Fed up with puzzling ambigous NMR, MS spectra and HPLC results for the identification of a new compound?

> Get the Molecular Structure from Crystallise!

Have you spent a lot of effort, time and money? And you have to close the project because the identification process fails?

Crystallise! will rescue your Investment

Contact us E-mail: info@crystallise.ch Tel: +41 44 55 83 400 www.crystallise.ch Skype: crystallise.ch or crystalliseag

X-ray Diffractometer

- State of the art technology We can measure crystals down to a size of $0.01 \times 0.01 \times -0.04 \text{ mm}^3$.
- Cu- μ source for absolute structure determination and light atom molecules.
- Cryo-stream for cooling the crystals down to -183°C (90 K) and for the crystallization of liquid samples.



State of the art Bruker D8-Venture X-ray Machine equipped with a CMOS (10 x 10 cm²) detector.

Chemical Laboratory

Crystallise! runs an independent fully equipped chemistry laboratory. We even have a high vacuum greaseless Schlenk line to avoid contamination of the samples.





Your Partner for Crystallization X-ray Structure Analysis

Grabenstrasse 11a. CH-8952 Schlieren, Switzerland

Crystallization of Liquids

The liquid sample is placed inside a sealed glass capillary and mounted in the X-ray machine. Using the cryo-stream the sample can be cooled down to -183 °C (90 K). Therefore the crystallization process takes place in situ.



Our Services

Crystallization

- Crystallization of Liquids
- Crystallization of Sensitive Compounds
- Crystallization of Stable Compounds

X-ray Measurement

Absolute Structure

For absolute structure determination, Sucrose is the "standard" system to be measured. It is a good test of experimental methods aimed at the determination of small molecules of biological origin (J. Appl. Cryst. 1991, 24, 352-354).

Crystallise! has grown at their laboratory single crystals Crystal Structure of Sucrose

of Sucrose. To "test" the settings and capabilities of their machine. A single crystal was picked and obtained @ Crystallise! AG meausured. R1 = 2.15%, wR2 = 5.21%, The results GooF 1.06, Flack -0.06(8) are as good (or even better) as the best standard structure reported in the literature (c.f. Cryst. Growth Des., 2009, 9, 3551-3561). Growing single crystals and picking the right one is essential for a good structure determination. Crystallise! AG



Single Crystal / Liquid Sample

Example of an in situ crystallization.

With the aid of a special technique that requires a laser,

Crystallise! is able to obtain a single crystal from pure liquid substances.



- Absolute Structure Determination
- Picking & Mounting Crystals under Inert Gas Atmosphere

Data Analysis

Unsolved Data Sets

Twinning and Disordered Structures

Do you need our services? Contact us at info@crystallise.ch

your smartest choice for better resutls!