BL-1A, BL-17A, AR-NE3A/2015G513 Structural analysis of the metal-substituted light-harvesting-reaction center core complexes

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1 Introduction

In purple photosynthetic bacteria, the light energy is absorbed by two types of light-harvesting complexes (LH1, LH2), and then is transferred efficiently to the reaction center (RC) where the primary charge separation takes place across the membrane and a cyclic electron transport chain occurs. The LH1 complex is located intimately around the RC with a fixed stoichiometric ratio to form the so-called core complex (LH1-RC). LH1 is a large oligomer of a basic structural unit composed of a heterodimer of two small integral membrane polypeptides (a and b, ca. 6 kDa) associated with bacteriochlorophyll (BChl) and carotenoid molecules. Thermochromatium (Tch.) tepidum is a thermophilic purple sulfur photosynthetic bacterium originally isolated from a hot spring in Yellowstone National Park. It grows anaerobically at optimum temperatures of 48 - 50 °C with an upper limit of 58 °C, and contains an unusual LH1 complex that absorbs maximally around 915 nm (Q_{y} transition). In a series of investigations, we found that the Tch. tepidum LH1 is highly stable at room temperature, and calcium ions are involved in both the enhanced thermal stability and the large red shift of the LH1 Q_y transition [1, 2].

The crystal structure of LH1-RC from *Tch. tepidum* has been determined [3], in which the Ca-binding sites have been identified. On the other hand, the Ca²⁺ ions in LH1 can be replaced by other divalent metal ions (e.g., Sr^{2+} and Ba^{2+}), resulting in a substituted LH1-RC with the LH1 Q_y in a range of 880 ~ 890 nm [1, 2]. Here, we report structure determination of the Sr- and Basubstituted LH1-RCs [4].

2 Experiment

Crystallization of the Sr- and Ba-LH1-RCs and postcrystallization treatment were carried out following the same method as describe previously [3]. X-ray diffraction data were collected at BL41XU and BL44XU of SPring-8, and BL1A, BL17A and NE3A of the Photon Factory at 100K. To identify the Sr²⁺- and Ba²⁺-binding sites, datasets were acquired at three wavelengths for the Sr-LH1-RC and two wavelengths for the Ba-LH1-RC.

3 Results and Discussion

Anomalous difference Fourier maps obtained at wavelengths of 0.76 Å and 1.5 Å revealed strong electron densities for the Sr- and Ba-LH1-RC crystals, respectively. The strong densities disappeared at 0.78 Å (remote) for the Sr-LH1-RC and largely reduced at 0.98 Å for the Ba-LH1-RC, confirming that they are truly from the Sr^{2+} and

 Ba^{2+} ions, respectively. Both the Sr^{2+} and Ba^{2+} have the same number and are located at essentially the same positions in the substituted LH1-RCs. These locations are clearly different from, though close to, the Ca-binding sites. Sixteen Sr²⁺ and Ba²⁺ ions are identified in the LH1 complexes. Conformational rearrangement induced by the substitution is limited to the metal-binding sites. Unlike the Ca-LH1-RC, only the α -polypeptides are involved in the Sr- and Ba-coordinations in LH1 (Figure 1). The difference in the thermostability between these complexes can be attributed to the different patterns of the network formed by metal-binding. The Sr- and Ba-LH1-RCs form a single-ring network formed by the LH1 α -polypeptides only, in contrast to the double-ring network composed of both α - and β -polypeptides in the Ca-LH1-RC. Based on the structural information, a combined effect of hydrogen-bonding, structural integrity and charge distribution is considered to influence the spectral properties of the core antenna complex.

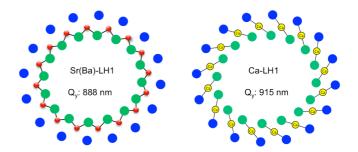


Fig. 1: Schematic illustration showing the difference between Sr(Ba)- and Ca-binding networks formed in LH1s [4].

<u>References</u>

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