

# Optimizing the Dose Distribution with Small Beams



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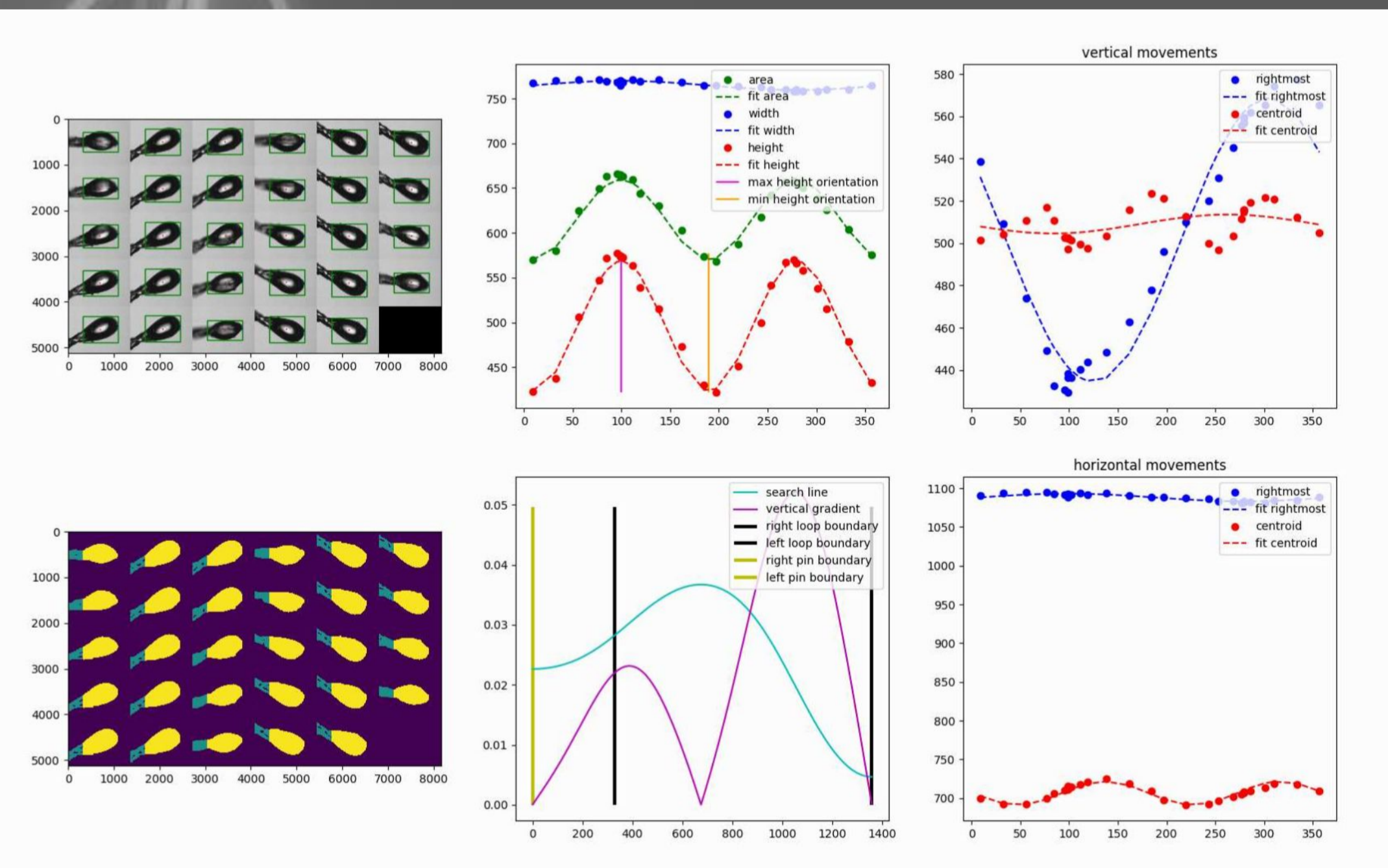
## Introduction

The fundamental variable governing radiation induced changes in crystals of biological macromolecules is the dose -- energy absorbed by the crystal during its lifetime. Ideally we should be able to control for it when designing the experiment. To do so we need to have a pretty good idea of the properties of the probe (photon energy, shape, intensity) and those of the sample (shape, composition) and their relative movements during the experiment.

## Reconstructing shape of sample environment

Series of optical sample images acquired over the rotational range of omega axis is used to determine key characteristics of the sample containing volume yielding

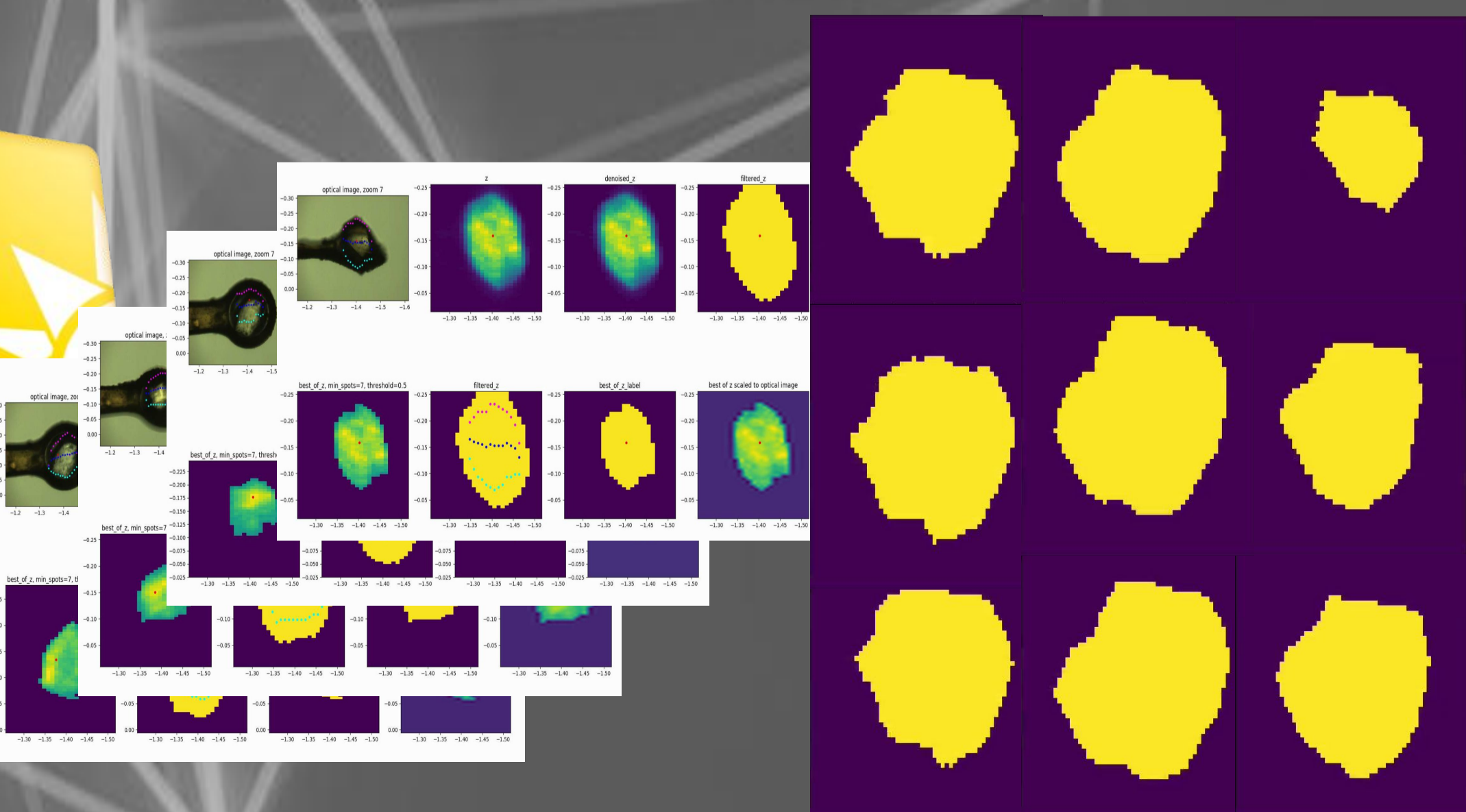
- loop bounding box at arbitrary orientation
- aligned center of the loop
- parameters of area to search with X-rays
- reconstructed volume enclosing the crystal



## Reconstructing crystal shape

Find the zones that diffract:

- crystal size and shape
- crystal quality and its variability
- crystal center curve model



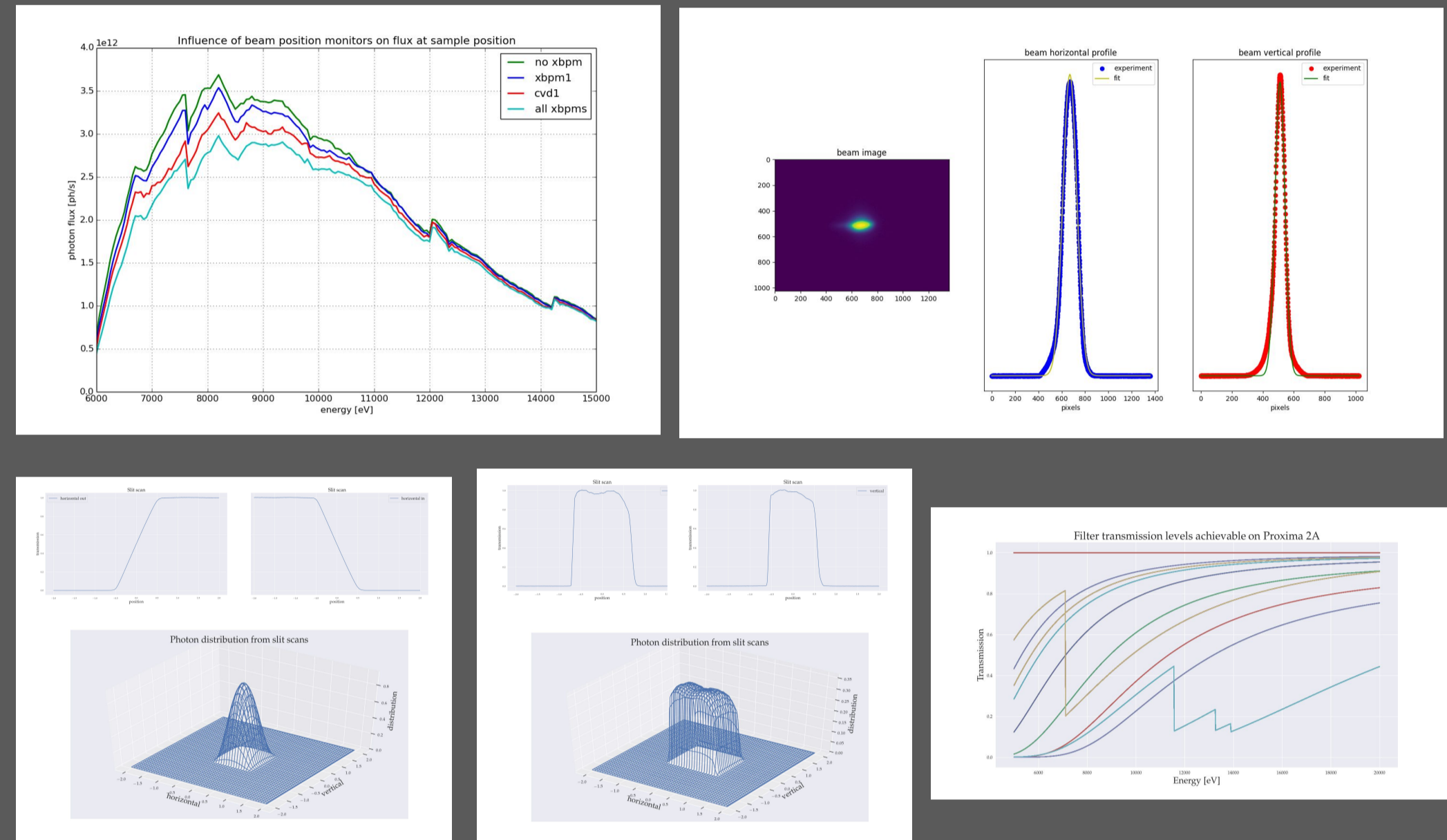
raster scans at four orientations

reconstructed crystal shape along rotational axis

## Controlling beam properties

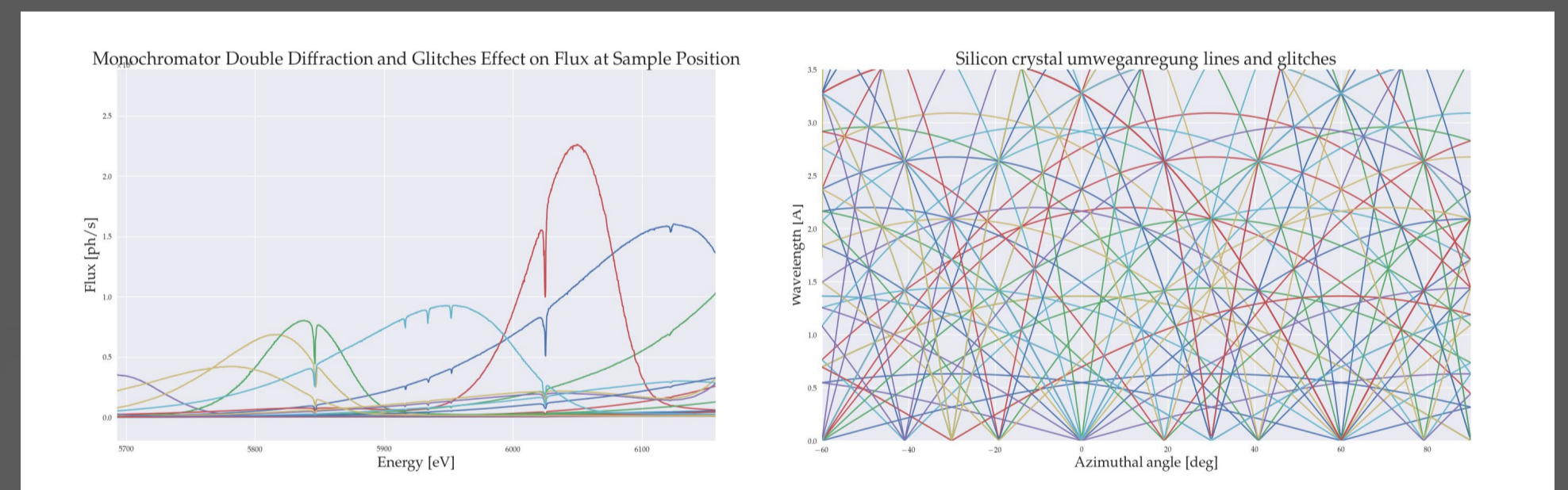
The x-ray probe

- size
- profile
- intensity
- photon energy
- influence of beamline components on flux at sample
- slits, filters, apertures, position monitors and monochromator



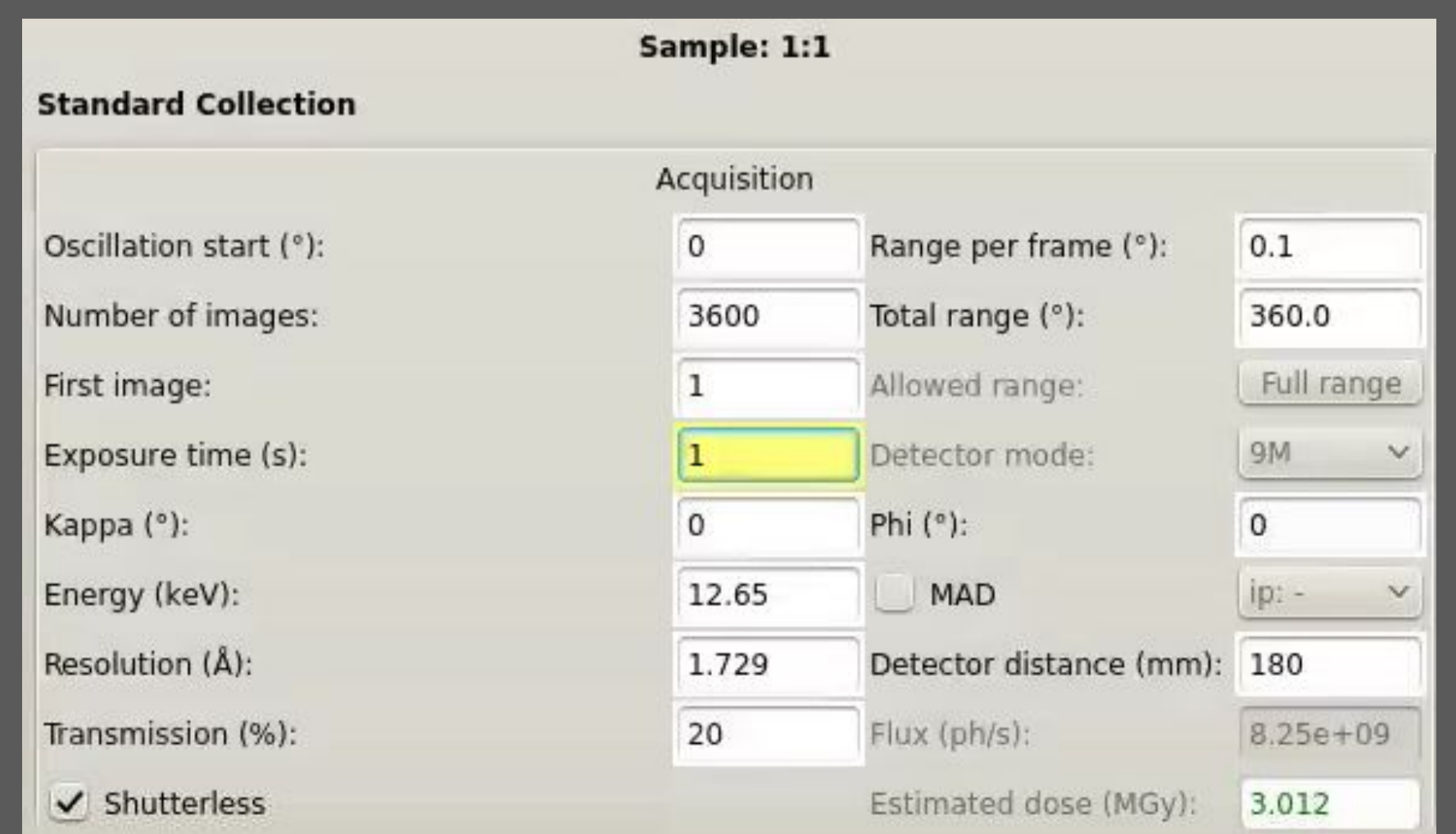
## Better look at the monochromator

- Beware of monochromator crystal double diffraction and glitches -- can be significant at specific energies



## Outlook

- sample shape from optical images alone
  - premis: reconstructing crystal shape purely from optical images is possible if sample is embedded in transparent medium
  - segment out support, mother liquor and crystal -- enhancing current pixel wise segmentation
  - dose as an experimental parameter



acquisition widget with the dose field