

PLATON(V-311206)-Run for: joer04 - SHELXL

TIME: Jun 9 13:21:22 2011

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Crystal Data

Input Cell (Lattice Type: P)		Temp = 200K	Reduced Cell		(Acta Cryst.(1976), A32, 297-298)
a =	15.327(3) Angstrom	alpha = 90 Degree	a =	15.163	alpha = 96.22 V = 3744.8
b =	15.163(3)	beta = 96.222(4)	b =	15.327	beta = 90.00
c =	16.209(3)	gamma = 90	c =	16.209	gamma = 90.00
V = 3744.8(13) Cubic-Angstrom		d(100) = 15.2367 Angstrom	Niggli Values		
		d(010) = 15.1630	229.917	234.917	262.732
Lambda(MoKa) = 0.71073 Angstrom		d(001) = 16.1135	-26.926	0.000	0.000

Orthogonalization Matrices

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

(X0) (15.32700 0 -1.75675) (X) , (X) (0.06524 0 0.00711) (X0) Orthogonal Axes A0, B0 and C0
 (Y0) = (0 15.16300 0)*(Y) , (Y) = (0 0.06595 0)*(Y0) are defined as:
 (Z0) (0 0 16.11352) (Z) , (Z) (0 0 0.06206) (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 P21/c No: 14, Laue: 2/m [Hall: -P 2ybc]

Lattice Type mP, Centric, Monoclinic, Order 4(2) [Schoenflies: C2h^5]

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

1	X ,	Y ,	Z
2	- X ,	1/2 + Y ,	1/2 - Z
3	- X ,	- Y ,	- Z
4	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 Br N Zn
 Cov.Rad(Ang): 0.68 0.35 1.21 0.68 1.45
 Atom Volume : 13.87 5.08 32.70 11.80 39.00
 Atom Number : 6 1 35 7 30
 Atom Weight : 12.010 1.008 79.90 14.01 65.39
 Scat.Fact.f0: 5.999 1.000 34.993 6.995 29.985
 Scat.Fact.f': 0.003 0.000 -0.290 0.006 0.284
 Scat.Fact.f'': 0.002 0.000 2.460 0.003 1.430
 Mu/Rho(MoKa): 0.58 0.37 75.37 0.84 53.97
 Elem. Type : -- -- HL -- TR

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
-	Br(51)	0.53700	0.17054	0.16725	7.9367	2.5859	2.6950	1	1	1	-	D/A
-	Br(52)	0.68762	0.35910	0.11714	10.3334	5.4450	1.8875	1	1	1	-	D/A
-	Br(53)	0.67609	0.24251	0.21931	9.9772	3.6772	3.5339	1	1	1	-	D/A
-	Br(54)	0.78782	0.13208	0.20226	11.7196	2.0027	3.2591	1	1	1	-	D/A

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	Br	N	Zn
1	1	0.672	0.226	0.176	319.60	1	4	-	-	4	-	-
Unit Cell Weight =					1278.40			0	0	16	0	0
Calculated Analysis (Percent) =								0.0	0.0100	0.0	0.0	0.0
Moiety_Formula =					Br4							
Sum_Formula =					Br4							
Formula_Weight =					319.60 [Note: Based on SHELXL97 Atomic Weights]							
Formula_Z =					4							
SpaceGroup_Z =					4 ==> Z' = 4 / 4 = 1.000							
Calculated Density =					0.5669(2) g cm-3 [= Mg m-3]				** WARNING **			
F(000) =					560.0 [556.64]				Please Check the Derived Crystal Data. They may be Incorrect for Disordered, Incomplete or Polymeric Structures.			
mu(MoKa) =					42.73 cm-1 = 4.273 mm-1							
Resonant Scattering =					10.00 Percent - (Girard,E. et al. (2003). Acta Cryst. D59, 1914-1922)							
Predicted Volume =					523.2[518.3] Ang**3, 298[200]K - (D.W.M. Hofmann (2002). Acta Cryst. B58, 489-493)							

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

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(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	Br51	0.02760						0.02760				
2	Br52	0.03222						0.03222				
3	Br53	0.03435						0.03435				
4	Br54	0.03848						0.03848				

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

where

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

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

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

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_{iso}] Averages per Element

	Non-H	C	H	Br	N	Zn
Average	0.0332	0.0000	0.0000	0.0332	0.0000	0.0000
Minimum	0.0276	0.0000	0.0000	0.0276	0.0000	0.0000
Maximum	0.0385	0.0000	0.0000	0.0385	0.0000	0.0000
Ratio	1.3942	0.0000	0.0000	1.3942	0.0000	0.0000
Number	4	0	0	4	0	0

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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)

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*** 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			
-	Br(51)	- Br(53)	-----	0	0	0	0.0	2.461	2.461	1	2			
-	Br(52)	- Br(53)	-----	0	0	0	0.0	2.442	2.442	1	2			
-	Br(53)	- Br(51)	Br(52) Br(54) -----	3	102	109	105.1	2.432	2.461	3	1			
-	Br(54)	- Br(53)	-----	0	0	0	0.0	2.432	2.432	1	2			

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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. 0.0002 Ang. , Abs. Angle Error 0.004 Deg.)
 - Bonds below with '>' or '<' Substituted for '-' have Distances that Deviate from Expected Values(Based on the hybridisations).
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Bond Lengths (Angstrom). - (Bonds are ordered on the first label, left to right and top to bottom) - su in last digit in ().

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Br(51) - Br(53)	2.4613	Br(52) - Br(53)	2.4419	Br(53) - Br(54)	2.4321
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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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Br(51) - Br(53) - Br(52)	102.25	Br(51) - Br(53) - Br(54)	104.49	Br(52) - Br(53) - Br(54)	108.53
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[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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