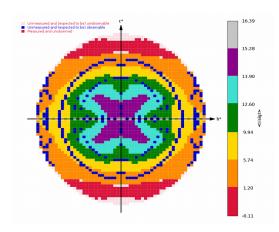
# ISPyB related work at Global Phasing

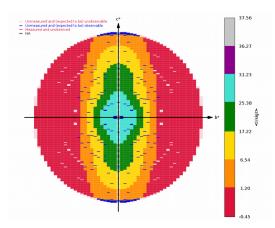
Clemens Vonrhein Global Phasing Ltd.

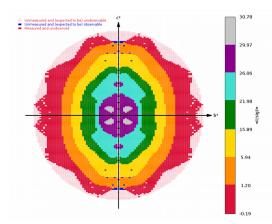
MXCube/ISPyB Meeting DLS 31.01.2018

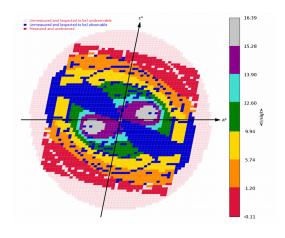


## **STARANISO**

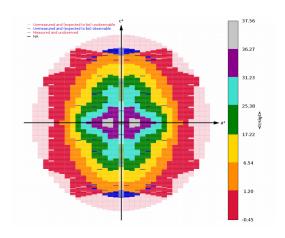








Plots of local <l/>for each r.l.p.





# autoPROC: merging statistics

□ autoPROC creates two types of merged data: traditional (isotropic) analysis of high-resolution limit: truncate.mtz new anisotropic analysis with STARANISO: staraniso alldata.mtz we tried to make it much clearer, what statistics relate to what output (merged) reflection data – especially in summary.html because autoPROC supports processing of multi-sweep datasets, these statistics are potentially given multiple times (persweep and final merging) since summary.html is also intended as a procedural record, statistics are given both during the scaling step and again at the end (producing final MTZ files) ... hopefully, this is not too confusing (we used colour to highlight) different types)



## autoPROC: "table 1" - traditional

#### 2.7. Finalising

Output file created (classical isotropic treatment of data) = <a href="mailto:truncate-unique.mtz">truncate-unique.mtz</a>

Spacegroup name P21

Unit cell parameters 50.68 89.80 60.47 90.000 94.778 90.000

Wavelength 0.97940 A

	0verall	InnerShell	OuterShell
Low resolution limit High resolution limit	89.800 1.937		
Rmerge (all I+ & I-) Rmerge (within I+/I-) Rmeas (within I+/I-) Rmeas (all I+ & I-) Rpim (within I+/I-) Rpim (all I+ & I-) Total number of observations Total number unique Mean(I)/sd(I) Completeness Multiplicity CC(1/2)	0.104 0.089 0.122 0.124 0.083 0.066 127299 37680 9.1 93.8 3.4 0.994		0.610 0.831 0.751 0.561 0.394 6869 1947 2.3
Anomalous completeness Anomalous multiplicity CC(ano)  DANO /sd(DANO)	83.2 1.8 0.121 0.890	84.2 1.8 0.479 1.402	1.9



# autoPROC: "table 1" - anisotropic

Final output file created (including analysis of anisotropy via <u>STARANISO</u>) = <u>staraniso\_alldata.mtz</u> (see above for detailed analysis of anisotropy.

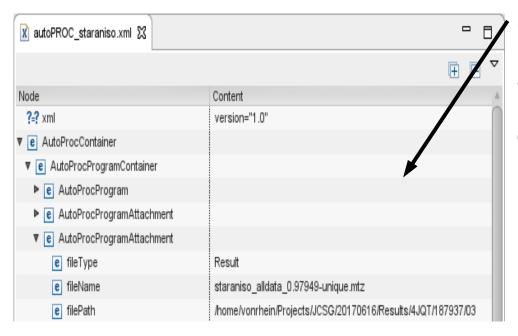
Spacegroup name P21 50.68 89.80 60.47 90.000 94.778 90.000 Unit cell parameters Wavelength 0.97940 A Resolution limits & eigenvectors of ellipsoid fitted to resolution cut-off surface: 2.163 0.9996 0.0000 0.0279 0.998 \_a\_\* - 0.066 \_c\_\* 1.658 0.0000 1.0000 0.0000 b \* 1.647 -0.0279 0.0000 0.9996  $-0.023 _a^* + _c^*$ Overall InnerShell OuterShell Low resolution limit 89.800 89.800 1.761 High resolution limit 1.647 5.075 1.647 Rmerge (all I+ & I-) 0.108 0.055 0.911 Rmerge (within I+/I-) 0.093 0.039 1.034 Rmeas (within I+/I-) 0.127 0.054 1.290 Rmeas (all I+ & I-) 0.129 0.066 1.082 0.085 0.038 0.766 Rpim (within I+/I-) Rpim (all I+ & I-) 0.069 0.035 0.576 Total number of observations 143816 7084 7292 Total number unique 43159 2150 2143 Mean(T)/sd(T)20.5 Completeness (spherical) 66.2 94.3 18.2 Completeness (ellipsoidal) 86.7 94.3 58.3 Multiplicity 3.3 3.3 3.4 CC(1/2)0.994 0.992 0.476 Anomalous completeness (spherical) 56.1 84.7 13.1 Anomalous completeness (ellipsoidal) 73.5 84.7 39.5 1.8 1.8 1.9 Anomalous multiplicity CC(ano) 0.112 0.410 -0.130 |DANO|/sd(DANO) 0.895 1.375 0.889



## autoPROC: STARANISO XML

XML file written as <a href="mailto:03/autoPROC.xml">03/autoPROC.xml</a> (for /home/vonrhein/Projects/JCSG/20170616/Results/4LII/177823/03/truncate-unique.mtz) isotropic analysis

XML file written as <u>03/autoPROC</u> <u>staraniso.xml</u> (for /home/vonrhein/Projects/JCSG/20170616/Results/4LII/177823 /03/staraniso alldata-unique.mtz)

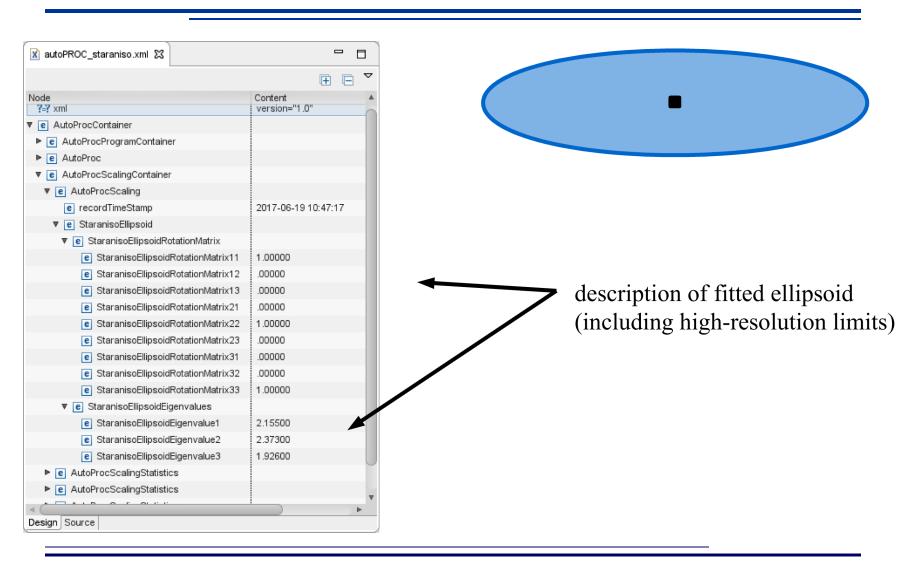


STARANISO mtz file (result)

all following statistics are based on **observations** (as defined by STARANISO)



## autoPROC: STARANISO XML





## autoPROC: STARANISO XML

