

checkCIF/PLATON report

Structure factors have been supplied for datablock(s) 3-Sm_hex

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: 3-Sm hex

Bond precision: C-C = 0.0117 Å

Wavelength=1.54178

```
Cell:      a=11.5391(8)
           alpha=85.410(4)
```

b=13.1434 (9) c=20.9935 (17)
beta=75.777 (4) gamma=67.979 (3)

Temperature: 120 K

	Calculated
Volume	2860.9 (4)
Space group	P -1
Hall group	-P 1
Moiety formula	C56 H89 O4 P4 Sm
Sum formula	C56 H89 O4 P4 Sm
Mr	1100.52
Dx, g cm ⁻³	1.278
Z	2
Mu (mm ⁻¹)	9.076
F000	1158.0
F000'	1152.35
h, k, lmax	14, 16, 25
Nref	11400
Tmin, Tmax	0.332, 0.520
Tmin'	0.138

Reported
2860.9(4)
P -1
-P 1
C56 H89 O4 P4 Sm
C56 H89 O4 P4 Sm
1100.50
1.278
2
9.076
1158.0

14,16,25
11259
0.286,0.754

```
Correction method= # Reported T Limits: Tmin=0.286 Tmax=0.754
AbsCorr = MULTI-SCAN
```

Data completeness= 0.988

$$\text{Theta (max)} = 72.725$$

R(reflections)= 0.0741(9612)

```
wR2 (reflections)=  
0.2030 ( 11259)
```

$$S = 1.026$$

Npar= 785

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

PLAT213_ALERT_2_B Atom C43 has ADP max/min Ratio 4.3 prolat

Author Response: This is due to the extensive disorder present in the model and overall quality of the dataset.

PLAT971_ALERT_2_B Check Calcd Resid. Dens. 0.85Ang From Sm1 2.84 eA-3

Author Response: The residualt density could not be modelled as any chemically sensible species and its presence is due to the proximity of a strong absorber (Sm)

PLAT971_ALERT_2_B Check Calcd Resid. Dens. 0.94Ang From Sm1 2.81 eA-3

Author Response: The residualt density could not be modelled as any chemically sensible species and its presence is due to the proximity of a strong absorber (Sm)

PLAT972_ALERT_2_B Check Calcd Resid. Dens. 0.95Ang From Sm1 -2.53 eA-3

Author Response: The residualt density could not be modelled as any chemically sensible species and its presence is due to the proximity of a strong absorber (Sm)



Alert level C

PLAT213_ALERT_2_C Atom Sm1 has ADP max/min Ratio 3.2 prolat

Author Response: This is due to the extensive disorder present in the model and overall quality of the dataset.

PLAT213_ALERT_2_C Atom P4 has ADP max/min Ratio 3.8 prolat

Author Response: This is due to the extensive disorder present in the model and overall quality of the dataset.

PLAT213_ALERT_2_C Atom P3 has ADP max/min Ratio 3.4 oblate

Author Response: This is due to the extensive disorder present in the model and overall quality of the dataset.

PLAT213_ALERT_2_C Atom O3 has ADP max/min Ratio 3.1 prolat

Author Response: This is due to the extensive disorder present in the model and overall quality of the dataset.

PLAT213_ALERT_2_C Atom O4 has ADP max/min Ratio 3.9 prolat

Author Response: This is due to the extensive disorder present in the model and overall quality of the dataset.

PLAT213_ALERT_2_C Atom C29 has ADP max/min Ratio 3.3 prolat

Author Response: This is due to the extensive disorder present in the model and overall quality of the dataset.

PLAT213_ALERT_2_C Atom C30 has ADP max/min Ratio 3.4 prolat

Author Response: This is due to the extensive disorder present in the model and overall quality of the dataset.

PLAT213_ALERT_2_C Atom C32 has ADP max/min Ratio 3.5 prolat

Author Response: This is due to the extensive disorder present in the model and overall quality of the dataset.

PLAT213_ALERT_2_C Atom C33 has ADP max/min Ratio 3.6 prolat

Author Response: This is due to the extensive disorder present in the model and overall quality of the dataset.

PLAT213_ALERT_2_C Atom C45 has ADP max/min Ratio 3.5 prolat

Author Response: This is due to the extensive disorder present in the model and overall quality of the dataset.

PLAT213_ALERT_2_C Atom C46 has ADP max/min Ratio 4.0 prolat

Author Response: This is due to the extensive disorder present in the model and overall quality of the dataset.

PLAT213_ALERT_2_C Atom C48 has ADP max/min Ratio 3.1 prolat

Author Response: This is due to the extensive disorder present in the model and overall quality of the dataset.

PLAT220_ALERT_2_C NonSolvent Resd 1 C Ueq(max)/Ueq(min) Range 3.8 Ratio
 PLAT222_ALERT_3_C NonSolvent Resd 1 H Uiso(max)/Uiso(min) Range 5.1 Ratio
 PLAT241_ALERT_2_C High 'MainMol' Ueq as Compared to Neighbors of 02 Check
 PLAT241_ALERT_2_C High 'MainMol' Ueq as Compared to Neighbors of 03 Check
 PLAT241_ALERT_2_C High 'MainMol' Ueq as Compared to Neighbors of 04 Check
 PLAT242_ALERT_2_C Low 'MainMol' Ueq as Compared to Neighbors of Sm1 Check
 PLAT242_ALERT_2_C Low 'MainMol' Ueq as Compared to Neighbors of C49 Check
 PLAT250_ALERT_2_C Large U3/U1 Ratio for <U(i,j)> Tensor(Resd 1) 2.7 Note
 PLAT342_ALERT_3_C Low Bond Precision on C-C Bonds 0.01175 Ang.
 PLAT601_ALERT_2_C Unit Cell Contains Solvent Accessible VOIDS of . 31 Ang**3
 PLAT911_ALERT_3_C Missing FCF Refl Between Thmin & STh/L= 0.600 41 Report
 -1 14 0, 1 15 0, 1-14 1, 0 1 1, 0 3 1, 0 4 1,
 1 4 1, 0 5 1, 1 7 1, -1 14 1, 0 14 1, 1 15 1,
 1 6 2, -1 14 2, 0 14 2, 2 15 2, 0 14 3, 1 14 3,
 2 15 3, 3 15 3, 2 1 4, 1 6 4, 1 13 4, 0 14 4,
 1 14 4, 2 15 4, -1 -4 5, 0 14 5, 2 15 5, 3 15 5,
 -10 -2 6, 0 14 6, 5 14 9, 5 15 9, 6 15 9, 4 14 10,
 5 14 10, 2 11 16, 6 -1 20, 9 1 20, 9 1 21,
 PLAT971_ALERT_2_C Check Calcd Resid. Dens. 0.99Ang From Sm1 1.93 eA-3

Author Response: The residualt density could not be modelled as any chemically sensible species and its presence is due to the proximity of a strong absorber (Sm)

PLAT971_ALERT_2_C Check Calcd Resid. Dens. 0.79Ang From Sm1 1.71 eA-3

Author Response: The residualt density could not be modelled as any chemically sensible species and its presence is due to the proximity of a strong absorber (Sm)

PLAT971_ALERT_2_C Check Calcd Resid. Dens. 1.12Ang From O2 1.57 eA-3

Author Response: The residualt density could not be modelled as any chemically sensible species and its presence is due to the proximity of a strong absorber (Sm)

PLAT972_ALERT_2_C Check Calcd Resid. Dens. 0.98Ang From Sm1 -2.50 eA-3

Author Response: The residual density could not be modelled as any chemically sensible species and its presence is due to the proximity of a strong absorber (Sm)

PLAT972_ALERT_2_C Check Calcd Resid. Dens. 0.82Ang From Sm1 -1.98 eA-3

Author Response: The residual density could not be modelled as any chemically sensible species and its presence is due to the proximity of a strong absorber (Sm)

PLAT975_ALERT_2_C Check Calcd Resid. Dens. 0.54Ang From O4 . 0.72 eA-3
 PLAT975_ALERT_2_C Check Calcd Resid. Dens. 0.75Ang From O3 . 0.58 eA-3

● Alert level G

PLAT002_ALERT_2_G	Number of Distance or Angle Restraints on AtSite	36	Note
PLAT003_ALERT_2_G	Number of Uiso or U(i,j) Restrained non-H Atoms	82	Report
PLAT007_ALERT_5_G	Number of Unrefined Donor-H Atoms	2	Report
	H3A H2		
PLAT072_ALERT_2_G	SHELXL First Parameter in WGHT Unusually Large	0.12	Report
PLAT083_ALERT_2_G	SHELXL Second Parameter in WGHT Unusually Large	10.32	Why ?
PLAT175_ALERT_4_G	The CIF-Embedded .res File Contains SAME Records	2	Report
PLAT178_ALERT_4_G	The CIF-Embedded .res File Contains SIMU Records	3	Report
PLAT187_ALERT_4_G	The CIF-Embedded .res File Contains RIGU Records	1	Report
PLAT188_ALERT_3_G	A Non-default SIMU Restraint Value has been used	0.0100	Report
PLAT188_ALERT_3_G	A Non-default SIMU Restraint Value has been used	0.0080	Report
PLAT188_ALERT_3_G	A Non-default SIMU Restraint Value has been used	0.0080	Report
PLAT301_ALERT_3_G	Main Residue Disorder(Resd 1)	28%	Note
PLAT412_ALERT_2_G	Short Intra XH3 .. XHn H12B ..H23F .	2.07	Ang.
	x,y,z =	1_555	Check
PLAT412_ALERT_2_G	Short Intra XH3 .. XHn H17 ..H26C .	2.11	Ang.
	x,y,z =	1_555	Check
PLAT412_ALERT_2_G	Short Intra XH3 .. XHn H31 ..H41A .	2.02	Ang.
	x,y,z =	1_555	Check
PLAT802_ALERT_4_G	CIF Input Record(s) with more than 80 Characters	2	Info
PLAT811_ALERT_5_G	No ADDSYM Analysis: Too Many Excluded Atoms	!	Info
PLAT860_ALERT_3_G	Number of Least-Squares Restraints	1457	Note
PLAT912_ALERT_4_G	Missing # of FCF Reflections Above STh/L= 0.600	100	Note
PLAT941_ALERT_3_G	Average HKL Measurement Multiplicity	4.6	Low
PLAT969_ALERT_5_G	The 'Henn et al.' R-Factor-gap value	3.009	Note
	Predicted wR2: Based on SigI**2 6.74 or SHELX Weight 19.79		
PLAT978_ALERT_2_G	Number C-C Bonds with Positive Residual Density.	0	Info

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

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

