

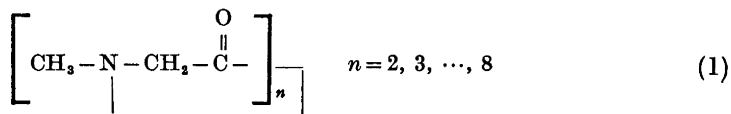
## Crystal Structure of Cyclooctasarcosyl

P. GROTH

*Department of Chemistry, University of Oslo, Oslo 3, Norway*

The crystals belong to the orthorhombic system with space group  $Pbca$  and cell dimensions  $a = 18.34_0$  Å,  $b = 18.27_0$  Å,  $c = 18.87_6$  Å. There are eight molecules in the unit cell. The phase problem was solved by direct methods. The  $R$ -value arrived at for 3471 observed reflections was 7.2 %. The ring conformation is surprisingly open with the inner volume filled by a cluster of four water molecules, which participate in a network of inter- and intra-molecular hydrogen bond bridges. Each of the four pairs of diametrically placed amino-acid residues is related by an approximate two-fold axis of symmetry. The conformation is *cis, cis, trans, trans, cis, cis, trans, trans*.

Cyclic oligopeptides of sarcosine of the general formula (I) have been studied by Dale and Titlestad.<sup>1</sup> To account for the relatively high observed



resistance to ring inversion, transannular interactions between N and C (carbonyl) were suggested. Such interactions have been reported to exist in certain cyclic aminoketones.<sup>2</sup> In the 10-ring lactone of 6-keto-9-hydroxy-nonanoic acid<sup>3</sup> there is strong evidence for transannular donor-acceptor attraction between the "ether" oxygen of the ester group and the carbonyl carbon atom.

In order to establish whether transannular N...C (carbonyl) attractions actually are stabilising the oligomers (I), and to obtain detailed information of the ring conformations and the geometries of the amino-acid residues, single crystals of some of these compounds are being examined by X-ray methods. Results for the cases  $n = 2$  and  $n = 4$  have been reported earlier.<sup>4,5</sup> In cyclo-tetrasarcosyl ( $n = 4$ ) the transannular N...C (carbonyl) distance was 3.08<sub>3</sub> Å, and no conclusion concerning stabilising transannular donor-acceptor attraction could be drawn. In the present paper the results of the crystal structure determination of cyclooctasarcosyl ( $n = 8$ ) are presented.

The crystals belong to the orthorhombic system and the systematic absences lead to the space group  $Pbca$ . The cell parameters, measured by means of a four circle diffractometer, and their estimated standard deviations are:

$$a = 18.340(3)\text{\AA}, \quad b = 18.279(3)\text{\AA}, \quad c = 18.875(2)\text{\AA}$$

With eight molecules per unit cell the calculated density is  $\rho_c = 1.20 \text{ g cm}^{-3}$ . The observed density,  $\rho_o = 1.34 \text{ g cm}^{-3}$ , corresponds to a difference in molecular weight of 72 which is accounted for by assuming the presence of four water molecules per asymmetric unit.

With  $2\theta(\text{max}) = 50^\circ$  and  $\text{MoK}\alpha$ -radiation, about 5000 independent reflections were measured on an automatic four-circle diffractometer. Using an observed-unobserved cutoff at  $2.5\sigma(I)$ , 3471 were recorded as observed. No corrections have been made for absorption or secondary extinction effects.

Table 1. Final fractional coordinates and anisotropic thermal vibration parameters with estimated standard deviations (multiplied by  $10^5$  for nonhydrogens and  $10^4$  for hydrogens). The symbols CC, CM and OV are used for carbonyl carbons, methyl carbons and water oxygens, respectively. Hn1 and Hn2 are bonded to Cn.

ATOM	X	Y	Z	B	B11	B22	B33	B12	B13	B23
O1	7692(19)	18723(28)	35285(28)		228( 12)	353( 15)	383( 15)	+152( 22)	138( 24)	+91( 28)
O2	55212(18)	28366(17)	29976(19)		256( 13)	217( 11)	311( 14)	+19( 17)	+12( 22)	173( 21)
O3	4318(20)	26792(18)	41380(19)		415( 15)	242( 13)	263( 13)	141( 15)	+213( 23)	+126( 21)
O4	25152(19)	34898(18)	39854(20)		333( 16)	177( 11)	396( 16)	+141( 20)	62( 24)	+4( 24)
O5	24735(17)	57814(19)	39979(19)		266( 12)	301( 13)	333( 14)	+83( 20)	+78( 22)	148( 24)
O6	44882(17)	55287(18)	31195(19)		198( 12)	289( 13)	307( 13)	21( 20)	+108( 21)	+179( 22)
O7	56598(28)	48143(18)	41377(19)		368( 16)	278( 13)	232( 13)	46( 23)	+118( 23)	72( 21)
O8	74555(28)	48383(19)	49883(21)		295( 14)	170( 11)	368( 14)	+69( 19)	+148( 23)	48( 23)
OV1	47653(24)	44883(22)	28788(22)		471( 18)	412( 16)	373( 17)	81( 29)	+131( 29)	+189( 27)
OV2	52358(29)	38892(23)	19468(26)		669( 23)	479( 19)	927( 22)	498( 35)	114( 37)	362( 34)
OV3	39965(22)	42211(23)	8128(22)		392( 16)	442( 17)	372( 16)	8( 27)	+93( 27)	+8( 28)
OV4	49238(23)	38748(23)	3794(24)		475( 18)	485( 17)	388( 17)	1( 30)	+87( 29)	+35( 29)
N1	85888(28)	12147(28)	36315(21)		166( 13)	168( 13)	268( 15)	20( 21)	7( 24)	38( 23)
N2	4848(28)	14147(28)	34818(23)		163( 13)	178( 13)	309( 17)	44( 21)	+16( 24)	+48( 24)
N3	37232(23)	31473(21)	31866(22)		264( 18)	228( 14)	258( 15)	198( 25)	+78( 26)	+144( 26)
N4	26332(23)	45816(28)	43437(22)		293( 16)	166( 13)	266( 16)	+43( 24)	189( 27)	+21( 24)
N5	84677(28)	83888(19)	38958(22)		191( 14)	147( 12)	317( 15)	38( 28)	64( 25)	29( 24)
N6	56477(28)	89228(21)	36929(22)		193( 13)	185( 13)	288( 15)	27( 22)	+25( 26)	+32( 24)
N7	62824(22)	42561(21)	31468(22)		244( 15)	193( 13)	259( 15)	114( 23)	+12( 26)	29( 25)
N8	71269(21)	29554(28)	43841(21)		237( 15)	171( 13)	278( 15)	7( 23)	+82( 25)	35( 24)
C1	58284(26)	11393(24)	38679(22)		193( 16)	127( 14)	282( 19)	+36( 25)	+7( 29)	54( 28)
C2	48744(27)	19853(27)	31333(26)		181( 17)	238( 17)	381( 20)	68( 29)	+71( 31)	+124( 33)
C3	36743(27)	38348(27)	38861(28)		258( 18)	188( 17)	292( 18)	59( 28)	+48( 33)	18( 31)
C4	33298(27)	86978(24)	43687(27)		266( 18)	139( 14)	241( 18)	23( 28)	+42( 31)	+25( 27)
C5	42816(27)	62845(24)	41399(29)		288( 17)	166( 15)	381( 20)	+18( 27)	+8( 31)	+97( 29)
C6	59282(25)	85894(26)	32387(27)		155( 16)	281( 16)	283( 19)	22( 27)	39( 30)	+18( 30)
C7	63628(28)	35898(28)	34749(29)		273( 19)	163( 16)	314( 21)	89( 28)	+77( 34)	17( 31)
C8	68769(38)	23279(28)	43298(33)		288( 19)	191( 16)	271( 21)	3( 38)	9( 34)	86( 34)
CM1	88728(38)	8696(29)	31219(31)		263( 20)	246( 20)	395( 22)	86( 33)	+11( 35)	+222( 35)
CM2	42978(31)	8698(38)	39737(32)		257( 20)	272( 20)	367( 23)	+188( 34)	155( 36)	58( 37)
CM3	33328(35)	38825(32)	24836(32)		393( 25)	369( 23)	252( 21)	282( 48)	+254( 39)	+78( 41)
CM4	21368(34)	45811(32)	47466(35)		364( 26)	296( 23)	489( 27)	+18( 38)	395( 44)	13( 42)
CM5	32819(31)	69291(29)	34494(34)		283( 20)	186( 17)	413( 25)	88( 32)	188( 38)	225( 36)
CM6	58818(32)	62977(28)	34883(34)		315( 23)	328( 23)	391( 26)	+88( 38)	+234( 48)	+279( 48)
CM7	86738(32)	43997(38)	24489(38)		328( 23)	326( 22)	224( 20)	114( 37)	163( 35)	+8( 37)
CM8	77972(32)	29954(38)	48336(31)		281( 21)	278( 21)	344( 23)	+48( 34)	+292( 37)	128( 37)
CC1	69988(27)	17998(26)	37865(27)		189( 17)	181( 16)	218( 19)	55( 28)	+16( 38)	66( 29)
CC2	83828(27)	15888(26)	34838(29)		188( 17)	148( 15)	257( 19)	25( 27)	19( 38)	+88( 29)
CC3	4868(26)	28128(29)	35389(28)		198( 17)	218( 19)	237( 19)	59( 38)	9( 32)	+86( 32)
CC4	29968(29)	38951(26)	39838(28)		253( 19)	151( 16)	216( 18)	67( 29)	+25( 38)	57( 30)
CC5	38654(28)	57222(26)	39844(27)		281( 18)	186( 16)	284( 18)	16( 29)	56( 39)	+13( 29)
CC6	47464(28)	58952(26)	36873(28)		168( 16)	167( 15)	232( 18)	8( 28)	+31( 36)	1( 29)
CC7	59458(28)	47547(27)	35485(28)		185( 17)	186( 17)	284( 18)	13( 28)	+85( 31)	+8( 29)
CC8	78287(29)	35388(26)	39787(28)		231( 18)	156( 16)	223( 19)	36( 29)	+14( 31)	+74( 38)
H11	5748(24)	1286(23)	4381(27)	1,82(1,13)						
H12	5712(25)	663(25)	3848(24)	1,84(1,18)						
H21	3562(26)	1783(28)	3178(25)	1,88(1,18)						
H22	4278(25)	1981(23)	2645(27)	2,93(1,88)						
H31	3686(25)	4244(28)	3194(25)	2,77(1,18)						
H32	4185(27)	3914(28)	3877(25)	4,88(1,11)						
H41	3883(26)	4946(24)	4171(24)	4,88(1,13)						
H42	3354(25)	5288(24)	4966(26)	3,42(1,11)						
H51	4362(25)	6885(24)	4595(26)	1,62(1,18)						
H52	4448(24)	6728(26)	4222(24)	2,38(1,11)						
H61	6429(26)	3678(24)	5229(28)	3,18(1,11)						
H62	5714(25)	8489(21)	2953(25)	4,44(1,12)						
H71	6393(25)	3196(25)	3885(25)	3,35(1,11)						
H72	5874(27)	3458(25)	3773(25)	2,39(1,18)						
H81	6227(27)	2486(28)	4244(25)	3,81(1,15)						
H82	6689(25)	2119(28)	4825(27)	2,79(1,19)						











$$\exp - (B_{11}h^2 + B_{22}k^2 + B_{33}l^2 + B_{12}hk + B_{13}hl + B_{23}kl)$$

A comparison between observed and calculated structure factors is presented in Table 2.

The principal axes of the thermal vibration ellipsoids for oxygen, nitrogen, and carbon atoms were calculated from the temperature parameters of Table 1. Maximum root mean square amplitudes range from about 0.20 Å for carbonyl carbons to about 0.30 Å for methyl carbon atoms and water oxygens. Due to the size of the molecule, no rigid-body analysis of translational and librational motion has been carried out.

Interatomic distances, bond angles, and dihedral angles are given in Table 3. The standard deviations, given in parentheses, are estimated from the correlation matrix of the last least squares refinement cycle. Fig. 1 shows the molecule viewed along [001].

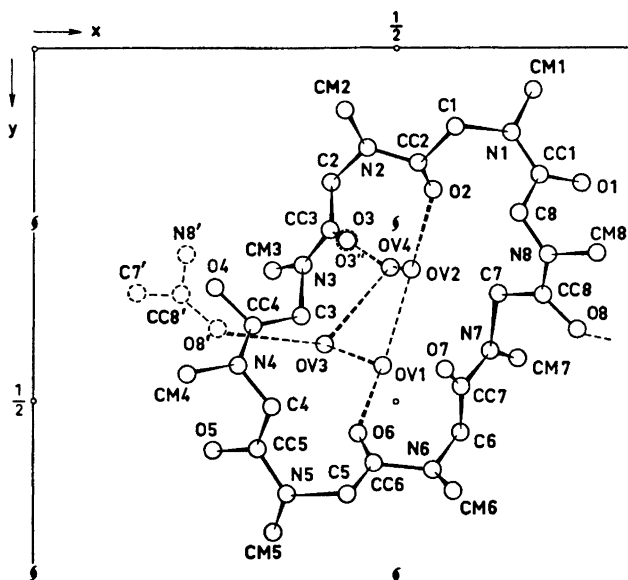


Fig. 1. The molecule viewed along [001].

By averaging bond distances of Table 3, and comparing with the results of the corresponding dimeric<sup>4</sup> and tetrameric<sup>5</sup> compounds, no significant differences are observed:

Distance	Cyclodisarcosyl	Cyclotetrasarcosyl	Cyclooctasarcosyl
CC - CM	1.506 Å	1.531 Å	1.527 Å
CC - N	1.348 Å	1.358 Å	1.344 Å
CC - O	1.234 Å	1.225 Å	1.228 Å
C - N	1.455 Å	1.458 Å	1.456 Å
CM - N	1.475 Å	1.467 Å	1.483 Å



Table 3. Interatomic distances, bond angles and dihedral angles with estimated standard deviations.

DISTANCE	(Å)	DISTANCE	(Å)	DISTANCE	(Å)
O1 = C1	1,227(5)	O2 = C2	1,234(5)	O3 = C3	1,231(5)
O4 = C4	1,222(5)	O5 = C5	1,219(5)	O6 = C6	1,233(5)
O7 = C7	1,227(5)	O8 = C8	1,230(5)	N1 = C1	1,488(5)
N2 = C2	1,479(6)	N3 = C3	1,488(6)	N4 = C4	1,483(6)
N5 = C5	1,481(6)	N6 = C6	1,487(6)	N7 = C7	1,487(6)
N8 = C8	1,482(6)	N1 = C1	1,484(6)	N2 = C2	1,482(6)
N3 = C3	1,464(6)	N4 = C4	1,448(6)	N5 = C5	1,451(6)
N6 = C6	1,466(6)	N7 = C7	1,452(6)	N8 = C8	1,447(6)
N1 = C1	1,352(6)	N2 = C2	1,352(6)	N3 = C3	1,339(6)
N4 = C4	1,348(6)	N5 = C5	1,358(6)	N6 = C6	1,338(6)
N7 = C7	1,345(6)	N8 = C8	1,338(6)	C1 = C2	1,519(7)
C2 = C3	1,520(7)	C3 = C4	1,530(7)	C4 = C5	1,531(7)
C5 = C6	1,520(7)	C6 = C7	1,526(7)	C7 = C8	1,529(7)
C8 = C1	1,531(7)	OV1 = OV2	2,787(6)	OV1 = OV3	2,743(6)
OV2 = O2	2,814(5)	OV3 = O8	2,868(6)	OV2 = OV4	2,928(7)
OV4 = O3	2,883(6)	O6 = OV1	2,791(5)	OV3 = OV4	2,884(6)

ANGLE	(°)	ANGLE	(°)
O1 = C1 = N1	121,94(48)	O2 = C2 = N2	121,79(46)
O3 = C3 = N3	122,85(47)	O4 = C4 = N4	121,98(48)
O5 = C5 = N5	122,93(46)	O6 = C6 = N6	121,99(45)
O7 = C7 = N7	121,72(47)	O8 = C8 = N8	121,76(46)
O1 = C1 = C8	128,64(46)	O2 = C2 = C1	121,26(43)
O3 = C3 = C2	128,92(48)	O4 = C4 = C3	128,86(46)
O5 = C5 = C4	129,71(45)	O6 = C6 = C5	128,92(43)
O7 = C7 = C6	121,16(47)	O8 = C8 = C7	128,51(45)
C1 = N1 = C1	116,87(41)	C2 = N2 = C2	118,99(43)
C3 = N3 = C3	116,31(43)	C4 = N4 = C4	116,68(44)
C5 = N5 = C5	116,28(41)	C6 = N6 = C6	118,88(41)
C7 = N7 = C7	119,69(43)	C8 = N8 = C8	117,33(43)
C1 = N1 = CC1	119,15(39)	C2 = N2 = CC2	123,68(43)
C3 = N3 = CC3	124,64(43)	C4 = N4 = CC4	118,37(43)
C5 = N5 = CC5	119,18(42)	C6 = N6 = CC6	122,83(43)
C7 = N7 = CC7	124,27(42)	C8 = N8 = CC8	118,28(42)
C8 = C1 = N1	117,36(42)	C1 = C2 = N2	119,88(44)
C2 = C3 = N3	116,53(44)	C3 = C4 = N4	117,24(45)
C4 = C5 = N5	116,23(43)	C5 = C6 = N6	117,88(44)
C6 = C7 = N7	117,11(43)	C7 = C8 = N8	117,73(44)
C1 = N1 = C1	123,68(42)	C2 = N2 = C2	116,76(44)
CC3 = N3 = C3	118,92(43)	CC4 = N4 = C4	125,83(45)
CC5 = N5 = C5	122,72(42)	CC6 = N6 = C6	116,57(42)
CC7 = N7 = C7	116,92(44)	CC8 = N8 = C8	123,79(44)
N1 = C1 = CC2	112,19(42)	N2 = C2 = CC3	111,86(43)
N3 = C3 = CC4	118,61(44)	N4 = C4 = CC5	112,44(44)
N5 = C5 = CC6	112,36(44)	N6 = C6 = CC7	118,38(42)
N7 = C7 = CC8	118,88(44)	N8 = C8 = CC1	111,86(44)
CC2 = O2 = OV2	149,97(33)	O2 = OV2 = OV1	132,15(22)
OV2 = OV1 = OV3	85,23(18)	OV1 = OV3 = O8	115,48(19)
OV2 = OV4 = O3	148,88(22)	OV4 = O3 = CC3	144,95(36)
OV4 = OV2 = O2	134,41(22)	OV4 = OV2 = OV1	91,68(17)
O6 = OV1 = OV2	141,19(20)	O6 = OV1 = OV3	127,46(19)
OV4 = OV3 = O8	121,82(19)	OV3 = OV4 = O3	111,78(18)
OV1 = OV3 = OV4	93,75(17)	OV3 = OV4 = OV2	88,88(16)

DIHEDRAL ANGLE	(°)	DIHEDRAL ANGLE	(°)
CC1 = N1 = C1 = C2	-77,58(68)	N1 = C1 = C2 = N2	-167,51(43)
C1 = C2 = N2 = C2	-169,88(44)	C2 = N2 = C2 = C3	76,99(61)
N2 = C2 = C3 = N3	-167,69(44)	C2 = C3 = N3 = C3	-178,28(46)
CC3 = N3 = C3 = C4	93,18(57)	N3 = C3 = C4 = N4	-179,39(43)
C3 = C4 = N4 = C4	-1,88(75)	CC4 = N4 = C4 = C5	-186,89(56)
N4 = C4 = C5 = N5	179,76(42)	C4 = C5 = N5 = C5	-15,88(69)
CC5 = N5 = C5 = C6	-72,38(63)	N5 = C5 = C6 = N6	-167,84(43)
C5 = C6 = N6 = C6	-171,72(45)	CC6 = N6 = C6 = C7	77,83(55)
N6 = C6 = C7 = N7	-173,83(41)	C6 = C7 = N7 = C7	-173,94(43)
CC7 = N7 = C7 = C8	83,87(57)	N7 = C7 = C8 = N8	-172,52(44)
C7 = C8 = N8 = C8	-5,68(73)	CC8 = N8 = C8 = C1	-91,88(58)
N8 = C8 = C1 = N1	-173,36(42)	C8 = C1 = N1 = C1	-9,88(67)

The geometry of the *cis* and *trans* amide groups, respectively, is roughly the same as for cyclotetrasarcosyl:

Angle	Cyclotetrasarcosyl	Cyclooctasarcosyl
(CM - N - CC) <i>cis</i>	119.8°	118.7°
(CM - N - CC) <i>trans</i>	124.3°	123.9°
(C - N - CC) <i>cis</i>	123.9°	123.8°
(C - N - CC) <i>trans</i>	120.1°	117.2°

The corresponding angles in cyclodisarcosyl, where the amide group has the *cis* conformation, are 119.7° and 124.6°, respectively.

As may be seen from Fig. 1, the ring has an open conformation with the inner volume filled by a cluster of four water molecules which participate in a network of inter- as well as intra-molecular hydrogen bond bridges. The four water oxygens are situated at the corners of a somewhat distorted square with  $OV-OV-OV$  angles of about  $81^\circ$ ,  $91^\circ$ ,  $85^\circ$ , and  $93^\circ$ , and  $OV\dots OV$  bondlengths ranging from 2.743 Å to 2.920 Å. The two  $OV\dots O$  bond distances of the intra-molecular hydrogen-bond bridge are 2.791 Å and 2.814 Å, respectively. The  $OV\dots O$  distances of the two inter-molecular bridges, linking different symmetry related molecules, are somewhat longer:  $OV3\dots O8' = 2.868$  Å,  $OV4\dots O3'' = 2.883$  Å. The  $OV-OV-O$  angles are distributed over a wide range (from  $112^\circ$  to  $150^\circ$ ).

There are no direct transannular interactions to be held responsible, as originally thought,<sup>1</sup> for the rigidity of this rather large ring. Since, from IR-spectra, it can be stated that the same conformation persists in  $CHCl_3$  solution, where no water molecules are present to form transannular bridges, the explanation must be sought in the intrinsic conformation of the peptide chain itself.<sup>8</sup>

Fig. 1 clearly shows that the ring conformation is *cis,cis,trans,trans,cis,cis,trans,trans*. It might also be seen that each of the four pairs of diametrically placed amino-acid residues is related by an approximate two fold axis of symmetry.

Apart from the hydrogen bonds, no short intermolecular distances are observed.

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