

checkCIF/PLATON report

Structure factors have been supplied for datablock(s) SAIOC-1@G12

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: SAIOC-1@G12

Bond precision:	C-C = 0.0121 Å	Wavelength=1.34050	
Cell:	a=29.9771(3) alpha=90	b=29.9771(3) beta=90	c=19.4854(3) gamma=90
Temperature:	100 K		
	Calculated	Reported	
Volume	17510.1(4)	17510.1(4)	
Space group	P 4/n	P 4/n	
Hall group	-P 4a	-P 4a	
Moiety formula	C216 H360 Al32 N12 O110 Pb6 S12, 3(C10 H8 N2) [+ solvent]	C216 H360 Al32 N12 O110 Pb6 S12, 3(C10 H8 N2)	
Sum formula	C246 H384 Al32 N18 O110 Pb6 S12 [+ solvent]	C246 H384 Al32 N18 O110 Pb6 S12	
Mr	7845.01	7844.91	
Dx, g cm ⁻³	1.488	1.488	
Z	2	2	
Mu (mm ⁻¹)	5.082	5.225	
F000	7932.0	7932.0	
F000'	7911.62		
h, k, lmax	35, 35, 22	32, 35, 22	
Nref	14929	14897	
Tmin, Tmax		1.000, 1.000	
Tmin'			
Correction method=	# Reported T Limits: Tmin=1.000 Tmax=1.000		
AbsCorr =	SPHERE		
Data completeness=	0.998	Theta(max)= 52.048	

R(reflections)= 0.0582(12193)

wR2(reflections)=
0.1635(14897)

S = 1.038

Npar= 967

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

PLAT242_ALERT_2_B Low 'MainMol' Ueq as Compared to Neighbors of C24 Check

Author Response: These alerts are generated because of a slight unresolved disorder caused by free rotation of the flexible chain at the end of the ligand. Some refi commands such as DELU, SIMU and ISOR were used to address such abnormal Ueq value, but failed. After inspection, this does not affect the correct assign of atom types.



Alert level C

RADNW01_ALERT_1_C The radiation wavelength lies outside the expected range
for the supplied radiation type. Expected range 1.34130-1.34150
Wavelength given = 1.34050

THETM01_ALERT_3_C The value of sine(theta_max)/wavelength is less than 0.590
Calculated sin(theta_max)/wavelength = 0.5882

PLAT094_ALERT_2_C	Ratio of Maximum / Minimum Residual Density	2.58	Report
PLAT213_ALERT_2_C	Atom O24 has ADP max/min Ratio	3.5	prolat
PLAT215_ALERT_3_C	Disordered C54 has ADP max/min Ratio	3.9	Note
PLAT220_ALERT_2_C	NonSolvent Resd 1 C Ueq(max)/Ueq(min) Range	4.3	Ratio
PLAT220_ALERT_2_C	NonSolvent Resd 1 O Ueq(max)/Ueq(min) Range	5.7	Ratio
PLAT222_ALERT_3_C	NonSolvent Resd 1 H Uiso(max)/Uiso(min) Range	6.9	Ratio
PLAT241_ALERT_2_C	High 'MainMol' Ueq as Compared to Neighbors of	C40	Check
PLAT241_ALERT_2_C	High 'MainMol' Ueq as Compared to Neighbors of	C51	Check
PLAT242_ALERT_2_C	Low 'MainMol' Ueq as Compared to Neighbors of	S2	Check

Author Response: These alerts are generated because of a slight unresolved disorder caused by free rotation of the flexible chain at the end of the ligand. Some refi commands such as DELU, SIMU and ISOR were used to address such abnormal Ueq value, but failed. After inspection, this does not affect the correct assign of atom types.

PLAT242_ALERT_2_C Low 'MainMol' Ueq as Compared to Neighbors of

S3 Check

Author Response: These alerts are generated because of a slight unresolved disorder caused by free rotation of the flexible chain at the end of the ligand. Some refi commands such as DELU, SIMU and ISOR were used to address such abnormal Ueq value, but failed. After inspection, this does not affect the correct assign of atom types.

PLAT242_ALERT_2_C Low 'MainMol' Ueq as Compared to Neighbors of N1 Check

Author Response: These alerts are generated because of a slight unresolved disorder caused by free rotation of the flexible chain at the end of the ligand. Some refi commands such as DELU, SIMU and ISOR were used to address such abnormal Ueq value, but failed. After inspection, this does not affect the correct assign of atom types.

PLAT242_ALERT_2_C Low 'MainMol' Ueq as Compared to Neighbors of N2 Check

Author Response: These alerts are generated because of a slight unresolved disorder caused by free rotation of the flexible chain at the end of the ligand. Some refi commands such as DELU, SIMU and ISOR were used to address such abnormal Ueq value, but failed. After inspection, this does not affect the correct assign of atom types.

PLAT242_ALERT_2_C Low 'MainMol' Ueq as Compared to Neighbors of N6 Check

Author Response: These alerts are generated because of a slight unresolved disorder caused by free rotation of the flexible chain at the end of the ligand. Some refi commands such as DELU, SIMU and ISOR were used to address such abnormal Ueq value, but failed. After inspection, this does not affect the correct assign of atom types.

PLAT242_ALERT_2_C Low 'MainMol' Ueq as Compared to Neighbors of C18 Check

Author Response: These alerts are generated because of a slight unresolved disorder caused by free rotation of the flexible chain at the end of the ligand. Some refi commands such as DELU, SIMU and ISOR were used to address such abnormal Ueq value, but failed. After inspection, this does not affect the correct assign of atom types.

PLAT242_ALERT_2_C Low 'MainMol' Ueq as Compared to Neighbors of C21 Check

Author Response: These alerts are generated because of a slight unresolved disorder caused by free rotation of the flexible chain at the end of the ligand. Some refi commands such as DELU, SIMU and ISOR were used to address such abnormal Ueq value, but failed. After inspection, this does not affect the correct assign of atom types.

PLAT242_ALERT_2_C Low 'MainMol' Ueq as Compared to Neighbors of C27 Check

Author Response: These alerts are generated because of a slight unresolved disorder caused by free rotation of the flexible chain at the end of the ligand. Some refi commands such as DELU, SIMU and ISOR were used to address such abnormal Ueq value, but failed. After inspection, this does not affect the correct assign of atom types.

PLAT242_ALERT_2_C Low 'MainMol' Ueq as Compared to Neighbors of C34 Check

Author Response: These alerts are generated because of a slight unresolved disorder caused by free rotation of the flexible chain at the end of the ligand. Some refi commands such as DELU, SIMU and ISOR were used to address such abnormal Ueq value, but failed. After inspection, this does not affect the correct assign of atom types.

PLAT250_ALERT_2_C	Large U3/U1 Ratio for <U(i,j)> Tensor(Resd	2)	2.3	Note
PLAT260_ALERT_2_C	Large Average Ueq of Residue Including	Pb1	0.107	Check
PLAT260_ALERT_2_C	Large Average Ueq of Residue Including	N4	0.114	Check
PLAT342_ALERT_3_C	Low Bond Precision on C-C Bonds		0.01215	Ang.
PLAT360_ALERT_2_C	Short C(sp3)-C(sp3) Bond C15 - C17		1.43	Ang.
PLAT906_ALERT_3_C	Large K Value in the Analysis of Variance		2.626	Check
PLAT911_ALERT_3_C	Missing FCF Refl Between Thmin & STh/L=	0.588	30	Report
	-1 3 0, -2 4 1, -4 5 1, 4 6 1, 3 17 1, 0 1 2,			
	1 2 2, 0 3 2, -2 5 2, 3 5 2, 5 7 2, -1 9 2,			
	-2 10 2, 3 10 2, -5 11 2, 0 2 3, 6 10 3, -7 12 3,			
	2 5 4, -2 7 4, -2 14 4, -2 4 6, -9 11 6, 0 12 6,			
	-4 14 6, 3 11 18, 3 12 18, 3 13 21, 4 13 21, 5 13 21,			
PLAT971_ALERT_2_C	Check Calcd Resid. Dens.	1.30Ang From Pb1	2.22	eA-3
PLAT971_ALERT_2_C	Check Calcd Resid. Dens.	1.05Ang From Pb2	2.16	eA-3
PLAT971_ALERT_2_C	Check Calcd Resid. Dens.	1.12Ang From Pb2	1.96	eA-3
PLAT971_ALERT_2_C	Check Calcd Resid. Dens.	0.95Ang From Pb1	1.88	eA-3
PLAT971_ALERT_2_C	Check Calcd Resid. Dens.	1.09Ang From Pb3	1.81	eA-3
PLAT971_ALERT_2_C	Check Calcd Resid. Dens.	0.89Ang From O19	1.74	eA-3
PLAT971_ALERT_2_C	Check Calcd Resid. Dens.	0.24Ang From Pb1	1.74	eA-3
PLAT971_ALERT_2_C	Check Calcd Resid. Dens.	1.11Ang From O2	1.73	eA-3
PLAT971_ALERT_2_C	Check Calcd Resid. Dens.	1.09Ang From Pb3	1.68	eA-3
PLAT971_ALERT_2_C	Check Calcd Resid. Dens.	0.79Ang From Pb2	1.53	eA-3
PLAT973_ALERT_2_C	Check Calcd Positive Resid. Density on	Pb2	1.39	eA-3
PLAT973_ALERT_2_C	Check Calcd Positive Resid. Density on	Pb3	1.30	eA-3



Alert level G

ABSMU01_ALERT_1_G Calculation of _exptl_absorpt_correction_mu

not performed for this radiation type.							
PLAT002_ALERT_2_G	Number of Distance or Angle Restraints on AtSite	21 Note					
PLAT003_ALERT_2_G	Number of Uiso or U(i,j) Restrained non-H Atoms	20 Report					
PLAT007_ALERT_5_G	Number of Unrefined Donor-H Atoms	2 Report					
H3	H21						
PLAT051_ALERT_1_G	Mu(calc) and Mu(CIF) Ratio Differs from 1.0 by .	2.74 %					
PLAT083_ALERT_2_G	SHELXL Second Parameter in WGHT Unusually Large	28.55 Why ?					
PLAT172_ALERT_4_G	The CIF-Embedded .res File Contains DFIX Records	15 Report					
PLAT177_ALERT_4_G	The CIF-Embedded .res File Contains DELU Records	4 Report					
PLAT178_ALERT_4_G	The CIF-Embedded .res File Contains SIMU Records	4 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	0.0100 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	0.0100 Report					
PLAT192_ALERT_3_G	A Non-default DELU Restraint Value for SecondPar	0.0200 Report					
PLAT192_ALERT_3_G	A Non-default DELU Restraint Value for SecondPar	0.0200 Report					
PLAT192_ALERT_3_G	A Non-default DELU Restraint Value for SecondPar	0.0200 Report					
PLAT192_ALERT_3_G	A Non-default DELU Restraint Value for SecondPar	0.0200 Report					
PLAT299_ALERT_4_G	Atom Site Occupancy Constrained at	0.5 Check					
C53	C54	H53A	H53B	H53C	H54A	H54B	H54C
PLAT300_ALERT_4_G	Atom Site Occupancy of C7				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C43				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C6				Constrained at		0.25 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C42				Constrained at		0.25 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H5A				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H5B				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H7A				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H7B				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H7C				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H41A				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H41B				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H43A				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H43B				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H43C				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H52A				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H52B				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H6A				Constrained at		0.25 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H6B				Constrained at		0.25 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H6C				Constrained at		0.25 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H42A				Constrained at		0.25 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H42B				Constrained at		0.25 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H42C				Constrained at		0.25 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H52C				Constrained at		0.25 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H52D				Constrained at		0.25 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of N4				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of N5				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C58				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C59				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C60				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C61				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C62				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C63				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C64				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C65				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C66				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of C67				Constrained at		0.75 Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H58				Constrained at		0.75 Check

PLAT300_ALERT_4_G	Atom Site Occupancy of H59	Constrained at	0.75	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H61	Constrained at	0.75	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H62	Constrained at	0.75	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H63	Constrained at	0.75	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H65	Constrained at	0.75	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H66	Constrained at	0.75	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of H67	Constrained at	0.75	Check
PLAT301_ALERT_3_G	Main Residue Disorder	(Resd 1)	3%	Note
PLAT302_ALERT_4_G	Anion/Solvent/Minor-Residue Disorder	(Resd 2)	100%	Note
PLAT413_ALERT_2_G	Short Inter XH3 .. XHn	H53C ..H57C	1.73	Ang.
	-1/2+y,1-x,1-z =		7_566	Check
PLAT605_ALERT_4_G	Largest Solvent Accessible VOID in the Structure		72	A**3
PLAT722_ALERT_1_G	Angle Calc 108.00, Rep 109.50 Dev...		1.50	Degree
	H42A -C42 -H42B	1_555 1_555 1_555	# 581	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -PB1 4_565	1_555 1_555	# 182	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -PB1 3_655	1_555 1_555	# 183	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -PB1 2_665	1_555 1_555	# 184	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -AL1 3_655	1_555 4_565	# 185	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -AL1 3_655	1_555 3_655	# 186	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -AL1 4_565	1_555 3_655	# 187	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -AL1 2_665	1_555 3_655	# 188	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -AL1 4_565	1_555 1_555	# 189	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -AL1 2_665	1_555 4_565	# 190	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -AL1 4_565	1_555 2_665	# 191	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -AL1 4_565	1_555 4_565	# 192	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -AL1 3_655	1_555 2_665	# 193	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -AL1 3_655	1_555 1_555	# 194	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -AL1 2_665	1_555 1_555	# 195	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -AL1 2_665	1_555 2_665	# 196	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -O2 3_655	1_555 2_665	# 197	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -O2 4_565	1_555 3_655	# 198	Check
PLAT779_ALERT_4_G	Suspect or Irrelevant (Bond) Angle(s) in CIF ...		0.00	Deg.
	O2 -O2 -O2 4_565	1_555 2_665	# 199	Check
PLAT794_ALERT_5_G	Tentative Bond Valency for Pb1	(II)	2.01	Info
PLAT794_ALERT_5_G	Tentative Bond Valency for Pb3	(II)	2.11	Info
PLAT794_ALERT_5_G	Tentative Bond Valency for Al1	(III)	2.76	Info
PLAT794_ALERT_5_G	Tentative Bond Valency for Al2	(III)	2.75	Info
PLAT794_ALERT_5_G	Tentative Bond Valency for Al4	(III)	2.76	Info
PLAT794_ALERT_5_G	Tentative Bond Valency for Al6	(III)	2.82	Info
PLAT794_ALERT_5_G	Tentative Bond Valency for Al7	(III)	2.81	Info

PLAT794_ALERT_5_G	Tentative Bond Valency for Al8	(III)	.	2.83	Info
PLAT802_ALERT_4_G	CIF Input Record(s) with more than 80 Characters			2	Info
PLAT860_ALERT_3_G	Number of Least-Squares Restraints		127	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	!
PLAT909_ALERT_3_G	Percentage of I>2sig(I) Data at Theta(Max)	Still		57%	Note
PLAT910_ALERT_3_G	Missing # of FCF Reflection(s) Below Theta(Min).			2	Note
	1 1 0, 0 0 1,				
PLAT913_ALERT_3_G	Missing # of Very Strong Reflections in FCF		1	Note
	1 1 0,				
PLAT933_ALERT_2_G	Number of HKL-OMIT Records in Embedded .res File			23	Note
	-1 3 0, 0 3 2, 0 1 2, -2 10 2, 1 2 2, 3 5 2,				
	-2 7 4, -5 11 2, -2 5 2, 0 2 3, -2 4 1, -1 9 2,				
	2 5 4, -9 11 6, 3 17 1, -2 4 6, -7 12 3, -4 5 1,				
	0 12 6, -4 14 6, 5 7 2, 3 10 2, -2 14 4,				
PLAT941_ALERT_3_G	Average HKL Measurement Multiplicity		3.4	Low
PLAT950_ALERT_5_G	Calculated (ThMax) and CIF-Reported Hmax Differ			3	Units
PLAT969_ALERT_5_G	The 'Henn et al.' R-Factor-gap value		3.449	Note
	Predicted wr2: Based on SigI**2	4.74	or SHELX Weight	15.76	
PLAT978_ALERT_2_G	Number C-C Bonds with Positive Residual Density.			0	Info
PLAT984_ALERT_1_G	The C-f' =	0.0147	Deviates from the B&C-Value	0.0137	Check
PLAT984_ALERT_1_G	The N-f' =	0.0253	Deviates from the B&C-Value	0.0241	Check
PLAT984_ALERT_1_G	The O-f' =	0.0412	Deviates from the B&C-Value	0.0389	Check
PLAT984_ALERT_1_G	The Pb-f' =	-4.2646	Deviates from the B&C-Value	-4.4950	Check
PLAT985_ALERT_1_G	The Al-f" =	0.1843	Deviates from the B&C-Value	0.1873	Check
PLAT985_ALERT_1_G	The Pb-f" =	7.1982	Deviates from the B&C-Value	6.8412	Check
PLAT985_ALERT_1_G	The S-f" =	0.4242	Deviates from the B&C-Value	0.4295	Check

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

12 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
 20 ALERT type 3 Indicator that the structure quality may be low
 70 ALERT type 4 Improvement, methodology, query or suggestion
 11 ALERT type 5 Informative message, check

checkCIF publication errors

Alert level A

PUBL004_ALERT_1_A The contact author's name and address are missing,
 _publ_contact_author_name and _publ_contact_author_address.

PUBL005_ALERT_1_A _publ_contact_author_email, _publ_contact_author_fax and
 _publ_contact_author_phone are all missing.
 At least one of these should be present.

PUBL006_ALERT_1_A _publ_requested_journal is missing
 e.g. 'Acta Crystallographica Section C'

PUBL008_ALERT_1_A _publ_section_title is missing. Title of paper.

PUBL009_ALERT_1_A _publ_author_name is missing. List of author(s) name(s).

PUBL010_ALERT_1_A _publ_author_address is missing. Author(s) address(es).

PUBL012_ALERT_1_A _publ_section_abstract is missing.
Abstract of paper in English.

7 **ALERT level A** = Data missing that is essential or data in wrong format
0 **ALERT level G** = General alerts. Data that may be required is missing

Publication of your CIF

You should 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 nature of your study may justify the reported deviations from journal submission requirements and the more serious of these should 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.

If level A alerts remain, which you believe to be justified deviations, and you intend to submit this CIF for publication in a journal, you should additionally insert an explanation in your CIF using the Validation Reply Form (VRF) below. This will allow your explanation to be considered as part of the review process.

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_PUBL004_GLOBAL
;
PROBLEM: The contact author's name and address are missing,
RESPONSE: ...
;
_vrf_PUBL005_GLOBAL
;
PROBLEM: _publ_contact_author_email, _publ_contact_author_fax and
RESPONSE: ...
;
_vrf_PUBL006_GLOBAL
;
PROBLEM: _publ_requested_journal is missing
RESPONSE: ...
;
_vrf_PUBL008_GLOBAL
;
PROBLEM: _publ_section_title is missing. Title of paper.
RESPONSE: ...
;
```



```
_vrf_PUBL009_GLOBAL
;
PROBLEM: _publ_author_name is missing. List of author(s) name(s).
RESPONSE: ...
;
_vrf_PUBL010_GLOBAL
;
PROBLEM: _publ_author_address is missing. Author(s) address(es).
RESPONSE: ...
;
_vrf_PUBL012_GLOBAL
;
PROBLEM: _publ_section_abstract is missing.
RESPONSE: ...
;
# end Validation Reply Form
```

If you wish to submit your CIF for publication in Acta Crystallographica Section C or E, you should upload your CIF via the web. If you wish to submit your CIF for publication in IUCrData you should upload your CIF via the web. If your CIF is to form part of a submission to another IUCr journal, you will be asked, either during electronic submission or by the Co-editor handling your paper, to upload your CIF via our web site.

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