

Metadata for raw data from X-ray diffraction and other structural techniques  
Rovinj, Croatia, 2015

# X-ray Origins; Protection or Paranoia?

Natalie Johnson  
Newcastle University

# X-ray Origins

- Storing raw diffraction images could provide a ‘greater hurdle’ against false structure solutions.<sup>1</sup>
- This is only the case if the manufacture of raw data is sufficiently difficult.
- This talk covers:
  - The need for a method to validate our diffraction frames.
  - Suggestion of a method to ensure that the diffraction frames archived are from genuine experimental data.

## Anesthesiologist Fabricates 172 Papers

A researchers in Japan faked patient data on nearly 200 studies over the past 2 decades, according to an investigating committee.

By Jef Akst | July 3, 2012

NATURE NEWS BLOG

## Former chemistry grad student found guilty of fraud

**BBC NEWS**

## S Korea cloning research was fake

Research by South Korea's top human cloning scientist - hailed as a breakthrough earlier this year - was fabricated

Nature retracts epigenetics paper by author who lost two Science papers last year

Posted by Ivan Oransky

## Grad student blamed for research misconduct at Utah

13 November 2014

Rebecca Trager

Suspicious Data in a JACS Paper from 2009

November 7th, 2013

## Retraction of "Bimetallic Effects for Enhanced Polar Co-Enchainment Selectivity in Catalytic Ethylene Polymerization"

Brandon A. Rodriguez, Massimiliano Delferro, and Tobin J. Marks\*

*J. Am. Chem. Soc.* **2009**, *131*, 5902–5919. DOI: 10.1021/ja900257k

The authors have been unable to reproduce the synthesis and spectroscopic characterization of the ethylene/acrylate copolymer described in this article. Accordingly, the authors are retracting this publication due to concerns over the validity of the aforementioned data. The authors regret any confusion that may have been created by publication of this work.

In contrast, the results reported in related publications regarding the synthesis of a binuclear catalyst and its ethylene polymerization are unaffected and remain valid.

## Retraction: Histone methylation by the *Drosophila* epigenetic transcriptional regulator Ash1

Christian Beisel, Axel Imhof, Jaime Greene, Elisabeth Kremmer & Frank Sauer

*Nature* (2015) | doi:10.1038/nature14421

Published online 15 April 2015

 PDF  Citation  Reprints  Rights & permissions  Article metrics

*Nature* **419**, 857–862 (2002); doi:10.1038/nature01126

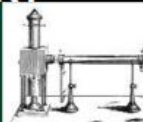
The authors and the University of California Riverside wish to retract this Letter owing to inappropriate image manipulation in the published figures. The figure panels affected are Figure 1b, d, Figure 2b, e, Figure 3a and Figure 4d. *Nature* has not received a response from Frank Sauer to approve this retraction.

Addition/Correction

## COMMENTARY

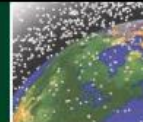
Helical spin order

338



Space junk

340



Bottling blue sky

344

LETTERS | POLICY FORUM | ESSAYS | BOOKS | PERSPECTIVES

## LETTERS

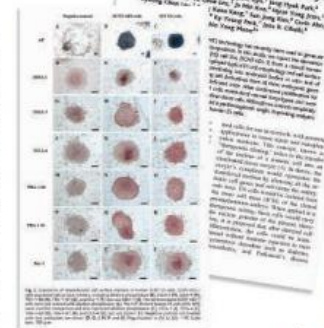
edited by Etta Kavanagh

### Editorial Retraction

#### INVESTIGATION COMMITTEE

(IC) has concluded that the authors of the paper in *Science* (2, 3) have provided data that are inconsistent with the published results and that the papers contain errors. The IC also found that the data showing the derivation of the pluripotent human embryonic stem cell line NT-1 is invalid because they are the result of a cloning procedure that is not consistent with the IC's findings. The IC also found that the authors of the paper in *Science* (2, 3) that 11 patient-specific pluripotent human embryonic stem cell lines were derived from a cloned blastocyst. According to the report of the IC, the laboratory "does not possess any scientific basis for the claim that the final product is a significant amount of the data presented in both papers. We believe that an immediate and unconditional retraction of both papers is warranted and advise the scientific community that the data presented in these papers are invalid."

All 15 authors of Hwang *et al.*, 2004 (2) have agreed to retract their paper. All 11 authors of Hwang *et al.*, 2005 (3) have agreed to retract their paper. The IC also found that the authors of the paper in *Science* (2, 3) that 11 patient-specific pluripotent human embryonic stem cell lines were derived from a cloned blastocyst. According to the report of the IC, the laboratory "does not possess any scientific basis for the claim that the final product is a significant amount of the data presented in both papers. We believe that an immediate and unconditional retraction of both papers is warranted and advise the scientific community that the data presented in these papers are invalid."



advocating not pure bottom-up divergence but a new Constitution replacing the American Confederation with an overarching institutional framework to govern the world.

Victor *et al.* also argue that the plurilateral approach advocated by the Kyoto and Wiener (1). This involves building a coalition of nations to create parallel institutions of international emission trading. It would not be a "top-down" approach, nor would it approach the degree of centralization in Madison's version of federalism. It would avoid the difficulties of establishing a single universal trading system under the Kyoto Protocol and focus on cooperation among major emitting countries, as advocated by Victor *et al.*

The plurilateral approach would foster a variety of trading systems to encourage precisely the kind of experimentation in policy and practice that Victor *et al.* favor. But it would also avoid binding mutual commitments by parties to the agreement. In contrast to both Madisonian federalism and plurilateral trading, Victor *et al.* favor a bottom-up local policies that they

advocate. The plurilateral approach would foster a variety of trading systems to encourage precisely the kind of experimentation in policy and practice that Victor *et al.* favor. But it would also avoid binding mutual commitments by parties to the agreement.

In contrast to both Madisonian federalism and plurilateral trading, Victor *et al.* favor a bottom-up local policies that they

## addenda and errata



Acta Crystallographica Section D

**Biological  
Crystallography**

ISSN 0907-4449

## addenda and errata

### Retraction of articles by H. M. Krishna Murthy *et al.*

Two papers by H. M. Krishna Murthy *et al.* are retracted.

Two papers by H. M. Krishna Murthy *et al.* (Krishna Murthy *et al.* 1999; Urs *et al.*, 1999) are retracted by the journal. This follows investigation by the University of Alabama at Birmingham, Alabama USA, of structures deposited by H. M. Krishna Murthy. Krishna Murthy has noted that he is not in agreement with the retraction:

#### References

Krishna Murthy, H. M., Judge, K., DeLucas, L., Clum, S. & Padmanabhan, (1999). *Acta Cryst. D55*, 1370–1372.  
Urs, U. K., Murali, R. & Krishna Murthy, H. M. (1999). *Acta Cryst. D55*, 1977.



Acta Crystallographica Section E

**Structure Reports  
Online**

ISSN 1600-5368

### Retraction of articles by T. Liu *et al.*

T. Liu,<sup>a\*</sup> Y.-X. Wang,<sup>b</sup> Z.-W. Wang,<sup>a</sup> Z.-P. Xie<sup>a,c</sup> and J. Y. Zhu<sup>d</sup>

<sup>a</sup>College of Engineering, Jingtangshan University, Jian 343009, People's Republic of China, <sup>b</sup>College of Mathematics and Physics, Jingtangshan University, Jian 343009, People's Republic of China, <sup>c</sup>Department of Chemistry, Jiangxi University of Science and Technology, Ganzhou 341000, People's Republic of China, and <sup>d</sup>Department of Information Engineering, Jiangxi University of Science and Technology, Nanchang 330013, People's Republic of China  
Correspondence e-mail: taoliu07@126.com

Received 20 November 2009; accepted 15 December 2009

A series of 29 papers by Liu *et al.* are retracted.

As a result of problems with the data sets and incorrect atom assignments, 29 papers by Liu *et al.* are retracted. Full details of all the articles are given in Table 1.

#### Table 1

Details of articles to be retracted, in order of publication.

Title	Reference	DOI	Refcode
<i>Tetrakis</i> (pyrazine- <i>c</i> , <i>N</i> )bis(thiocyanato- <i>c</i> , <i>N</i> )manganese(II)	Liu & Xie (2007a)	10.1107/S1600536807026852	EDUMAS
<i>(Dihydroxyglyoxime-<math>\kappa^2</math>,<i>N,N'</i>)bis(1,10-phenanthroline-<math>\kappa^2</math>,<i>N,N'</i>)copper(II) dinitrate dihydrate</i>	Liu, Wang, Wang & Xie (2007b)	10.1107/S1600536807026255	EDUVAB
<i>Tetrakis</i> (pyrazine- <i>c</i> , <i>N</i> )bis(thiocyanato- <i>c</i> , <i>N</i> )zinc(II)	Liu & Xie (2007b)	10.1107/S1600536807028735	RIGQAA
<i>Tetrakis</i> ( $\mu$ -2-pyridylglyoxacetato)bis(1,10-phenanthroline)(2-pyridylglyoxacetato)-lanthanum(III)	Liu, Wang, Wang & Xie (2007c)	10.1107/S1600536807030917	UDUMIQ
<i>Polymeric KNOF<sub>2</sub></i>	Liu, Wang, Wang & Xie (2007a)	10.1107/S1600536807027195	ICSD 240891

## addenda and errata

Acta Crystallographica Section E

**Structure Reports  
Online**

ISSN 1600-5368

### Retraction of articles by H. Zhong *et al.*

H. Zhong,<sup>a\*</sup> S.-H. Duan,<sup>a</sup> Y.-P. Hong,<sup>a</sup> M.-L. Li,<sup>a</sup> Y.-Q. Liu,<sup>a</sup> C.-J. Luo,<sup>a</sup> Q.-Y. Luo,<sup>a</sup> S.-Z. Xiao,<sup>a</sup> H.-L. Xie,<sup>a</sup> Y.-P. Xu,<sup>a</sup> X.-M. Yang,<sup>b,a</sup> X.-R. Zeng<sup>a</sup> and Q. Y. Zhong<sup>c</sup>

<sup>a</sup>College of Chemistry and Chemical Engineering, Provincial Key Laboratory of Coordination Chemistry, Jingtangshan University, Jian 343009, People's Republic of China, <sup>b</sup>Institute of Applied Materials, Jiangxi University of Finance and Economics, Nanchang 330032, People's Republic of China, and <sup>c</sup>Jian Training School, Jian 343000, People's Republic of China  
Correspondence e-mail: huazhong06@126.com

Received 20 November 2009; accepted 15 December 2009

A series of 41 papers by H. Zhong *et al.* are retracted.

As a result of problems with the data sets and incorrect atom assignments, 41 papers by H. Zhong *et al.* are retracted. Full details of all the articles are given in Table 1.

#### Table 1

Details of articles to be retracted, in order of publication.

Title	Reference	DOI	Refcode
<i>Aquachlorobis(1,10-phenanthroline)cobalt(II) chloride thiourea solvate</i>	Zhong, Zeng, Liu & Luo (2006a)	10.1107/S1600536806041122	KERQEE
<i>cis-Dichlorobis(1,10-phenanthroline)cobalt(II)</i>	Zhong, Zeng & Luo (2006)	10.1107/S1600536806047295	MEQFOE
<i>Tris(quinolin-8-olato-<math>\kappa^2</math>,<i>N,O</i>)cobalt(III) glyoxal hemisolvate monohydrate</i>	Zhong, Zeng, Liu & Luo (2006b)	10.1107/S1600536806050240	MEQHEW
<i>(8-Quinolino-<math>\kappa^2</math>,<i>N,O</i>)bis(8-quinolinolato-<math>\kappa^2</math>,<i>N,O</i>)nickel(II) glyoxal hemisolvate monohydrate</i>	Zhong, Zeng, Liu & Luo (2007)	10.1107/S1600536806053232	METVUD
<i>Aquachlorobis(1,10-phenanthroline)cobalt(II) chloride thioacetamide solvate</i>	Zhong, Zeng & Luo (2007)	10.1107/S1600536806053530	METQIM
<i>(8-Quinolino-<math>\kappa^2</math>,<i>N,O</i>)bis(8-quinolinolato-<math>\kappa^2</math>,<i>N,O</i>)zinc(II) glyoxal hemisolvate monohydrate</i>	Zhong, Zeng, Luo, Li & Xiao (2007)	10.1107/S1600536807001171	DEXTEG
<i>(Dimethylglyoxime-<math>\kappa^2</math>,<i>N,N'</i>)bis(1,10-phenanthroline-<math>\kappa^2</math>,<i>N,N'</i>)nickel(II) dinitrate dihydrate</i>	Zhong, Zeng, Yang, Luo & Li (2007a)	10.1107/S1600536807004102	YEYGOZ

- Discovered during the testing of *checkCIF*.
- Structure factor files were discovered to be almost identical.
- One set of intensities were taken and refined, unit cell size and elements within were manually changed to produce data for other 'structures'.



editorial

Acta Crystallographica Section E  
Structure Reports  
Online  
ISSN 1600-5368

**Editorial**

**William T. A. Harrison,<sup>a</sup> Jim Simpson<sup>b</sup> and Matthias Weil<sup>c</sup>**

<sup>a</sup>Department of Chemistry, University of Aberdeen, Aberdeen AB24 3UE, Scotland, <sup>b</sup>Department of Chemistry, University of Otago, PO Box 56, Dunedin, New Zealand, and <sup>c</sup>Institute of Chemical Technologies and Analytics, Division of Structural Chemistry, Vienna University of Technology, Getreidemarkt 9/164-SC, Austria

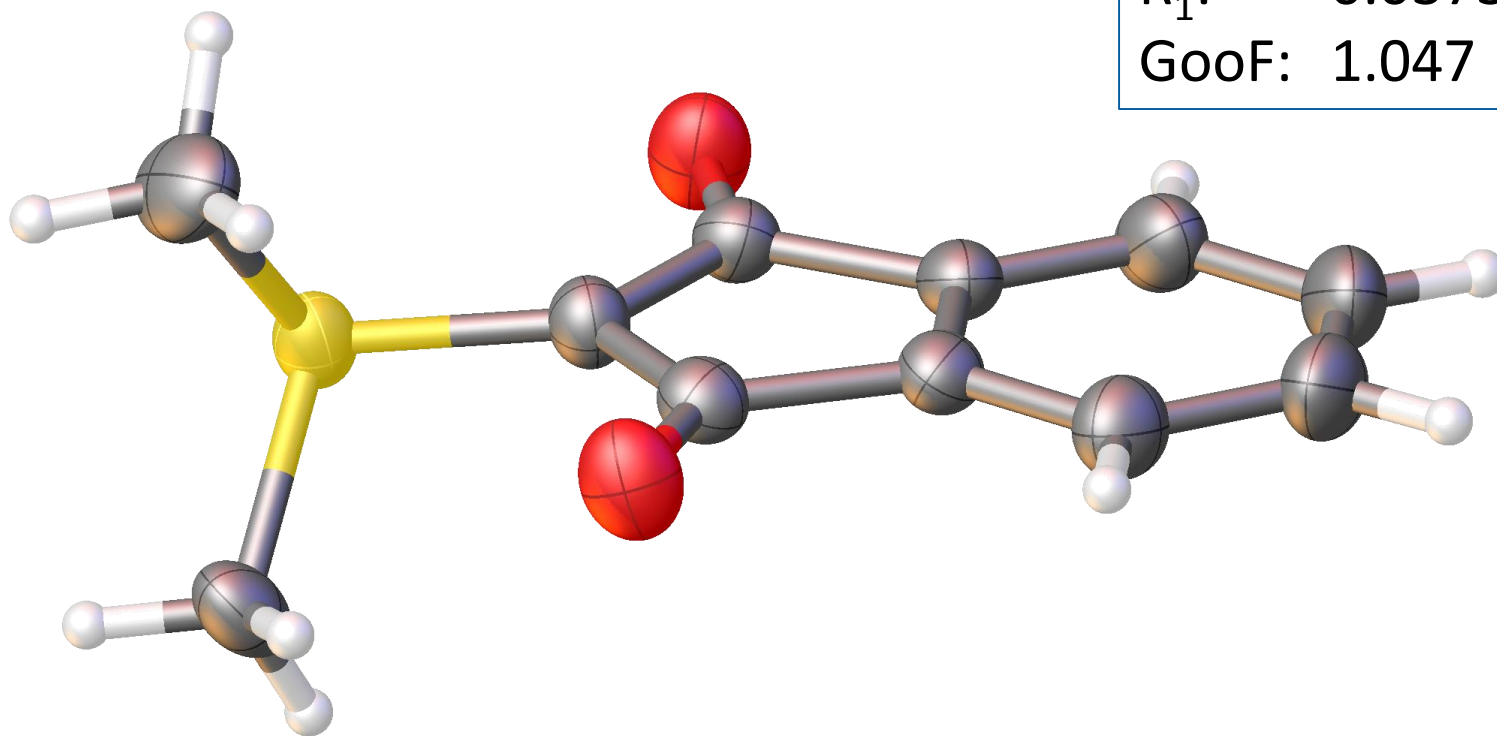
Regrettably, this editorial is to alert readers and authors of *Acta Crystallographica Section E* and the wider scientific community to the fact that we have recently uncovered evidence for an extensive series of scientific frauds involving papers published in the journal, principally during 2007. Although several thousands of structures published in *Acta Crystallographica Section E* every year will continue to reflect results of serious scientific work, the extent of these problems is significant with at least 70 structures demonstrated to be falsified and meanwhile acknowledged by the authors as such. Our work is ongoing and it is likely that this figure will rise further.

These problems were first discovered by Ton Spek during testing of the checking programs for the journal. Testing is routinely carried out using cifs and structure-factor files from back issues of *Acta Crystallographica Sections E* or *C*. Initially, unexplained Hirshfeld rigid-bond alerts and unusual metal–ligand donor-atom distances led to the discovery that metal atoms had been transposed and that more than one structure had been ‘determined’ using identical sets of data. Investigation of these cases sparked a search of papers written by the correspondence authors involved.

A program written by Toine Schreurs of Utrecht University that can examine and compare two structure-factor files was then used to examine the data deposited for the structures under investigation. For all of the problem structures, the program revealed that the data sets used to refine two or more supposedly unique structures were in fact identical, but with the cell parameters apparently manually altered by the authors concerned.

The falsified structures have many features in common: in each case, a *bona fide* set of intensity data, usually on a compound whose structure had been correctly determined and reported in the literature, was used to produce a number of papers, with the authors changing one or more atoms in the structure to produce what appeared to be a genuine structure determination of a new compound. The worst example generated no fewer than 18 supposedly original structures from a single common set of data. There is nothing to suggest that the authors of the original papers describing the real structures are in any way aware of, or complicit in, this fraud.

W. T. A. Harrison, Jim Simpson, and  
Matthias Weil, *Acta Crystallogr., Sect.  
E*, 2010, **66**, e1-e2



$R_1$ : 0.0373  
GooF: 1.047

R(reflections) = 0.0279 ( 1749)      wR2(reflections) = 0.0787 ( 1799)

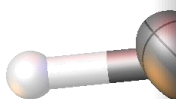
S = 1.100

Npar = 129

**Alert level G**

PLAT912 ALERT 4 G Missing # of FCF Reflections Above STh/L= 0.600      111 Note

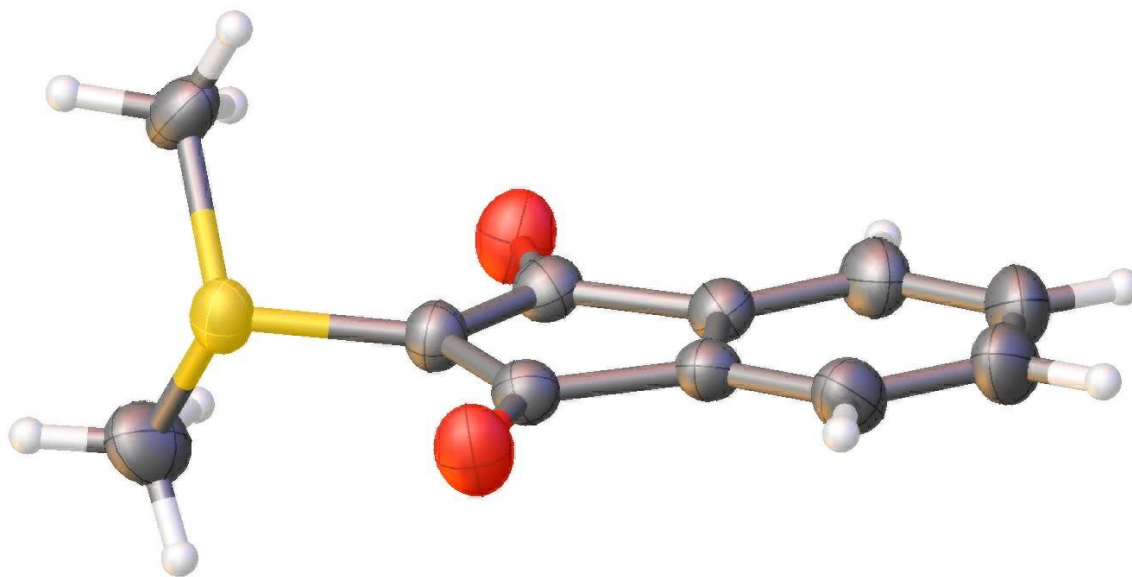
- 0 ALERT level A = Most likely a serious problem - resolve or explain
  - 0 ALERT level B = A potentially serious problem, consider carefully
  - 0 ALERT level C = Check. Ensure it is not caused by an omission or oversight
  - 1 ALERT level G = General information/check it is not something unexpected
- 
- 0 ALERT type 1 CIF construction/syntax error, inconsistent or missing data
  - 0 ALERT type 2 Indicator that the structure model may be wrong or deficient
  - 0 ALERT type 3 Indicator that the structure quality may be low
  - 1 ALERT type 4 Improvement, methodology, query or suggestion
  - 0 ALERT type 5 Informative message, check

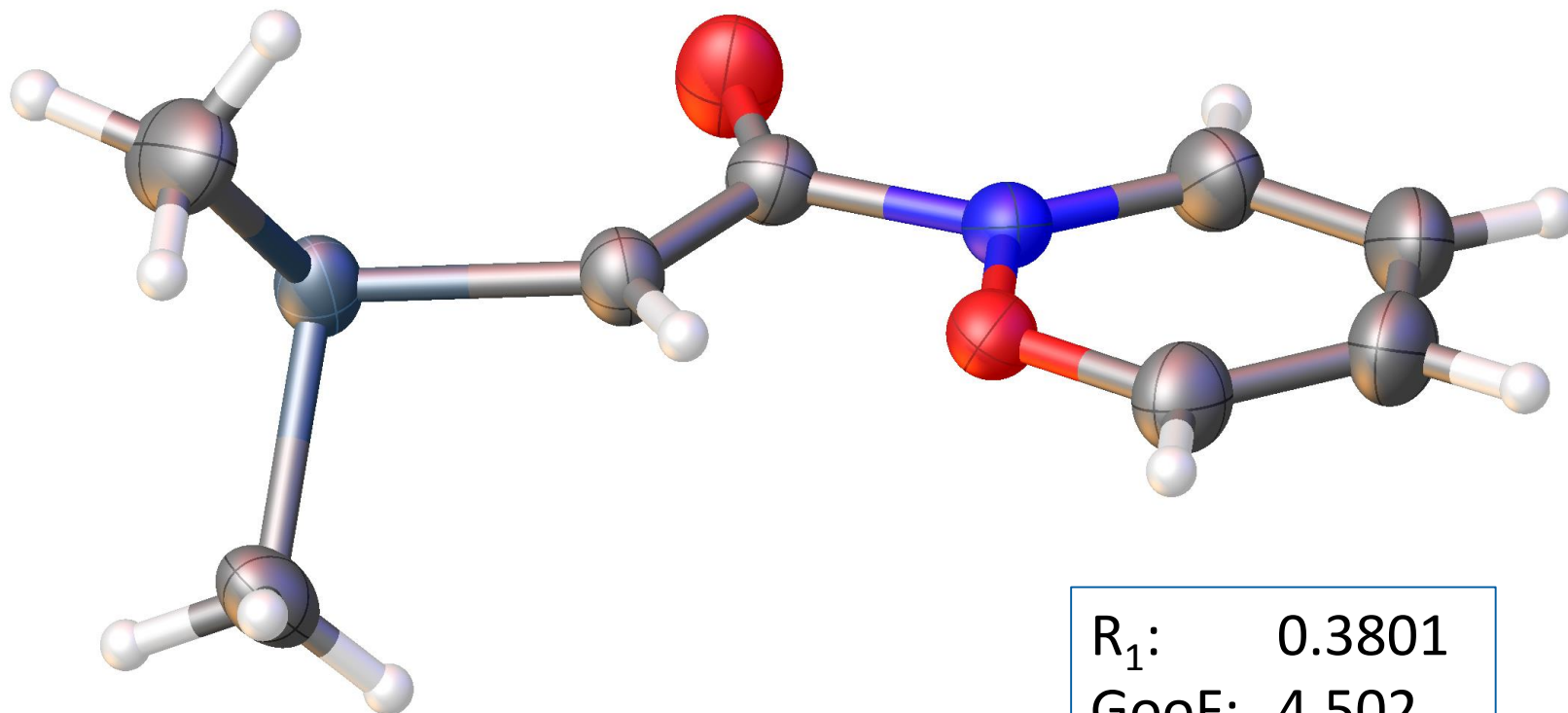


It is advisable to attempt to resolve as many as possible of the alerts in all categories. Often the minor alerts point to easily fixed oversights, errors and omissions in your CIF or refinement strategy, so attention to these fine details can be worthwhile. In order to resolve some of the more serious problems it may be necessary to carry out additional measurements or structure refinements. However, the purpose of your study may justify the reported deviations and the more serious of these should normally be commented upon in the discussion or experimental section of a paper or in the "special\_details" fields of the CIF. checkCIF was carefully designed to identify outliers and unusual parameters, but every test has its limitations and alerts that are not important in a particular case may appear. Conversely, the absence of alerts does not guarantee there are no aspects of the results needing attention. It is up to the individual to critically assess their own results and, if necessary, seek expert advice.





File Edit View Structure Mode Tools Model Select Help





$R_1$ :	0.3801
GooF:	4.502



---

32 **ALERT level A** = Most likely a serious problem - resolve or explain  
22 **ALERT level B** = A potentially serious problem, consider carefully  
72 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight  
5 **ALERT level G** = General information/check it is not something unexpected

18 **ALERT type 1** CIF construction/syntax error, inconsistent or missing data  
60 **ALERT type 2** Indicator that the structure model may be wrong or deficient  
12 **ALERT type 3** Indicator that the structure quality may be low  
41 **ALERT type 4** Improvement, methodology, query or suggestion  
0 **ALERT type 5** Informative message, check

---



$R_1$ :	0.3801
GooF:	4.502

Sample Instrument Windows Help

lations\ECM25\Worsyld\_try375\_yld\_01\_0215.sfrm

Setup  
Evaluate

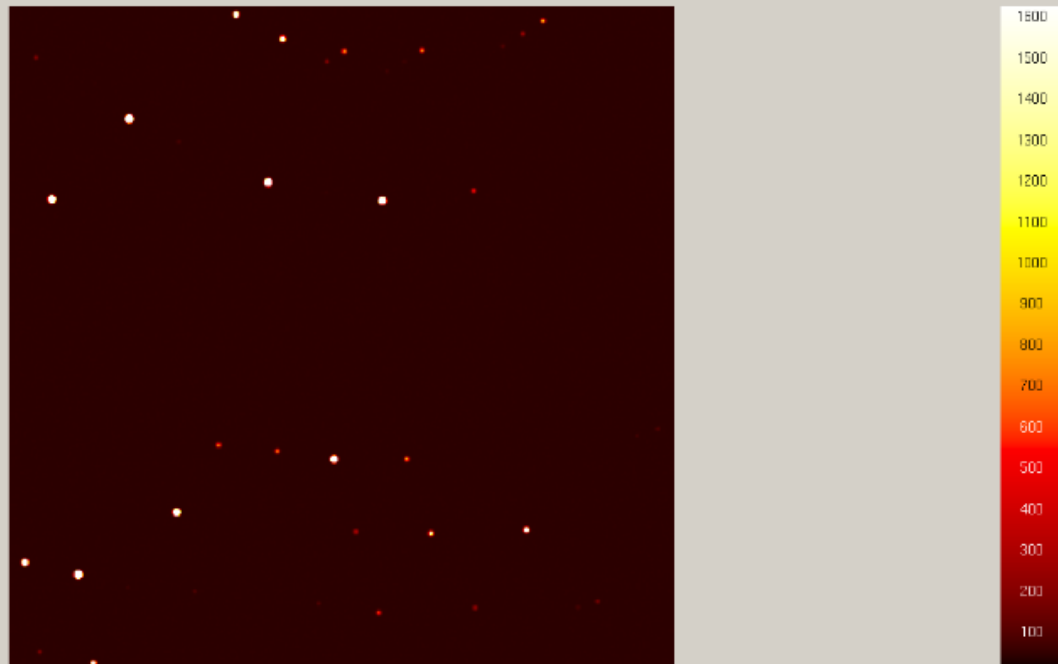
Determine Unit Cell

Transform Unit Cell

Compare Unit Cells

Reciprocal Lattice Viewer

View Images



Automatic Mode

Start at: Collect Data

Stop after: Search

Run

Manual Mode

Collect Data

Harvest Spots

Index

Traverse

Refine

Search

Unit cells

a = 5.95Å, α = 90.00°, V = 966Å<sup>3</sup>

b = 9.02Å, β = 90.00°, Orthorhombic P

c = 10.36Å, γ = 90.00°

Edit...  
Delete  
Delete All

Reflections:

Group 0: 8021 reflections

Edit...  
Delete  
Delete All

Expected resolution:

Exposure time [s]	Resolution [Å]
1	5.0
2	20.0
3	60.0
4	120.0
5	600.0

Crystal Mosaicity [°] 0.72

Cursor

Position [mm]	30.60	-15.95
Position [pixels]	511	122
Intensity [counts]	33	
HKL index	-2.00	0.07
Resolution [Å]	0.70	
2Theta [°]	47.35	

Collect

Integrate

Scale

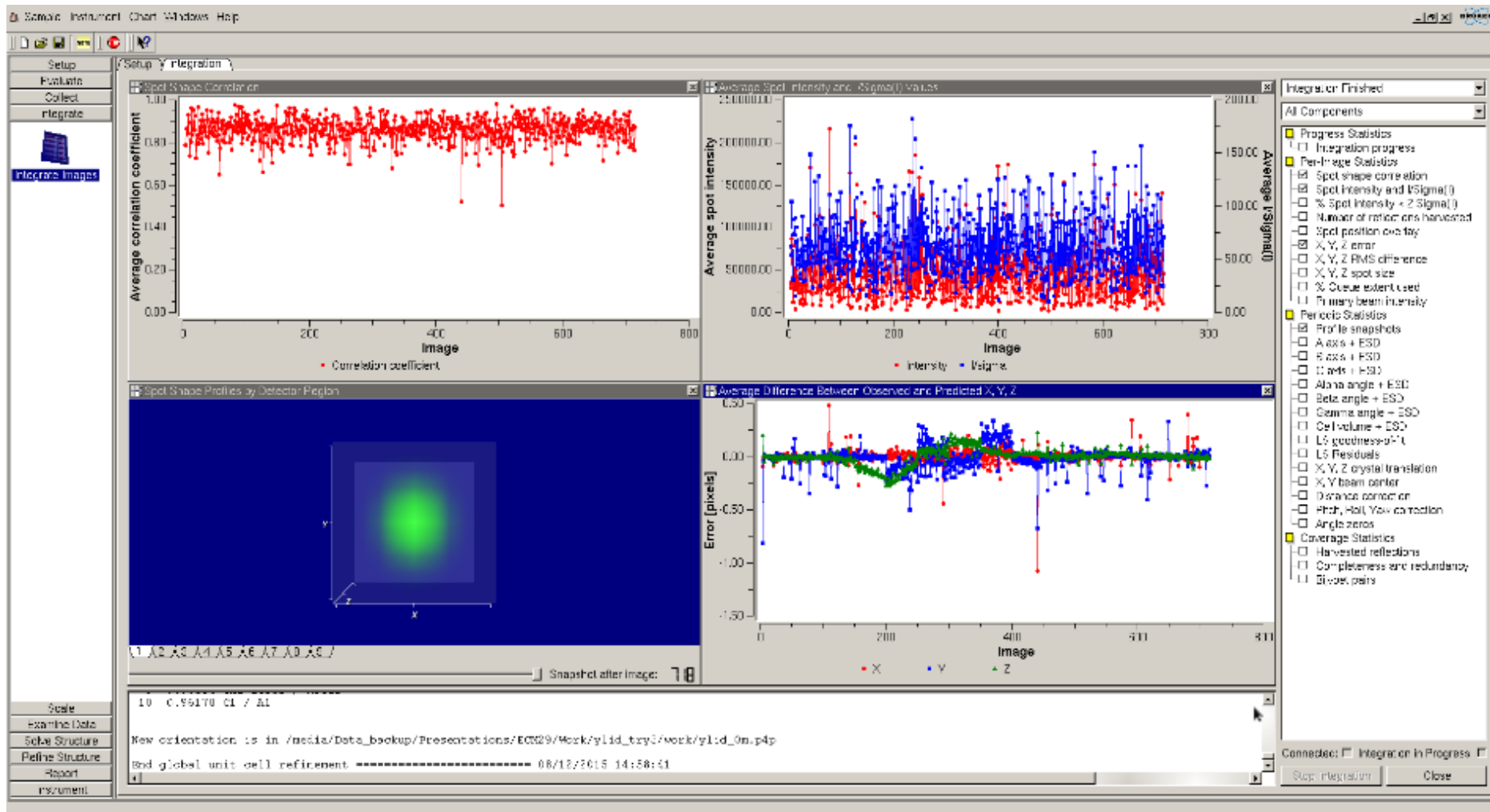
Examine Data

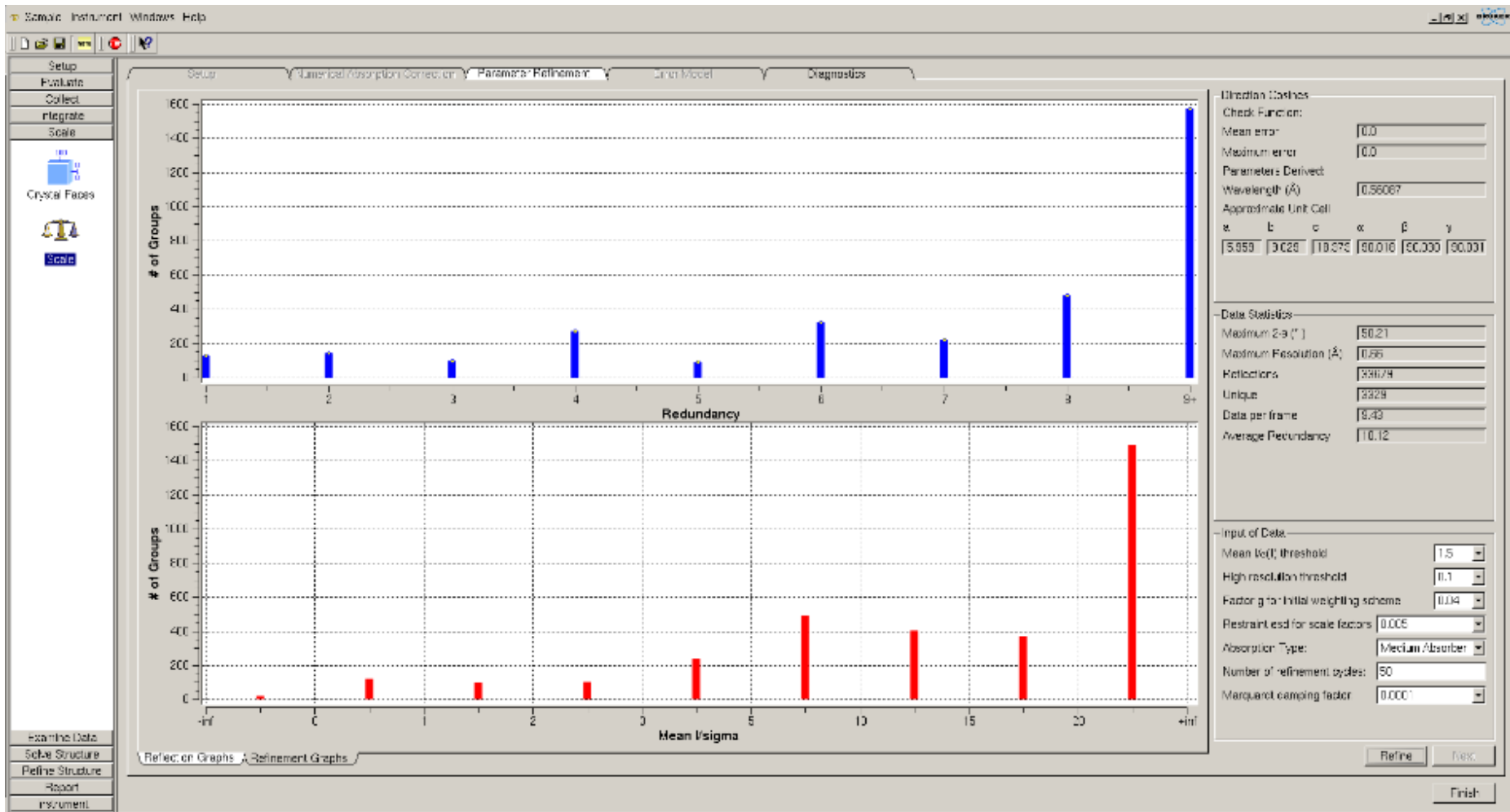
Solve Structure

Refine Structure

Report

Instrument





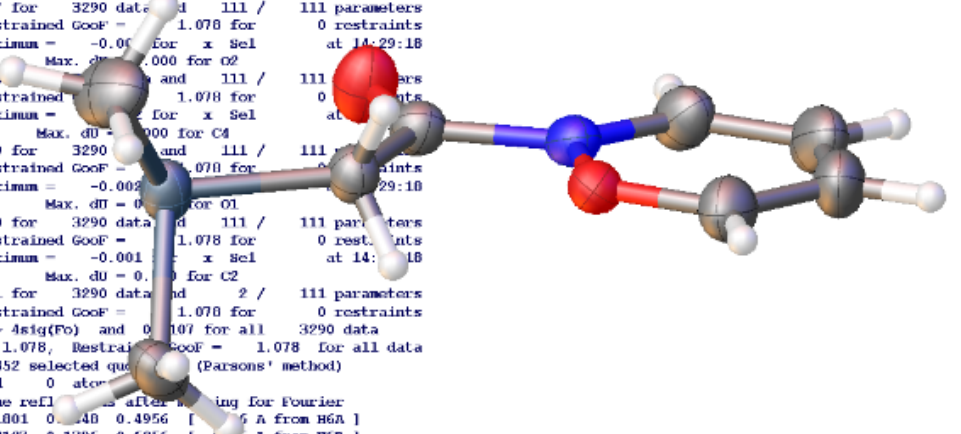
Olex2

File Edit View Structure Mode Tools Model Select Help

```

wR2 = 0.0352 before cycle 3 for 3290 data and 111 / 111 parameters
Goof = S = 1.077; Restrained Goof = 1.077 for 0 restraints
Mean shift/esd = 0.000 Maximum = -0.003 for z C8
Max. shift = 0.000 A for H8C Max. du = 0.000 for C8
wR2 = 0.0353 before cycle 4 for 3290 data and 111 / 111 parameters
Goof = S = 1.078; Restrained Goof = 1.078 for 0 restraints
Mean shift/esd = 0.000 Maximum = -0.001 for x Se1
Max. shift = 0.000 A for H7B Max. du = 0.000 for C1
wR2 = 0.0352 before cycle 5 for 3290 data and 111 / 111 parameters
Goof = S = 1.077; Restrained Goof = 1.077 for 0 restraints
Mean shift/esd = 0.000 Maximum = -0.001 for x Se1
Max. shift = 0.000 A for H7B Max. du = 0.000 for C8
wR2 = 0.0353 before cycle 6 for 3290 data and 111 / 111 parameters
Goof = S = 1.078; Restrained Goof = 1.078 for 0 restraints
Mean shift/esd = 0.000 Maximum = -0.001 for x Se1
Max. shift = 0.000 A for O2 Max. du = 0.000 for C8
wR2 = 0.0353 before cycle 7 for 3290 data and 111 / 111 parameters
Goof = S = 1.078; Restrained Goof = 1.078 for 0 restraints
Mean shift/esd = 0.000 Maximum = -0.001 for x Se1
Max. shift = 0.000 A for H7A Max. du = 0.000 for O2
wR2 = 0.0353 before cycle 8 for 3290 data and 111 / 111 parameters
Goof = S = 1.078; Restrained Goof = 1.078 for 0 restraints
Mean shift/esd = 0.000 Maximum = -0.001 for x Se1
Max. shift = 0.000 A for C4 Max. du = 0.000 for C4
wR2 = 0.0353 before cycle 9 for 3290 data and 111 / 111 parameters
Goof = S = 1.078; Restrained Goof = 1.078 for 0 restraints
Mean shift/esd = 0.000 Maximum = -0.002 for x Se1
Max. shift = 0.000 A for H7B Max. du = 0.000 for O1
wR2 = 0.0353 before cycle 10 for 3290 data and 111 / 111 parameters
Goof = S = 1.078; Restrained Goof = 1.078 for 0 restraints
Mean shift/esd = 0.000 Maximum = -0.001 for x Se1
Max. shift = 0.000 A for H8A Max. du = 0.000 for C2
wR2 = 0.0353 before cycle 11 for 3290 data and 111 / 111 parameters
Goof = S = 1.078; Restrained Goof = 1.078 for 0 restraints
R1 = 0.0107 for 3263 Fo > 4sig(Fo) and 0.0107 for all 3290 data
wR2 = 0.0353, Goof = S = 1.078, Restrained Goof = 1.078 for all data
Flack x = -0.002(1) from 1352 selected quot (Parsons' method)
0 atoms may be split and 0 atoms
R1 = 0.0114 for 1918 unique refls after merging for Fourier
Highest peak 0.49 at 0.1001 0.440 0.4956 [ 0.6 A from H6A ]
Deepest hole -0.47 at 0.2127 0.1396 0.6256 [ 0.5 A from H6B ]
+-----+
+ t5_ylid_0m_b finished at 14:29:18 Total elapsed time: 0.25 secs +
+-----+
Checking absolute structure...
Boofft y: 0.0140(13)
Flack x: -0.002(1)
OK
Skipping 'space_group_crystal_system'
Skipping 'space_group_name_H_M_all'
Refinement CIF file has been merged with the meta-data cif file
>>|

```



t5\_ylid\_0m\_b P2<sub>1</sub>2<sub>1</sub>1

/media/Data\_backup/Presentations/EC...ork/ylid\_try3/work/t5\_ylid\_0m\_b.ins

C<sub>8</sub>H<sub>12</sub>NO<sub>2</sub>Se

a = 9.0257(4) α = 90° Z = 4  
 b = 18.3726(9) β = 90° Z' = 1  
 c = 5.9550(3) γ = 90° V = 987.49(8)

R1 = 1.07%

Shift/Std = -0.001 Max Peak 0.5 Min Peak -0.5 Goof 1.078

Home Work View Tools Info

Solve Refine Report

Refinement Program XL Method Least Squares

Reflection File t5\_ylid\_0m\_b.hkl

Max. refinement cycles and peaks Cycles 10 Peaks 20

Weight: 0.026(0.026) | 0.031(0.031) Auto-update when R1 < 15.0 %

Extinction correction n/a CONF, MORE -1, Bond SH, ACTA

Use solvent mask Recalculate mask

Refinement Settings Extra

Toolbox Work

Labels Labels OFF/ON

C H N O Se ... Add H ...

Z = 1

Split atoms you click next with No Restraints SADF ISOB SIMU

Select group or atom(s) and then Split Fit Split or Move with SHIFT key

Electron Density Map

Peak & Uiso Sliders

Growing

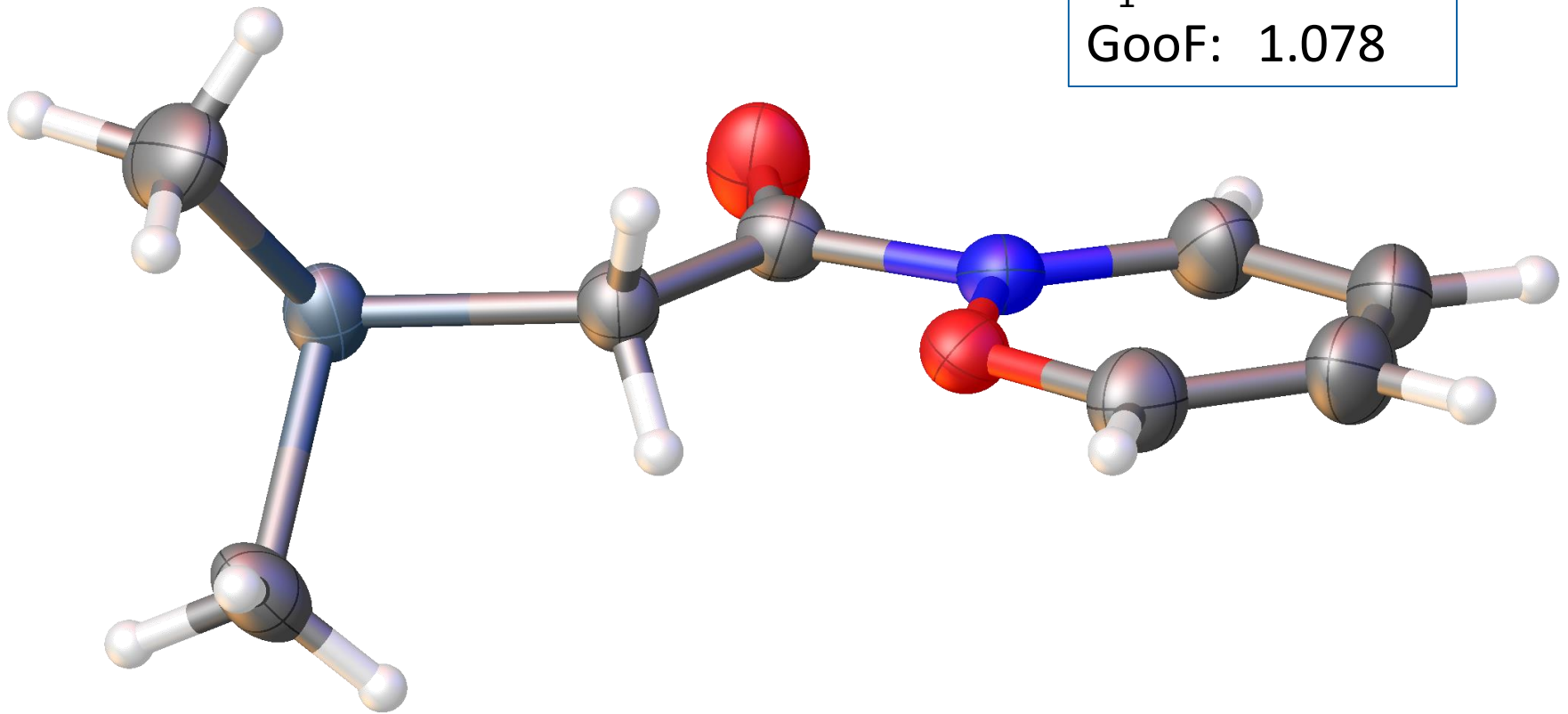
Finishing

History

Select

Naming

Sorting



$R_1$ : 0.0107  
GooF: 1.078



$R_1$ : 0.0107  
Goof: 1.078

**Alert level C**

PLAT911 ALERT 3 C Missing # FCF Refl Between THmin & STh/L= 0.600 9 Report

- 
- 0 ALERT level A = Most likely a serious problem - resolve or explain
  - 0 ALERT level B = A potentially serious problem, consider carefully
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  - 0 ALERT type 2 Indicator that the structure model may be wrong or deficient
  - 1 ALERT type 3 Indicator that the structure quality may be low
  - 0 ALERT type 4 Improvement, methodology, query or suggestion
  - 0 ALERT type 5 Informative message, check
-

# How is it done?

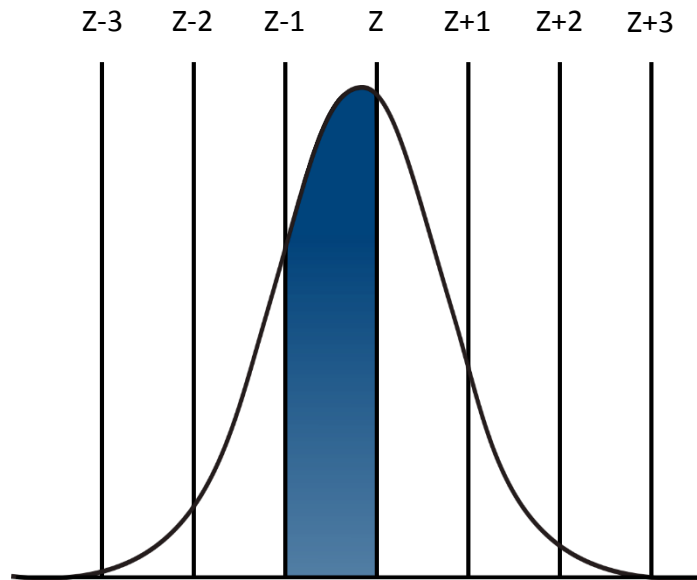
# .raw file

Integrated Intensity

X, Y, Z coordinates of spots

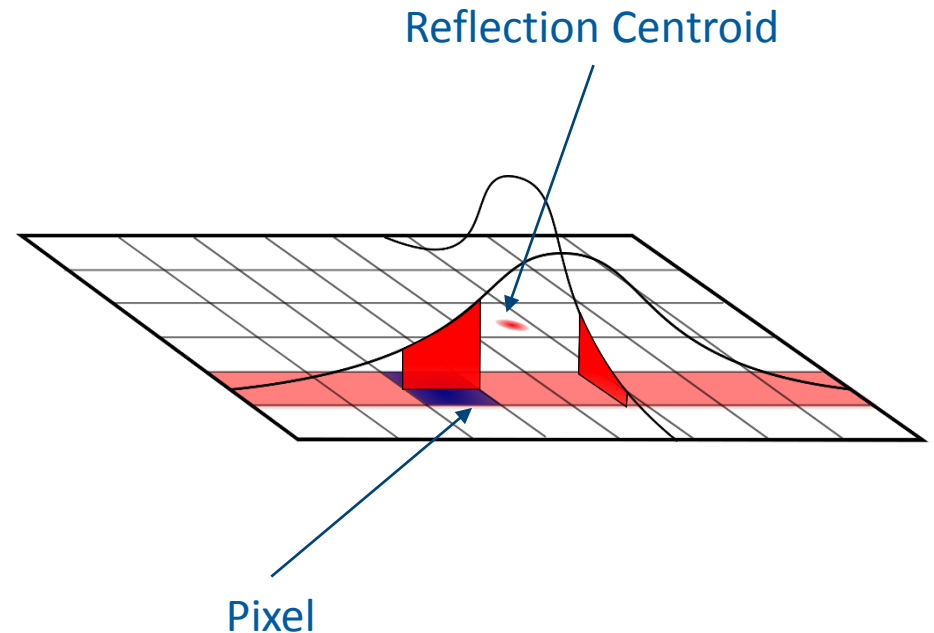
0	0	107.342	7.680	1	0.99166-0.98834-0.11839	0.11839	0.02248	0.02248	0	530.00	259.88	236.68	329.94	259.61	236.74	0.046	0.75	0.33	10.00	16.09	3	316	537	1352	118.37	1	0.984	536346112	2	0.044	76.51	3.90	→	
0	0	-311882.	1679.	1	0.98734-0.99399-0.10571	0.10571	0.04497	0.04497	0	411.00	245.00	228.83	410.94	245.09	228.87	0.091	0.89	0.32	10.00	12.16	3	633	609844	1401	114.44	1	0.992	536346113	-2	0.088	157.51	-10.98	→	
0	0	329459.	1278.	1	0.99206-0.98542-0.12249	0.12249	0.04497	0.04497	0	303.87	264.13	239.38	303.66	264.33	239.36	0.092	0.81	0.33	10.00	17.44	3	633	622508	1352	119.68	1	0.977	536346113	2	0.088	50.38	8.15	→	
0	0	-310.665	21.830	1	0.98486-0.99483-0.10137	0.10137	0.06745	0.06745	0	439.02	240.02	226.31	439.11	240.01	226.25	0.137	0.91	0.32	10.00	10.91	3	949	518	1309	113.12	1	1.000	536346114	-2	0.133	185.53	-15.96	→	
0	0	3	54.538	8.462	1	0.99194-0.98198-0.12655	0.12655	0.06745	0.06745	0	277.84	268.98	241.83	277.54	269.03	241.99	0.138	0.75	0.34	10.00	18.67	3	949	151	1353	120.99	1	0.969	536346114	2	0.133	24.35	13.00	→
0	0	-4	985093.	2184.	1	0.98186-0.99515-0.09700	0.09700	0.08994	0.08994	0	468.00	235.00	223.58	468.12	234.74	223.62	0.183	0.90	0.31	10.00	9.54	3	1265	943725	1341	111.81	1	1.000	536346115	-2	0.176	214.51	-20.98	→
0	0	4	906564.	3250.	1	0.99131-0.97803-0.13044	0.13044	0.08994	0.08994	0	251.87	273.89	244.58	251.46	273.74	244.61	0.185	0.80	0.34	10.00	20.04	3	1265	839969	1252	122.31	1	0.988	536346115	2	0.176	-1.62	17.91	→
0	0	-5	964.800	52.778	1	0.97834-0.99495-0.09252	0.09252	0.11242	0.11242	0	498.02	228.99	220.91	498.14	229.27	220.98	0.229	0.90	0.31	10.00	8.21	3	1582	833	1326	110.49	1	0.986	536346116	-2	0.220	244.53	-26.99	→
0	0	5	340.585	29.445	1	0.99015-0.97354-0.13438	0.13438	0.11242	0.11242	0	225.01	278.86	247.26	225.29	278.50	247.25	0.233	0.71	0.34	10.00	21.38	3	1582	363	1362	123.63	1	0.996	536346116	2	0.220	-28.48	22.88	→
0	0	6	1.07166	3212.99	1	0.98853-0.96854-0.13779	0.13779	0.13493	0.13493	0	199.00	283.12	249.83	198.92	283.34	249.88	0.281	0.89	0.35	10.00	22.66	3	1898	648111	1329	124.94	1	0.976	536346117	2	0.263	-54.49	27.14	→
0	0	7	342.115	32.955	1	0.98630-0.96298-0.14161	0.14161	0.15741	0.15741	0	172.01	288.13	252.50	172.22	288.27	252.53	0.331	0.85	0.35	10.00	24.00	3	2215	314	1321	126.27	1	0.997	536346118	2	0.306	-81.48	32.15	→
0	0	8	9404.6	219.2	1	0.98354-0.95698-0.14535	0.14535	0.17990	0.17990	0	145.01	293.06	255.24	145.05	293.34	255.19	0.382	0.88	0.36	10.00	25.37	3	2531	4426	1417	127.60	1	0.999	536346119	2	0.348	-108.48	37.08	→
0	0	9	0.21114	5.9597	1	0.98023-0.95025-0.14902	0.14902	0.20239	0.20239	0	117.35	298.81	257.91	117.29	298.59	257.86	0.434	0.02	0.36	10.00	26.71	3	2848	104	1302	128.93	1	0.992	536346120	2	0.390	-136.14	42.83	→
0	0	10	2.16366	792.28	1	0.97637-0.94305-0.15266	0.15266	0.22488	0.22488	0	88.99	304.00	260.52	88.77	304.05	260.55	0.487	0.91	0.36	10.00	28.01	3	3164	771529	1415	130.27	1	1.000	536346121	2	0.431	-164.50	48.02	→
0	0	11	1040.6	78.7	1	0.97194-0.93530-0.15620	0.15620	0.24737	0.24737	0	59.02	309.95	263.27	59.31	309.78	263.25	0.543	0.87	0.37	10.00	29.39	3	3480	497	1329	131.62	1	0.999	536346122	2	0.472	-194.47	53.97	→
0	0	12	524050.	2095.	1	0.96695-0.92697-0.15969	0.15969	0.26985	0.26985	0	28.99	316.00	265.89	28.74	315.83	265.96	0.600	0.91	0.37	10.00	30.70	3	3797	147096	1335	132.98	1	0.974	536346123	2	0.511	-224.50	60.02	→
-2	-1	32201.3	183.2	1	0.98603-0.98271-0.10331	0.10331	0.05748	0.10245	0	311.88	165.00	227.51	311.67	165.02	227.41	0.077	0.87	0.32	10.00	11.51	3	1185	71051	1334	113.71	1	0.998	536348160	-1	0.074	58.39	-90.98	→	
-2	-1	32040.8	183.2	1	0.87794-0.88125-0.02672	0.18906	0.41213	0.36718	0	402.95	347.00	184.74	402.65	347.03	184.59	0.077	0.90	0.26	10.00	350.12	3	1185	71024	1328	92.29	1	0.998	536348160	1	0.074	149.46	91.02	→	
-2	-2	370173.	564.	1	0.99165-0.99829-0.11831	0.11831	0.04400	0.02209	0.11202	0	393.98	148.00	236.55	393.78	147.85	236.69	0.063	0.87	0.33	10.00	16.02	3	1305	994526	1358	118.35	1	1.000	536348161	-1	0.061	140.49	-107.98	→
-2	-2	327346.	741.	1	0.99010-0.98345-0.11218	0.05014	0.01102	0.10096	0	285.89	170.01	232.91	285.68	170.19	232.83	0.123	0.93	0.32	10.00	14.20	3	1305	459665	1348	116.42	1	0.993	536348161	1	0.117	32.40	-85.97	→	
-2	-2	331804.	742.	1	0.93123-0.93787-0.05489	0.21720	0.29050	0.20056	0	430.00	343.00	199.82	430.38	343.05	199.74	0.122	0.93	0.28	10.00	-2.34	3	1305	467672	1377	99.87	1	0.996	536348161	1	0.117	176.51	87.02	→	
-2	-2	367906.	1140.	1	0.85300-0.84633-0.19622	0.35854	0.54554	0.45559	0	319.99	360.00	301.54	319.88	359.63	301.52	0.064	0.85	0.42	10.00	48.52	3	1306	990454	1364	150.76	1	1.000	536348161	1	0.061	66.50	104.02	→	
-2	-3	17153.7	158.6	1	0.98820-0.99817-0.10747	0.05485	0.03580	0.09911	0	421.01	141.02	229.84	421.36	141.49	229.94	0.108	0.83	0.32	10.00	12.67	3	1485	27291	1356	114.97	1	1.000	536348162	-1	0.104	167.52	-114.96	→	
-2	-3	15759.5	189.8	1	0.99167-0.98170-0.11841	0.04390	0.02263	0.11228	0	260.00	175.00	236.80	259.65	175.15	236.76	0.169	0.93	0.33	10.00	16.15	3	1485	16552	1386	118.38	1	1.000	536348162	-1	0.161	6.51	-80.98	→	
-2	-3	13464.3	174.5	1	0.94706-0.95703-0.06520	0.22752	0.24355	0.10864	0	459.00	339.00	205.49	459.02	339.15	205.43	0.168	0.98	0.29	10.00	0.50	3	1485	13833	1303	102.71	1	1.000	536348162	1	0.161	205.51	83.02	→	
-2	-3	14394.0	146.1	1	0.93532-0.92533-0.17516	0.33748	0.37582	0.24090	0	294.00	364.00	279.07	293.83	364.20	279.05	0.118	0.87	0.39	10.00	37.29	3	1485	23046	1307	139.53	1	1.000	536348162	1	0.104	40.51	108.02	→	
-2	-4	11410.4	153.8	1	0.98422-0.99751-0.10036	0.06196	0.07262	0.10726	0	450.00	134.99	225.57	449.95	134.67	225.64	0.154	0.95	0.31	10.00	10.54	3	1705	12724	1302	112.82	1	1.000	536348163	-1	0.147	196.51	-120.99	→	
-2	-4	10687.7	177.9	1	0.99208-0.97879-0.12368	0.03863	0.05157	0.12831	0	233.95	180.00	240.22	233.57	179.93	240.13	0.216	0.92	0.33	10.00	17.86	3	1705	8658	1368	120.07	1	0.984	536348163	-1	0.204	-19.54	-75.98	→	
-2	-4	7205.8	143.6	1	0.95248-0.96577-0.06906	0.23138	0.22550	0.04562	0	488.99	335.01	207.60	488.65	335.32	207.59	0.214	0.88	0.29	10.00	1.55	3	1705	5997	1325	103.80	1	1.000	536348163	1	0.204	235.50	79.03	→	
-2	-4	11258.4	153.9	1	0.95567-0.94235-0.16621	0.31315	0.13324	0	267.97	369.00	271.29	267.63	368.97	271.25	0.156	0.93	0.38	10.00	33.39	3	1705	12577	1316	135.62	1	1.000	536348163	1	0.148	14.48	113.02	→		
-2	-5	27332.0	272.3	1	0.97990-0.99651-0.09444	0.06788	0.10284	0.12201	0	479.98	127.01	222.05	479.55	127.39	222.11	0.201	0.88	0.31	10.00	8.77	3	1951	23663	1352	111.05	1	1.000	536348164	-1	0.190	226.49	-128.97	→	
-2	-5	25292.1	301.1	1	0.99170-0.97509-0.12848	0.03384	0.07827	0.14658	0	207.00	184.95	243.29	207.34	184.56	243.25	0.265	0.75	0.34	10.00	19.40	3	1951	16801	1365	121.63	1	0.998	536348164	-1	0.247	-46.49	-71.03	→	
-2	-5	28749.4	281.5	1	0.96241-0.94576-0.16249	0.32481	0.28830	0.06343	0	241.00	374.00	268.19	241.30	373.99	268.21	0.204	0.91	0.37	10.00	31.84	3	1951	24793	1425	134.10	1	1.000	536348164	1	0.191	-12.49	118.02	→	
-2	-6	22953.1	311.7	1	0.99066-0.97072-0.13288	0.02944	0.10380	0.16602	0	180.99	189.00	246.31	180.84	189.08	246.24	0.314	0.97	0.34	10.00	20.90	3	2215	12738	1373	123.12	1	1.000	536348165	-1	0.290	-72.50	-66.98	→	

- .raw files contain information on position and intensity of diffraction spots.
- Positional information can be used in conjunction with  $F^2$  values, which are used as the total intensity values.
- Diffraction frames are all well documented, this knowledge allows us to produce frames that can be read within existing processing software.



- The sum of the intensity of each spot is calculated for each frame.
- Cumulative distribution is calculated between each pair of consecutive frames.
- Total intensity of a reflection on frame  $Z$  is the cumulative distribution of the Gaussian between  $Z$  and  $Z-1$ .

- Cumulative distribution is calculated for each row of pixels around the centroid.
- Then the cumulative distribution calculated for each pixel within the row.
- This process takes place for each reflection on a frame.
- After this is complete, a background is added to the image.



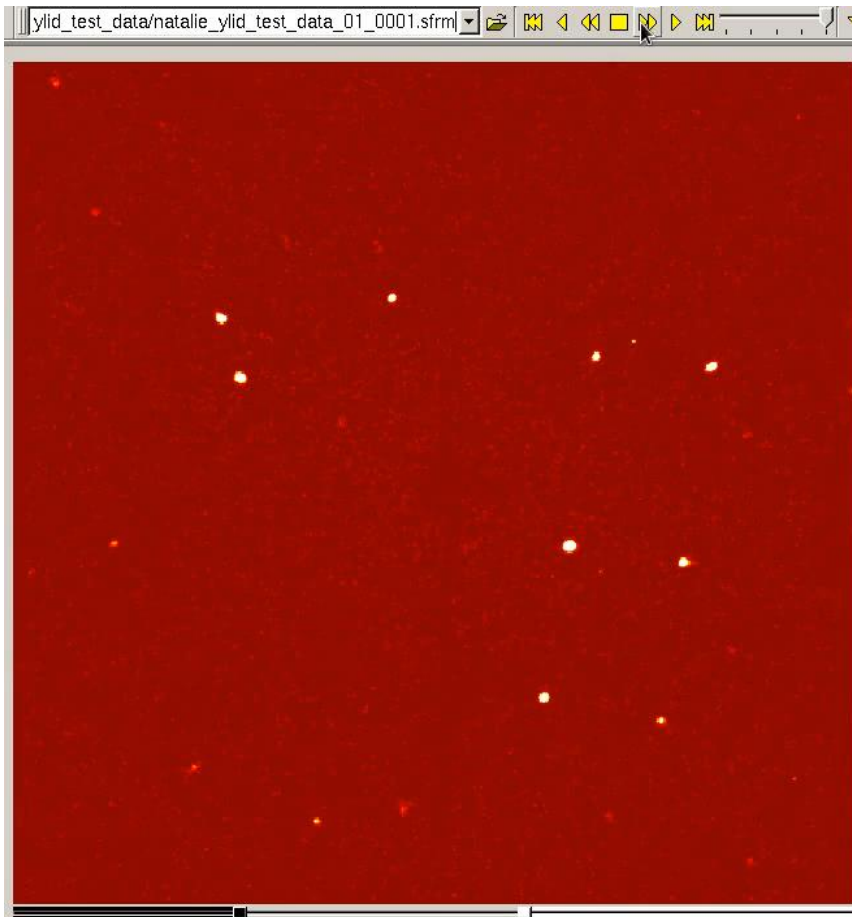
- Frames are then manufactured.
- Header information is recycled from real frames – with minor edits.
- Image is encoded in the required format.

```

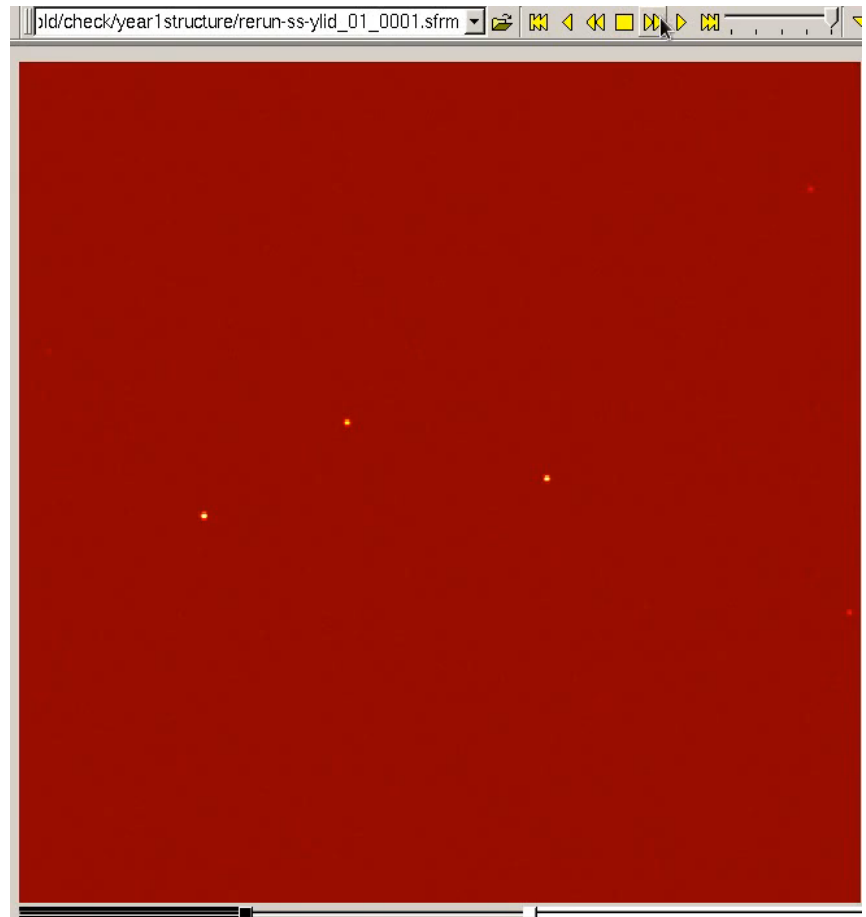
FORMAT :100
VERSION:15
HDRBLKS:15
TYPE :Phi Scan Image
SITE :XIPHOSIIA
MODEL :D85 [08/10-2338] with 4-CIRCLE (HUBER)
USER :BrukerAdministrator
SAMPLE :
SETNAME:
RUN :1
SAMPNUM:0
TITLE :
TITLE :
TITLE :
TITLE :
TITLE :
TITLE :
TITLE :
TITLE :
TITLE :
TITLE :
NCOUNTS:9286403
NOVERFL:-1
MINIMUM:26
MAXIMUM:12316
NONTIME:-2
NLATE :-1
FILENAM:\XIPHOSIIa\frames\guest\ylid_28_02_12\ylid_28_02_12_01_0025.sfrm
CREATED:02/28/12
CUMULAT:2.999000
ELAPSDR:3.000000
ELAPSDA:2.999000
OSCILLA:0
NSTEPS :1
RANGE :0.500000
START :193.000000
INCREME:0.500000
NUMBER :25
CSIZE :
DNSMET :
DARK :0482_00512_00003._dk
AUTORNG:0.000000E+000 0.000000E+000 0.000000E+000 0.000000E+000 1.251000E+000
ZEROADJ:0.000000 0.000000 0.000000 0.000000
XTRANS :0.000000 0.000000 0.000000 0.000000
HKL&XY :0.000000 0.000000 0.000000 0.000000 0.000000
AXES2 :0.000000 0.000000 0.000000 1.000000
ENDING2:0.000000 0.000000 0.000000 1.000000
FILTER2:0.000000 0.000000 0.000000 4.170000
LEPTOS :
CFR: HDR: IMG:
.....J!# " !" !!!#
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*8)#!$$ $(!" " #!"##### "$#" #!"!"# #!"#"$#$" "$ &#" "### # %$ " " !!
!("# !"# # $# # $#!" !%"! " ! # $& " ! $!" !# %&(! """"!"$ " !&#"%)%"$!"3$#!$$#1$ !
# ! # %!#" "$ ! %!!!!!"!"$!#!*$ " !#! $!"#&#(#!"'"( " #&!!!!" "%!!"$(%!"!"

```

# Frame Comparison



Real



Replica

# How Could We Detect Fraud Within Raw Diffraction Frames?



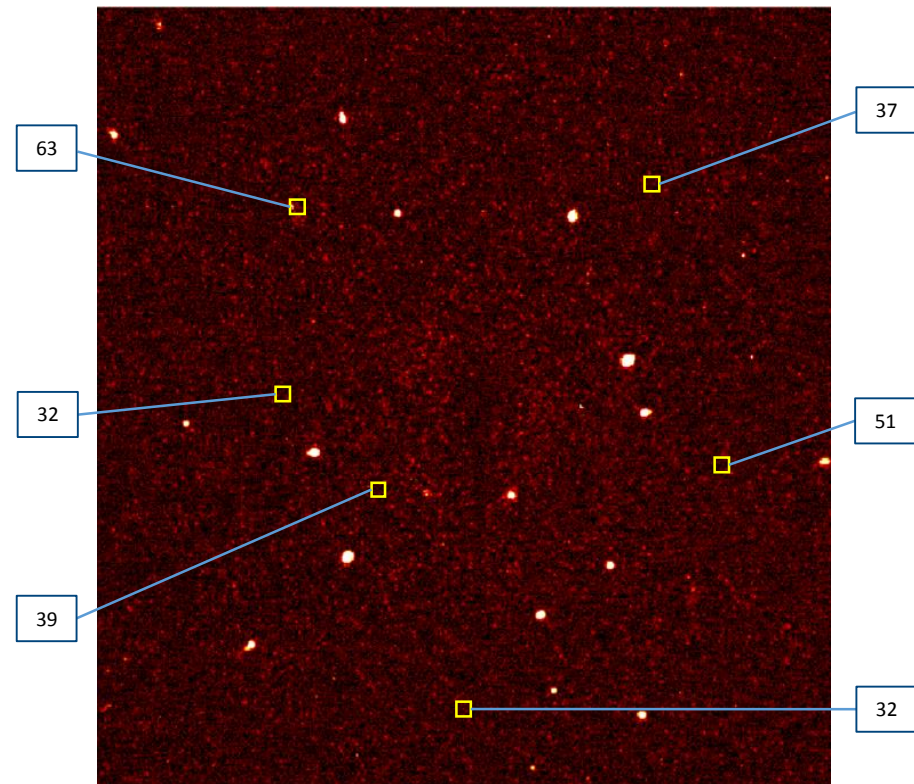
# Encrypt Entire Frame

- Fraudsters would have to crack the encryption and be able to re-encrypt files.
- Could slow down diffraction software as the decryptions need to be performed.
- Could hamper the development of current and new diffraction frame formats.



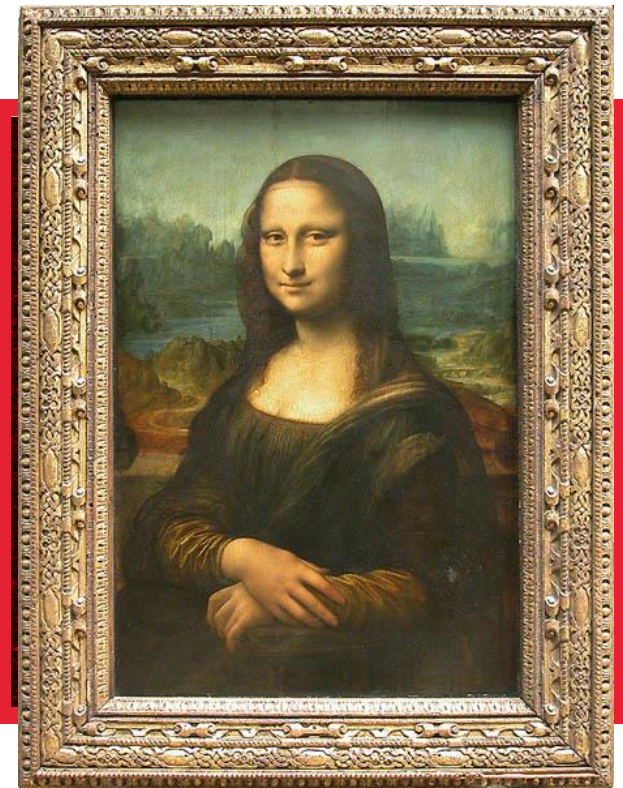
# Encryptions within frame header

- Using information from the diffraction image to create an encrypted section within the header.
- Encryption check required when file is submitted.



# Addition of information

- Fraud is prevented by photographing and cataloguing famous artwork without its frame.
- In a similar way, extra values or information could be added around the image, within the image binary itself.
- Fraudsters would have to know of its existence.

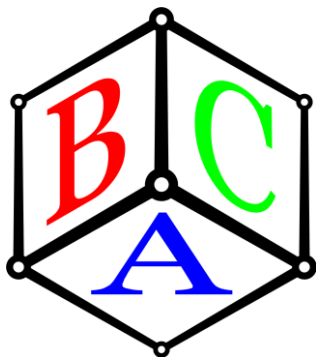


# Certification

- An additional file produced during the diffraction process, which contains encrypted information about the frames.
- Encryption check needed when raw images are submitted.



# Acknowledgements



Thank you for listening

# Should we rely on the honesty of our peers?

Has anything like this already been done?