

ProjectID

Scientist / Research Group

Protein / Molecule

Goal

- salt detection
- dataset < ____ Å (anomalous? element ____)
- flour scan (expected element ____)

remarks

Prior knowledge	
Spacegroup	_____
	_____ / _____ / _____ Å
Beamline	_____ / _____ / _____ °
	Synchrotron _____
	Beamline _____
	Detector _____
	Date _____

Crystal position by priority (cane id and position e.g. Msch1 / b+1) - **please tick of measured crystals!**

Datasets

basic folder: _____

collection	processing	comment
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1. prefix _____

λ _____ Å	Δφ _____ °	_____	_____ Å resolution
dist _____ mm	exp _____ s	_____ / _____ / _____ Å	_____ % completeness
φ _____ - _____ °		_____ / _____ / _____ °	anisotropic? <input type="checkbox"/> Y/ <input type="checkbox"/> N
Transmission _____	Dose _____ MGy		anomalous? <input type="checkbox"/> Y/ <input type="checkbox"/> N

2. prefix _____

λ _____ Å	Δφ _____ °	_____	_____ Å resolution
dist _____ mm	exp _____ s	_____ / _____ / _____ Å	_____ % completeness
φ _____ - _____ °		_____ / _____ / _____ °	anisotropic? <input type="checkbox"/> Y/ <input type="checkbox"/> N
Transmission _____	Dose _____ MGy		anomalous? <input type="checkbox"/> Y/ <input type="checkbox"/> N

3. prefix _____

λ _____ Å	Δφ _____ °	_____	_____ Å resolution
dist _____ mm	exp _____ s	_____ / _____ / _____ Å	_____ % completeness
φ _____ - _____ °		_____ / _____ / _____ °	anisotropic? <input type="checkbox"/> Y/ <input type="checkbox"/> N
Transmission _____	Dose _____ MGy		anomalous? <input type="checkbox"/> Y/ <input type="checkbox"/> N

4. prefix _____

λ _____ Å	Δφ _____ °	_____	_____ Å resolution
dist _____ mm	exp _____ s	_____ / _____ / _____ Å	_____ % completeness
φ _____ - _____ °		_____ / _____ / _____ °	anisotropic? <input type="checkbox"/> Y/ <input type="checkbox"/> N
Transmission _____	Dose _____ MGy		anomalous? <input type="checkbox"/> Y/ <input type="checkbox"/> N

