checkCIF/PLATON report

You have not supplied any structure factors. As a result the full set of tests cannot be run.

Datablock: 4b

Bond precision: C-C = 0.0029 A Wavelength=0.71073

Cell: a=15.5497(7) b=14.4945(7) c=12.9298(6)

alpha=90 beta=102.3006(6) gamma=90

Temperature: 173 K

 Calculated
 Reported

 Volume
 2847.3(2)
 2847.3(2)

 Space group
 P 21/c
 P 21/c

 Hall group
 -P 2ybc
 -P 2ybc

Moiety formula C20 H29 N5, 2(C F3 O3 S) C20 H29 N5, 2(C F3 O3 S)

Sum formula C22 H29 F6 N5 O6 S2 C22 H29 F6 N5 O6 S2

Mr 637.64 637.62 Dx,g cm-3 1.487 1.487 Z 4 4 Mu (mm-1) 0.272 0.272 F000 1320.0 1320.0

F000' 1321.86

h,k,lmax 19,18,16 19,18,16 Nref 5837 5831

Tmin, Tmax 0.880, 0.989 0.883, 0.989

Tmin' 0.880

Correction method= INTEGRATION

Data completeness= 0.999 Theta(max)= 26.400

R(reflections) = 0.0361(4762) wR2(reflections) = 0.0968(5831)

S = 1.021 Npar= 372

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 C

PLAT220_ALERT_2_C Large Non-Solvent C Ueq(max)/Ueq(min) ... 3.7 Ratio PLAT242_ALERT_2_C Check Low Ueq as Compared to Neighbors for C15 PLAT242_ALERT_2_C Check Low Ueq as Compared to Neighbors for C18

Alert level G PLAT005_ALERT_5_G No _iucr_refine_instructions_details in CIF ? PLAT432_ALERT_2_G Short Inter X...Y Contact 05 ... C6 ... 2.98 Ang. PLAT432_ALERT_2_G Short Inter X...Y Contact 05 ... C4 ... 2.99 Ang. O ALERT level A = Most likely a serious problem - resolve or explain O ALERT level B = A potentially serious problem, consider carefully 3 ALERT level C = Check. Ensure it is not caused by an omission or oversight 3 ALERT level G = General information/check it is not something unexpected O ALERT type 1 CIF construction/syntax error, inconsistent or missing data 5 ALERT type 2 Indicator that the structure model may be wrong or deficient O ALERT type 3 Indicator that the structure quality may be low

Datablock: 5b

1 ALERT type 5 Informative message, check

Bond precision: C-C = 0.0031 A Wavelength=0.71073

O ALERT type 4 Improvement, methodology, query or suggestion

Cell: a=10.1647(5) b=10.4449(5) c=13.1034(6)

alpha=76.4620(5) beta=72.6898(5) gamma=78.6303(5)

Temperature: 173 K

	Calculated	Reported		
Volume	1279.03(11)	1279.03(11)		
Space group	P -1	P -1		
Hall group	-P 1	-P 1		
Moiety formula	C22 H30 I2 N5 O2 Rh	C22 H30 I2 N5 O2 Rh		
Sum formula	C22 H30 I2 N5 O2 Rh	C22 H30 I2 N5 O2 Rh		
Mr	753.22	753.22		
Dx,g cm-3	1.956	1.956		
Z	2	2		
Mu (mm-1)	3.109	3.109		
F000	728.0	728.0		
F000′	724.08			
h,k,lmax	13,13,17	13,13,17		
Nref	5855	5814		
Tmin,Tmax	0.449,0.856	0.487,0.855		
Tmin'	0.415			

Correction method= NUMERICAL

Data completeness= 0.993 Theta(max)= 27.480

R(reflections) = 0.0172(5305) wR2(reflections) = 0.0436(5814)

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 C

PLAT411_ALERT_2_C Short Inter H...H Contact H5 .. H20 .. 2.14 Ang.

Alert level G

PLAT005_ALERT_5_G No _iucr_refine_instructions_details in CIF ?

PLAT232_ALERT_2_G Hirshfeld Test Diff (M-X) I2 -- Rh .. 12.7 su

PLAT232_ALERT_2_G Hirshfeld Test Diff (M-X) Rh -- O1 .. 5.0 su

PLAT232_ALERT_2_G Hirshfeld Test Diff (M-X) Rh -- C4 .. 5.6 su

- 0 ALERT level A = Most likely a serious problem resolve or explain
- 0 ALERT level B = A potentially serious problem, consider carefully
- 1 ALERT level C = Check. Ensure it is not caused by an omission or oversight
- 4 ALERT level G = General information/check it is not something unexpected
- 0 ALERT type 1 CIF construction/syntax error, inconsistent or missing data
- 4 ALERT type 2 Indicator that the structure model may be wrong or deficient
- O ALERT type 3 Indicator that the structure quality may be low
- 0 ALERT type 4 Improvement, methodology, query or suggestion
- 1 ALERT type 5 Informative message, check

Datablock: 7b

Bond precision: C-C = 0.0214 A Wavelength=0.71073

Cell: a=30.695(19) b=17.496(11) c=15.149(9)

alpha=90 beta=119.530(7) gamma=90

Temperature: 173 K

Volume Space group Hall group	Calculated 7079(8) C 2/c -C 2yc	Reported 7079(7) C2/c -C 2yc		
Moiety formula	C20 H28 I3 N5 Pd, C2 H3 N	C20 H28 I3 N5 Pd, 4(C2 H3 N)		
Sum formula	C22 H31 I3 N6 Pd	C28 H40 I3 N9 Pd		
Mr	866.63	989.79		
Dx,g cm-3	1.626	1.857		
Z	8	8		
Mu (mm-1)	3.158	3.173		
F000	3280.0	3808.0		
F000′	3261.53			
h,k,lmax	38,21,18	38,21,18		
Nref	7250	7143		
Tmin,Tmax	0.664,0.752	0.252,0.757		
Tmin'	0.148			

Correction method= NUMERICAL

Data completeness= 0.985 Theta(max)= 26.400

R(reflections) = 0.0734(4484) wR2(reflections) = 0.2186(7143)

S = 1.080 Npar= 288

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

 ${\tt CHEMW03_ALERT_2_A~ALERT:~The~ratio~of~given/expected~molecular~weight~as~calculated~from~the~_atom_site*~data~lies~outside}$

the range 0.90 <> 1.10

From the CIF: _cell_formula_units_Z 8
From the CIF: _chemical_formula_weight 989.79

TEST: Calculate formula weight from _atom_site_*

mass num sum C 12.01 22.00 264.24 1.01 31.00 Η 31.25 14.01 6.00 84.04 N Pd 106.42 1.00 106.42 3.00 380.71 126.90

Calculated formula weight 866.67

Author Response: Attempts to refine peaks of residual electron density as disordered or partial-occupancy solvent acetonitrile nitrogen or carbon atoms were unsuccessful. The data were corrected for disordered electron density through use of the SQUEEZE procedure (Sluis, P. van der; Spek, A. L. Acta Crystallogr. 1990, A46, 194-201) as implemented in PLATON (Spek, A. L. Acta Crystallogr. 1990, A46, C34; Spek, A. L. J. Appl. Cryst. 2003, 36, 7-13. PLATON - a multipurpose crystallographic tool. Utrecht University, Utrecht, The Netherlands). A total solvent-accessible void volume of 1470 cubic Angstroms with a total electron count of 519(consistent with 24 molecules of solvent acetonitrile, or 3 moleculesper formula unit of the palladium complex) was found in the unit cell.

PLAT043_ALERT_1_A Check Reported Molecular Weight 989.79

Author Response: see response to CHEMW03.

Alert level B PLAT342_ALERT_3_B Low Bond Precision on C-C Bonds	0.0214 Ang
Alert level C PLAT041_ALERT_1_C Calc. and Reported SumFormula Strings Differ	?
Author Response: see response to CHEMW03.	
PLAT042_ALERT_1_C Calc. and Reported MoietyFormula Strings Differ	?
Author Response: see response to CHEMW03.	
PLAT068_ALERT_1_C Reported F000 Differs from Calcd (or Missing)	?

Author Response: see response to CHEMW03.

PLAT234_ALERT_4_C Large Hirshfeld Difference N5 C9	0.16 Ang.
PLAT234_ALERT_4_C Large Hirshfeld Difference C13 C14	0.19 Ang.
PLAT242_ALERT_2_C Check Low	C18
PLAT244_ALERT_4_C Low 'Solvent' Ueq as Compared to Neighbors of	C1S
PLAT250_ALERT_2_C Large U3/U1 Ratio for Average U(i,j) Tensor	2.2
PLAT366_ALERT_2_C Short? C(sp?)-C(sp?) Bond C9 - C10	1.37 Ang.
PLAT366_ALERT_2_C Short? C(sp?)-C(sp?) Bond C10 - C11	1.37 Ang.

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:C28 H40 I3 N9 Pd1

Atom count from the _atom_site data: C22 H31 I3 N6 Pd1

Author Response: see response to CHEMW03.

Author Response: see response to CHEMW03.

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CELLZ01_ALERT_1_G ALERT: Large difference may be due to a
          symmetry error - see SYMMG tests
          From the CIF: _cell_formula_units_Z
          From the CIF: _chemical_formula_sum    C28 H40 I3 N9 Pd
          TEST: Compare cell contents of formula and atom_site data
                 Z*formula cif sites diff
          atom
                 224.00 176.00 48.00
          C
                 320.00 248.00 72.00
          Т
                  24.00
                           24.00 0.00
                  72.00
                           48.00 24.00
          N
          Pd
                   8.00
                            8.00 0.00
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Author Response: see response to CHEMW03.

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PLAT002_ALERT_2_G Number of Distance or Angle Restraints on AtSite 6
PLAT005_ALERT_5_G No _iucr_refine_instructions_details in CIF ... ?
PLAT072_ALERT_2_G SHELXL First Parameter in WGHT Unusually Large. 0.12
PLAT301_ALERT_3_G Note: Main Residue Disorder ...... 10 Perc.
PLAT606_ALERT_4_G VERY LARGE Solvent Accessible VOID(S) in Structure !
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Author Response: see response to CHEMW03.

Author Response: see response to CHEMW03.

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

6 ALERT type 1 CIF construction/syntax error, inconsistent or missing data
8 ALERT type 2 Indicator that the structure model may be wrong or deficient
3 ALERT type 3 Indicator that the structure quality may be low
8 ALERT type 4 Improvement, methodology, query or suggestion
1 ALERT type 5 Informative message, check
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Datablock: 10b

Cell: a=13.782(3) b=15.694(3) c=22.394(5)

alpha=79.906(3) beta=88.010(3) gamma=86.666(3)

Temperature: 173 K

Calculated Reported Volume 4759.2(17) 4759.3(17)

Space group P -1 P -1 Hall group -P 1 -P 1

Moiety formula 2(C38 H43 I2 N5 P Pd), C38 H43 I2 N5 P Pd, 2(C2

3(C2 H3 N), 2(I) H3 N), I

Sum formula C82 H95 I6 N13 P2 Pd2 C42 H49 I3 N7 P Pd

Mr2298.851169.95Dx,g cm-31.6041.633Z24Mu (mm-1)2.4032.405F0002236.02280.0

F000' 2227.31 h,k,lmax 17,20,29 17,20,29 Nref 22166 21838

Tmin, Tmax 0.450, 0.866 0.393, 0.877

Tmin' 0.312

Correction method= NUMERICAL

Data completeness= 0.985 Theta(max)= 27.640

R(reflections) = 0.0351(16825) wR2(reflections) = 0.0924(21838)

S = 1.031 Npar= 951

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.

风 Alert level B

PLAT220_ALERT_2_B Large Non-Solvent C Ueq(max)/Ueq(min) ... 4.4 Ratio

Alert level C

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 4
From the CIF: _chemical_formula_weight 1169.95

TEST: Calculate formula weight from _atom_site_*

 atom
 mass
 num
 sum

 C
 12.01
 41.00
 492.45

 H
 1.01
 47.50
 47.88

 N
 14.01
 6.50
 91.05

P	30.97	1.00	30.97	
Pd	106.42	1.00	106.42	
I	126.90	3.00	380.71	
Calcul	1149.49			

Author Response: Attempts to refine peaks of residual electron density as disordered or partial-occupancy solvent acetonitrile nitrogen or carbon atoms were unsuccessful. The data were corrected for disordered electron density through use of the SQUEEZE procedure (Sluis, P. van der; Spek, A. L. Acta Crystallogr. 1990, A46, 194-201) as implemented in PLATON (Spek, A. L. Acta Crystallogr. 1990, A46, C34; Spek, A. L. J. Appl. Cryst. 2003, 36, 7-13. PLATON - a multipurpose crystallographic tool. Utrecht University, Utrecht, The Netherlands). A total solvent-accessible void volume of 199 cubic Angstroms with a total electron count of 45(consistent with 2 molecule of solvent acetonitrile, or 0.5 molecule performula unit of the palladium complex) was found in the unit cell.

PLAT041_ALERT_1_C Calc. and Reported SumFormula Strings Differ ?

Author Response: see response to CHEMW03.

PLAT042_ALERT_1_C Calc. and Reported MoietyFormula Strings Differ ?

Author Response: see response to CHEMW03.

PLAT043_ALERT_1_C Check Reported Molecular Weight 1169.95

Author Response: see response to CHEMW03.

PLAT045_ALERT_1_C Calculated and Reported Z Differ by	0.50 Ratio
PLATO68 ALERT 1 C Reported FOOD Differs from Calcd (or Missing)	2

Author Response: see response to CHEMW03.

PLAT220_ALERT_2_C Large	Non-Solvent	C	Ueq(r	nax)/Ueq(min)		3.5 Ratio
PLAT242_ALERT_2_C Check	Low Ue	q as	Compared	to	Neighbors	for	C18A
PLAT242_ALERT_2_C Check	Low Ue	q as	Compared	to	Neighbors	for	C15B
PLAT244_ALERT_4_C Low	'Solvent' Ued	q as	Compared	to	Neighbors	of	C1S
PLAT244_ALERT_4_C Low	'Solvent' Ue	q as	Compared	to	Neighbors	of	C3S

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:C42 H49 I3 N7 P1 Pd1

Atom count from the _atom_site data: C41 H47.5 I3 N6.5 P1 Pd1

Author Response: see response to CHEMW03.

Author Response: see response to CHEMW03.

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CELLZ01_ALERT_1_G ALERT: Large difference may be due to a
          symmetry error - see SYMMG tests
          From the CIF: _cell_formula_units_Z
          From the CIF: _chemical_formula_sum    C42 H49 I3 N7 P Pd
          TEST: Compare cell contents of formula and atom_site data
                 Z*formula cif sites diff
          atom
          C
                 168.00 164.00 4.00
                 196.00 190.00 6.00
         Η
         I
                  12.00
                          12.00 0.00
                  28.00
                          26.00 2.00
         N
          Ρ
                   4.00
                           4.00 0.00
                   4.00
                           4.00 0.00
          Рd
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Author Response: see response to CHEMW03.

Author Response: see response to CHEMW03.

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PLAT710_ALERT_4_G Delete 1-2-3 or 2-3-4 Linear Torsion Angle ... #
          I1A -PDA -P1A -C21A 176.50 0.30 1.555 1.555 1.555 1.555
PLAT710_ALERT_4_G Delete 1-2-3 or 2-3-4 Linear Torsion Angle ... #
                                                               6
          I1A -PDA -P1A -C31A -60.80 0.40 1.555 1.555 1.555 1.555
PLAT710_ALERT_4_G Delete 1-2-3 or 2-3-4 Linear Torsion Angle ... #
                                                               9
          I1A -PDA -P1A -C41A 58.30 0.40 1.555 1.555 1.555 1.555
PLAT710_ALERT_4_G Delete 1-2-3 or 2-3-4 Linear Torsion Angle ... #
                                                               23
          I2A -PDA -C1A -N1A -165.20 1.20 1.555 1.555 1.555 1.555
PLAT710_ALERT_4_G Delete 1-2-3 or 2-3-4 Linear Torsion Angle ... #
                                                               26
          I2A -PDA -C1A -N2A 9.50 1.70 1.555 1.555 1.555
PLAT710_ALERT_4_G Delete 1-2-3 or 2-3-4 Linear Torsion Angle ... #
                                                               115
          I1B -PDB -P1B -C21B 149.90 0.30 1.555 1.555 1.555 1.555
PLAT710_ALERT_4_G Delete 1-2-3 or 2-3-4 Linear Torsion Angle ... #
                                                               118
          I1B -PDB -P1B -C31B -87.10 0.30 1.555 1.555 1.555 1.555
PLAT710_ALERT_4_G Delete 1-2-3 or 2-3-4 Linear Torsion Angle ... #
                                                               121
          I1B -PDB -P1B -C41B 32.40 0.30 1.555 1.555 1.555 1.555
PLAT710_ALERT_4_G Delete 1-2-3 or 2-3-4 Linear Torsion Angle ... #
                                                               135
          I2B -PDB -C1B -N1B -162.30 0.80 1.555 1.555 1.555
PLAT710_ALERT_4_G Delete 1-2-3 or 2-3-4 Linear Torsion Angle ... #
                                                               138
          I2B -PDB -C1B -N2B 11.60 1.40 1.555 1.555 1.555
                                                               1.555
PLAT720_ALERT_4_G Number of Unusual/Non-Standard Labels .....
                                                               36
PLAT869_ALERT_4_G ALERTS Related to the use of SQUEEZE Suppressed
                                                                 !
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Author Response: see response to CHEMW03.

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O ALERT level A = Most likely a serious problem - resolve or explain

1 ALERT level B = A potentially serious problem, consider carefully

11 ALERT level C = Check. Ensure it is not caused by an omission or oversight

25 ALERT level G = General information/check it is not something unexpected

8 ALERT type 1 CIF construction/syntax error, inconsistent or missing data

10 ALERT type 2 Indicator that the structure model may be wrong or deficient

0 ALERT type 3 Indicator that the structure quality may be low

18 ALERT type 4 Improvement, methodology, query or suggestion

1 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*, 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 19/04/2012; check.def file version of 14/04/2012







