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Time-resolved kinetic refolding of PI3K SH3 in the presence of 45% ethylene glycol at pH 6

Yoshitaka Matsumura¹, Masaji Shinjo¹, Nobuyuki Okishio² and Hiroshi Kihara¹ Department of Physics, Kansai Medical University, 18-89 Uyama-Higashi Hirakata, Osaka 573-1136, Japan

²Hokkaido Red Cross Blood Center, Yamanote 2-2, Nishi-ku, Sapporo 063-0002, Japan

Introduction

We have investigated kinetic protein folding study of src SH3 domain and its mutant, A45G and Fyn SH3 domain, by various probes including X-ray solution scattering combined with cryo-stoppoed-flow method [1, 2]. We have done kinetic refolding experiments of another SH3 domain, PI3K SH3 domain. This protein is also mainly composed of β -sheets as src SH3 domain [3]. In the present study, we performed kinetic refolding of PI3K SH3 domain by a concentration jump of guanidine hydrochloride (GuHCl) from 5 M to 0.7 M. The experiments were done in the presence of 45% ethylene glycol at pH 6 at -5°C.

Experimental

X-ray scattering experiments were done at the beamline of 15A, keeping the sample-to-detector-distance at c.a. 1.3 m with a CCD-based X-ray detector (Hamamatsu Photonics, C7300). The obtained data were corrected for image distortion, non-uniformity of sensitivity, and the contrast reduction on X-ray image intensifier.

Results and Discussion

Figure 1 shows kinetic refolding of PI3K SH3 domain SAXS measurements by of time-resolved stopped-flow in the presence of 45% ethylene glycol at pH 6 at -5 °C. The figure does not show time-dependent change on Rg within the experimental errors. The averaged Rg value of the fitting line was $17.7 \pm 1.0 \text{ Å}$. The Rg of the native state of the PI3K SH3 domain is 15 Å. In contrast, the Rg of the unfolded state of the protein is 27.5 Å. Dobson et al reported that the slow refolding took place with rate of 0.0565 and 0.0140 s⁻¹ [4]. Taken into consideration of these native and denatured Rg values, it is thought that the obtained Rg value we observed is a kinetic transient intermediate, of which size is slightly bigger than the native structure.

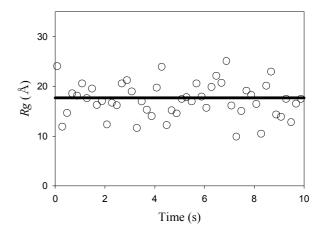


Fig. 1 Kinetic refolding of PI3K SH3 domain of time-resolved SAXS measurements by the stopped-flow in the presence of 45% ethylene glycol at pH 6 at -5 $^{\circ}$ C. The black line is fitting line. The Rg was 17.7 Å.

References

- [1] Li et al. (2007) Biochemistry. 46, 5072-5082.
- [2] Li et al. (2007) J. Mol. Biol., 372, 747-755.
- [3] Koyama et al. (1993) Cell, 72, 945-952.
- [4] Dobson et al. (1998) J. Mol. Biol., 276, 657-667

^{*} kihara@makino.kmu.ac.jp