

Automated Crystal Structure Validation

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Why Automated Structure Validation ?

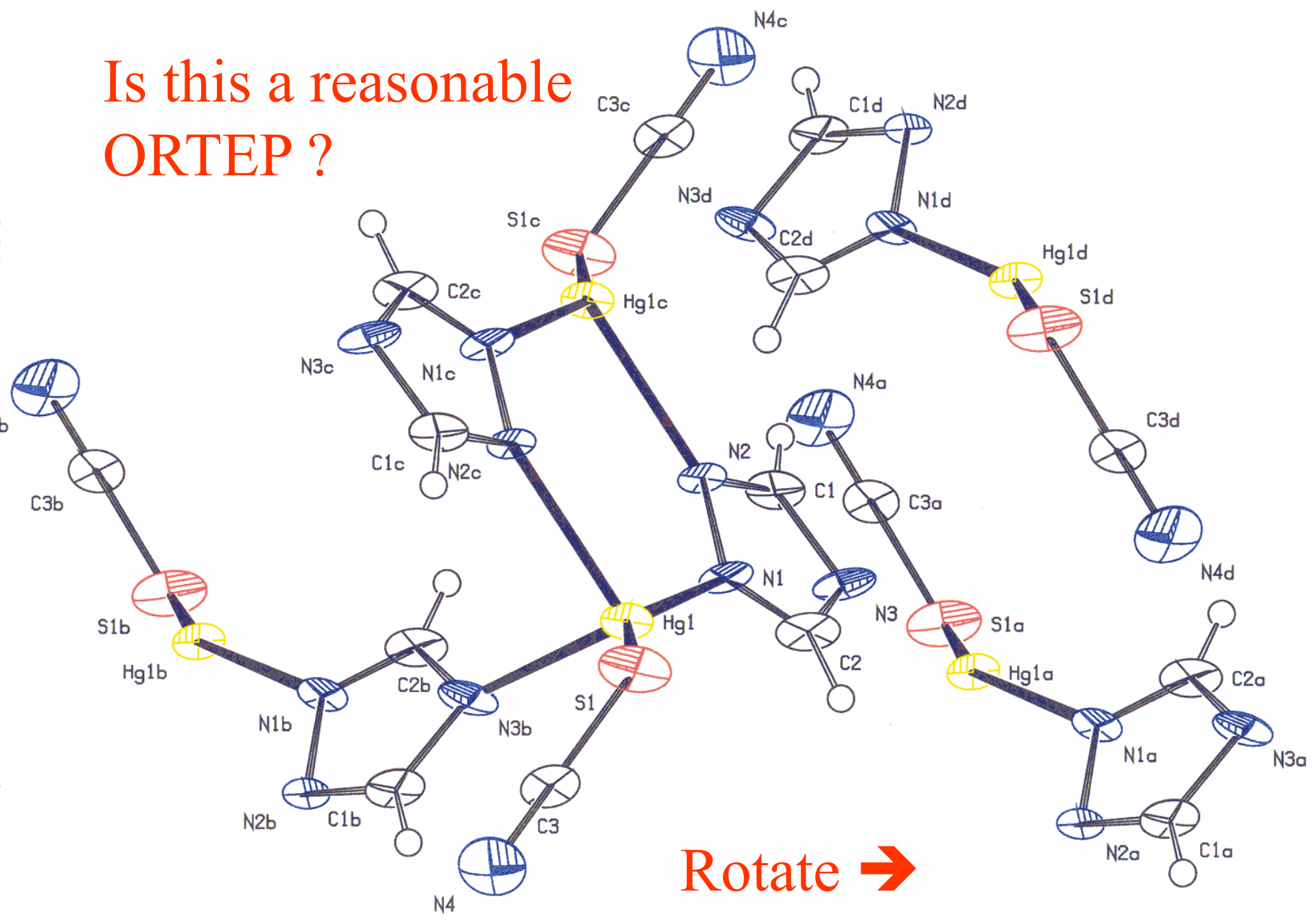
- It is easy to miss problems with a structure as a busy author or as a referee
- Increasingly: Black-Box style analyses done by non-experts
- Limited number of referees & experts available
- It is easy to hide problems with a ball-and-stick style illustration
- Sadly, Fraudulous results and structures published and in the CSD
- Even an ORTEP can hide problems =>

81 Y
PLATON-Dec 18 16:50:21 2006 - (171206)

NOMOVE FORCED

PROBA= 50

Is this a reasonable
ORTEP ?



Rotate →

Z 53

84-g

R = 0.07

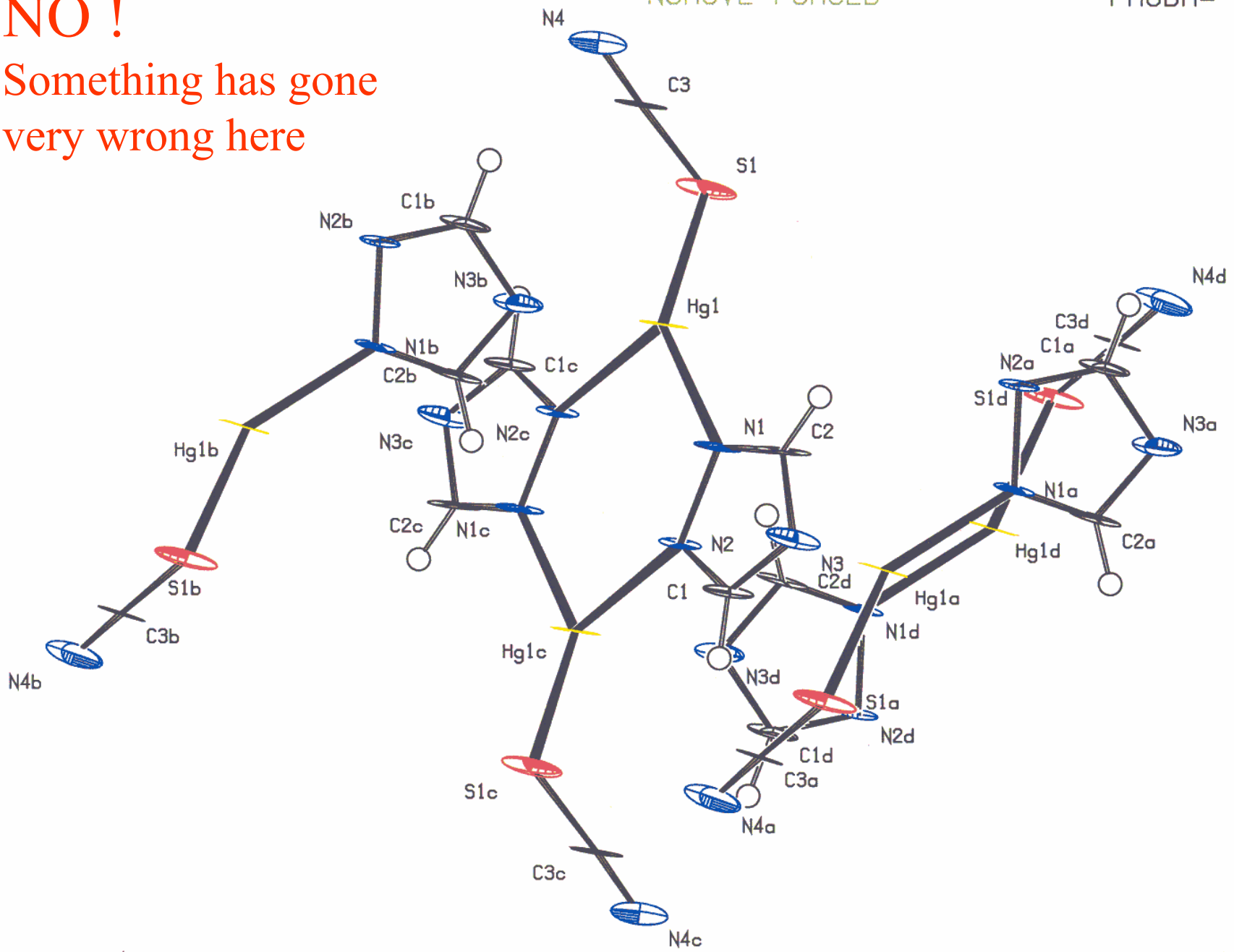
RES= 0 -141 X

-41 Y
- (171206)
PLATON-Dec 18 16:50:21 2006
Z -67 84-g

NO !
Something has gone
very wrong here

NOMOVE FORCED

PROBA= 50



R = 0.07

RES= 0 15 X

The CIF Standard Solution

- The IUCr Automated Structure Validation project was pioneered and ‘pushed’ by Syd Hall, at that time section editor of Acta Cryst C. by:
- The creation of the CIF Standard for data archival and exchange (Hall et al., (1991) Acta Cryst., A47, 655-685.
- Having CIF adopted by Sheldrick for SHELXL93
- Making CIF the Acta Cryst. submission standard
- Setting up early CIF checking procedures for Acta
- Inviting me to include PLATON checking tools such as ADDSYM and VOID search.

WHAT ARE THE VALIDATION QUESTIONS ?

Single Crystal Structure Validation addresses three simple but important questions:

- 1 – Is the reported information complete?
- 2 – What is the quality of the analysis?
- 3 – Is the Structure Correct?

How is Validation Currently Implemented ?

- Validation checks on CIF data can be executed at any time, both in-house (PLATON/CHECK) or through the WEB-based IUCr CHECKCIF server.
- A file, **check.def**, defines the issues that are tested (currently more than 400) with levels of severity and associated explanation and advise.
(www.cryst.chem.uu.nl/platon/CIF-VALIDATION.pdf)
- Most non-trivial tests on the IUCr CheckCIF server are executed with routines in the program PLATON. (Identified as PLATxyz)

VALIDATION ALERT LEVELS

CheckCIF/PLATON creates a report in the form of a list of ALERTS with the following ALERT levels:

- ALERT A – Serious Problem
- ALERT B – Potentially Serious Problem
- ALERT C – Check & Explain
- ALERT G – Verify or Take Notice

VALIDATION ALERT TYPES

- 1 - CIF Construction/Syntax errors,
Missing or Inconsistent Data.
- 2 - Indicators that the Structure Model
may be Wrong or Deficient.
- 3 - Indicators that the quality of the results
may be low.
- 4 – Info, Cosmetic Improvements, Queries and
Suggestions.

PLATON/CHECK CIF + FCF Results

=====
>>> The Following Model and Quality ALERTS were generated - (Acta-Mode) <<<
=====

Format: alert-number_ALERT_alert-type_alert-level text

148_ALERT_3_B	su on the	a	-	Axis is Too Large (x 1000)	.	10	Ang.
148_ALERT_3_B	su on the	b	-	Axis is Too Large (x 1000)	.	8	Ang.
148_ALERT_3_B	su on the	c	-	Axis is Too Large (x 1000)	.	30	Ang.
230_ALERT_2_B	Hirshfeld Test Diff for	01	--	C1	..	26.27	su
230_ALERT_2_B	Hirshfeld Test Diff for	02	--	C1	..	11.10	su
242_ALERT_2_B	Check Low	Ueq	as Compared to Neighbors for			C1	
420_ALERT_2_B	D-H Without Acceptor	01	-	H1	...	?	

#=====
230_ALERT_2_C Hirshfeld Test Diff for C1 -- C2 .. 6.87 su
242_ALERT_2_C Check Low Ueq as Compared to Neighbors for C11
911_ALERT_3_C Missing # FCF Refl Between THmin & STh/L= 0.595 11
913_ALERT_3_C Missing # of Very Strong Reflections in FCF 1

#=====
860_ALERT_3_G Note: Number of Least-Squares Restraints 2
#=====
#=====

=====
>>> The Following Improvement and Query ALERTS were generated - (Acta-Mode) <<<
=====

926_ALERT_1_B	Reported and Calculated	R1 * 100.0	Differ by	.	-0.81
927_ALERT_1_B	Reported and Calculated	wR2 * 100.0	Differ by	.	-2.26

#=====
042_ALERT_1_C Calc. and Reported MoietyFormula Strings Differ ?
790_ALERT_4_C Centre of Gravity not Within Unit Cell; Resd. # 2
C8 H6 O4
928_ALERT_1_C Reported and Calculated S value Differ by . -0.13
#=====
#=====

128_ALERT_4_G Non-standard setting of Space-group P21/c P21/n
#=====
#=====

checkCIF/PLATON report (publication check)

No syntax errors found.
Please wait while processing

[CIF dictionary](#)
[Interpreting this report](#)

Datablock: I

Bond precision:	C-C = 0.0157 Å	Wavelength=0.71073	
Cell:	a=7.6336(15)	b=27.725(6)	c=12.051(2)
	alpha=90	beta=98.80(3)	gamma=90
Temperature:	153 K		
	Calculated	Reported	
Volume	2520.5(9)	2520.5(9)	
Space group	P 21/n	P 21/n	
Hall group	-P 2yn	-P 2yn	
Moiety formula	C28 H32 Br0.97 Cl1.03 N2 Ru	C28 H32 Br0.97 Cl1.03 N2 Ru	
Sum formula	C28 H32 Br0.97 Cl1.03 N2 Ru	C28 H32 Br0.97 Cl1.03 N2 Ru	
Mr	611.69	611.69	
Dx,g cm-3	1.612	1.612	
Z	4	4	
Mu (mm-1)	2.290	2.290	
F000	1237.9	1237.9	
F000'	1232.67		
h,k,lmax	9,33,14	9,33,14	
Nref	4452	4449	
Tmin,Tmax	0.726,0.955	0.581,0.955	
Tmin'	0.546		
Correction method=	MULTI-SCAN		
Data completeness=	0.999	Theta(max)=	25.030
R(reflections)=	0.0891(3288)	wR2(reflections)=	0.2547(4449)
S =	1.037	Npar=	306

The following ALERTS were generated. Each ALERT has the format

test-name_ALERT_alert-type_alert-level.

Click on the hyperlinks for more details of the test.

Alert level B

[PLAT083_ALERT_2_B](#) SHELXL Second Parameter in WGHT unusually Large. 37.00

Alert level C

[RFACR01_ALERT_3_C](#) The value of the weighted R factor is > 0.25
Weighted R factor given 0.255
[PLAT301_ALERT_3_C](#) Main Residue Disorder 6.00 Perc.

Done

Which Key Validation Issues are Addressed

- Missed Space Group symmetry (“being Marshded”)
- Wrong chemistry (Mis-assigned atom types).
- Too many, too few or misplaced H-atoms.
- Unusual displacement parameters.
- Hirshfeld Rigid Bond test violations.
- Missed solvent accessible voids in the structure.
- Missed Twinning.
- Absolute structure
- Data quality and completeness.

Some Relevant ALERTS

Wrong atom type assignments generally cause:

- Serious Hirshfeld Rigid Bond Violation ALERTS
- Larger than expected difference map minima and maxima.
- $wR2 \gg 2 * R1$
- High values for the SHELXL refined weight parameter

Evaluation and Performance

- The validation scheme has been very successful for Acta Cryst. C & E in setting standards for quality and reliability.
- The missed symmetry problem has been solved for the IUCr journals (unfortunately not generally yet: There are still numerous ‘Marshable’ structures).
- Most major chemical journals currently have now some form of a validation scheme implemented.
- Recently included: FCF validation

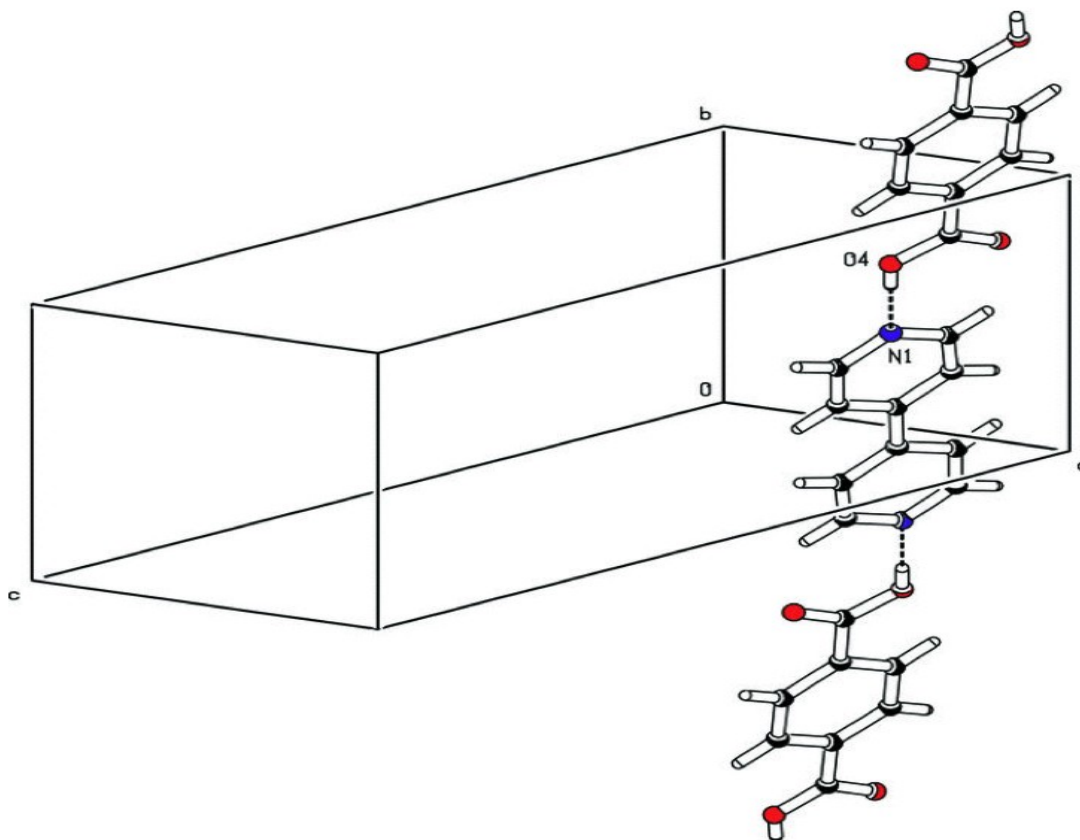
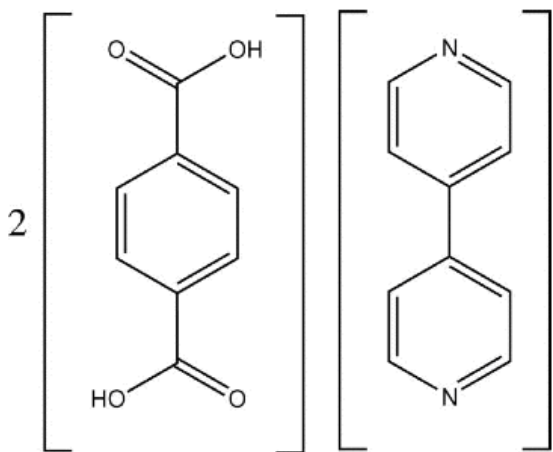
FCF-VALIDATION

- Check of the CIF & FCF data Consistency (including R-values, cell dimensions)
- Check of Completeness of the reflection data set.
- Automatic Detection of ignored twinning
- Detection of Applied Twinning Correction without having been Reported in the paper.
- Validity check of the reported Flack parameter value against the Hooft parameter value.
- Analysis of the details of the Difference Density Fourier Map for unreported features.

Sloppy, Novice or Fraudulent ?

- Errors are easily made and unfortunately not always discernable from fraud.
- Wrong element type assignments can be caused as part of an incorrect analysis of an unintended reaction product.
- Alternative element types can be (and have been) substituted deliberately to create a ‘new publishable’ structures.
- Reported and calculated R-values differing in the first relevant digit !?
- FCF Validation is the tool to sort out this type of issues.

Published structure is claimed to form an infinite hydrogen bonded chain



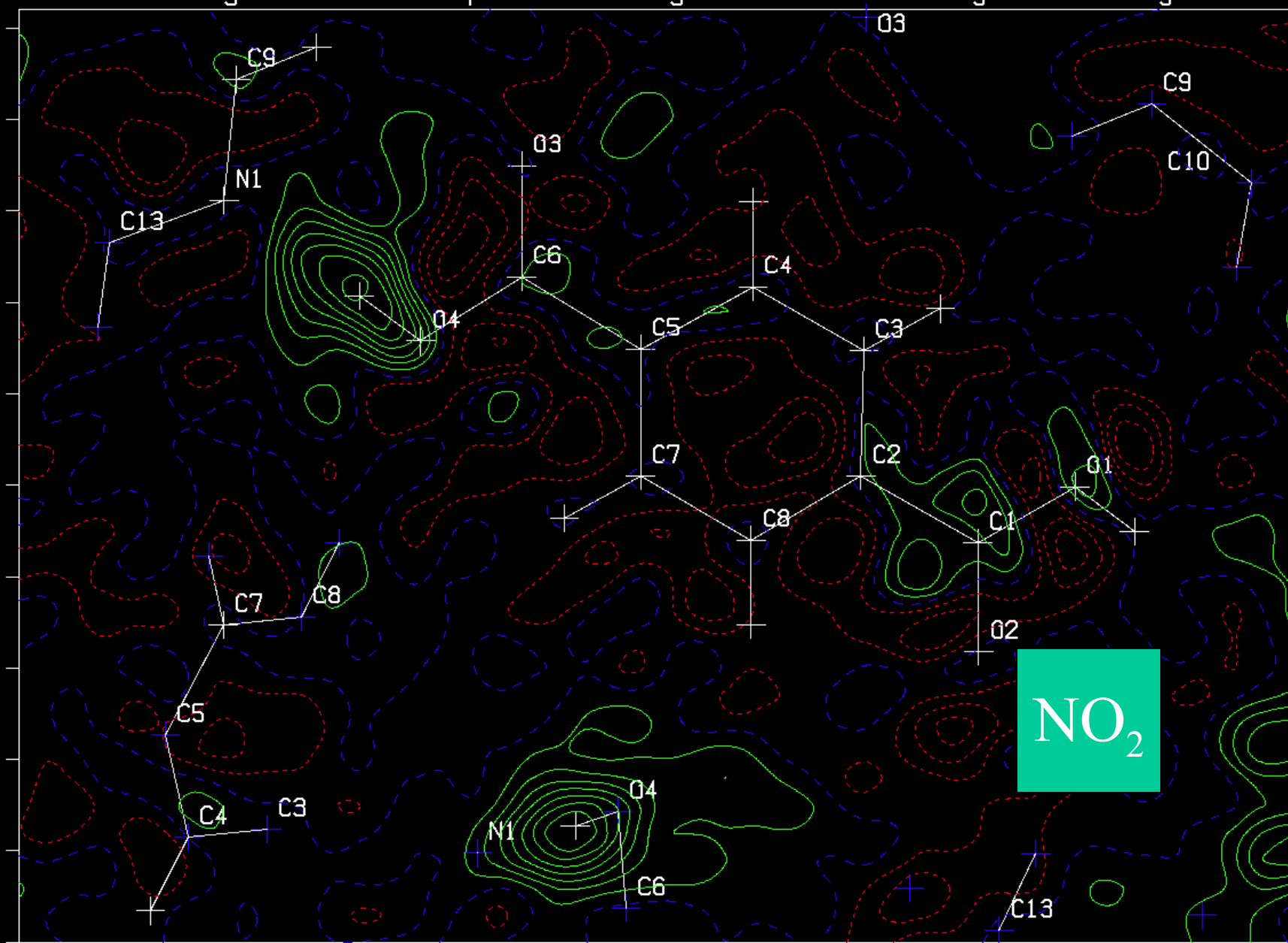
However: This structure does not include a dicarboxylic acid but the previously published para-nitrobenzoic acid.

PROOF: Difference map calculated without the 2 carboxylic H-atoms



Plane: -6.5083x 3.1226y -5.6013z = -3.6642 Cont-Lev(eA-3): -0.15 0.35 0.05 Dlf-Map
Tot = 1.5 Ang Step = 0.3000 Ang Resolution 25.0 Deg. Omlt 2*SlgI

PLATON-Sep 13 16:54:16 2009 - (90909)



I P 21/n R = 0.05 Ang

There are clear ALERTS ! But apparently ignored

```
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=====
```

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```
148_ALERT_3_B su on the      a - Axis is Too Large (x 1000) .      10 Ang.  
148_ALERT_3_B su on the      b - Axis is Too Large (x 1000) .      8 Ang.  
148_ALERT_3_B su on the      c - Axis is Too Large (x 1000) .     30 Ang.  
230_ALERT_2_B Hirshfeld Test Diff for 01 -- C1 .. 26.27 su  
230_ALERT_2_B Hirshfeld Test Diff for 02 -- C1 .. 11.10 su  
242_ALERT_2_B Check Low      Ueq as Compared to Neighbors for C1  
420_ALERT_2_B D-H Without Acceptor 01 - H1 ... ?
```

```
#=====  
230_ALERT_2_C Hirshfeld Test Diff for C1 -- C2 .. 6.87 su  
242_ALERT_2_C Check Low      Ueq as Compared to Neighbors for C11  
911_ALERT_3_C Missing # FCF Refl Between THmin & STh/L= 0.595 11  
913_ALERT_3_C Missing # of Very Strong Reflections in FCF .... 1
```

```
#=====  
860_ALERT_3_G Note: Number of Least-Squares Restraints ..... 2  
#=====  
#=====  
#=====
```

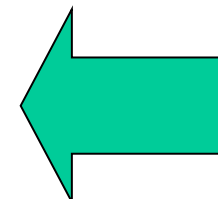
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C8 H6 O4
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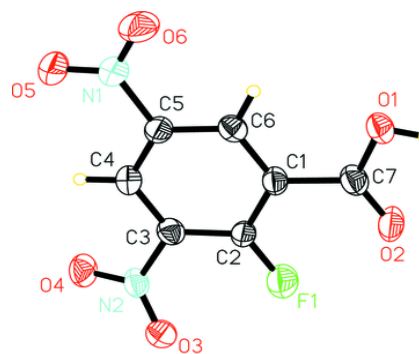
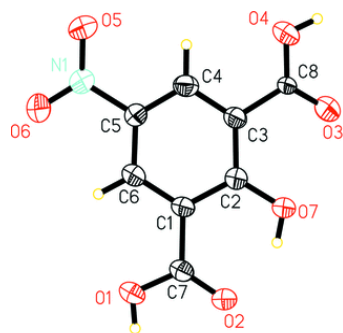
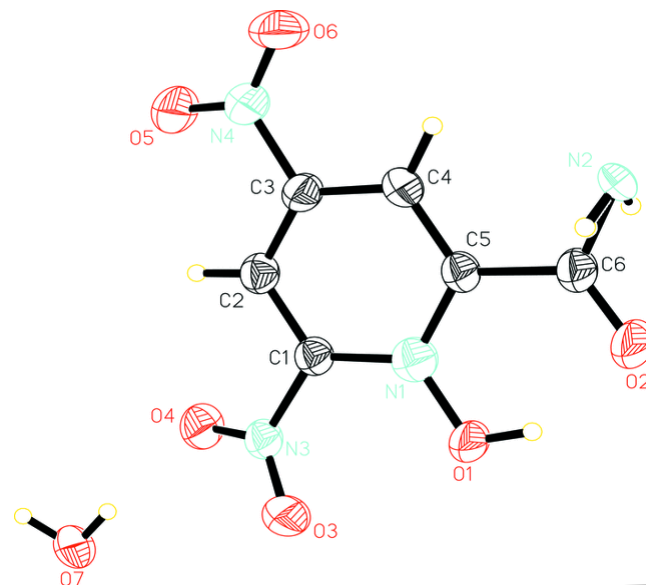
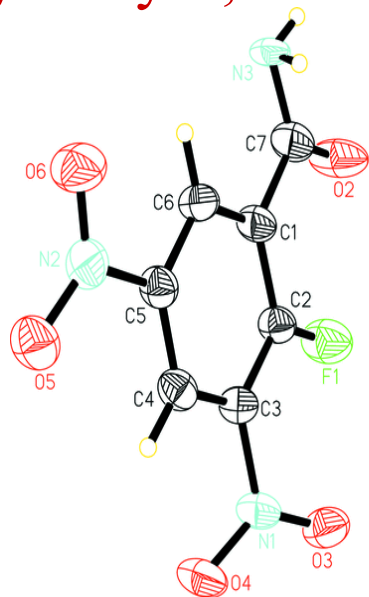
```
#=====  
128_ALERT_4_G Non-standard setting of Space-group P21/c .... P21/n  
#=====  
#=====
```



The Ultimate Shame

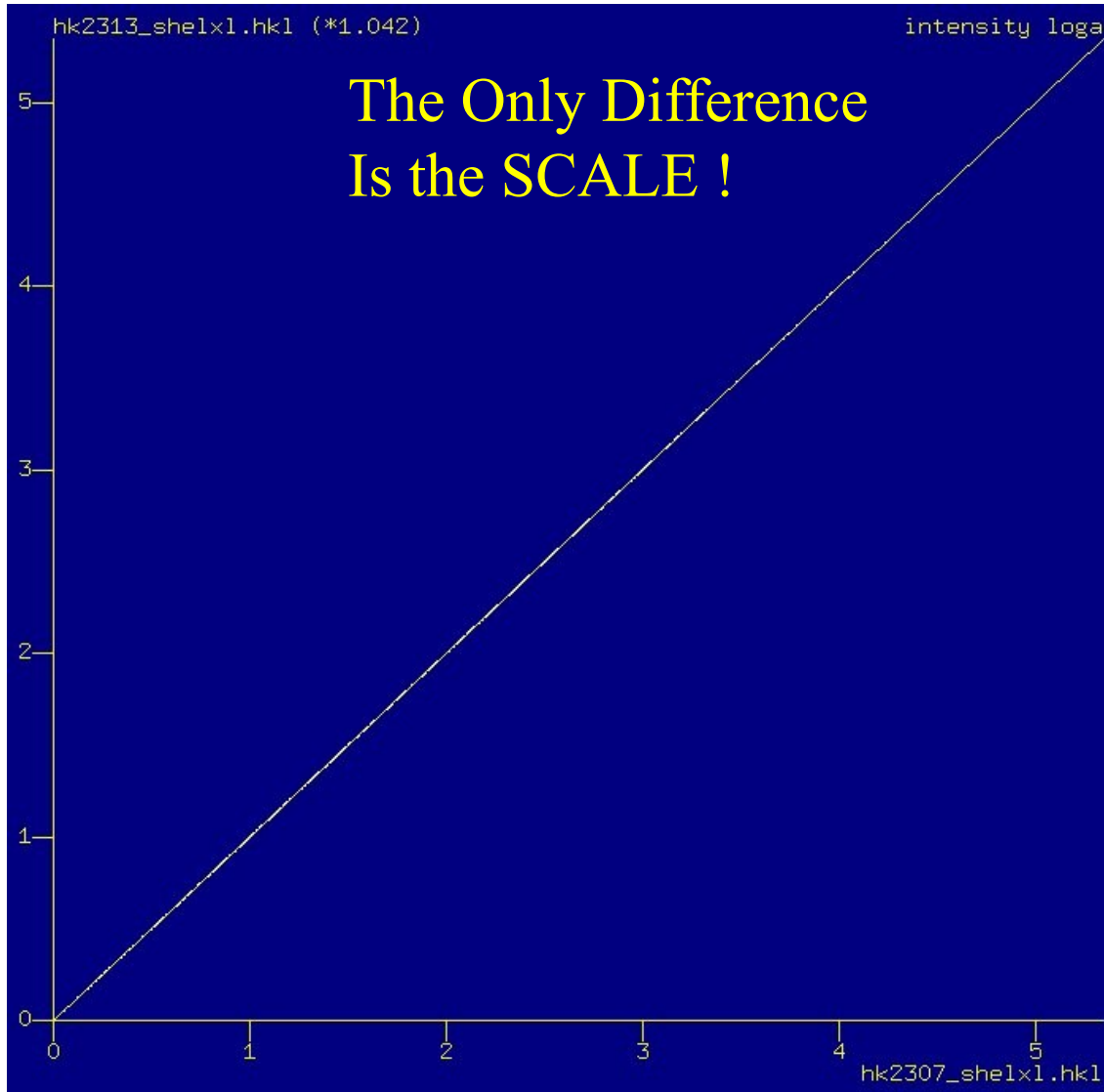
- Recently a whole series of ‘isomorphous’ substitutions was detected for an already published structure.
- Similar series have now been detected for coordination complexes (Transition metals and lanthanides)
- How could referees let those pass ?
- **Over 100 structures now retracted**
- Fraud detected by looking at all papers of the same authors of a ‘strange’ structure (and their institutions)

Bogus Variations (with Hirshfeld ALERTS) on the Published Structure 2-hydroxy-3,5-nitrobenzoic acid (ZAJGUM)



Comparison of the Observed data for two 'isomorphous' compounds.

Tool: platon -d name1.fcf name2.fcf



Conclusion
The Same
Data !

SLOPPY

Or

FRAUD ?

Info

www.cryst.chem.uu.nl/platon/CIF-VALIDATION.pdf

www.cryst.chem.uu.nl/platon/FCF-VALIDATION.pdf

- Papers on structure validation:
- A.L.Spek (2003). *J. Appl. Cryst.* 36, 7-13.
- A.L.Spek (2009). *Acta Cryst.* D65, 148-155.

