

checkCIF () running

Checking for embedded fcf data in CIF ...

Found embedded fcf data in CIF. Extracting fcf data from uploaded CIF, please wait . .

```
+++++++ + SHELXL - CRYSTAL STRUCTURE
REFINEMENT - MULTI-CPU VERSION + + Copyright(C) George M. Sheldrick 1993-2014 Version 2014/7 + + infile_sx
started at 05:18:56 on 04-Aug-2016 + ++++++++
Command line parameters: infile_sx -a50000 -b3000 -c624 -t8 -a sets the approximate maximum number of atoms
including hydrogens. -b sets the maximum number of full-matrix parameters (leave unchanged for CGLS). For
example -b9000 allows refinement of 1000 anisotropic atoms or 3000 with BLOC 1. For a 32-bit version, -b times the
square root of the number of threads should not exceed about 65500. -c sets the reflection buffer size. This depends
on the CPU cache size but will rarely need changing. -t sets the number of threads, otherwise the multi-CPU version
sets this equal to the number of available CPUs. For optimal performance on systems with hyperthreading, usually the
hyperthreading should be switched off or -t used to halve the number of threads; e.g. -t4 rather than -t8 for an Intel
i7 processor. Running 8 threads on 8 processors Read instructions and data ** ATOM NAMES NOT ALLOWED **
+++++++ + infile_sx finished at
05:18:56 Total elapsed time: 0.00 secs +
+++++++ . Create .fcf from SHELXL
.ins & .hkl. Substituted SHELXL Instruction:LS. 1 Inserted SHELXL Instruction:BLOC ** PROBLEM to Recreate FCF for
this ENTRY ** Entry # 1 - 0A, 0B, 1C, 1G-Alerts, BP C-C = 0.0038 :: CheckCIF out on :infile.chk . . . . .
. . . . .
```

checkCIF/PLATON (full publication check)

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

No syntax errors found.

Please wait while processing

[CIF dictionary](#)

[Interpreting this report](#)

Datablock: I

Bond precision: C-C = 0.0038 A Wavelength=0.71073

Cell: a=8.717(3) b=16.325(5) c=21.043(7)

alpha=90 beta=90 gamma=90

Temperature: 90 K

	Calculated	Reported
Volume	2994.5(17)	2994.5(17)
Space group	P b c a	P b c a
Hall group	-P 2ac 2ab	-P 2ac 2ab
Moiety formula	C36 H30 S4	C36 H30 S4
Sum formula	C36 H30 S4	C36 H30 S4
Mr	590.84	590.84
Dx, g cm-3	1.311	1.311
Z	4	4
Mu (mm-1)	0.342	0.342
F000	1240.0	1240.0
F000'	1242.35	
h, k, lmax	10, 19, 25	10, 19, 25
Nref	2646	2641
Tmin, Tmax	0.902, 0.934	0.659, 0.934
Tmin'	0.887	

Correction method= # Reported T Limits: Tmin=0.659 Tmax=0.934 AbsCorr = EMPIRICAL

Data completeness= 0.998

Theta (max)= 25.020


```

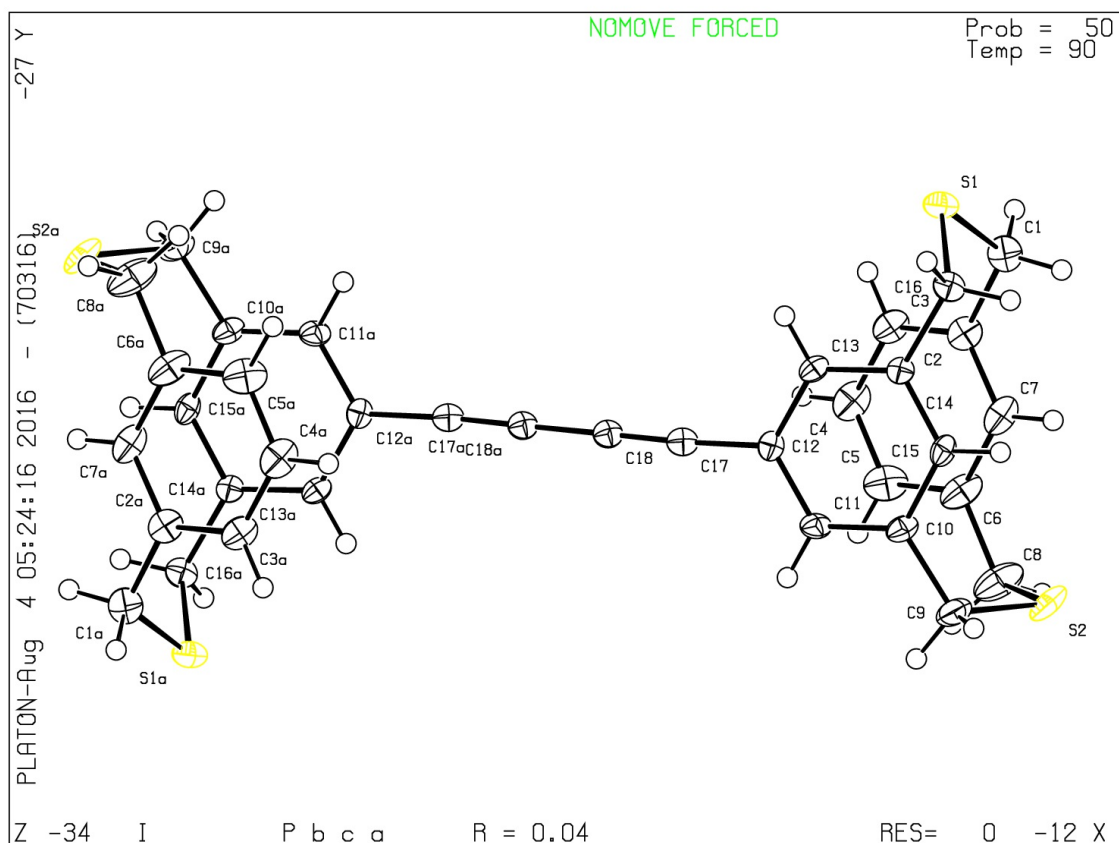
;
_vrf_PUBL010_GLOBAL
;
PROBLEM: _publ_author_address is missing. Author(s) address(es).
RESPONSE: ...
;
# end Validation Reply Form

```

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.

PLATON version of 08/07/2016; check.def file version of 05/07/2016

Datablock I – ellipsoid plot



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