フォトンファクトリーでの高圧蛋白質結晶実験 Photon Factory and High-Pressure Macromolecular Crystallography

Leonard M.G. Chavas¹、永江峰幸²、渡邊信久²、平木雅彦¹、山田悠介¹、 五十嵐教之¹、松垣直宏¹、若槻壮市¹

1構造生物学研究センター、2名古屋大学大学院工学研究科

Recent developments in instrumentation for high-pressure studies of macromolecular crystals have been undertaken at various synchrotron sites. These new implementations have allowed the collection of complete diffraction data with high accuracy from samples subjected to extreme conditions. Thus, previous reports highlighted the analysis of protein crystals at pressures higher than 1 GPa with no variation the clear in crystals' diffraction power. As а direct consequence, increasing interests in high-pressure macromolecular crystallography nowadays are emerging, notably in specific areas of detailed thermodynamic and kinetics. A non-exhaustive list of the principal investigations of interest includes diverse applications to biotechnology developments, the clarification of the effects of high-pressure in deep-sea organisms, or the studies of the influence of compression on proteins' flexibility, possible regulatory factor

of biological functions such as oligomerization enzymatic or activities. Although numerous innovative projects are emerging from these methodologies, they remain poorly investigated because of the limited access to synchrotron facilities providing high-pressure equipment. Taking advantage of the present set-up at PF AR-NW12A. studies high-pressure of macromolecular crystals have been made possible by adapting optional already tools to the installed equipment, such as a diamond anvil cell anchored to the goniometer head, a bigger light for a proper or visualization of the sample within the compressed cell for crystal centering. In this presentation, we will introduce the potential for high-pressure studies Some at AR-NW12A. concrete examples will be discussed for which full data sets of macromolecular crystals submitted to extreme stress have been collected and analyzed.