

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: LK15

Bond precision: C-C = 0.0155 A

Wavelength=0.71073

Cell: a=16.2248(14) b=20.2271(17) c=28.297(2)
 alpha=103.560(2) beta=92.408(2) gamma=104.177(2)
Temperature: 100 K

	Calculated	Reported
Volume	8703.9(12)	8703.8(13)
Space group	P -1	P -1
Hall group	-P 1	-P 1
	C147 H101.50 Cl8 Dy7 N10	
Moiety formula	O42.50, N O3, C2 H3 N, 2.75(C H4 O)	?
Sum formula	C151.75 H115.50 Cl8 Dy7 N12 O48.25	C3.99 H3.04 Cl0.21 Dy0.18 N0.32 O1.27
Mr	4300.17	113.16
Dx,g cm-3	1.641	1.641
Z	2	76
Mu (mm-1)	3.167	3.167
F000	4188.0	4188.0
F000'	4189.53	
h,k,lmax	20,24,34	20,24,34
Nref	34264	34234
Tmin,Tmax	0.691,0.729	0.450,0.742
Tmin'	0.383	

Correction method= # Reported T Limits: Tmin=0.450 Tmax=0.742
AbsCorr = ?

Data completeness= 0.999

Theta(max)= 26.022

R(reflections)= 0.0631(25380)

wR2(reflections)= 0.1788(34234)

S = 1.052

Npar= 2079

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

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 6.902
Additional refinement cycles may be required.

Author Response: Multiple disorders and significant residual electron density around th

PLAT080_ALERT_2_A Maximum Shift/Error 6.90 Why ?

Author Response: Multiple disorders and significant residual electron density around th

PLAT417_ALERT_2_A Short Inter D-H..H-D H32A ..H53 . 0.97 Ang.
x,y,z = 1_555 Check

Author Response: Multiple disorders and significant residual electron density around th

PLAT602_ALERT_2_A VERY LARGE Solvent Accessible VOID(S) in Structure ! Info

Author Response: Multiple solvent disorders - SQUEEZE (part of PLATON was used.

Alert level B

PLAT201_ALERT_2_B Isotropic non-H Atoms in Main Residue(s) 3 Report
O42 O43 C109
PLAT220_ALERT_2_B NonSolvent Resd 1 C Ueq(max) / Ueq(min) Range 7.0 Ratio
PLAT420_ALERT_2_B D-H Without Acceptor O39 --H39B . Please Check
PLAT420_ALERT_2_B D-H Without Acceptor O47 --H47' . Please Check
PLAT420_ALERT_2_B D-H Without Acceptor O54 --H54 . Please Check

Alert level C

ABSMU01_ALERT_1_C The ratio of given/expected absorption coefficient lies
outside the range 0.99 <> 1.01
Calculated value of mu = 3.100
Value of mu given = 3.167
PLAT041_ALERT_1_C Calc. and Reported SumFormula Strings Differ Please Check
PLAT052_ALERT_1_C Info on Absorption Correction Method Not Given Please Do !
PLAT094_ALERT_2_C Ratio of Maximum / Minimum Residual Density 2.06 Report
PLAT213_ALERT_2_C Atom C11 has ADP max/min Ratio 3.1 prolat
PLAT215_ALERT_3_C Disordered O44 has ADP max/min Ratio 3.2 Note
PLAT215_ALERT_3_C Disordered N13 has ADP max/min Ratio 3.3 Note
PLAT220_ALERT_2_C NonSolvent Resd 1 O Ueq(max) / Ueq(min) Range 5.5 Ratio
PLAT222_ALERT_3_C NonSolvent Resd 1 H Uiso(max)/Uiso(min) Range 7.9 Ratio
PLAT242_ALERT_2_C Low MainMol Ueq as Compared to Neighbors of O12 Check
PLAT260_ALERT_2_C Large Average Ueq of Residue Including O45 0.134 Check
PLAT329_ALERT_4_C Carbon Atom Hybridisation Unclear for C109 Check
PLAT342_ALERT_3_C Low Bond Precision on C-C Bonds 0.01548 Ang.
PLAT369_ALERT_2_C Long C(sp2)-C(sp2) Bond C124 - C130 . 1.53 Ang.
PLAT369_ALERT_2_C Long C(sp2)-C(sp2) Bond C131 - C132 . 1.53 Ang.

PLAT410_ALERT_2_C	Short Intra H...H Contact	H80	..H83	.	1.99 Ang.
			x,y,z =		1_555 Check
PLAT410_ALERT_2_C	Short Intra H...H Contact	H119	..H122	.	1.94 Ang.
			x,y,z =		1_555 Check

● Alert level G

CELLZ01_ALERT_1_G Difference between formula and atom_site contents detected.

CELLZ01_ALERT_1_G ALERT: Large difference may be due to a

symmetry error - see SYMMG tests

From the CIF: _cell_formula_units_Z 76

From the CIF: _chemical_formula_sum C3.99 H3.04 Cl0.21 Dy0.18 N0.32 O1

TEST: Compare cell contents of formula and atom_site data

atom	Z*formula	cif sites	diff
C	303.24	303.50	-0.26
H	231.04	231.00	0.04
Cl	15.96	16.00	-0.04
Dy	13.68	14.00	-0.32
N	24.32	24.00	0.32
O	96.52	96.50	0.02

PLAT002_ALERT_2_G	Number of Distance or Angle Restraints on AtSite		8	Note
PLAT003_ALERT_2_G	Number of Uiso or Uij Restrained non-H Atoms ...		218	Report
PLAT007_ALERT_5_G	Number of Unrefined Donor-H Atoms		19	Report
PLAT012_ALERT_1_G	N.O.K. _shelx_res_checksum Found in CIF			Please Check
PLAT013_ALERT_1_G	N.O.K. _shelx_hkl_checksum Found in CIF			Please Check
PLAT045_ALERT_1_G	Calculated and Reported Z Differ by a Factor ...		0.03	Check
PLAT083_ALERT_2_G	SHELXL Second Parameter in WGHT Unusually Large		79.47	Why ?
PLAT154_ALERT_1_G	The s.u.'s on the Cell Angles are Equal ..(Note)		0.002	Degree
PLAT172_ALERT_4_G	The CIF-Embedded .res File Contains DFIX 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		1	Report
PLAT300_ALERT_4_G	Atom Site Occupancy of O44	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of N10	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of N13	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H10A	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H10B	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H10C	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O45	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O55	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O56	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of N14	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O50	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O51	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O52	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of N12	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O47	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C2'	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H15G	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H15H	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H15I	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H47'	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O49	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C154	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H15M	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H15N	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H15O	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H49'	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O54	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C1'	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H1'1	Constrained at	0.5	Check

PLAT300_ALERT_4_G	Atom Site Occupancy of H1'2	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H1'3	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H54	Constrained at	0.5	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of O46	Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C129	Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H12A	Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H12B	Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H12C	Constrained at	0.25	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H46	Constrained at	0.25	Check
PLAT301_ALERT_3_G	Main Residue Disorder(Resd 1)		1%	Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 2)		100%	Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder (Resd 3)		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 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 in (Resd 9)		1.50	Check
PLAT333_ALERT_2_G	Large Aver C6-Ring C-C Dist C1 -C10		1.42	Ang.
PLAT333_ALERT_2_G	Large Aver C6-Ring C-C Dist C112 -C121		1.43	Ang.
PLAT412_ALERT_2_G	Short Intra XH3 .. XHn H01C ..H10A		1.67	Ang.
	x,y,z =	1_555	Check	
PLAT413_ALERT_2_G	Short Inter XH3 .. XHn H150 ..H69		1.77	Ang.
	x,l+y,z =	1_565	Check	
PLAT413_ALERT_2_G	Short Inter XH3 .. XHn H1'3 ..H41		2.11	Ang.
	1+x,y,z =	1_655	Check	
PLAT414_ALERT_2_G	Short Intra D-H..H-X H10B ..H39A		1.98	Ang.
	x,y,z =	1_555	Check	
PLAT417_ALERT_2_G	Short Inter D-H..H-D H25B ..H49'		0.89	Ang.
	x,y,z =	1_555	Check	

Author Response: Multiple disorders and significant residual electron density around th

PLAT431_ALERT_2_G	Short Inter HL..A Contact C14 ..O42		3.09	Ang.
	1-x,-y,1-z =	2_656	Check	
PLAT432_ALERT_2_G	Short Inter X...Y Contact C16 ..C2'		3.22	Ang.
	1-x,1-y,2-z =	2_667	Check	
PLAT432_ALERT_2_G	Short Inter X...Y Contact C42 ..C92		3.19	Ang.
	-1+x,y,z =	1_455	Check	
PLAT720_ALERT_4_G	Number of Unusual/Non-Standard Labels		8	Note
PLAT721_ALERT_1_G	Bond Calc 0.98000, Rep 1.11360 Dev...		0.13	Ang.
	C1' -H1'1 1.555 1.555	#	13	Check
PLAT721_ALERT_1_G	Bond Calc 0.97000, Rep 1.10930 Dev...		0.14	Ang.
	C1' -H1'2 1.555 1.555	#	14	Check
PLAT721_ALERT_1_G	Bond Calc 0.99000, Rep 1.11780 Dev...		0.13	Ang.
	C1' -H1'3 1.555 1.555	#	15	Check
PLAT721_ALERT_1_G	Bond Calc 0.94000, Rep 0.97350 Dev...		0.03	Ang.
	O46 -H46 1.555 1.555	#	131	Check
PLAT721_ALERT_1_G	Bond Calc 0.95000, Rep 0.96050 Dev...		0.01	Ang.
	O54 -H54 1.555 1.555	#	142	Check
PLAT721_ALERT_1_G	Bond Calc 0.98000, Rep 1.00560 Dev...		0.03	Ang.
	C109 -H10A 1.555 1.555	#	342	Check
PLAT721_ALERT_1_G	Bond Calc 0.98000, Rep 1.00600 Dev...		0.03	Ang.
	C109 -H10B 1.555 1.555	#	343	Check
PLAT721_ALERT_1_G	Bond Calc 0.98000, Rep 1.00600 Dev...		0.03	Ang.
	C109 -H10C 1.555 1.555	#	344	Check
PLAT721_ALERT_1_G	Bond Calc 0.98000, Rep 0.99780 Dev...		0.02	Ang.
	C129 -H12A 1.555 1.555	#	377	Check
PLAT721_ALERT_1_G	Bond Calc 0.99000, Rep 1.00690 Dev...		0.02	Ang.
	C129 -H12B 1.555 1.555	#	378	Check
PLAT721_ALERT_1_G	Bond Calc 0.97000, Rep 1.00390 Dev...		0.03	Ang.

C129	-H12C		1.555	1.555	#	379	Check
PLAT722_ALERT_1_G	Angle	Calc	109.00,	Rep	123.50	Dev...		14.50 Degree
O54	-C1'	-H1'1	1.555	1.555	1.555	#	56	Check
PLAT722_ALERT_1_G	Angle	Calc	110.00,	Rep	124.10	Dev...		14.10 Degree
O54	-C1'	-H1'2	1.555	1.555	1.555	#	57	Check
PLAT722_ALERT_1_G	Angle	Calc	110.00,	Rep	92.30	Dev...		17.70 Degree
H1'1	-C1'	-H1'2	1.555	1.555	1.555	#	58	Check
PLAT722_ALERT_1_G	Angle	Calc	109.00,	Rep	124.20	Dev...		15.20 Degree
O54	-C1'	-H1'3	1.555	1.555	1.555	#	59	Check
PLAT722_ALERT_1_G	Angle	Calc	109.00,	Rep	91.80	Dev...		17.20 Degree
H1'1	-C1'	-H1'3	1.555	1.555	1.555	#	60	Check
PLAT722_ALERT_1_G	Angle	Calc	110.00,	Rep	91.50	Dev...		18.50 Degree
H1'2	-C1'	-H1'3	1.555	1.555	1.555	#	61	Check
PLAT722_ALERT_1_G	Angle	Calc	109.00,	Rep	54.50	Dev...		54.50 Degree
DY6	-O4	-H4'	1.555	1.555	1.555	#	436	Check
PLAT722_ALERT_1_G	Angle	Calc	111.00,	Rep	51.80	Dev...		59.20 Degree
DY2	-O4	-H4'	1.555	1.555	1.555	#	437	Check
PLAT722_ALERT_1_G	Angle	Calc	113.00,	Rep	44.90	Dev...		68.10 Degree
DY3	-O8	-H8''	1.555	1.555	1.555	#	448	Check
PLAT722_ALERT_1_G	Angle	Calc	108.00,	Rep	60.90	Dev...		47.10 Degree
DY6	-O8	-H8''	1.555	1.555	1.555	#	449	Check
PLAT722_ALERT_1_G	Angle	Calc	121.00,	Rep	117.50	Dev...		3.50 Degree
C129	-O46	-H46	1.555	1.555	1.555	#	502	Check
PLAT722_ALERT_1_G	Angle	Calc	119.00,	Rep	120.60	Dev...		1.60 Degree
C154	-O49	-H49'	1.555	1.555	1.555	#	505	Check
PLAT722_ALERT_1_G	Angle	Calc	108.00,	Rep	20.60	Dev...		87.40 Degree
C48	-N2	-H2'	1.555	1.555	1.555	#	513	Check
PLAT722_ALERT_1_G	Angle	Calc	112.00,	Rep	8.30	Dev...		103.70 Degree
C61	-N5	-H5'	1.555	1.555	1.555	#	522	Check
PLAT722_ALERT_1_G	Angle	Calc	112.00,	Rep	8.70	Dev...		103.30 Degree
C79	-N6	-H6'	1.555	1.555	1.555	#	525	Check
PLAT722_ALERT_1_G	Angle	Calc	109.00,	Rep	111.80	Dev...		2.80 Degree
O12	-C109	-H10A	1.555	1.555	1.555	#	874	Check
PLAT722_ALERT_1_G	Angle	Calc	109.00,	Rep	112.00	Dev...		3.00 Degree
O12	-C109	-H10B	1.555	1.555	1.555	#	875	Check
PLAT722_ALERT_1_G	Angle	Calc	109.00,	Rep	106.80	Dev...		2.20 Degree
H10A	-C109	-H10B	1.555	1.555	1.555	#	876	Check
PLAT722_ALERT_1_G	Angle	Calc	110.00,	Rep	112.00	Dev...		2.00 Degree
O12	-C109	-H10C	1.555	1.555	1.555	#	877	Check
PLAT722_ALERT_1_G	Angle	Calc	109.00,	Rep	106.90	Dev...		2.10 Degree
H10A	-C109	-H10C	1.555	1.555	1.555	#	878	Check
PLAT722_ALERT_1_G	Angle	Calc	110.00,	Rep	106.90	Dev...		3.10 Degree
H10B	-C109	-H10C	1.555	1.555	1.555	#	879	Check
PLAT722_ALERT_1_G	Angle	Calc	109.00,	Rep	113.70	Dev...		4.70 Degree
O46	-C129	-H12B	1.555	1.555	1.555	#	938	Check
PLAT722_ALERT_1_G	Angle	Calc	109.00,	Rep	107.80	Dev...		1.20 Degree
H12A	-C129	-H12B	1.555	1.555	1.555	#	939	Check
PLAT722_ALERT_1_G	Angle	Calc	110.00,	Rep	112.40	Dev...		2.40 Degree
O46	-C129	-H12C	1.555	1.555	1.555	#	940	Check
PLAT722_ALERT_1_G	Angle	Calc	110.00,	Rep	106.70	Dev...		3.30 Degree
H12A	-C129	-H12C	1.555	1.555	1.555	#	941	Check
PLAT722_ALERT_1_G	Angle	Calc	110.00,	Rep	106.50	Dev...		3.50 Degree
H12B	-C129	-H12C	1.555	1.555	1.555	#	942	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond)	Angle(s) in CIF	#				448	Check
DY3	-O8	-H8''	1.555	1.555	1.555		44.90	Deg.
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond)	Angle(s) in CIF	#				513	Check
C48	-N2	-H2'	1.555	1.555	1.555		20.60	Deg.
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond)	Angle(s) in CIF	#				522	Check
C61	-N5	-H5'	1.555	1.555	1.555		8.30	Deg.
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond)	Angle(s) in CIF	#				525	Check
C79	-N6	-H6'	1.555	1.555	1.555		8.70	Deg.
PLAT790_ALERT_4_G	Centre of Gravity not Within Unit Cell:	Resd.	#				8	Note

C H4 O

PLAT794_ALERT_5_G	Tentative Bond Valency for Dy4	(III)	.	3.01	Info
PLAT794_ALERT_5_G	Tentative Bond Valency for Dy5	(III)	.	2.88	Info
PLAT794_ALERT_5_G	Tentative Bond Valency for Dy6	(III)	.	3.02	Info
PLAT802_ALERT_4_G	CIF Input Record(s) with more than 80 Characters			3	Info
PLAT860_ALERT_3_G	Number of Least-Squares Restraints			3396	Note
PLAT883_ALERT_1_G	No Info/Value for _atom_sites_solution_primary				Please Do !

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

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

