

PLATON(V-311206)-Run for: xray02 - SHELXL

TIME: Mar 4 13:35:29 2010

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Crystal Data

Input Cell (Lattice Type: C) - Temp = 200K		Reduced Cell (Acta Cryst.(1976), A32, 297-298)	
a = 30.333(18) Angstrom	alpha = 90 Degree	a = 6.101	alpha = 105.01 V = 1741.5
b = 6.101(3)	beta = 105.313(11)	b = 15.470	beta = 90.00
c = 19.513(11)	gamma = 90	c = 19.513	gamma = 101.37
V = 3483(3) Cubic-Angstrom		Niggli Values	
d(100) = 29.2561 Angstrom		37.221	239.328 380.757
d(010) = 6.1010		-78.157	0.000 -18.611
d(001) = 18.8202			
Lambda(MoKa) = 0.71073 Angstrom			

Orthogonalization Matrices

(See e.g. J.D.Dunitz, Xray Analysis and Structure Determination of Organic Molecules, Cornell Univ. Press, 1979, P236)

(X0) (30.33300 0 -5.15323) (X) , (X) (0.03297 0 0.00903) (X0) Orthogonal Axes A0, B0 and C0
 (Y0) = (0 6.10100 0)*(Y) , (Y) = (0 0.16391 0)*(Y0) are defined as:
 (Z0) (0 0 18.82024) (Z) , (Z) (0 0 0.05313) (Z0) A0 // A, C0 // C*, B0 // C0 X A0

Space Group Symmetry

(See e.g. G. Burns & A.M. Glazer, Space Groups for Solid State Scientists, Academic Press, 1990 or Int. Tables A)

Space Group C2/c No: 15, Laue: 2/m [Hall: -C 2yc]

Lattice Type mC, Centric, Monoclinic, Order 8(2) [Schoenflies: C2h^6]

Nr ***** Symmetry Operation(s) *****

1	X ,	Y ,	Z
2	- X ,	Y ,	1/2 - Z
3	- X ,	- Y ,	- Z
4	X ,	- Y ,	1/2 + Z
5	1/2 + X ,	1/2 + Y ,	Z
6	1/2 - X ,	1/2 + Y ,	1/2 - Z
7	1/2 - X ,	1/2 - Y ,	- Z
8	1/2 + X ,	1/2 - Y ,	1/2 + Z

(Not-Rule Rounded) Coordinates of Unique Residue(s) Identified. Standard Deviations in the Last Digit are in Parentheses.

Site = Site Symmetry; SSN = Site Symmetry Number; SSOF = SHELX Site Occupation Factor (= S.O.F / SSN).
 ***** Move = Transformation on Input Data: N.IJK (N = SymOp, IJK = Translation) i.e. 1.555 = nomove
 SYMBOLS: Type = D/A = Potential Donor or Acceptor atom, D-H = H on Donor atom, MET = Metal.
 ***** El Type = AK = Alkali Metal, AE = Alkaline Earth, HL = Halogen, AN = Actinide, LN = Lanthanide, TR = Transition Element.
 ARU = Asymmetric Residue Unit encoded as sklm.nn, with s = symmetry op, klm = translation, nn = residue #.
 RESIDUE = collection of ARU's constituting an isolated unit (= molecule, ion).
 FLAGS : d = determined, c = calculated, R = riding G = group

Atom Types : C H S
 Cov.Rad(Ang): 0.68 0.35 1.04
 Atom Volume : 13.87 5.08 25.20
 Atom Number : 6 1 16
 Atom Weight : 12.010 1.008 32.07
 Scat.Fact.f0: 5.999 1.000 16.000
 Scat.Fact.f': 0.003 0.000 0.125
 Scat.Fact.f": 0.002 0.000 0.123
 Mu/Rho(MoKa): 0.58 0.37 9.99
 Elem. Type : -- -- --

Sources - Cov. Radii : Manual Cambridge Crystallographic Data Base
 - Atom Volume: D.W.M. Hofmann (2002). Acta Cryst. B58, 489-493
 - Atomic Wt. : SHELXL97
 - Scat. Fact.: SHELXL97 (International Tables)
 - mu/rho : International Tables C, table 4.2.4.2, p193-199

NOMOVE

Flags	Label	Fractional Coordinates (x,y,z)			Orthogonal Coordinates (X0,Y0,Z0)			Site	SSN*SSOF =	S.O.F	Move	Type
-	S(501)	0.33589	0.03489	0.13729	9.4809	0.2129	2.5838	1	1	1	-	D/A
-	S(502)	0.09445	0.04914	0.11878	2.2528	0.2998	2.2355	1	1	1	-	D/A
-	C(501)	0.18942	0.32181	0.19330	4.7496	1.9634	3.6379	1	1	1	-	-
-	C(502)	0.18542	0.11637	0.16082	4.7956	0.7099	3.0267	1	1	1	-	-
-	C(503)	0.25119	0.20852	0.11822	7.0101	1.2722	2.2249	1	1	1	-	-
-	C(504)	0.21726	0.06609	0.12240	5.9595	0.4032	2.3036	1	1	1	-	-
-	C(505)	0.22398	0.46665	0.18802	5.8250	2.8470	3.5387	1	1	1	-	-
-	C(506)	0.25480	0.41361	0.15225	6.9441	2.5235	2.8654	1	1	1	-	-
-	C(507)	0.37103	-0.05456	0.08141	10.8348	-0.3329	1.5321	1	1	1	-	-
-	C(508)	0.14998	-0.04501	0.16840	3.6814	-0.2746	3.1694	1	1	1	-	-
-	C(509)	0.28501	0.14715	0.07404	8.2638	0.8977	1.3934	1	1	1	-	-
-	C(510)	0.05383	-0.12568	0.14236	0.8992	-0.7668	2.6792	1	1	1	-	-
-	C(511)	0.00731	-0.04581	0.10878	-0.3389	-0.2795	2.0473	1	1	1	-	-
-	C(512)	0.38261	0.13846	0.04031	11.3979	0.8447	0.7586	1	1	1	-	-
-	C(513)	0.34563	-0.24123	0.03254	10.3164	-1.4718	0.6124	1	1	1	-	-
-	C(514)	0.41322	-0.14448	0.12508	11.8896	-0.8815	2.3540	1	1	1	-	-
-	C(515)	0.05058	-0.33417	0.08632	1.0894	-2.0388	1.6247	1	1	1	-	-
-	C(516)	0.06904	-0.38377	0.15390	1.3012	-2.3414	2.8965	1	1	1	-	-
-	C(517)	0.05828	-0.11638	0.22562	0.6051	-0.7101	4.2463	1	1	1	-	-

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Ordered Structure                               Unit Cell Contents (Based on Contents of Atom List, that may be Incomplete)
=====
Resd Site  X(cen) Y(cen) Z(cen)  Mol.Wt S.O.F  Z      C      H      S
=====
  1      1  0.208  0.027  0.128   268.31    1    8     17     -     2
=====
Unit Cell Weight =      2146.48              136      0     16

Calculated Analysis (Percent) =              76.1    0.0 23.9

Moiety_Formula = C17 S2

Sum_Formula = C17 S2

Formula_Weight =      268.31 [Note: Based on SHELXL97 Atomic Weights]

Formula_Z =          8

SpaceGroup_Z =          8 ==> Z' = 8 / 8 = 1.000

Calculated Density = 1.0234(9) g cm-3 [= Mg m-3]

F(000) =      1072.0 [      1074.33]

mu(MoKa) =      2.89 cm-1 = 0.289 mm-1

Resonant Scattering =      0.73 Percent - (Girard,E. et al. (2003). Acta Cryst. D59, 1914-1922)

Predicted Volume =      2289.5[ 2268.2] Ang**3, 298[200]K - (D.W.M. Hofmann (2002). Acta Cryst. B58, 489-493)

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Note on F000: The first number is a pure electron count, the second number between [] is calculated from f, f' & f''

** WARNING **
Please Check the Derived Crystal Data.
They may be Incorrect for Disordered,
Incomplete or Polymeric Structures.

(An)isotropic, Equivalent and Main Axes Displacement Parameters - Unusual Values Marked with a # - [Optional Coordinate Split-up]

Atom	Label	U11 or Uiso	U22	U33	U23	U13	U12	Ueq(sUeq)	U1	U2	U3	U3/U1
1	S501	0.07664						0.07664				
2	S502	0.08103						0.08103				
3	C501	0.05520						0.05520				
4	C502	0.04868						0.04868				
5	C503	0.05200						0.05200				
6	C504	0.05310						0.05310				
7	C505	0.05468						0.05468				
8	C506	0.05960						0.05960				
9	C507	0.06012						0.06012				
10	C508	0.07668						0.07668				
11	C509	0.07468						0.07468				
12	C510	0.06273						0.06273				
13	C511	0.12295						0.12295				
14	C512	0.11581						0.11581				
15	C513	0.16172						0.16172				
16	C514	0.15687						0.15687				
17	C515	0.21525						0.21525				
18	C516	0.21540						0.21540				
19	C517	0.24976						0.24976				

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The Displacement Factor has the Form of $\text{Exp}(-T)$

where

$T = 8 \cdot (\pi^2) \cdot U_{\text{iso}} \cdot \sin(\theta/\lambda)^2$, for Isotropic Atoms,

$T = 2 \cdot (\pi^2) \cdot (U_{11} \cdot (h \cdot a)^2 + U_{22} \cdot (k \cdot b)^2 + U_{33} \cdot (l \cdot c)^2 + 2 \cdot U_{23} \cdot k \cdot l \cdot b \cdot c + 2 \cdot U_{13} \cdot h \cdot l \cdot a \cdot c + 2 \cdot U_{12} \cdot h \cdot k \cdot a \cdot b)$, for Anisotr. Atoms

$U_{\text{eq}} = 1/3 \text{ Sum}(i,j) (U_{ij} \cdot a(i) \cdot a(j) \cdot a(i) \cdot a(j))$

U1, U2, U3 are the three Main Axes Components of U_{ij}

Reference U_{eq} : R.X. Fischer & E. Tillmanns, Acta Cryst. (1988). C44, 775-776

U_{eq} [or $U(\text{iso})$] Averages per Element

	Non-H	C	H	S
Average	0.1049	0.1080	0.0000	0.0788

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Minimum  0.0487 0.0487 0.0000 0.0766
Maximum  0.2498 0.2498 0.0000 0.0810
Ratio    5.1306 5.1306 0.0000 1.0573
Number      19    17      0      2

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Analysis of Bond Distance and Angle Values - Identification of Chiral Center(s) and Their (R/S)-Configuration (Cahn-Ingold-Prelog)

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The Following Tests are done. Faults are Marked Under Note

-- V : Valency Check Fault for H, C

-- S : Bond Too Short

-- A : Unusual Bond Angle Values (PLEASE CHECK)

*** PLEASE NOTE: R/S ASSIGNMENTS ARE TENTATIVE *** (CIP Special rules NOT Implemented)

*** See Angew.Chem.Intern. Ed. Eng.,(1966),5,385 & (1982),21,567 for Authoritative Details for Special Cases

Flag	Label	- Connected to (May be Incomplete for Polymeric Structures)				=A.N.G.L.E.S=				=B.O.N.D.S=				Hyb	RS	Note
						nra	min	max	Aver	min	max	nrb	tnr			
-	S(501) - C(507)	C(509)				1	103	103	103.5	1.799	1.835	2	14			
-	S(502) - C(508)	C(510)				1	106	106	106.4	1.780	1.801	2	15			
-	C(501) - C(502)	C(505)				1	121	121	120.8	1.395	1.395	2	3			?
-	C(502) - C(501)	C(504)	C(508)			3	117	122	120.0	1.395	1.494	3	7	sp2		
-	C(503) - C(504)	C(506)	C(509)			3	120	120	120.0	1.366	1.550	3	5	sp2		
-	C(504) - C(502)	C(503)				1	122	122	121.9	1.366	1.404	2	4			?
-	C(505) - C(501)	C(506)				1	122	122	121.6	1.345	1.395	2	1			?
-	C(506) - C(503)	C(505)				1	119	119	118.7	1.345	1.407	2	2			?
-	C(507) - S(501)	C(512)	C(513)	C(514)		6	108	113	109.5	1.445	1.799	4	10	sp3		
-	C(508) - S(502)	C(502)				1	109	109	109.4	1.494	1.801	2	13			?
-	C(509) - S(501)	C(503)				1	106	106	106.2	1.550	1.835	2	11			?
-	C(510) - S(502)	C(511)	C(515)	C(516)	C(517)	10	47	132	102.9	1.473	1.780	5	12			A
-	C(511) - C(510)					0	0	0	0.0	1.473	1.473	1	8			?
-	C(512) - C(507)					0	0	0	0.0	1.517	1.517	1	6			?
-	C(513) - C(507)					0	0	0	0.0	1.553	1.553	1	6			?
-	C(514) - C(507)					0	0	0	0.0	1.445	1.445	1	6			?
-	C(515) - C(510)	C(516)				1	65	65	65.4	1.324	1.663	2	9			A
-	C(516) - C(510)	C(515)				1	67	67	67.3	1.324	1.640	2	9			A
-	C(517) - C(510)					0	0	0	0.0	1.595	1.595	1	8			?

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Analysis of the IntraMolecular Geometry in Terms of Unique Molecule(s)/Ions, with Bond Criterium: d(i-j) < R(i) + R(j) + Tol

-- Tol = 0.40 Ang. for Normal Bonds + 0.70 for (Earth)alkali-NonMetal Contacts and adjusted by -.40 Ang. for Metal-Metal Distances

-- The Bond Distance and Angle su's have been Incremented to Include the Effect of the Unit-cell su.
(Rel.Error in Dist. 0.0006 Ang. , Abs. Angle Error 0.011 Deg.)

-- Bonds below with '>' or '<' Substituted for '-' have Distances that Deviate from Expected Values(Based on the hybridisations).

Bond Lengths (Angstrom). - (Bonds are ordered on the first label, left to right and top to bottom) - su in last digit in ().

=====

S(501) - C(507)	1.7992	S(501) - C(509)	1.8351	S(502) - C(508)	1.8009	S(502) - C(510)	1.7795
C(501) - C(502)	1.3952	C(501) - C(505)	1.3955	C(502) - C(504)	1.4041	C(502) - C(508)	1.4937
C(503) - C(504)	1.3657	C(503) - C(506)	1.4073	C(503) - C(509)	1.5503	C(505) - C(506)	1.3455
C(507) - C(512)	1.5173	C(507) - C(513)	1.5530	C(507) - C(514)	1.4453	C(510) - C(511)	1.4730
C(510) - C(515)	1.6632	C(510) - C(516)	1.6396	C(510) - C(517)	1.5954	C(515) - C(516)	1.3244

Bond/Valence Angles (Degrees) - (Angles are ordered on the middle label, left to right and top to bottom) - su in last digit in ().

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C(507) - S(501) - C(509)	103.48	C(508) - S(502) - C(510)	106.44	C(502) - C(501) - C(505)	120.82
C(501) - C(502) - C(504)	116.69	C(501) - C(502) - C(508)	121.71	C(504) - C(502) - C(508)	121.57
C(504) - C(503) - C(506)	120.23	C(504) - C(503) - C(509)	119.96	C(506) - C(503) - C(509)	119.79
C(502) - C(504) - C(503)	121.90	C(501) - C(505) - C(506)	121.62	C(503) - C(506) - C(505)	118.72
S(501) - C(507) - C(512)	109.99	S(501) - C(507) - C(513)	108.51	S(501) - C(507) - C(514)	109.39
C(512) - C(507) - C(513)	113.02	C(512) - C(507) - C(514)	108.28	C(513) - C(507) - C(514)	107.58
S(502) - C(508) - C(502)	109.39	S(501) - C(509) - C(503)	106.17	S(502) - C(510) - C(511)	109.52
S(502) - C(510) - C(515)	102.31	S(502) - C(510) - C(516)	114.97	S(502) - C(510) - C(517)	111.34
C(511) - C(510) - C(515)	94.42	C(511) - C(510) - C(516)	125.49	C(511) - C(510) - C(517)	104.76
C(515) - C(510) - C(516)	47.27	C(515) - C(510) - C(517)	132.15	C(516) - C(510) - C(517)	87.09
C(510) - C(515) - C(516)	65.43	C(510) - C(516) - C(515)	67.30		

Torsion/Dihedral Angles (Deg.) - Klyne & Prelog Convention (Dunitz, p241) - (Excl. Minor Disorder & Embedded Bond Angl. > 160. Deg.)

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C(505) C(501) C(502) C(504)	-0.84	C(505) C(501) C(502) C(508)	177.29	C(502) C(501) C(505) C(506)	-0.60
C(501) C(502) C(504) C(503)	1.48	C(508) C(502) C(504) C(503)	-176.65	C(501) C(502) C(508) S(502)	71.09
C(504) C(502) C(508) S(502)	-110.87	C(506) C(503) C(504) C(502)	-0.72	C(509) C(503) C(504) C(502)	-178.94
C(504) C(503) C(506) C(505)	-0.74	C(509) C(503) C(506) C(505)	177.47	C(504) C(503) C(509) S(501)	-96.65
C(506) C(503) C(509) S(501)	85.13	C(501) C(505) C(506) C(503)	1.39	C(512) C(507) S(501) C(509)	60.22
C(513) C(507) S(501) C(509)	-63.88	C(514) C(507) S(501) C(509)	179.02	C(502) C(508) S(502) C(510)	-170.50
C(503) C(509) S(501) C(507)	175.01	C(511) C(510) S(502) C(508)	174.78	C(515) C(510) S(502) C(508)	-85.95
C(516) C(510) S(502) C(508)	-37.59	C(517) C(510) S(502) C(508)	59.40	S(502) C(510) C(515) C(516)	112.74
C(511) C(510) C(515) C(516)	-136.17	C(517) C(510) C(515) C(516)	-21.68	S(502) C(510) C(516) C(515)	-83.69
C(511) C(510) C(516) C(515)	58.01	C(517) C(510) C(516) C(515)	164.08		

Statistics of Bond Length per Bond Type (NOTE: A Indicates 10 Occurrences, B Indicates 11, Etc. and * more than 35)

[illegible]

Selected Bond Lengths (Angstrom) - see M.F.C. Ladd & R.A. Palmer, Structure Determination by X-Ray Crystallography (1985)

Formal single bonds

C4-C4 1.54	C4-C3 1.52	C4-C2 1.46	C4-N3 1.47	C4-N2 1.47	C4-O2 1.43	C3-C3 1.46	C3-C2 1.45	C3-N3 1.40	C3-N2 1.40
C3-O2 1.36	C2-C2 1.38	C2-N3 1.33	C2-N2 1.33	C2-O2 1.36	N3-N3 1.45	N3-N2 1.45	N3-O2 1.36	N2-N2 1.45	N2-O2 1.41

Formal double bonds

C3-C3 1.34 C3-C2 1.31 C3-N2 1.32 C3-O1 1.22 C2-C2 1.28 C2-N2 1.32 C2-O1 1.16 N3-O1 1.24 N2-N2 1.25 N2-O1 1.21

Formal triple bonds

Aromatic bonds

C2-C2 1.20 C2-N1 1.16 C3-C3 1.40 C2-N2 1.34 N2-N2 1.35

The notation in the table indicates the connectivity of the atoms

For more detailed standard bond distance tabulations see: J. Chem. Soc. Perkin II, (1987), S1-S19; J. Chem. Soc. Dalton Trans. (1989), S1 - S83 or International Tables C, (1992), 707-791.

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