

checkCIF/PLATON report

Structure factors have been supplied for datablock(s) cu_20240708a_1_a

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: cu_20240708a_1_a

Bond precision:	C-C = 0.0186 Å	Wavelength=1.54178	
Cell:	a=30.283(1) alpha=90	b=32.1268(12) beta=94.511(2)	c=36.5647(14) gamma=90
Temperature:	150 K		
	Calculated	Reported	
Volume	35464(2)	35463(2)	
Space group	P 21/n	P 1 21/n 1	
Hall group	-P 2yn	-P 2yn	
Moiety formula	2(C144 H140 N8 O12 Rh6), 7(C F3 O3 S) [+ solvent]	2(C144 H140 N8 O12 Rh6), 7(C F3 O3 S), 11.5[CF3SO3], 1[CH3OH]	
Sum formula	C295 H280 F21 N16 O45 Rh12 S7 [+ solvent]	C307.50 H284 F55.50 N16 O80.50 Rh12 S18.50	
Mr	6627.71	8374.02	
Dx, g cm ⁻³	1.241	1.568	
Z	4	4	
Mu (mm ⁻¹)	5.369	6.342	
F000	13452.0	16882.0	
F000'	13504.08		
h, k, lmax	33, 35, 40	33, 35, 40	
Nref	51466	51070	
Tmin, Tmax	0.444, 0.386		
Tmin'	0.336		

Correction method= Not given

Data completeness= 0.992 Theta (max)= 59.256

R(reflections)= 0.1055(21206)

wR2 (reflections)=
0.2598(51070)

S = 1.094

Npar= 3409

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

RINTA01_ALERT_3_B	The value of Rint is greater than 0.18	
	Rint given	0.186
THETM01_ALERT_3_B	The value of sine(theta_max)/wavelength is less than 0.575	
	Calculated sin(theta_max)/wavelength =	0.5574
PLAT020_ALERT_3_B	The Value of Rint is Greater Than 0.12	0.186 Report
PLAT260_ALERT_2_B	Large Average Ueq of Residue Including	S3 0.384 Check
PLAT260_ALERT_2_B	Large Average Ueq of Residue Including	S5 0.327 Check

Alert level C

PLAT026_ALERT_3_C	Ratio Observed / Unique Reflections (too) Low ..	42% Check
PLAT052_ALERT_1_C	Info on Absorption Correction Method Not Given	Please Do !
PLAT082_ALERT_2_C	High R1 Value	0.11 Report
PLAT084_ALERT_3_C	High wR2 Value (i.e. > 0.25)	0.26 Report
PLAT220_ALERT_2_C	NonSolvent Resd 1 C Ueq(max)/Ueq(min) Range	4.3 Ratio
PLAT220_ALERT_2_C	NonSolvent Resd 2 C Ueq(max)/Ueq(min) Range	4.1 Ratio
PLAT222_ALERT_3_C	NonSolvent Resd 1 H Uiso(max)/Uiso(min) Range	5.3 Ratio
PLAT222_ALERT_3_C	NonSolvent Resd 2 H Uiso(max)/Uiso(min) Range	5.1 Ratio
PLAT242_ALERT_2_C	Low 'MainMol' Ueq as Compared to Neighbors of	Rh1 Check
PLAT242_ALERT_2_C	Low 'MainMol' Ueq as Compared to Neighbors of	Rh3 Check
PLAT242_ALERT_2_C	Low 'MainMol' Ueq as Compared to Neighbors of	Rh8 Check
PLAT242_ALERT_2_C	Low 'MainMol' Ueq as Compared to Neighbors of	Rh12 Check
PLAT242_ALERT_2_C	Low 'MainMol' Ueq as Compared to Neighbors of	N7 Check
PLAT242_ALERT_2_C	Low 'MainMol' Ueq as Compared to Neighbors of	N15 Check
PLAT242_ALERT_2_C	Low 'MainMol' Ueq as Compared to Neighbors of	Rh2 Check
PLAT242_ALERT_2_C	Low 'MainMol' Ueq as Compared to Neighbors of	Rh7 Check
PLAT242_ALERT_2_C	Low 'MainMol' Ueq as Compared to Neighbors of	Rh9 Check
PLAT244_ALERT_4_C	Low 'Solvent' Ueq as Compared to Neighbors of	S1 Check
PLAT244_ALERT_4_C	Low 'Solvent' Ueq as Compared to Neighbors of	S2 Check
PLAT244_ALERT_4_C	Low 'Solvent' Ueq as Compared to Neighbors of	S3 Check
PLAT244_ALERT_4_C	Low 'Solvent' Ueq as Compared to Neighbors of	S4 Check
PLAT244_ALERT_4_C	Low 'Solvent' Ueq as Compared to Neighbors of	S5 Check
PLAT244_ALERT_4_C	Low 'Solvent' Ueq as Compared to Neighbors of	S6 Check
PLAT244_ALERT_4_C	Low 'Solvent' Ueq as Compared to Neighbors of	S7 Check
PLAT250_ALERT_2_C	Large U3/U1 Ratio for <U(i,j)> Tensor(Resd 8)	2.7 Note
PLAT260_ALERT_2_C	Large Average Ueq of Residue Including	Rh1 0.144 Check
PLAT260_ALERT_2_C	Large Average Ueq of Residue Including	Rh2 0.126 Check
PLAT260_ALERT_2_C	Large Average Ueq of Residue Including	S1 0.220 Check
PLAT260_ALERT_2_C	Large Average Ueq of Residue Including	S2 0.251 Check
PLAT260_ALERT_2_C	Large Average Ueq of Residue Including	S4 0.239 Check
PLAT260_ALERT_2_C	Large Average Ueq of Residue Including	S6 0.213 Check
PLAT260_ALERT_2_C	Large Average Ueq of Residue Including	S7 0.240 Check
PLAT342_ALERT_3_C	Low Bond Precision on C-C Bonds	0.01859 Ang.
PLAT369_ALERT_2_C	Long C(sp2)-C(sp2) Bond C232 - C258 ..	1.53 Ang.
PLAT410_ALERT_2_C	Short Intra H...H Contact H216 ..H230 ..	1.99 Ang.

	x,y,z =	1_555 Check
PLAT434_ALERT_2_C Short Inter HL..HL Contact F15	..F20	. 2.50 Ang.
	3/2-x,-1/2+y,3/2-z =	2_646 Check
PLAT905_ALERT_3_C Negative K value in the Analysis of Variance ...		-14.687 Report
PLAT905_ALERT_3_C Negative K value in the Analysis of Variance ...		-0.216 Report
PLAT911_ALERT_3_C Missing FCF Refl Between Thmin & STh/L=	0.557	395 Report
2 0 0, 1 1 0, 1 2 0, 2 2 0, 3 2 0, 5 2 0,		
6 3 0, 33 7 0, 1 10 0, 32 11 0, 15 32 0, 13 33 0,		
7 35 0, -31 0 1, 1 0 1, -5 1 1, -3 1 1, 1 1 1,		
3 1 1, -2 2 1, -1 2 1, 0 2 1, 5 2 1, 9 3 1,		
2 5 1, 4 5 1, 33 6 1, -33 7 1, 11 9 1, -31 14 1,		
30 16 1, 29 18 1, 15 32 1, -13 33 1, -7 35 1, 7 35 1,		
0 0 2, 2 0 2, -2 1 2, -1 1 2, 0 1 2, 1 2 2,		
-3 3 2, -1 3 2, 0 3 2, 4 4 2, 6 4 2, 33 5 2,		
33 6 2, -31 14 2, -28 20 2, -7 35 2, -1 0 3, 1 0 3,		
33 0 3, -2 1 3, -3 2 3, 0 2 3, 2 2 3, 13 2 3,		
-1 3 3, 4 3 3, 33 5 3, 32 10 3, 31 13 3, -31 14 3,		
-28 20 3, 18 30 3, -13 33 3, 10 34 3, 0 0 4, -4 3 4,		
33 3 4, 3 4 4, 33 4 4, 32 9 4, -31 14 4, 28 19 4,		
-15 32 4, 6 35 4, 33 0 5, -7 1 5, 33 1 5, 1 2 5,		
33 2 5, -33 7 5, -31 14 5, 29 17 5, 16 31 5, 12 33 5,		
-10 34 5, -6 35 5, -5 1 6, -1 1 6, 8 3 6, -33 7 6,		
PLAT977_ALERT_2_C Check Negative Difference Density on H28H	.	-0.34 eA-3

● 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: C307.5 H284 F55.5 N16 O80.5 R
 Atom count from _chemical_formula_moiety: C306.5 H280 F55.5 N16 O79.5 R

FORMU01_ALERT_2_G There is a discrepancy between the atom counts in the _chemical_formula_sum and the formula from the _atom_site* data.
 Atom count from _chemical_formula_sum: C307.5 H284 F55.5 N16 O80.5 Rh12
 Atom count from the _atom_site data: C295 H280 F21 N16 O45 Rh12 S7

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 4
 From the CIF: _chemical_formula_sum C307.50 H284 F55.50 N16 O80.50 Rh1
 TEST: Compare cell contents of formula and atom_site data

atom	Z*formula	cif	sites	diff
C	1230.00	1180.00	50.00	
H	1136.00	1120.00	16.00	
F	222.00	84.00	138.00	
N	64.00	64.00	0.00	
O	322.00	180.00	142.00	
Rh	48.00	48.00	0.00	
S	74.00	28.00	46.00	

PLAT002_ALERT_2_G Number of Distance or Angle Restraints on AtSite	161 Note
PLAT003_ALERT_2_G Number of Uiso or U(i,j) Restrained non-H Atoms	396 Report
PLAT041_ALERT_1_G Calc. and Reported SumFormula Strings Differ	Please Check
Calc: C295 H280 F21 N16 O45 Rh12 S7	
Rep.: C307.50 H284 F55.50 N16 O80.50 Rh12 S18.50	
PLAT042_ALERT_1_G Calc. and Reported MoietyFormula Strings Differ	Please Check
Calc: 2(C144 H140 N8 O12 Rh6), 7(C F3 O3 S)	

Rep.: 2(C144 H140 N8 O12 Rh6), 7(C F3 O3 S), 11.5[CF3SO3], 1[CH3OH]

PLAT191_ALERT_3_G	A	Non-default SADI Restraint Value has been used	0.0400	Report
PLAT191_ALERT_3_G	A	Non-default SADI Restraint Value has been used	0.0400	Report
PLAT191_ALERT_3_G	A	Non-default SADI Restraint Value has been used	0.0400	Report
PLAT244_ALERT_4_G	Low	'Solvent' Ueq as Compared to Neighbors of	C9	Check
PLAT244_ALERT_4_G	Low	'Solvent' Ueq as Compared to Neighbors of	C73	Check
PLAT244_ALERT_4_G	Low	'Solvent' Ueq as Compared to Neighbors of	C147	Check
PLAT244_ALERT_4_G	Low	'Solvent' Ueq as Compared to Neighbors of	C289	Check
PLAT244_ALERT_4_G	Low	'Solvent' Ueq as Compared to Neighbors of	C293	Check
PLAT244_ALERT_4_G	Low	'Solvent' Ueq as Compared to Neighbors of	C145	Check
PLAT244_ALERT_4_G	Low	'Solvent' Ueq as Compared to Neighbors of	C291	Check
PLAT344_ALERT_2_G	Unusual	sp3 Angle Range in Solvent/Ion for	C73	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C2	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C245	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C273	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C233	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C257	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C33	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C285	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C237	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C59	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C61	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C253	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C265	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C277	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C69	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C71	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C235	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C117	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C247	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C243	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C123	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C255	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C127	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C267	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C263	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C133	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C275	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C137	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C283	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C143	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C193	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C5	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C7	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C11	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C217	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C21	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C205	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C27	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C43	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C197	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C49	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C51	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C213	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C225	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C87	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C85	Check
PLAT380_ALERT_4_G	Incorrectly?	Oriented X(sp2)-Methyl Moiety	C195	Check

PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety C97 Check
 PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety C207 Check
 PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety C203 Check
 PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety C103 Check
 PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety C215 Check
 PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety C107 Check
 PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety C227 Check
 PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety C223 Check
 PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety C113 Check
 PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety C175 Check
 PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety C169 Check
 PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety C171 Check
 PLAT380_ALERT_4_G Incorrectly? Oriented X(sp2)-Methyl Moiety C173 Check
 PLAT432_ALERT_2_G Short Inter X...Y Contact F11 ..C34 . 2.86 Ang.
 x,y,z = 1_555 Check
 PLAT432_ALERT_2_G Short Inter X...Y Contact F19 ..C139 . 2.86 Ang.
 -1/2+x,1/2-y,1/2+z = 4_566 Check
 PLAT432_ALERT_2_G Short Inter X...Y Contact F19 ..C287 . 2.95 Ang.
 -1/2+x,1/2-y,1/2+z = 4_566 Check
 PLAT432_ALERT_2_G Short Inter X...Y Contact O15 ..C183 . 2.99 Ang.
 x,1+y,z = 1_565 Check
 PLAT434_ALERT_2_G Short Inter HL..HL Contact F5 ..F6 . 2.73 Ang.
 x,y,z = 1_555 Check
 PLAT606_ALERT_4_G Solvent Accessible VOID(S) in Structure ! Info
 PLAT860_ALERT_3_G Number of Least-Squares Restraints 6161 Note
 PLAT868_ALERT_4_G ALERTS Due to the Use of _smtbx_masks Suppressed ! Info
 PLAT883_ALERT_1_G No Info/Value for _atom_sites_solution_primary . Please Do !
 PLAT933_ALERT_2_G Number of HKL-OMIT Records in Embedded .res File 53 Note
 -7 1 5, -7 4 9, -5 1 1, -5 1 6, -4 3 4, -3 1 1,
 -3 2 3, -3 3 2, -2 1 2, -2 1 3, -2 1 7, -2 2 1,
 -1 0 3, -1 1 2, -1 1 6, -1 2 1, -1 3 2, -1 3 3,
 0 0 2, 0 0 4, 0 1 2, 0 2 1, 0 2 3, 0 3 2,
 0 10 6, 1 0 1, 1 0 3, 1 1 0, 1 1 1, 1 2 0,
 1 2 2, 1 2 5, 1 10 0, 2 0 0, 2 0 2, 2 2 0,
 2 2 3, 2 5 1, 3 1 1, 3 2 0, 3 4 4, 4 3 3,
 4 4 2, 4 5 1, 5 2 0, 5 2 1, 6 3 0, 6 4 2,
 7 4 8, 8 3 6,
 PLAT969_ALERT_5_G The 'Henn et al.' R-Factor-gap value 1.847 Note
 Predicted wR2: Based on SigI**2 14.07 or SHELX Weight 23.76
 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
 5 **ALERT level B** = A potentially serious problem, consider carefully
 40 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight
 145 **ALERT level G** = General information/check it is not something unexpected

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

PLATON version of 13/05/2024; check.def file version of 04/05/2024

Datablock cu_20240708a_1_a - ellipsoid plot

