.jp