

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

Bond precision: Si- O = 0.0060 Å Wavelength=0.71073

Cell: a=12.7351(10) b=12.7351(10) c=18.0797(14)
 alpha=90 beta=90 gamma=90
Temperature: 293 K

	Calculated	Reported
Volume	2932.2(5)	2932.2(4)
Space group	P -4 n 2	P -4 n 2
Hall group	P -4 -2n	P -4;-2n
Moiety formula	Mo O40 Si W11, 30(O)	?
Sum formula	Mo O70 Si W11	H64 Mo1 O70 Si1 W11
Mr	3266.31	3330.82
Dx,g cm-3	3.700	3.773
Z	2	2
Mu (mm-1)	21.840	21.842
F000	2860.0	2860.0
F000'	2842.54	
h,k,lmax	15,15,22	15,15,22
Nref	3025[1662]	3013
Tmin,Tmax	0.084,0.174	0.500,0.800
Tmin'	0.033	

Correction method= # Reported T Limits: Tmin=0.500 Tmax=0.800
AbsCorr = MULTI-SCAN

Data completeness= 1.81/1.00 Theta(max)= 26.420

R(reflections)= 0.0229(2677) wR2(reflections)= wR= 0.0276(3013)

S = 0.430 Npar= 175

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

GEOM006_ALERT_1_A `_geom_angle_atom_site_label_2` is missing
Label identifying the atom site 2.
GEOM007_ALERT_1_A `_geom_angle_atom_site_label_3` is missing
Label identifying the atom site 3.
WEIGH01_ALERT_1_A Unit weights are not acceptable for submissions to Acta
Crystallographica Section C.
n.b. unit is however a legal CIF keyword.
PLAT211_ALERT_2_A ADP of Atom O6 is N.P.D. or (nearly) 2D ... Please Check
PLAT213_ALERT_2_A Atom O1 has ADP max/min Ratio 9.4 prolat
PLAT213_ALERT_2_A Atom O5 has ADP max/min Ratio 5.6 prolat

Alert level B

GOODF01_ALERT_2_B The least squares goodness of fit parameter lies
outside the range 0.60 <> 4.00
Goodness of fit given = 0.430
PLAT043_ALERT_1_B Calculated and Reported Mol. Weight Differ by .. 64.51 Check
PLAT214_ALERT_2_B Atom Ow2 (Anion/Solvent) ADP max/min Ratio 5.7 prolat
PLAT220_ALERT_2_B Non-Solvent Resd 1 O Ueq(max)/Ueq(min) Range 6.7 Ratio
PLAT241_ALERT_2_B High 'MainMol' Ueq as Compared to Neighbors of O5 Check
PLAT306_ALERT_2_B Isolated Oxygen Atom (H-atoms Missing ?) Ow1 Check
PLAT306_ALERT_2_B Isolated Oxygen Atom (H-atoms Missing ?) Ow2 Check
PLAT306_ALERT_2_B Isolated Oxygen Atom (H-atoms Missing ?) Ow3 Check
PLAT306_ALERT_2_B Isolated Oxygen Atom (H-atoms Missing ?) Ow4 Check
PLAT306_ALERT_2_B Isolated Oxygen Atom (H-atoms Missing ?) Ow5 Check
PLAT306_ALERT_2_B Isolated Oxygen Atom (H-atoms Missing ?) Ow6 Check
PLAT306_ALERT_2_B Isolated Oxygen Atom (H-atoms Missing ?) Ow7 Check
PLAT306_ALERT_2_B Isolated Oxygen Atom (H-atoms Missing ?) Ow8 Check

Alert level C

ABSTY02_ALERT_1_C An `_exptl_absorpt_correction_type` has been given without
a literature citation. This should be contained in the
`_exptl_absorpt_process_details` field.
Absorption correction given as multi-scan
CHEMW03_ALERT_2_C The ratio of given/expected molecular weight as
calculated from the `_atom_site*` data lies outside
the range 0.99 <> 1.01
From the CIF: `_cell_formula_units_Z` 2
From the CIF: `_chemical_formula_weight` 3330.82
TEST: Calculate formula weight from `_atom_site*`
atom mass num sum
H 1.01 0.00 0.00
Mo 95.94 1.00 95.90
O 16.00 70.00 1119.93
Si 28.09 1.00 28.09
W 183.85 11.00 2022.42
Calculated formula weight 3266.34
STRVA01_ALERT_4_C Flack test results are ambiguous.
From the CIF: `_refine_ls_abs_structure_Flack` 0.500
From the CIF: `_refine_ls_abs_structure_Flack_su` 0.030
PLAT041_ALERT_1_C Calc. and Reported SumFormula Strings Differ Please Check
PLAT127_ALERT_1_C Implicit Hall Symbol Inconsistent with Explicit P -4;-2n
PLAT202_ALERT_3_C Isotropic non-H Atoms in Anion/Solvent 1 Check
PLAT214_ALERT_2_C Atom Ow5 (Anion/Solvent) ADP max/min Ratio 4.9 prolat

PLAT241_ALERT_2_C	High	'MainMol'	Ueq as Compared to Neighbors of	01	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	06	Check

● Alert level G

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: H64 Mo1 O70 Si1 W11
 Atom count from the _atom_site data: Mol.0046 O70 Si1 W11.

CELLZ01_ALERT_1_G Difference between formula and atom_site contents detected.

CELLZ01_ALERT_1_G WARNING: H atoms missing from atom site list. Is this intentional?
 From the CIF: _cell_formula_units_Z 2
 From the CIF: _chemical_formula_sum H64 Mo1 O70 Si1 W11
 TEST: Compare cell contents of formula and atom_site data

atom	Z*formula	cif sites	diff
H	128.00	0.00	128.00
Mo	2.00	2.00	0.00
O	140.00	140.00	0.00
Si	2.00	2.00	0.00
W	22.00	22.00	0.00

PLAT004_ALERT_5_G	Polymeric Structure Found with Maximum Dimension	2	Info
PLAT005_ALERT_5_G	No Embedded Refinement Details found in the CIF		Please Do !
PLAT199_ALERT_1_G	Reported _cell_measurement_temperature (K)	293	Check
PLAT200_ALERT_1_G	Reported _diffrn_ambient_temperature (K)	293	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of >W1 is Constrained at	0.9167	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of >W2 is Constrained at	0.9167	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of >W3 is Constrained at	0.9167	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of <Mo1 is Constrained at	0.0833	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of <Mo2 is Constrained at	0.0833	Check
PLAT300_ALERT_4_G	Atom Site Occupancy of <Mo3 is Constrained at	0.0833	Check
PLAT301_ALERT_3_G	Main Residue Disorder Percentage =	21	Note
PLAT720_ALERT_4_G	Number of Unusual/Non-Standard Labels	8	Note
PLAT792_ALERT_1_G	The Model has Chirality at Si1 (Polar SPGR)	R	Verify
PLAT808_ALERT_5_G	No Parseable SHELXL Style Weighting Scheme Found		Please Check

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

12 ALERT type 1 CIF construction/syntax error, inconsistent or missing data
 21 ALERT type 2 Indicator that the structure model may be wrong or deficient
 2 ALERT type 3 Indicator that the structure quality may be low
 8 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.

