

The Crystal and Molecular Structure of Dimedone

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The crystal and molecular structure of dimedone, 5,5-dimethyl-1,3-cyclohexanedione, has been determined from three-dimensional single crystal X-ray diffraction data. The space group is $P2_1/c$, with cell dimensions $a = 10.040(1)$, $b = 6.823(1)$, $c = 12.984(2)$ Å, $\beta = 116.20(1)^\circ$. There are four molecules in the unit cell. The structure has been refined by full matrix least-squares methods to a weighted R -factor of 0.046 for the 1428 independent reflections. The estimated standard deviations in bond lengths involving non-hydrogen atoms are 0.002 Å. The molecules crystallize in the enol form and are hydrogen bonded to each other forming infinite helices parallel to the b axis. The hydrogen bond length is 2.593 Å.

Dimedone (5,5-dimethyl-1,3-cyclohexanedione) is a white crystalline solid of melting point 148–149°C. It is commonly used as a reagent for separation and identification of aldehydes.¹ Unlike acyclic β -dicarbonyl compounds, enolization of dimedone gives a *trans*-enol where *intra*-molecular hydrogen bonding is impossible.^{2,3} Nevertheless, dimedone is believed to exist largely in the enol form both in solution⁴ and in the solid state.⁴ In solvents incapable of forming hydrogen bonds the comparatively high enol content has been explained by assuming a hydrogen bonded dimeric enol form.^{2,3} Several X-ray investigations of acyclic β -diketones have been carried out,^{5–8} but there is a paucity of structural data for the cyclic analogs. An X-ray investigation of dimedone has therefore been undertaken.

EXPERIMENTAL

A commercial sample of dimedone was purified and crystallized from methanol. The observed extinctions, $h0l$ for $l = 2n + 1$ and $0k0$ for $k = 2n + 1$ uniquely determined the space group

as $P2_1/c$. The cell dimensions were determined at room temperature on a manual four circle diffractometer using CuK radiation. The crystal used for the collection of intensity data was of nearly spherical form with diameter 0.35 mm. Data were collected on a SYNTEX PI diffractometer by the $\omega/2\theta$ scan technique using a graphite monochromator and MoK α radiation ($= 0.71069$ Å) at 18°C. Each reflection was scanned from $2\theta_{\alpha 1} - 0.85^\circ$ to $2\theta_{\alpha 2} + 0.85^\circ$, at a variable scan rate ranging from 1°/min to 12°/min depending on intensity. Stationary-counter-stationary-crystal background counts were made at the upper and lower extremities of the 2θ scan range for each reflection. A rejection level was also specified to avoid measurement of reflections with intensities less than the threshold value. The intensities of three standard reflections, measured every 50' reflections, showed no change during the data collection. A total of 1548 independent intensities were recorded, out to $2\theta < 70^\circ$. Of these, 1428 reflections had intensities greater than twice their standard deviations. The intensities and their standard deviations were corrected for Lorentz and polarization effects and a 2 % uncertainty in diffractometer stability was included in the standard deviations. No corrections were made for absorption since intensity errors from this source are less than 1.5 % ($\mu = 0.09$ mm⁻¹). All programs applied are written or revised for CDC 3300 by Dahl *et al.*¹¹

CRYSTAL DATA

Dimedone, $C_8H_{12}O_2$, F.W. 140.2, monoclinic; $a = 10.040(1)$, $b = 6.823(1)$, $c = 12.984(2)$ Å, $\beta = 116.20(1)^\circ$, $V = 798.3$ Å³, $F(000) = 444$, $Z = 4$, $\rho_{obs} = 1.15$ g cm⁻³, $\rho_{calc} = 1.15$ g cm⁻³. Space group $P2_1/c$.

SOLUTION AND REFINEMENT

The phase problem was solved three-dimensionally by a computer program based on

symbolic addition methods. By permuting the signs of two structure invariant reflections with large unitary structure amplitudes, four sets of 350 signs were determined. From one of the resulting Fourier maps, peaks corresponding to all the heavy atoms were located. The trial structure was refined by full-matrix least-squares technique. A difference Fourier synthesis led to the location of the hydrogen atoms with the exception of those of the methyl groups. The coordinates for these atoms were calculated assuming staggered positions of the methyl groups relative to C5. Refinement of all positional and thermal parameters including a common isotropic thermal parameter for the methyl hydrogen atoms terminated at $R_w = 0.046$ and $R = 0.040$. Unobserved reflections were excluded throughout the refinement. Weight analysis showed that weighting based on standard deviations from counter statistics was satisfactory. No correction for secondary extinction was necessary. Atomic form factors were those of Hanson *et al.*⁹ except for hydrogen.¹⁰ A final difference Fourier map contained no larger density fluctuations than $\pm 0.3 \text{ e/}\text{\AA}^3$. The parameters obtained in the last cycle of refinement are given in Table 1 together with their estimated standard deviations. Table 2 contains the observed and calculated structure factors.

Rigid-body analysis of translational, librational, and screw motion¹¹ of the molecule gave a r.m.s. value of 0.0033 Å² when all the non-

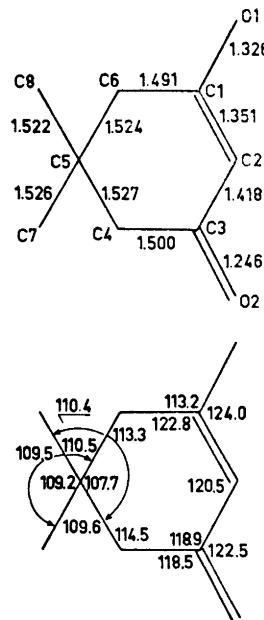


Fig. 1. Schematic drawing of the molecule showing bond distances and angles. (Bond distances uncorrected for librational motion).

hydrogen atoms were included. When the two methyl carbon atoms were excluded, the r.m.s. value dropped to 0.0022 Å.² The cyclohexene ring and the two oxygen atoms may therefore be regarded as a rigid body. Corrections in bond lengths generally exceed the standard devia-

Table 1. Fractional atomic coordinates and thermal parameters with estimated standard deviations ($\times 10^6$), isotropic temperature factor and positional parameters ($\times 10^4$) for the hydrogen atoms. The anisotropic temperature factor is given by $\exp - (B_{11}h^2 + B_{22}k^2 + B_{33}l^2 + B_{12}hk + B_{13}hl + B_{23}kl)$.

ATOM	X	Y	Z	B	611	822	B33	B12	B13	B23
O 1	26132(12)	65079(16)	35714(11)	1361(15)	2793(29)	833(18)	=818(33)	764(21)	549(26)	
O 2	=883(11)	19773(15)	35972(8)	1272(14)	2557(27)	831(9)	=971(31)	621(19)	=386(23)	
C 1	21184(15)	51711(19)	48534(12)	1151(17)	2024(32)	732(12)	=11(39)	879(24)	=9(29)	
C 2	8335(15)	41651(21)	34848(12)	1125(17)	2247(33)	628(11)	=77(48)	632(22)	=15(38)	
C 3	3224(15)	28588(19)	40777(11)	1149(17)	1840(29)	734(11)	=45(48)	899(23)	=366(29)	
C 4	12484(16)	25816(23)	53431(12)	1448(21)	2869(34)	699(11)	=337(45)	1881(25)	=187(38)	
C 5	29911(17)	29271(21)	57624(12)	1294(19)	2246(32)	597(18)	=74(41)	584(22)	=66(28)	
C 6	30788(17)	49165(23)	53854(12)	1268(28)	2492(39)	691(12)	=688(46)	651(25)	=232(33)	
C 7	36671(24)	29562(34)	70718(14)	1971(38)	3517(53)	664(13)	=797(69)	585(32)	=83(41)	
C 8	35889(23)	13281(29)	53343(17)	1632(26)	3869(47)	1849(17)	1259(63)	944(35)	189(47)	
H 1	229(15)	4352(28)	2677(12)	3,8(-3)						
H 2	835(17)	3444(22)	5751(13)	4,8(-3)						
H 3	1861(16)	1272(24)	5543(12)	4,4(-3)						
H 4	2833(16)	5968(22)	5733(12)	4,8(-3)						
H 5	4899(19)	5129(23)	6438(12)	5,1(-4)						
H 6	2818(21)	6664(25)	2665(16)	5,9(-4)						
H 7	4728(24)	3197(27)	7379(17)	7,8(-2)						
H 8	3484(21)	1789(29)	7382(15)	7,8(-8)						
H 9	3222(19)	4857(28)	7347(15)	7,8(-8)						
H10	4623(24)	1511(27)	5676(16)	7,8(-8)						
H11	30952(21)	1211(25)	4486(17)	7,8(-8)						
H12	3413(20)	27(28)	5684(15)	7,8(-8)						

Table 2. Observed and calculated structure factors. The columns are l , $10F_0$, $10F_c$ respectively.

	$0, K_a$	0	6	101	97	6	289	261	*	1	32	33	2	89	51	$0, K_a$	8	*	8	9	10	9	31	32				
2	486	489	7	37	33	6	202	195	*	0	82	85	3	182	154	*	0, K _a	8	*	8	15	11	28	25				
4	302	293	8	41	38	4	251	250	1	49	68	4	102	103	0	0, K _a	30	*	3	83	31	14	20	16				
6	16	16	9	24	24	3	242	241	2	19	19	0	37	38	0, K _a	30	*	2	33	31	14	16	2					
8	20	14	10	56	54	2	795	822	5	47	64	7	125	126	*	3, K _a	23	*	1	14	15	18	13	15				
10	66	79	11	31	31	1	155	153	6	107	110	8	111	110	6	16	11	0	39	39	16	37	35					
12	30	23	13	47	46	1	353	352	7	113	114	6	61	61	0, K _a	3	*	1	55	56	13	33	36					
14	21	27	14	25	21	1	357	352	7	113	114	14	39	38	*	18	26	2	3	78	79	-12	19	23				
16	63	54	15	22	22	2	233	220	8	33	33	0, K _a	3	*	12	23	19	4	27	27	*11	38	40					
18	34	30	He	0, K _a	5	3	117	114	11	56	56	0, K _a	5	*	14	56	56	8	60	61	7	30	32	9	67	89		
He	0, K _a	0	*	12	25	27	4	409	389	0, K _a	6	*	14	55	54	8	60	61	7	30	32	*	8	54	53			
*17	26	24	11	42	42	5	246	232	*	9	51	54	0, K _a	12	*	6	118	112	11	22	22	*	8	54	53			
*14	18	15	6	40	39	6	49	46	*	8	28	27	0, K _a	10	*	8	4	26	26	13	22	22	*	7	55	54		
*10	37	36	*	8	23	24	7	67	65	*	5	59	60	10	15	16	*	2, K _a	501	498	He	3, K _a	5	*	6	139	132	
*8	15	15	*	7	125	126	8	97	98	*	3	30	29	0, K _a	10	*	0, K _a	480	476	*	12	19	0	5	19			
*6	5	5	*	8	48	48	10	30	41	*	2	57	55	0, K _a	9	*	2	97	205	10	31	33	*	24	26			
*6	344	342	7	27	27	18	21	*	1	89	92	*	7	29	26	4	140	135	*	7	15	15	20	*	3	285		
*5	70	68	*	4	57	58	13	15	15	0	61	62	0, K _a	6	*	6	63	54	8	76	76	*	5	16	15			
*4	153	146	*	3	116	118	16	31	28	1	66	69	*	8	124	121	10	56	54	*	4	61	62	*	1	40		
*3	245	242	*	2	59	62	He	1, K _a	2	3	29	30	4	316	307	14	30	29	*	3	20	19	0	124	119			
*2	143	147	*	1	139	140	15	22	22	5	34	33	*	3	259	259	He	3, K _a	1	*	2	62	61	1	172	164		
*1	475	484	139	140	12	19	22	6	39	38	*	2	271	269	16	58	53	*	1	78	78	2	18	15				
1	473	484	2	62	62	10	63	68	11	20	21	*	1	113	105	9	84	86	1	79	80	3	17	18				
2	143	147	3	118	118	9	31	33	He	1, K _a	7	0	63	61	8	16	16	3	59	59	4	31	31					
3	245	242	4	59	57	8	49	53	*	9	28	21	1	104	103	*	7	63	85	4	24	21	5	65	65			
4	154	145	5	27	29	7	101	105	*	4	30	27	0, K _a	165	167	5	68	65	6	28	25	7	32	30				
5	70	68	6	46	49	9	11	11	*	3	24	26	3	112	108	*	4	229	220	7	29	27	8	19				
6	345	342	7	126	126	29	31	34	*	4	41	40	0, K _a	44	44	4	44	44	0, K _a	51	50	10	16	14				
7	9	8	*	24	24	4	34	34	*	4	48	46	0, K _a	92	94	*	1	13	9	*	10	35	35	51	51			
8	14	16	9	41	39	*	3	116	114	2	26	27	6	53	50	0	310	314	*	10	35	35	51	51				
10	36	36	11	42	42	*	2	55	55	He	1, K _a	3	0	60	64	*	7	33	33	33	33	He	4, K _a	3				
13	23	24	12	24	27	1	22	22	5	27	25	8	46	49	2	58	54	*	6	33	33	33	33	He	4, K _a	3		
17	26	24	13	19	17	0	222	227	5	21	20	10	37	38	3	28	28	*	4	34	34	*	18	26				
He	0, K _a	2	15	15	11	1	215	217	He	1, K _a	8	13	50	47	4	26	26	*	4	10	9	*	12	39	40			
*14	36	34	He	0, K _a	6	2	17	13	*	10	24	21	14	38	35	5	55	55	*	2	117	116	*	11	82	83		
13	39	36	*	10	13	15	3	24	24	*	5	21	19	He	2, K _a	4	*	6	67	67	*	1	47	47	*	10	72	75
*12	19	18	*	8	21	21	4	94	93	*	4	29	29	9	34	37	7	55	59	0	30	30	*	9	111	114		
*8	22	22	*	7	27	27	5	24	25	*	1	65	60	8	118	121	9	33	35	1	23	25	*	8	36	36		
*7	53	57	*	5	33	34	6	11	11	*	0	12	12	7	29	29	11	27	26	4	40	40	*	124	126			
*6	31	31	*	4	23	25	4	44	44	*	0	15	15	6	59	59	3	35	35	3	30	30	*	6	79	75		
*4	149	149	*	3	39	39	8	101	101	*	3	45	42	0, K _a	57	57	He	3, K _a	1	*	4	55	53	*	4	10	4	
*4	110	10	*	47	47	9	42	42	*	5	20	19	0, K _a	4	*	4	64	60	*	17	30	28	*	24	20	200		
*3	10	7	0	46	47	11	30	32	6	32	30	*	3	105	101	101	*	13	67	66	6	75	74	*	2	100	106	
*2	2	2	1	46	47	11	30	32	6	32	30	*	3	80	78	0, K _a	12	*	23	22	7	34	33	*	1	32	30	
*1	255	249	2	9	11	16	23	19	He	1, K _a	9	*	1	74	71	*	11	39	40	He	3, K _a	7	0	56	61			
0	436	426	3	37	38	He	1, K _a	3	*	6	16	17	1	32	32	0, K _a	24	*	11	54	49	1	202	202				
1	256	249	4	24	26	*	9	180	187	2	35	34	2	56	55	*	9	13	20	*	9	15	15	2	30	28		
2	82	78	5	30	34	8	169	174	4	14	17	3	44	46	*	8	183	187	*	6	41	45	3	46	43			
3	11	7	6	13	16	7	125	132	He	2, K _a	0	4	24	25	7	189	185	186	*	5	14	13	4	21	20			
4	113	110	7	27	27	5	129	138	142	*	14	36	37	5	78	78	6	173	172	172	*	17	17	17	17	17		
5	150	149	23	21	21	12	129	128	121	*	12	62	62	9	37	36	0, K _a	55	55	55	55	55	55	55	55			
6	35	31	5	25	25	3	156	153	163	*	6	12	12	0, K _a	52	52	52	52	52	52	52	52						
7	23	25	He	0, K _a	9	*	54	68	4	52	46	11	31	28	*	10	110	109	*	2	152	149	2	31	33			
3	185	186	*	18	10	36	37	3	85	85	0, K _a	6	*	9	91	92	0	92	93	3	14	11	11	11				
4	30	32	*	2	37	37	9	73	77	*	2	84	83	1	31	32	*	8	38	39	2	146	147	4	13			
5	25	25	*	1	26	26	5	103	104	*	1	56	56	0, K _a	15	16	*	7	13	13	4	264	275					
6	102	110	1	28	26	7	130	134	0	284	278	9	52	51	*	6	34	37	6	150	159	7	13					
7	67	69	3	37	37	5	55	55	1	20	22	6	67	67	*	7	202	197	197	6	18	17	17					
8	24	21	1	45	44	4	32	32	30	35	35	0, K _a	55	55	55	55	55	55	55	55	55	55	55					
9	12	16	*	12	13	63	4	180	185	129	121	*	17	25	*	3	32	32	32	32	32	32	32					
13	32	31	*	8	159	163	1	21	19	4	17	17	3	40	38	*	2	78	71	*	25	21	21	13				
15	33	25	*	6	546	518	0	92	93	6	15	14	0	89	87	*	0	154	147	*	16	37	37	10				
He	0, K _a	4	*	4	145	144	1	119	119	6	97	94	1	37	34	*	1	55	53	*	15	13	13	34				
*14	23	21	*	2	78	77	2	20	18	7	14	8	2	24	22	*	2	28	24	*	13	37	43	7				
*13	47	46	0	374	368	3	11	16	12	40	40	3	84	83	4	68	68	*	12	143	147	4	6					
*11	30	31	2	29	25	4	37	33	14	47	41	4	75	76	5	64	66	*	11	51	52	*	5					
*10	57	54	4	69	68	5	129	125	16	23	24	5	69	68	6</td													

Table 2. Continued.

* 5	59	60	6	39	39	+11	11	5	+ 9	10	0	+ 4	13	15	H _e	8, K _e	6	3	19	20	+ 6	35	35			
* 4	12	11	10	25	25	+10	11	79	82	+ 6	46	+ 3	23	31	H _e	2, K _e	25	16	H _e	6, K _e	+ 4	29	32			
* 2	76	81	H _e	5, K _e	4	+ 9	69	71	+ 6	56	59	+ 1	32	31	H _e	23	18	+ 12	28	25	+ 3	30	31			
* 1	134	130	+14	16	13	+ 8	60	65	+ 2	45	20	+ 2	20	21	H _e	35	18	+ 8	51	51	+ 2	17	18			
* 0	62	61	+10	44	43	+ 5	45	43	+ 2	24	55	+ 3	71	H _e	24	+ 11	22	22	+ 2	32	29	+ 0	37			
* 3	39	39	+ 9	25	26	+ 8	60	60	+ 0	73	71	+ 7	31	28	+ 8	21	19	H _e	9, K _e	7	1	21	16			
* 2	43	42	+ 8	40	66	+ 4	43	46	+ 2	24	69	+ 2	13	13	H _e	44	39	+ 9	35	33	+ 1	19	18			
* 3	23	24	+ 7	64	63	+ 2	69	59	+ 4	60	59	+ 3	55	50	+ 7	44	39	+ 8	35	33	+ 7	14	14			
* 4	23	25	+ 6	81	78	+ 1	33	35	+ 6	16	18	+ 2	12	12	+ 5	45	47	+ 7	43	41	+ 5	12	12			
* 5	22	19	+ 4	114	115	+ 0	57	63	+ 8	46	46	+ 3	13	13	+ 11	+ 2	25	27	+ 1	18	15	+ 2	17	16		
* 7	23	21	+ 4	155	157	+ 1	62	62	H _e	7, K _e	1	H _e	7, K _e	8	+ 2	13	12	H _e	10, K _e	0	+ 0	19	18			
* 9	16	16	H _e	4, K _e	7	+ 2	64	64	+ 3	92	93	+ 16	18	18	+ 9	18	17	H _e	23	18	+ 10	33	32	H _e	11, K _e	3
* 8	22	22	+ 1	120	123	+ 4	49	49	+ 15	25	24	+ 5	21	21	+ 3	18	14	H _e	41	35	+ 8	58	58	+ 0	17	15
* 6	28	29	+ 1	58	63	+ 5	18	17	+ 14	20	21	+ 2	18	18	+ 13	41	35	+ 7	37	37	+ 2	23	25			
* 4	18	17	+ 2	58	57	+ 6	32	30	+ 13	32	32	H _e	8, K _e	1	+ 1	23	20	+ 8	31	26	+ 3	19	19			
* 5	25	23	+ 4	31	30	H _e	5, K _e	5	+ 11	57	57	+ 12	57	50	+ 10	61	62	+ 6	19	21	H _e	10, K _e	1	+ 0	37	33
* 1	42	38	+ 10	14	13	+ 14	14	14	+ 9	57	56	+ 8	35	34	+ 6	23	23	+ 10	18	16	+ 2	19	18			
* 2	29	28	+ 11	25	27	+ 12	24	25	+ 8	25	23	+ 6	68	67	+ 3	20	20	+ 14	27	28	H _e	11, K _e	4			
* 3	21	21	+ 12	23	20	+ 11	45	45	+ 7	63	63	+ 4	61	62	+ 4	45	20	+ 13	27	25	+ 16	20	17			
* 4	27	28	H _e	5, K _e	5	+ 10	51	55	+ 6	62	59	+ 2	58	58	H _e	8, K _e	1	+ 7	23	20	+ 12	37	36	+ 13	20	
* 5	32	31	+ 14	16	19	+ 9	46	46	+ 4	185	164	H _e	8, K _e	1	+ 7	23	20	+ 10	31	31	+ 11	15	16			
* 6	4, K _e	8	+ 12	27	25	+ 8	26	31	+ 3	98	101	+ 15	42	41	H _e	9, K _e	6	+ 10	61	62	+ 10	32	31			
* 12	22	19	+ 11	36	35	+ 7	16	18	+ 2	114	116	+ 14	17	16	+ 16	39	38	+ 9	61	62	+ 10	32	31			
* 2	23	20	+ 9	20	27	+ 5	20	70	+ 1	72	76	+ 13	29	29	+ 14	42	40	+ 8	67	66	+ 8	29	24			
* 1	34	34	+ 8	34	36	+ 4	29	29	+ 0	91	90	+ 12	40	41	+ 10	12	7	+ 2	26	27	+ 6	26	24			
* 2	21	19	+ 7	51	59	+ 3	49	48	+ 1	17	17	+ 10	27	26	+ 2	24	24	+ 6	35	32	+ 3	33	32			
* 3	29	27	+ 5	72	72	+ 2	42	42	+ 12	42	42	+ 6	60	61	+ 1	60	61	+ 5	65	65	+ 2	26	25			
* 4	21	22	+ 5	61	63	+ 9	13	13	+ 3	23	23	+ 8	82	82	+ 2	23	23	+ 7	44	66	+ 6	30	31			
* 5	H _e	4, K _e	+ 3	159	161	+ 1	47	50	+ 4	86	86	+ 5	71	71	+ 21	24	24	+ 9	37	41	H _e	11, K _e	5			
* 6	20	19	+ 2	34	37	+ 2	67	64	+ 5	33	36	+ 6	95	93	+ 4	29	29	+ 2	39	41	H _e	11, K _e	5			
* 7	H _e	5, K _e	+ 0	55	53	+ 4	43	43	+ 6	41	42	+ 4	60	59	+ 6	15	13	+ 1	54	53	+ 17	23	20			
* 8	21	19	+ 1	49	51	+ 6	21	19	+ 8	38	35	+ 3	21	23	H _e	9, K _e	1	+ 0	27	25	+ 13	23	22			
* 9	62	55	+ 3	61	62	+ 10	21	18	+ 9	33	30	+ 2	53	51	+ 14	29	30	+ 1	18	20	+ 10	25	21			
* 10	66	67	+ 4	68	70	H _e	6, K _e	2	+ 15	22	23	+ 1	75	75	+ 12	64	60	H _e	10, K _e	2	+ 4	19	18			
* 8	111	110	H _e	5, K _e	6	+ 14	29	25	+ 15	16	17	+ 2	66	67	+ 7	62	61	+ 13	29	27	+ 4	20	23			
* 6	278	271	+ 5	39	35	+ 3	13	26	+ 14	33	30	+ 6	39	39	+ 6	53	54	+ 12	16	16	+ 2	20	21			
* 4	63	59	+ 1	24	24	+ 12	23	23	+ 12	24	22	+ 2	44	44	+ 4	85	85	+ 6	19	10	+ 6	19	20			
* 5	30	30	+ 3	53	53	+ 10	45	45	+ 10	17	17	+ 17	17	17	+ 2	44	46	+ 5	35	36	+ 2	12, K _e	0			
* 2	26	27	+ 5	33	33	+ 9	102	109	+ 9	36	36	+ 10	24	24	+ 20	+ 1	17	19	+ 2	43	45	+ 16	19	17		
* 4	54	59	H _e	5, K _e	7	+ 8	28	28	+ 7	74	76	H _e	5, K _e	0	+ 0	46	46	+ 7	22	23	+ 14	25	22			
* 5	51	51	+ 13	24	17	+ 7	12	12	+ 7	55	56	+ 16	38	36	+ 3	14	14	+ 2	30	26	+ 10	50	54			
* 8	12	10	+ 11	25	22	+ 5	108	110	+ 6	126	125	+ 19	28	29	+ 4	19	19	+ 5	20	15	+ 5	31	32			
* 10	99	95	+ 9	14	15	+ 4	152	157	+ 5	28	27	+ 14	55	55	+ 5	20	23	H _e	10, K _e	3	+ 5	65	65			
* 17	17	15	+ 2	25	26	+ 3	133	135	+ 3	54	50	+ 11	67	66	+ 6	17	19	+ 14	27	24	+ 2	47	42			
* 18	15	15	+ 4	26	25	+ 1	83	87	+ 2	33	29	+ 9	34	37	+ 14	34	32	+ 9	20	18	H _e	12, K _e	1			
* 15	18	21	H _e	5, K _e	8	+ 0	38	40	+ 0	37	44	+ 8	14	13	+ 13	34	35	+ 7	29	27	+ 11	28	27			
* 14	16	18	H _e	5, K _e	8	+ 1	68	72	+ 1	37	33	+ 7	91	92	+ 1	35	35	+ 9	37	36	+ 9	44	43			
* 12	122	125	+ 2	24	19	+ 2	67	93	+ 104	104	104	+ 6	102	101	+ 10	101	101	+ 1	15	18	+ 8	16	14			
* 11	71	68	+ 20	15	16	+ 3	69	70	+ 2	35	35	+ 5	16	16	+ 49	49	49	+ 9	52	53	+ 1	15	15			
* 10	27	25	+ 5	12	10	+ 4	48	47	+ 12	24	24	+ 2	27	27	+ 19	55	55	+ 5	25	24	+ 4	16	12			
* 9	47	47	+ 4	43	42	+ 4	40	45	+ 4	47	47	+ 1	25	27	+ 27	65	65	+ 5	25	24	+ 4	16	12			
* 8	47	47	+ 4	43	42	+ 4	40	45	+ 4	47	47	+ 1	25	27	+ 27	65	65	+ 5	25	24	+ 4	16	12			
* 8	17	117	+ 1	30	29	+ 6	26	26	+ 1	14	14	+ 12	22	22	+ 2	22	22	+ 17	36	35	+ 15	25	21			
* 7	52	52	+ 1	19	14	+ 8	21	21	+ 1	14	14	+ 10	27	27	+ 17	36	35	+ 15	25	21	+ 10	16	13			
* 6	66	62	+ 3	26	26	+ 6	13	14	+ 11	24	24	+ 27	30	27	+ 7	24	24	+ 15	36	34	+ 15	16	13			
* 5	29	27	+ 8	23	23	+ 5	32	35	+ 1	41	41	+ 13	33	37	+ 10	23	19	+ 15	21	21	+ 9	20	17			
* 4	45	47	+ 6	179	180	+ 3	135	138	+ 3	25	27	+ 11	44	46	+ 6	50	H _e	9, K _e	3	+ 14	26	23	+ 0	30	27	
* 8	15	14	+ 4	31	35	+ 2	83	80	+ 7	74	74	+ 3	41	42	+ 1	42	38	+ 6	43	43	+ 4	14	16			
* 9	40	39	+ 1	24	24	+ 1	28	26	+ 16	15	15	+ 9	40	44	+ 10	36	36	+ 9	29	29	+ 1	19	17			
* 10	43	39	+ 1	8	0	+ 0	48	46	+ 14	26	26	+ 7	73	76	+ 9	28	31	+ 1	12	14	H _e	13, K _e	1			
* 1	100	99	+ 12	15	15	+ 5	4	20	+ 3	22	22	+ 6	32	32	+ 7	42	46	+ 1	20	18	+ 14	16	14			
* 0	102	95	+ 11	20	19	+ 1	21	21	+ 2	10	12	+ 1	20	19	+ 6	29	27	+ 2	33	25	+ 10	37	35			
* 4	72	74	+ 10	12	10	+ 1	21	21	+ 2	0	31	+ 9	17	19	H _e	9, K _e	4	+ 3	20	20	H _e	13, K _e	3			
* 5	24	23	+ 9	65	64	+ 5	28	29	+ 2	89	90	+ 7	31	29	+ 14	24	19	H _e	11, K _e	0	+ 17	15	9			
* 6	62	62	+ 8	13	9	H _e	6, K _e	7	+ 4	22	22	+ 6	23	23	+ 9	15	12	+ 16	23	22	+ 1	32	28			
* 5	3	+ 7	45	44	+ 11	20	17	H _e	7, K _e	5	+ 5	19	22	+ 7	32	3										

Table 3. Bond distances, bond angles and hydrogen bond lengths and angles. Estimated standard deviations in bond lengths between heavy atoms are 0.002 Å, in angles 0.1°, for bonds and angles involving hydrogen atoms 0.02 Å and 1.0°, respectively. Distances in parenthesis are corrected for librational motion.

Bond distances (Å)		Bond angles (°)	
C1—C2	1.351 (1.357)	C6—C1—C2	122.8
C2—C3	1.418 (1.425)	C1—C2—C3	120.5
C3—C4	1.500 (1.510)	C2—C3—C4	118.9
C4—C5	1.527 (1.534)	C3—C4—C5	114.5
C5—C6	1.524 (1.530)	C4—C5—C6	107.7
C5—C7	1.526 (1.534)	C4—C5—C7	109.5
C5—C8	1.522 (1.531)	C4—C5—C8	110.3
C6—C1	1.491 (1.495)	C5—C6—C1	113.3
C1—O1	1.326 (1.329)	C6—C5—C7	109.5
C3—O2	1.246 (1.249)	C6—C5—C8	110.5
C2—H1	0.96	C7—C5—C8	109.2
C4—H2	1.00	C2—C1—O1	123.9
C4—H3	0.96	C6—C1—O1	113.2
C6—H4	1.00	C2—C3—O2	122.5
C6—H5	0.97	C4—C3—O2	118.5
C7—H7	0.97		
C7—H8	0.99		
C7—H9	1.01		
C8—H10	0.96		
C8—H11	0.99		
C8—H12	0.99		

Hydrogen bond lengths (Å)	Hydrogen bond angles (°)
O1...O2	2.593
O1—H6	0.87
O2...H6	1.74
	O1—H6...O2 177
	C1—O1...O2 112.7
	C1—O1—H6 111
	C3—O2...O1 129.7
	C3—O2...H6 129

tions as shown in Table 3. The r.m.s. translational amplitudes corresponding to the discrepancy of 0.0022 Å are 0.22, 0.20, and 0.18 Å and the r.m.s. librational amplitudes are 5.5, 3.9, and 2.9°.

DISCUSSION

Bond lengths and angles are given in Table 3 and Fig. 1. The molecular arrangement in the unit cell is shown in Fig. 2. The compound crystallizes in the *trans*-enol form as expected from spectroscopic investigations,^{2,4} and there is an *inter*-molecular hydrogen bond (2.593 Å) between the hydrogen atom of the enol hydroxyl

and the oxygen atom of the carbonyl group. These bonds tie the molecules in helices along the two-fold screw axes. The molecules thus form polymers rather than dimers as has been found in solution. However, formation of a dimer with two hydrogen bridges as suggested,^{2,3} seems to be incompatible with normal hydrogen bonds if reasonable van der Waals distances are assumed. The O...O distance is significantly larger than corresponding distances found in analogous *cis*-enolized β-diketones (2.43–2.50 Å).^{6–8} In the open *trans* arrangement in dimedone, *inter*-molecular nonbonded repulsions may preferably give an expansion of the O...O distance, whereas in the *cis*-enolized β-diketones investigated,^{6–8} there are varying amounts of *intra*-molecular nonbonded repulsions giving the opposite effect. The hydrogen atom (H6) is unambiguously located in an asymmetric position near O1 (O1—H6 = 0.87 Å), and is situated very close to the least-squares plane through the conjugated enol system. The angle O1—H6...O2 is 177°.

The distances in the enol system reveal a considerable degree of conjugation, although not as much as in the *cis*-analogues.^{6–8} Similar conjugation effects have also been found in ascorbic,¹³ dialuric,¹⁴ and α-methyltetronic acids.¹⁵ Changes in bond lengths due to hydrogen bond formation in the conjugated system in barbiturates have been discussed by Craven *et al.*¹⁶ It is also reasonable to assume that formation of hydrogen bonds is partly responsible for the relatively large amount of conjugation in the above mentioned systems.

The conformation of the molecule might be described as an envelope. The carbon atom C5 is displaced by 0.61 Å from a least-squares plane through the other carbon atoms in the ring and the oxygen atoms (Table 4). The deviations from the plane are probably significant. The oxygen atoms and C4, and the other carbon atoms are displaced on opposite sides of the plane. The deviations thus give the cyclohexene ring a slightly curved boat-shaped form. The small deviation of O1 from a plane through C2, C1, and C6 is possibly associated with the formation of the hydrogen bond. The other deviations may be due to non-bonded interactions, especially between O2 and C1' across the two-fold screw axis in *x*=0 and O2 and H1 across the centre of symmetry in *x*=0. Very

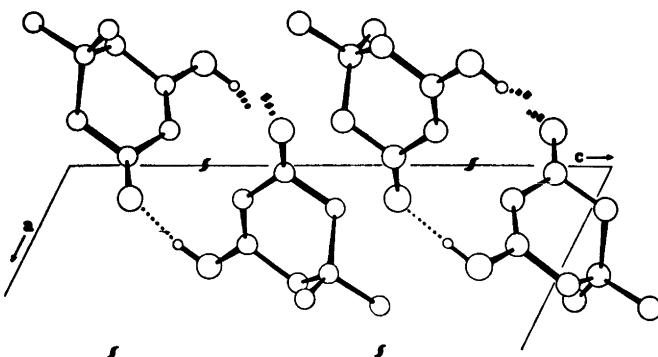


Fig. 2. The structure viewed along the *b* axis.

Table 4. Deviation (\AA) of atoms from least-squares plane through parts of the ring system in dimedone. Distances to atoms not defining the plane in parenthesis.

O1	0.035
O2	0.025
C1	-0.024
C2	-0.049
C3	-0.017
C4	0.017
C6	-0.007
C5	(-0.61)
H1	(-0.063)
H6	(-0.087)

similar deviations are found in the conformation of related six-membered rings.¹⁷

The sp^2-sp^3 and sp^3-sp^3 bonds have fairly normal values.¹⁸ The angles inside the ring progressively decrease from values close to 120° near the double bond to a tetrahedral value at C5. The arrangements around C5, C7, and C8 are very regular, none of the angles departing much from tetrahedral values, and the hydrogen atoms at C4 and C6 are in staggered positions relative to the methyl groups at C7 and C8.

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