

checkCIF/PLATON report

You have not supplied any structure factors. As a result the full set of tests cannot be run.

THIS REPORT IS FOR GUIDANCE ONLY. IF USED AS PART OF A REVIEW PROCEDURE FOR PUBLICATION, IT SHOULD NOT REPLACE THE EXPERTISE OF AN EXPERIENCED CRYSTALLOGRAPHIC REFEREE.

No syntax errors found. CIF dictionary Interpreting this report

Datablock: fl206_a

Bond precision:	C-C = 0.0114 A	Wavelength=0.71073	
Cell:	a=21.131(4)	b=17.391(4)	c=21.069(4)
	alpha=90	beta=107.00(3)	gamma=90
Temperature:	293 K		
	Calculated	Reported	
Volume	7404(3)	7404(3)	
Space group	C 2/c	C 1 2/c 1	
Hall group	-C 2yc	-C 2yc	
Moiety formula	2(C63 H54 Fe2 N18), 4(B0.40 F1.60), 4(B F4), 4(C0.50 H0.75 N0.5	C63 H54 Fe2 N18, 2(B0.4 F0.8), 2(B F4), 2(B0.6 F3.2), 2(C0.5 H0	
Sum formula	C130 H114 B8 F32 Fe4 N39	C65 H57 B4 F16 Fe2 N19.50	
Mr	3140.49	1570.24	
Dx, g cm ⁻³	1.409	1.409	
Z	2	4	
Mu (mm ⁻¹)	0.486	0.486	
F000	3198.0	3198.0	
F000'	3202.45		
h,k,lmax	26,21,26	26,21,26	
Nref	7678	6995	
Tmin,Tmax	0.994,0.995		
Tmin'	0.990		

Correction method= Not given

Data completeness= 0.911 Theta(max)= 26.499

R(reflections)= 0.1035(4686) wR2(reflections)= 0.3220(6995)

S = 1.096 Npar= 630

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 A

PLAT029_ALERT_3_A _diffrn_measured_fraction_theta_full value Low . 0.917 Note

Author Response: 'The MX1 beam line of the Australian Synchrotron is a single circle instrument, leaving gaps in the coverage of the reciprocal space.'

Alert level C

PLAT084_ALERT_3_C High wR2 Value (i.e. > 0.25) 0.32 Report
PLAT094_ALERT_2_C Ratio of Maximum / Minimum Residual Density 2.47 Report
PLAT234_ALERT_4_C Large Hirshfeld Difference N3B -- C4B .. 0.23 Ang.
PLAT234_ALERT_4_C Large Hirshfeld Difference N5A -- C20A .. 0.16 Ang.
PLAT250_ALERT_2_C Large U3/U1 Ratio for Average U(i,j) Tensor 2.2 Note
PLAT341_ALERT_3_C Low Bond Precision on C-C Bonds 0.01136 Ang.

Alert level G

FORMU01_ALERT_1_G There is a discrepancy between the atom counts in the
_chemical_formula_sum and _chemical_formula_moiety. This is
usually due to the moiety formula being in the wrong format.
Atom count from _chemical_formula_sum: C65 H57 B4 F16 Fe2 N19.5
Atom count from _chemical_formula_moiety:C64 H55.5 B4 F16 Fe2 N19

PLAT002_ALERT_2_G Number of Distance or Angle Restraints on AtSite 13 Note
PLAT003_ALERT_2_G Number of Uiso or Uij Restrained non-H Atoms ... 42 Report
PLAT007_ALERT_5_G Number of Unrefined Donor-H Atoms 4 Report
PLAT012_ALERT_1_G No _shelx_res_checksum found in CIF Please Check
PLAT042_ALERT_1_G Calc. and Reported MoietyFormula Strings Differ Please Check
PLAT045_ALERT_1_G Calculated and Reported Z Differ by a Factor ... 0.50 Check
PLAT072_ALERT_2_G SHELXL First Parameter in WGHT Unusually Large 0.13 Report
PLAT083_ALERT_2_G SHELXL Second Parameter in WGHT Unusually Large 69.22 Why ?
PLAT171_ALERT_4_G The CIF-Embedded .res File Contains EADP Records 1 Report
PLAT172_ALERT_4_G The CIF-Embedded .res File Contains DFIX Records 2 Report
PLAT176_ALERT_4_G The CIF-Embedded .res File Contains SADI Records 1 Report
PLAT177_ALERT_4_G The CIF-Embedded .res File Contains DELU Records 1 Report
PLAT187_ALERT_4_G The CIF-Embedded .res File Contains RIGU Records 6 Report
PLAT199_ALERT_1_G Reported _cell_measurement_temperature (K) 293 Check
PLAT200_ALERT_1_G Reported _diffrn_ambient_temperature (K) 293 Check
PLAT232_ALERT_2_G Hirshfeld Test Diff (M-X) Fe01 -- N1A .. 5.2 s.u.
PLAT244_ALERT_4_G Low 'Solvent' Ueq as Compared to Neighbors of B01A Check
PLAT300_ALERT_4_G Atom Site Occupancy of N1B' is Constrained at 0.5 Check
PLAT300_ALERT_4_G Atom Site Occupancy of N1B is Constrained at 0.5 Check
PLAT300_ALERT_4_G Atom Site Occupancy of N2B' is Constrained at 0.5 Check
PLAT300_ALERT_4_G Atom Site Occupancy of N2B is Constrained at 0.5 Check
PLAT300_ALERT_4_G Atom Site Occupancy of C1 is Constrained at 0.5 Check
PLAT300_ALERT_4_G Atom Site Occupancy of C7A' is Constrained at 0.5 Check
PLAT300_ALERT_4_G Atom Site Occupancy of C1B is Constrained at 0.5 Check
PLAT300_ALERT_4_G Atom Site Occupancy of C2 is Constrained at 0.5 Check
PLAT300_ALERT_4_G Atom Site Occupancy of C8A' is Constrained at 0.5 Check
PLAT300_ALERT_4_G Atom Site Occupancy of C2B is Constrained at 0.5 Check
PLAT300_ALERT_4_G Atom Site Occupancy of C3 is Constrained at 0.5 Check
PLAT300_ALERT_4_G Atom Site Occupancy of C9A' is Constrained at 0.5 Check
PLAT300_ALERT_4_G Atom Site Occupancy of C3B is Constrained at 0.5 Check

PLAT300_ALERT_4_G	Atom Site Occupancy of C1AA	is Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C2AA	is Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H2AB	is Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H2AC	is Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H2AD	is Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of N01L	is Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C4	is Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C0AA	is Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H4D	is Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H4E	is Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H4F	is Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of F00H	is Constrained at	0.8	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of F00T	is Constrained at	0.8	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of F00Y	is Constrained at	0.8	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of B01C	is Constrained at	0.6	Check
PLAT301_ALERT_3_G	Main Residue Disorder	(Resd 1)..	46	% Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder	(Resd 2)..	100	% Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder	(Resd 4)..	100	% Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder	(Resd 5)..	100	% Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder	(Resd 6)..	100	% Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder	(Resd 7)..	100	% Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder	(Resd 8)..	100	% Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder	(Resd 9)..	100	% Note
PLAT304_ALERT_4_G	Non-Integer Number of Atoms (1.75)	in Resd. #	4	Check
PLAT304_ALERT_4_G	Non-Integer Number of Atoms (1.50)	in Resd. #	5	Check
PLAT304_ALERT_4_G	Non-Integer Number of Atoms (0.80)	in Resd. #	6	Check
PLAT304_ALERT_4_G	Non-Integer Number of Atoms (0.80)	in Resd. #	7	Check
PLAT304_ALERT_4_G	Non-Integer Number of Atoms (0.80)	in Resd. #	8	Check
PLAT304_ALERT_4_G	Non-Integer Number of Atoms (0.60)	in Resd. #	9	Check
PLAT367_ALERT_2_G	Long? C(sp?)-C(sp?) Bond C8B	- C11B ..	1.51	Ang.
PLAT432_ALERT_2_G	Short Inter X...Y Contact F00A	.. C0AA ..	2.95	Ang.
PLAT432_ALERT_2_G	Short Inter X...Y Contact F00Y	.. C20A ..	2.93	Ang.
PLAT720_ALERT_4_G	Number of Unusual/Non-Standard Labels		40	Note
PLAT811_ALERT_5_G	No ADDSYM Analysis: Too Many Excluded Atoms		!	Info
PLAT860_ALERT_3_G	Number of Least-Squares Restraints		218	Note

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

6 **ALERT type 1** CIF construction/syntax error, inconsistent or missing data
10 **ALERT type 2** Indicator that the structure model may be wrong or deficient
5 **ALERT type 3** Indicator that the structure quality may be low
112 **ALERT type 4** Improvement, methodology, query or suggestion
2 **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.

Publication of your CIF in IUCr journals

A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section C* or *E* or *IUCrData*, you should make sure that full publication checks are run on the final version of your CIF prior to submission.

Publication of your CIF in other journals

Please refer to the *Notes for Authors* of the relevant journal for any special instructions relating to CIF submission.

Validation response form

Please find below a validation response form (VRF) that can be filled in and pasted into your CIF.

```
# start Validation Reply Form
_vrf_PLAT084_fl206_a
;
PROBLEM: High wR2 Value (i.e. > 0.25) ..... 0.32 Report
RESPONSE: ...
;
_vrf_PLAT094_fl206_a
;
PROBLEM: Ratio of Maximum / Minimum Residual Density .... 2.47 Report
RESPONSE: ...
;
_vrf_PLAT234_fl206_a
;
PROBLEM: Large Hirshfeld Difference N3B -- C4B .. 0.23 Ang.
RESPONSE: ...
;
_vrf_PLAT250_fl206_a
;
PROBLEM: Large U3/U1 Ratio for Average U(i,j) Tensor .... 2.2 Note
RESPONSE: ...
;
_vrf_PLAT341_fl206_a
;
PROBLEM: Low Bond Precision on C-C Bonds ..... 0.01136 Ang.
RESPONSE: ...
;
# end Validation Reply Form
```

PLATON version of 27/03/2017; check.def file version of 24/03/2017

Datablock fl206_a - ellipsoid plot

