



CCTBX: AN INTRODUCTION

NOT QUITE FOR DUMMIES

Luc J. Bourhis

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Durham University, UK and Bruker AXS

THE BIG PICTURE

Modular design to minimise dependencies

phenix.refine

Olex2.refine

Macro-Molecular Toolbox (*mmtbx*) Small Molecule Toolbox (smtbx)

Input/Output Toolbox (smtbx)

Common Crystallography Toobox (cctbx)

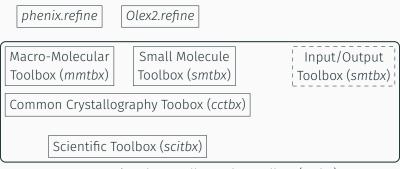
Scientific Toolbox (scitbx)

In Python:

from cctbx import sgtbx

THE BIG PICTURE

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Computational Crystallography Toolbox (cctbx)

In Python:

from cctbx import sgtbx

DOWNLOADS

- This tutorial: slides, code and data https://github.com/luc-j-bourhis/ rovinj-2015-cctbx-tutorial
- Source and binary packages: http://cci.lbl.gov/cctbx_build/ Not very up-to-date anymore but it will do for this tutorial!
 - Linux: choose CentOS 6.2 (64-bit) even if you don't wear a red hat source cctbx_build/setpaths.sh
 - Windows 7 (64-bit) or Windows XP (32-bit) cctbx_build/setpaths.bat
 - cctbx.python my_script.py
- · Source code repository (version control):
 - · Currently hosted on Sourceforge
 - · Plans to move to Github soon-ish

MODULES

From the foundations to the roofs:

- Python \leftrightarrow C++ bridge
 - · Python is slow but convenient
 - · C++ is fast but cumbersome
 - Boost.Python is a bridge between them and boost_adaptbx rules it
- scitbx: mathematics, non-crystallographic physics (rigid body)
 - · n-dimensional arrays, bridge with NumPy
 - · linear algebra, special functions, non-linear least-squares
- · cctbx: common crystallography
 - structure factors computation, and their derivatives
 - · space group toolbox
 - Fourier transforms
- smtbx: constraints
 - structure factors computation and derivatives

MODULES

Based on cctbx:

- · rstbx: indexing and integration of diffraction images
- xfel: X-ray free-electron lasers
- crys3d, gltbx: tools to write GUI displaying 3D objects (structures, Fourier maps)
 - $\cdot \longrightarrow Phenix$

SCITBX.MATRIX

Matrices in pure Python:

small matrices.

```
>>> from scitbx.matrix import rec, col
>>> A = rec(elems=(-1,0,0,0,-1,0,0,0,-1), n=(3,3))
>>> b = col((3,4,5))
>>> C = A * b
>>> print C
matrix.rec(elems=(-3, -4, -5), n=(3,1))
>>> abs(C)
7.0710678118654755
Matrix written by row: \begin{bmatrix} -1 & 0 & 0 \\ 0 & -1 & 0 \\ 0 & 0 & -1 \end{bmatrix} Use only for few operations on
```

SCITBX.MATRIX

Particularly useful in crystallography

Used a lot.

AND NOW, LADIES AND GENTLEMENT, THE MAIN FEATURE!

Let's triple the unit cell of a structure.

Why, oh, why?

Phase transition: unit cell before the transition (a, b, c, 90, 90, 90), and after (3a, b, 3c, 90, 90, 90).

To compare the two structures, it is useful to put the first one in a tripled unit cell, keeping the same symmetries.