

# 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: i-4

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Bond precision:    C-C = 0.0380 A                      Wavelength=0.71073

Cell:                      a=9.817(1)              b=9.817(1)              c=19.541(4)  
                                    alpha=90              beta=90              gamma=90

Temperature:              100 K

	Calculated	Reported
Volume	1883.2(5)	1883.3(5)
Space group	I -4	I -4
Hall group	I -4	I -4
Moiety formula	C32 H64 Li4 N4 O8	C32 H64 Li4 N4 O8
Sum formula	C32 H64 Li4 N4 O8	C32 H64 Li4 N4 O8
Mr	660.63	660.63
Dx,g cm-3	1.165	1.165
Z	2	2
Mu (mm-1)	0.080	0.080
F000	720.0	720.0
F000'	720.31	
h,k,lmax	12,12,24	12,12,24
Nref	2100[ 1086]	2072
Tmin,Tmax		0.369,0.429
Tmin'		

Correction method= # Reported T Limits: Tmin=0.369 Tmax=0.429  
AbsCorr = MULTI-SCAN

Data completeness= 1.91/0.99                      Theta(max)= 27.113

R(reflections)= 0.3023( 1057)                      wR2(reflections)= 0.6523( 2072)

S = 2.155    Npar= 51

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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.

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**Alert level A**

EXPT005\_ALERT\_1\_A \_exptl\_crystal\_description is missing  
Crystal habit description.  
The following tests will not be performed.  
CRYSR\_01

ATOM007\_ALERT\_1\_A \_atom\_site\_aniso\_label is missing  
Unique label identifying the atom site.

RINTA01\_ALERT\_3\_A The value of Rint is greater than 0.25  
Rint given 0.313

SHFSU01\_ALERT\_2\_A The absolute value of parameter shift to su ratio > 0.20  
Absolute value of the parameter shift to su ratio given 1.991  
Additional refinement cycles may be required.

PLAT020\_ALERT\_3\_A The Value of Rint is Greater Than 0.12 ..... 0.313 Report  
PLAT080\_ALERT\_2\_A Maximum Shift/Error ..... 1.99 Why ?  
PLAT082\_ALERT\_2\_A High R1 Value ..... 0.30 Report  
PLAT084\_ALERT\_3\_A High wR2 Value (i.e. > 0.25) ..... 0.65 Report  
PLAT097\_ALERT\_2\_A Large Reported Max. (Positive) Residual Density 2.53 eA-3  
PLAT201\_ALERT\_2\_A Isotropic non-H Atoms in Main Residue(s) ..... 12 Report  
PLAT203\_ALERT\_2\_A Negative Isotropic ADP for O1 ..... -0.001 Report  
PLAT203\_ALERT\_2\_A Negative Isotropic ADP for LI1 ..... -0.001 Report  
PLAT316\_ALERT\_2\_A Too many H on C in C=N Moiety in Main Residue .. C4 Check  
PLAT410\_ALERT\_2\_A Short Intra H...H Contact H4A ..H8A 1.30 Ang.  
PLAT410\_ALERT\_2\_A Short Intra H...H Contact H4A ..H8B 1.69 Ang.  
PLAT413\_ALERT\_2\_A Short Inter XH3 .. XHn H3B ..H7B 1.89 Ang.

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**Alert level B**

DIFMN02\_ALERT\_2\_B The minimum difference density is < -0.1\*ZMAX\*1.00  
\_refine\_diff\_density\_min given = -1.029  
Test value = -0.800

PLAT098\_ALERT\_2\_B Large Reported Min. (Negative) Residual Density -1.03 eA-3  
PLAT340\_ALERT\_3\_B Low Bond Precision on C-C Bonds ..... 0.038 Ang.  
PLAT360\_ALERT\_2\_B Short C(sp3)-C(sp3) Bond C7 - C8 1.32 Ang.  
PLAT411\_ALERT\_2\_B Short Inter H...H Contact H4B ..H7A 1.93 Ang.

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**Alert level C**

DIFMN03\_ALERT\_1\_C The minimum difference density is < -0.1\*ZMAX\*0.75  
The relevant atom site should be identified.

DIFMX02\_ALERT\_1\_C The maximum difference density is > 0.1\*ZMAX\*0.75  
The relevant atom site should be identified.

GOODF01\_ALERT\_2\_C The least squares goodness of fit parameter lies  
outside the range 0.80 <> 2.00  
Goodness of fit given = 2.155

STRVA01\_ALERT\_4\_C Flack parameter is too small  
From the CIF: \_refine\_ls\_abs\_structure\_Flack -4.100  
From the CIF: \_refine\_ls\_abs\_structure\_Flack\_su 1.000

PLAT053\_ALERT\_1\_C Minimum Crystal Dimension Missing (or Error) ... Please Check  
PLAT054\_ALERT\_1\_C Medium Crystal Dimension Missing (or Error) ... Please Check  
PLAT055\_ALERT\_1\_C Maximum Crystal Dimension Missing (or Error) ... Please Check  
PLAT087\_ALERT\_2\_C Unsatisfactory S value (Too High) ..... 2.15 Check  
PLAT094\_ALERT\_2\_C Ratio of Maximum / Minimum Residual Density .... 2.46 Report  
PLAT241\_ALERT\_2\_C High 'MainMol' Ueq as Compared to Neighbors of O1 Check  
PLAT241\_ALERT\_2\_C High 'MainMol' Ueq as Compared to Neighbors of C4 Check  
PLAT241\_ALERT\_2\_C High 'MainMol' Ueq as Compared to Neighbors of C6 Check  
PLAT241\_ALERT\_2\_C High 'MainMol' Ueq as Compared to Neighbors of C8 Check  
PLAT242\_ALERT\_2\_C Low 'MainMol' Ueq as Compared to Neighbors of C1 Check  
PLAT242\_ALERT\_2\_C Low 'MainMol' Ueq as Compared to Neighbors of C5 Check  
PLAT361\_ALERT\_2\_C Long C(sp3)-C(sp3) Bond C1 - C4 .. 1.67 Ang.  
PLAT410\_ALERT\_2\_C Short Intra H...H Contact H7A ..H8B 1.94 Ang.  
PLAT410\_ALERT\_2\_C Short Intra H...H Contact H7B ..H8A 1.92 Ang.

PLAT601\_ALERT\_2\_C Structure Contains Solvent Accessible VOIDS of . 54 Ang\*\*3  
PLAT790\_ALERT\_4\_C Centre of Gravity not Within Unit Cell: Resd. # 1 Note  
C32 H64 Li4 N4 O8

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### ● Alert level G

PLAT012_ALERT_1_G	No	_shelx_res_checksum Found in CIF . . . . .	Please Check
PLAT032_ALERT_4_G	Std. Uncertainty on Flack Parameter Value High .		1.000 Report
PLAT072_ALERT_2_G	SHELXL First Parameter in WGHT Unusually Large		0.20 Report
PLAT303_ALERT_2_G	Full Occupancy Atom H5B with # Connections		2.00 Check
PLAT343_ALERT_2_G	Unusual Angle Range in Main Residue for		C5 Check
PLAT343_ALERT_2_G	Unusual sp3 Angle Range in Main Residue for		C8 Check
PLAT367_ALERT_2_G	Long? C(sp?)-C(sp?) Bond C5 - C6 .		1.59 Ang.
PLAT764_ALERT_4_G	Overcomplete CIF Bond List Detected (Rep/Expd) .		1.31 Ratio
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle in CIF . . . #		21 Check
	01 -C1 -Li1 1.555 1.555 2.445		44.70 Deg.

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

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

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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.

Datablock i-4 - ellipsoid plot

