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B-XRD1043 - Powder crystal structure of organic crystals

Introduction

Previously, it was typical to conduct crystal structure analysis using the single crystal method. However, analysis of crystal structure using the single crystal method is difficult in the case of samples for which it is hard to produce single crystals, and samples for which crystal growth is difficult even if single crystals can be produced. In recent years, the precision of powder X-ray diffractometers has been improving, and thus it is becoming possible to conduct crystal structure analysis using powder samples. Here we show an example where a powder sample was measured, and structural analysis was conducted based on the obtained diffraction pattern.

Measurement and results

The crystal system of γ -indometacin crystals belongs to the triclinic type, and thus many diffraction lines overlap, and it is difficult to determine the lattice constants from the powder X-ray diffraction pattern. Figure 1 shows an overlay of the X-ray diffraction pattern obtained with a high-resolution convergent beam optical system, which enables linear focus using an elliptical surface multilayer mirror of monochromatized X-rays only on Cu Ka₁, in a Ge (111) Johansson-type curved crystal; and the X-ray diffraction pattern obtained using a conventional parallel beam optical system containing Cu Ka₁ and Ka₂ lines. With the data measured with the high-resolution convergent beam system, it can be easily confirmed that there are minute, strongly overlapping peaks which were difficult to discover with the conventional optical system. The structure was successfully determined due to this improvement in data quality.

The initial structure was determined using the direct-space method (parallel tempering method), and the structure was refined using the Rietveld method. The crystal structure obtained through refinement exhibited a good match with the structure obtained using single crystal structure analysis, as shown in Figure 2.

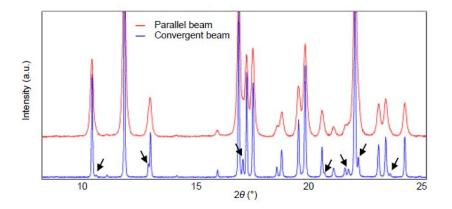


Figure 1: Measurement data for γ -indometacin obtained with a conventional parallel beam optical system (red, upper pattern) and a high-resolution convergent beam optical system (blue, lower pattern). The arrow marks \downarrow indicate peaks

that cannot be detected with a parallel beam optical system.



Figure 2: Crystal structure of y-indometacin obtained from powder crystal (yellow) and single crystal (orange)

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