

## ***Appendix 2***

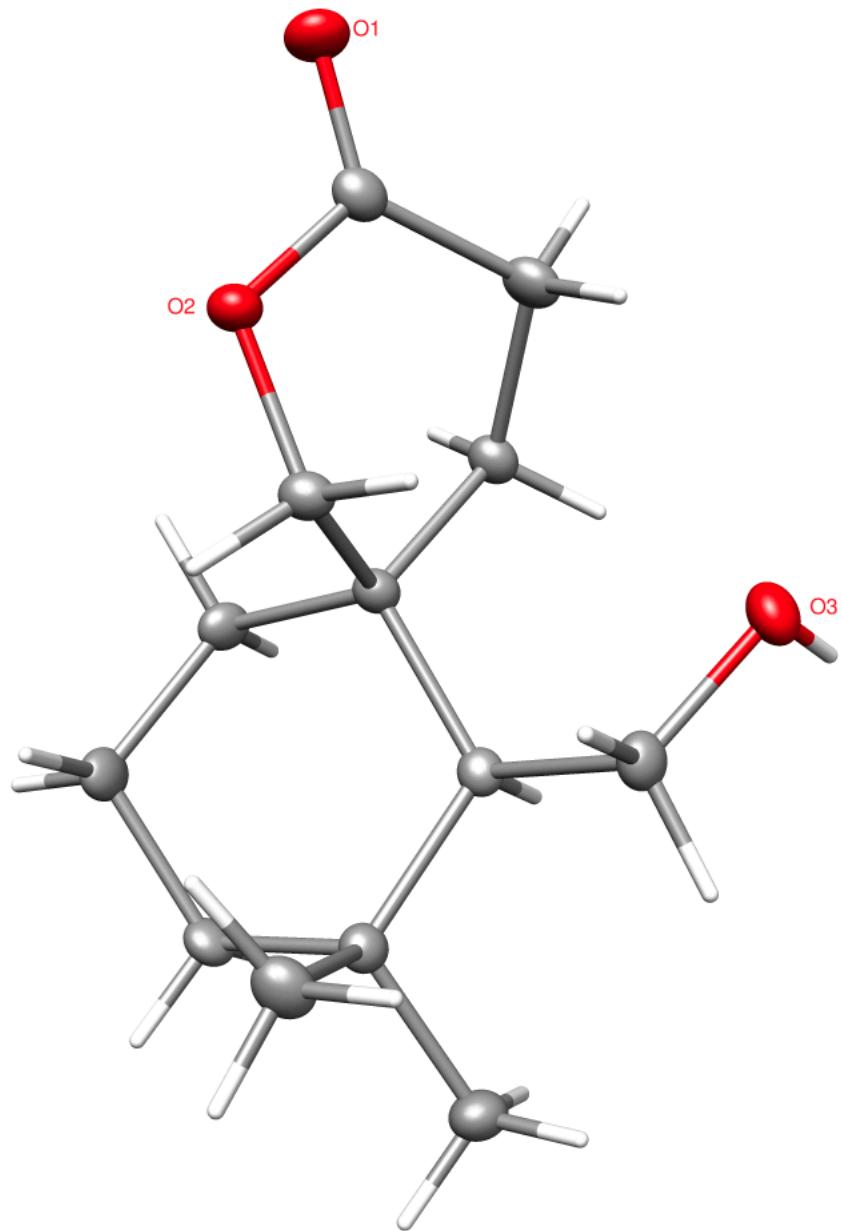
*X-Ray Crystallography Reports Relevant to Chapter 2:  
Total Synthesis of (–)-Maoecrystal Z<sup>†</sup>*

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<sup>†</sup> The work disclosed in this appendix for the X-ray crystallographic analysis of **109** and **127** was completed entirely by Mr. Larry Henling and Dr. Michael Day in the Caltech X-ray crystallography lab.

## A2.1 CRYSTAL STRUCTURE ANALYSIS OF SPIROLACTONE 109

**Figure A2.1.** Spirolactone **109**. Crystallographic data have been deposited at the CCDC, 12 Union Road, Cambridge CB2 1EZ, UK and copies can be obtained on request, free of charge, by quoting the publication citation and the deposition number 837088.



**Table A2.1.** Crystal data and structure refinement for spirolactone **109** (CCDC 837088).

Empirical formula	C <sub>13</sub> H <sub>22</sub> O <sub>3</sub>	
Formula weight	226.31	
Crystallization Solvent	Isooctane/dichloromethane	
Crystal Habit	Column	
Crystal size	0.22 x 0.15 x 0.07 mm <sup>3</sup>	
Crystal color	Colorless	

### Data Collection

Type of diffractometer	Bruker KAPPA APEX II		
Wavelength	0.71073 Å MoKα		
Data Collection Temperature	100(2) K		
θ range for 5677 reflections used in lattice determination	2.53	to 36.05°	
Unit cell dimensions	a = 8.3489(4) Å	$\alpha$ = 90°	b = 7.3054(3) Å c = 9.9944(5) Å $\beta$ = 105.627(3)° $\gamma$ = 90°
Volume	587.05(5) Å <sup>3</sup>		
Z	2		
Crystal system	Monoclinic		
Space group	P 2 <sub>1</sub>		
Density (calculated)	1.280 Mg/m <sup>3</sup>		
F(000)	248		
Data collection program	Bruker APEX2 v2009.7-0		
θ range for data collection	2.12	to 36.35°	
Completeness to θ = 36.35°	99.6 %		
Index ranges	-13 ≤ h ≤ 12, -12 ≤ k ≤ 12, -16 ≤ l ≤ 16		
Data collection scan type	ω scans; 9 settings		
Data reduction program	Bruker SAINT-Plus v7.66A		
Reflections collected	17473		
Independent reflections	5474 [R <sub>int</sub> = 0.0331]		
Absorption coefficient	0.089 mm <sup>-1</sup>		
Absorption correction	None		
Max. and min. transmission	0.9938 and 0.9807		

**Table A2.1 (cont.)****Structure solution and Refinement**

Structure solution program	SHELXS-97 (Sheldrick, 2008)
Primary solution method	Direct methods
Secondary solution method	Difference Fourier map
Hydrogen placement	Difference Fourier map
Structure refinement program	SHELXL-97 (Sheldrick, 2008)
Refinement method	Full matrix least-squares on $F^2$
Data / restraints / parameters	5474 / 1 / 233
Treatment of hydrogen atoms	Unrestrained
Goodness-of-fit on $F^2$	1.709
Final R indices [ $I > 2\sigma(I)$ , 4661 reflections]	$R_1 = 0.0408$ , $wR_2 = 0.0498$
R indices (all data)	$R_1 = 0.0512$ , $wR_2 = 0.0504$
Type of weighting scheme used	Sigma
Weighting scheme used	$w = 1/\sigma^2(F_{\text{o}}^2)$
Max shift/error	0.001
Average shift/error	0.000
Absolute structure determination	Unable to reliably determine
Absolute structure parameter	-0.2(6)
Largest diff. peak and hole	0.380 and -0.314 e. $\text{\AA}^{-3}$

**Special Refinement Details**

Crystals were mounted on a glass fiber using Paratone oil then placed on the diffractometer under a nitrogen stream at 100K.

Refinement of  $F^2$  against ALL reflections. The weighted R-factor ( $wR$ ) and goodness of fit (S) are based on  $F^2$ , conventional R-factors (R) are based on F, with F set to zero for negative  $F^2$ . The threshold expression of  $F^2 > 2\sigma(F^2)$  is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on  $F^2$  are statistically about twice as large as those based on F, and R-factors based on ALL data will be even larger.

All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds

involving l.s. planes.

**Table A2.2.** Atomic coordinates ( $\times 10^4$ ) and equivalent isotropic displacement parameters ( $\text{\AA}^2 \times 10^3$ ) for spirolactone **109** (CCDC 837088).  $U(\text{eq})$  is defined as the trace of the orthogonalized  $U^{ij}$  tensor.

	x	y	z	$U_{\text{eq}}$
O(1)	6715(1)	-573(1)	7277(1)	22(1)
O(2)	4260(1)	682(1)	7046(1)	18(1)
O(3)	4941(1)	6385(1)	5917(1)	21(1)
C(1)	5908(1)	820(1)	7204(1)	17(1)
C(2)	6586(1)	2720(1)	7332(1)	18(1)
C(3)	5836(1)	3874(1)	8286(1)	16(1)
C(4)	3897(1)	3793(1)	7901(1)	13(1)
C(5)	3322(1)	2395(1)	6732(1)	16(1)
C(6)	3432(1)	3115(2)	9204(1)	16(1)
C(7)	1601(1)	3218(1)	9127(1)	19(1)
C(8)	1023(1)	5195(1)	8884(1)	16(1)
C(9)	1284(1)	6031(1)	7544(1)	14(1)
C(10)	3106(1)	5731(1)	7462(1)	13(1)
C(11)	3280(1)	6376(1)	6038(1)	17(1)
C(12)	-36(1)	5250(2)	6284(1)	20(1)
C(13)	938(1)	8103(1)	7603(1)	21(1)

**Table A2.3.** Bond lengths [Å] and angles [°] for spirolactone **109** (CCDC 837088).

O(1)-C(1)	1.2118(12)	C(2)-C(3)-C(4)	113.43(8)
O(2)-C(1)	1.3455(11)	C(2)-C(3)-H(3A)	108.7(7)
O(2)-C(5)	1.4640(12)	C(4)-C(3)-H(3A)	109.8(6)
O(3)-C(11)	1.4239(12)	C(2)-C(3)-H(3B)	107.5(6)
O(3)-H(3)	0.780(12)	C(4)-C(3)-H(3B)	108.2(6)
C(1)-C(2)	1.4917(14)	H(3A)-C(3)-H(3B)	109.2(8)
C(2)-C(3)	1.5280(14)	C(5)-C(4)-C(6)	109.43(8)
C(2)-H(2A)	0.972(10)	C(5)-C(4)-C(3)	107.76(8)
C(2)-H(2B)	0.946(10)	C(6)-C(4)-C(3)	106.58(7)
C(3)-C(4)	1.5608(13)	C(5)-C(4)-C(10)	111.49(7)
C(3)-H(3A)	0.954(10)	C(6)-C(4)-C(10)	110.09(8)
C(3)-H(3B)	0.966(10)	C(3)-C(4)-C(10)	111.34(8)
C(4)-C(5)	1.5288(13)	O(2)-C(5)-C(4)	111.57(7)
C(4)-C(6)	1.5377(13)	O(2)-C(5)-H(5A)	105.0(6)
C(4)-C(10)	1.5737(13)	C(4)-C(5)-H(5A)	109.2(6)
C(5)-H(5A)	0.989(11)	O(2)-C(5)-H(5B)	107.9(6)
C(5)-H(5B)	0.968(9)	C(4)-C(5)-H(5B)	111.2(6)
C(6)-C(7)	1.5115(14)	H(5A)-C(5)-H(5B)	111.8(8)
C(6)-H(6A)	0.971(10)	C(7)-C(6)-C(4)	115.05(8)
C(6)-H(6B)	0.967(10)	C(7)-C(6)-H(6A)	108.3(6)
C(7)-C(8)	1.5213(14)	C(4)-C(6)-H(6A)	108.2(6)
C(7)-H(7A)	1.022(10)	C(7)-C(6)-H(6B)	112.0(7)
C(7)-H(7B)	1.006(11)	C(4)-C(6)-H(6B)	108.1(7)
C(8)-C(9)	1.5394(13)	H(6A)-C(6)-H(6B)	104.6(8)
C(8)-H(8A)	0.974(10)	C(6)-C(7)-C(8)	109.17(9)
C(8)-H(8B)	0.921(11)	C(6)-C(7)-H(7A)	112.5(6)
C(9)-C(12)	1.5420(13)	C(8)-C(7)-H(7A)	111.7(5)
C(9)-C(13)	1.5449(14)	C(6)-C(7)-H(7B)	110.1(6)
C(9)-C(10)	1.5613(13)	C(8)-C(7)-H(7B)	108.9(6)
C(10)-C(11)	1.5425(12)	H(7A)-C(7)-H(7B)	104.2(8)
C(10)-H(10)	0.960(9)	C(7)-C(8)-C(9)	113.76(8)
C(11)-H(11A)	1.026(10)	C(7)-C(8)-H(8A)	108.8(6)
C(11)-H(11B)	1.036(10)	C(9)-C(8)-H(8A)	109.6(6)
C(12)-H(12A)	0.961(12)	C(7)-C(8)-H(8B)	111.0(6)
C(12)-H(12B)	0.960(10)	C(9)-C(8)-H(8B)	109.0(6)
C(12)-H(12C)	0.938(12)	H(8A)-C(8)-H(8B)	104.2(8)
C(13)-H(13A)	0.952(11)	C(8)-C(9)-C(12)	109.05(8)
C(13)-H(13B)	0.931(11)	C(8)-C(9)-C(13)	106.48(8)
C(13)-H(13C)	0.991(10)	C(12)-C(9)-C(13)	107.08(8)
		C(8)-C(9)-C(10)	110.98(7)
C(1)-O(2)-C(5)	115.62(8)	C(12)-C(9)-C(10)	113.40(8)
C(11)-O(3)-H(3)	105.0(10)	C(13)-C(9)-C(10)	109.54(8)
O(1)-C(1)-O(2)	118.63(9)	C(11)-C(10)-C(9)	109.93(7)
O(1)-C(1)-C(2)	125.72(9)	C(11)-C(10)-C(4)	113.29(8)
O(2)-C(1)-C(2)	115.60(9)	C(9)-C(10)-C(4)	116.95(8)
C(1)-C(2)-C(3)	110.91(9)	C(11)-C(10)-H(10)	107.4(6)
C(1)-C(2)-H(2A)	109.9(6)	C(9)-C(10)-H(10)	101.7(6)
C(3)-C(2)-H(2A)	109.8(6)	C(4)-C(10)-H(10)	106.4(5)
C(1)-C(2)-H(2B)	107.2(6)	O(3)-C(11)-C(10)	114.57(8)
C(3)-C(2)-H(2B)	111.1(6)	O(3)-C(11)-H(11A)	107.1(6)
H(2A)-C(2)-H(2B)	107.8(8)	C(10)-C(11)-H(11A)	108.5(6)

O(3)-C(11)-H(11B)	110.9(6)	H(12B)-C(12)-H(12C)	109.2(9)
C(10)-C(11)-H(11B)	108.1(5)	C(9)-C(13)-H(13A)	110.4(6)
H(11A)-C(11)-H(11B)	107.5(7)	C(9)-C(13)-H(13B)	111.3(6)
C(9)-C(12)-H(12A)	113.5(6)	H(13A)-C(13)-H(13B)	110.6(9)
C(9)-C(12)-H(12B)	108.9(6)	C(9)-C(13)-H(13C)	110.9(6)
H(12A)-C(12)-H(12B)	108.8(9)	H(13A)-C(13)-H(13C)	104.7(8)
C(9)-C(12)-H(12C)	107.1(6)	H(13B)-C(13)-H(13C)	108.8(9)
H(12A)-C(12)-H(12C)	109.2(9)		

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**Table A2.4.** Anisotropic displacement parameters ( $\text{\AA}^2 \times 10^4$ ) for spiro lactone **109** (CCDC 837088). The anisotropic displacement factor exponent takes the form:  
 $-2p^2 [ h^2 a^{*2} U^{11} + \dots + 2 h k a^* b^* U^{12} ]$

	U <sup>11</sup>	U <sup>22</sup>	U <sup>33</sup>	U <sup>23</sup>	U <sup>13</sup>	U <sup>12</sup>
O(1)	188(4)	184(4)	282(4)	-9(3)	54(3)	43(3)
O(2)	152(4)	148(4)	242(3)	-22(3)	72(3)	9(3)
O(3)	218(4)	219(4)	233(4)	-10(3)	125(3)	-28(3)
C(1)	161(5)	199(5)	147(4)	-20(4)	55(3)	-6(4)
C(2)	125(5)	214(6)	217(5)	-39(4)	51(4)	-6(4)
C(3)	128(5)	180(5)	165(4)	-17(4)	39(3)	-9(4)
C(4)	115(5)	142(4)	132(4)	-7(3)	31(3)	3(4)
C(5)	135(5)	153(5)	174(4)	-29(4)	37(4)	5(4)
C(6)	186(5)	155(5)	139(4)	23(4)	42(4)	24(4)
C(7)	213(5)	182(5)	194(4)	39(4)	106(4)	7(4)
C(8)	153(5)	182(5)	168(4)	-19(4)	74(4)	7(4)
C(9)	143(5)	150(4)	141(4)	2(4)	40(3)	0(4)
C(10)	133(5)	131(4)	113(4)	-23(3)	29(3)	-17(4)
C(11)	184(5)	173(5)	155(4)	20(4)	62(4)	-11(4)
C(12)	136(5)	248(6)	193(5)	6(4)	11(4)	0(4)
C(13)	198(6)	177(5)	240(5)	24(4)	63(4)	38(5)

**Table A2.5.** Hydrogen coordinates ( $\times 10^4$ ) and isotropic displacement parameters ( $\text{\AA}^2 \times 10^3$ ) for spirolactone **109** (CCDC 837088).

	X	y	z	$U_{\text{iso}}$
H(3)	5343(16)	7264(17)	6308(13)	38(4)
H(2A)	6355(12)	3286(14)	6419(11)	18(3)
H(2B)	7754(13)	2629(13)	7687(10)	18(3)
H(3A)	6191(13)	5112(14)	8255(10)	19(3)
H(3B)	6263(12)	3407(14)	9218(11)	19(3)
H(5A)	2155(13)	2052(13)	6656(10)	16(3)
H(5B)	3458(12)	2861(13)	5863(10)	13(3)
H(6A)	4038(12)	3843(13)	9990(10)	16(3)
H(6B)	3868(13)	1889(14)	9408(11)	26(3)
H(7A)	898(12)	2361(13)	8392(10)	15(3)
H(7B)	1405(13)	2780(15)	10024(11)	28(3)
H(8A)	1609(12)	5927(15)	9680(10)	15(2)
H(8B)	-81(13)	5306(12)	8868(10)	12(3)
H(10)	3709(12)	6580(12)	8147(10)	10(3)
H(11A)	2608(12)	5509(14)	5287(10)	14(3)
H(11B)	2749(12)	7663(13)	5842(9)	14(3)
H(12A)	25(13)	3944(16)	6195(11)	27(3)
H(12B)	99(13)	5811(16)	5453(10)	20(3)
H(12C)	-1080(15)	5566(15)	6397(10)	27(3)
H(13A)	-115(13)	8302(14)	7776(10)	16(3)
H(13B)	990(13)	8688(14)	6789(11)	26(3)
H(13C)	1749(13)	8684(14)	8399(11)	22(3)

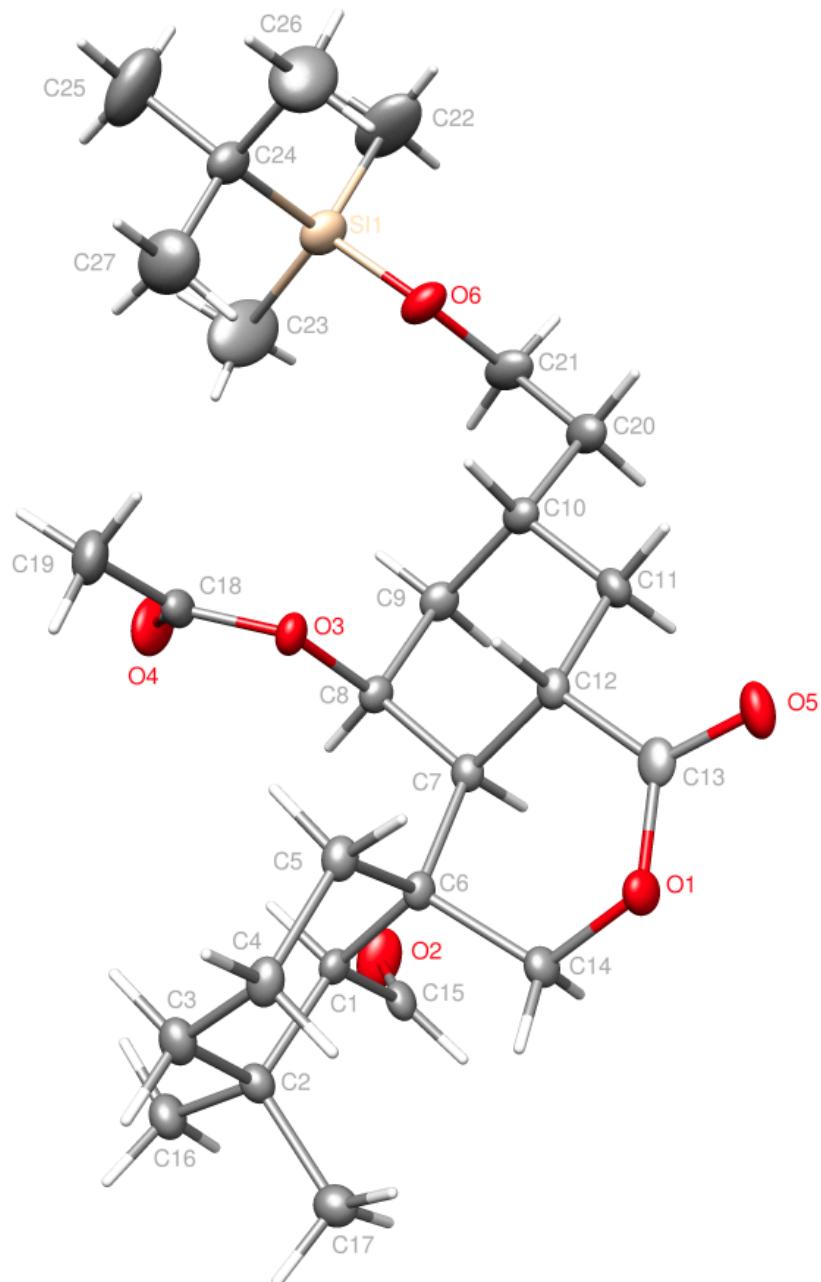
**Table A2.6.** Hydrogen bonds for spirolactone **109** (CCDC 837088) [Å and °].

D-H...A	d(D-H)	d(H...A)	d(D...A)	$\angle$ (DHA)
O(3)-H(3)...O(1)#1	0.780(12)	2.037(12)	2.8100(10)	171.5(13)

Symmetry transformations used to generate equivalent atoms:  
#1 x,y+1,z

## A2.2 CRYSTAL STRUCTURE ANALYSIS OF ALDEHYDE 127

**Figure A2.2.** Aldehyde 127. Crystallographic data have been deposited at the CCDC, 12 Union Road, Cambridge CB2 1EZ, UK and copies can be obtained on request, free of charge, by quoting the publication citation and the deposition number 830951.



**Table A2.7.** Crystal data and structure refinement for aldehyde **127** (CCDC 830951).

Empirical formula	C <sub>27</sub> H <sub>46</sub> O <sub>6</sub> Si	
Formula weight	494.73	
Crystallization Solvent	Ether	
Crystal Habit	Plate	
Crystal size	0.40 x 0.14 x 0.05 mm <sup>3</sup>	
Crystal color	Colorless	

## Data Collection

Type of diffractometer	Bruker KAPPA APEX II		
Wavelength	0.71073 Å MoKα		
Data Collection Temperature	100(2) K		
θ range for 9883 reflections used in lattice determination	2.56 to 25.95°		
Unit cell dimensions	a = 7.0008(2) Å	α = 90°	
	b = 14.2170(5) Å	β = 90°	
	c = 28.7340(9) Å	γ = 90°	
Volume	2859.91(16) Å <sup>3</sup>		
Z	4		
Crystal system	Orthorhombic		
Space group	P 2 <sub>1</sub> 2 <sub>1</sub> 2 <sub>1</sub>		
Density (calculated)	1.149 Mg/m <sup>3</sup>		
F(000)	1080		
Data collection program	Bruker APEX2 v2009.7-0		
θ range for data collection	2.02 to 30.56°		
Completeness to θ = 30.56°	99.8 %		
Index ranges	-10 ≤ h ≤ 6, -20 ≤ k ≤ 20, -41 ≤ l ≤ 39		
Data collection scan type	ω scans; 9 settings		
Data reduction program	Bruker SAINT-Plus v7.66A		
Reflections collected	62150		
Independent reflections	8741 [R <sub>int</sub> = 0.0589]		
Absorption coefficient	0.118 mm <sup>-1</sup>		
Absorption correction	Semi-empirical from equivalents		
Max. and min. transmission	0.7461 and 0.6633		

**Table A2.7 (cont.)****Structure solution and Refinement**

Structure solution program	SHELXS-97 (Sheldrick, 2008)
Primary solution method	Direct methods
Secondary solution method	Difference Fourier map
Hydrogen placement	Geometric positions
Structure refinement program	SHELXL-97 (Sheldrick, 2008)
Refinement method	Full matrix least-squares on $F^2$
Data / restraints / parameters	8741 / 0 / 315
Treatment of hydrogen atoms	Riding
Goodness-of-fit on $F^2$	1.797
Final R indices [ $I > 2\sigma(I)$ , 7186 reflections]	$R_1 = 0.0453$ , $wR_2 = 0.0579$
R indices (all data)	$R_1 = 0.0634$ , $wR_2 = 0.0594$
Type of weighting scheme used	Sigma
Weighting scheme used	$w = 1/\sigma^2(F_{\text{o}}^2)$
Max shift/error	0.001
Average shift/error	0.000
Absolute structure determination	Unknown
Absolute structure parameter	0.07(7)
Largest diff. peak and hole	0.592 and -0.387 e. $\text{\AA}^{-3}$

**Special Refinement Details**

Crystals were mounted on a glass fiber using Paratone oil then placed on the diffractometer under a nitrogen stream at 100K.

Refinement of  $F^2$  against ALL reflections. The weighted R-factor ( $wR$ ) and goodness of fit (S) are based on  $F^2$ , conventional R-factors (R) are based on F, with F set to zero for negative  $F^2$ . The threshold expression of  $F^2 > 2\sigma(F^2)$  is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on  $F^2$  are statistically about twice as large as those based on F, and R-factors based on ALL data will be even larger.

All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds

involving l.s. planes.

**Table A2.8.** Atomic coordinates ( $\times 10^4$ ) and equivalent isotropic displacement parameters ( $\text{\AA}^2 \times 10^3$ ) for aldehyde **127** (CCDC 830951). U(eq) is defined as the trace of the orthogonalized  $U^{ij}$  tensor.

	x	y	z	U <sub>eq</sub>
Si(1)	14159(1)	5918(1)	7588(1)	29(1)
O(1)	5263(1)	6642(1)	9945(1)	20(1)
O(2)	11944(1)	5324(1)	10481(1)	33(1)
O(3)	9511(1)	4778(1)	9005(1)	19(1)
O(4)	12207(1)	3915(1)	8990(1)	38(1)
O(5)	4978(1)	7610(1)	9361(1)	23(1)
O(6)	12638(1)	6504(1)	7909(1)	27(1)
C(1)	8905(2)	4690(1)	10257(1)	15(1)
C(2)	7988(2)	3911(1)	10569(1)	18(1)
C(3)	6711(2)	3307(1)	10254(1)	21(1)
C(4)	5222(2)	3885(1)	9995(1)	20(1)
C(5)	6176(2)	4609(1)	9679(1)	18(1)
C(6)	7543(2)	5282(1)	9939(1)	14(1)
C(7)	8769(2)	5904(1)	9609(1)	15(1)
C(8)	10377(2)	5446(1)	9330(1)	16(1)
C(9)	11508(2)	6173(1)	9062(1)	20(1)
C(10)	10287(2)	6773(1)	8735(1)	19(1)
C(11)	8627(2)	7207(1)	9004(1)	19(1)
C(12)	7490(2)	6485(1)	9285(1)	16(1)
C(13)	5864(2)	6962(1)	9531(1)	18(1)
C(14)	6387(2)	5989(1)	10226(1)	18(1)
C(15)	10245(2)	5291(1)	10541(1)	18(1)
C(16)	9628(2)	3300(1)	10760(1)	24(1)
C(17)	6838(2)	4271(1)	10987(1)	23(1)
C(18)	10579(2)	4026(1)	8878(1)	24(1)
C(19)	9408(2)	3359(1)	8593(1)	34(1)
C(20)	11434(2)	7545(1)	8496(1)	25(1)
C(21)	13182(2)	7208(1)	8238(1)	29(1)
C(22)	15483(3)	6733(1)	7199(1)	63(1)
C(23)	15881(2)	5281(2)	7969(1)	68(1)
C(24)	12623(2)	5082(1)	7256(1)	28(1)
C(25)	13809(3)	4506(1)	6914(1)	66(1)
C(26)	11102(3)	5619(1)	6984(1)	62(1)
C(27)	11605(2)	4425(1)	7594(1)	56(1)

**Table A2.9.** Bond lengths [Å] and angles [°] for aldehyde **127** (CCDC 830951).

Si(1)-O(6)	1.6355(9)	C(15)-C(1)-C(6)	112.88(10)
Si(1)-C(22)	1.8584(17)	C(2)-C(1)-C(6)	117.75(10)
Si(1)-C(23)	1.8635(17)	C(17)-C(2)-C(3)	110.03(10)
Si(1)-C(24)	1.8647(15)	C(17)-C(2)-C(16)	107.40(10)
O(1)-C(13)	1.3433(14)	C(3)-C(2)-C(16)	109.25(10)
O(1)-C(14)	1.4589(13)	C(17)-C(2)-C(1)	115.34(10)
O(2)-C(15)	1.2024(13)	C(3)-C(2)-C(1)	107.34(10)
O(3)-C(18)	1.3539(15)	C(16)-C(2)-C(1)	107.34(10)
O(3)-C(8)	1.4648(14)	C(4)-C(3)-C(2)	112.66(11)
O(4)-C(18)	1.1946(15)	C(3)-C(4)-C(5)	110.86(10)
O(5)-C(13)	1.2125(14)	C(4)-C(5)-C(6)	113.57(10)
O(6)-C(21)	1.4286(15)	C(14)-C(6)-C(5)	109.84(9)
C(1)-C(15)	1.5095(17)	C(14)-C(6)-C(7)	104.33(10)
C(1)-C(2)	1.5634(17)	C(5)-C(6)-C(7)	113.46(10)
C(1)-C(6)	1.5654(16)	C(14)-C(6)-C(1)	111.19(10)
C(2)-C(17)	1.5346(16)	C(5)-C(6)-C(1)	109.00(10)
C(2)-C(3)	1.5335(17)	C(7)-C(6)-C(1)	108.98(9)
C(2)-C(16)	1.5411(17)	C(8)-C(7)-C(12)	109.97(9)
C(3)-C(4)	1.5223(17)	C(8)-C(7)-C(6)	119.00(11)
C(4)-C(5)	1.5269(17)	C(12)-C(7)-C(6)	110.76(9)
C(5)-C(6)	1.5472(16)	O(3)-C(8)-C(9)	109.54(9)
C(6)-C(14)	1.5310(16)	O(3)-C(8)-C(7)	107.82(9)
C(6)-C(7)	1.5545(16)	C(9)-C(8)-C(7)	111.20(10)
C(7)-C(8)	1.5272(16)	C(8)-C(9)-C(10)	113.64(10)
C(7)-C(12)	1.5333(16)	C(20)-C(10)-C(9)	112.59(10)
C(8)-C(9)	1.5131(17)	C(20)-C(10)-C(11)	109.77(11)
C(9)-C(10)	1.5287(16)	C(9)-C(10)-C(11)	109.88(10)
C(10)-C(20)	1.5253(17)	C(12)-C(11)-C(10)	113.04(10)
C(10)-C(11)	1.5266(16)	C(13)-C(12)-C(11)	109.82(10)
C(11)-C(12)	1.5286(17)	C(13)-C(12)-C(7)	113.61(10)
C(12)-C(13)	1.5021(17)	C(11)-C(12)-C(7)	112.23(9)
C(18)-C(19)	1.4975(19)	O(5)-C(13)-O(1)	117.02(11)
C(20)-C(21)	1.5089(18)	O(5)-C(13)-C(12)	122.82(12)
C(24)-C(27)	1.523(2)	O(1)-C(13)-C(12)	120.06(11)
C(24)-C(26)	1.526(2)	O(1)-C(14)-C(6)	113.88(9)
C(24)-C(25)	1.525(2)	O(2)-C(15)-C(1)	124.07(12)
		O(4)-C(18)-O(3)	124.03(13)
O(6)-Si(1)-C(22)	110.23(7)	O(4)-C(18)-C(19)	125.78(13)
O(6)-Si(1)-C(23)	109.73(7)	O(3)-C(18)-C(19)	110.17(12)
C(22)-Si(1)-C(23)	109.49(9)	C(21)-C(20)-C(10)	114.84(11)
O(6)-Si(1)-C(24)	103.71(6)	O(6)-C(21)-C(20)	109.37(10)
C(22)-Si(1)-C(24)	112.19(7)	C(27)-C(24)-C(26)	107.83(13)
C(23)-Si(1)-C(24)	111.36(8)	C(27)-C(24)-C(25)	109.58(14)
C(13)-O(1)-C(14)	122.50(9)	C(26)-C(24)-C(25)	108.56(13)
C(18)-O(3)-C(8)	117.04(9)	C(27)-C(24)-Si(1)	109.61(10)
C(21)-O(6)-Si(1)	123.77(8)	C(26)-C(24)-Si(1)	110.22(11)
C(15)-C(1)-C(2)	110.26(9)	C(25)-C(24)-Si(1)	110.97(11)

**Table A2.10.** Anisotropic displacement parameters ( $\text{\AA}^2 \times 10^4$ ) for aldehyde **127** (CCDC 830951). The anisotropic displacement factor exponent takes the form:  
 $-2p^2 [ h^2 a^{*2} U^{11} + \dots + 2 h k a^* b^* U^{12} ]$

	U <sup>11</sup>	U <sup>22</sup>	U <sup>33</sup>	U <sup>23</sup>	U <sup>13</sup>	U <sup>12</sup>
Si(1)	247(2)	349(3)	278(2)	-58(2)	22(2)	4(2)
O(1)	170(5)	187(5)	247(5)	9(4)	21(4)	55(4)
O(2)	152(5)	329(6)	522(7)	-116(5)	-16(5)	-9(4)
O(3)	197(5)	142(5)	223(5)	-44(4)	13(4)	13(4)
O(4)	351(6)	327(7)	466(6)	-82(5)	-71(5)	188(5)
O(5)	211(5)	137(5)	352(5)	8(4)	-52(4)	25(4)
O(6)	278(5)	303(6)	223(5)	-96(5)	37(4)	-44(4)
C(1)	150(6)	133(6)	166(6)	-20(6)	22(5)	13(5)
C(2)	190(7)	141(7)	204(7)	16(6)	6(6)	-21(5)
C(3)	222(7)	148(7)	274(7)	5(6)	22(6)	-32(6)
C(4)	177(7)	150(7)	266(7)	-43(6)	14(6)	-36(5)
C(5)	160(7)	155(7)	216(7)	-26(6)	-17(5)	-1(5)
C(6)	122(6)	132(7)	180(6)	-14(6)	-2(5)	9(5)
C(7)	137(6)	129(6)	176(6)	-15(6)	-11(5)	-17(5)
C(8)	146(6)	166(7)	180(6)	-21(6)	6(5)	6(5)
C(9)	159(7)	215(8)	214(7)	-27(6)	15(6)	-25(5)
C(10)	231(7)	153(7)	177(6)	-25(6)	20(6)	-44(6)
C(11)	249(7)	133(7)	199(7)	18(6)	-22(6)	-5(5)
C(12)	153(6)	121(7)	192(6)	-29(6)	-23(6)	-14(5)
C(13)	143(7)	146(7)	254(7)	-48(6)	-59(6)	-34(6)
C(14)	167(7)	158(7)	200(7)	17(6)	-4(5)	23(6)
C(15)	188(7)	127(7)	215(7)	22(6)	-26(6)	14(6)
C(16)	286(8)	174(7)	264(7)	11(6)	-12(6)	-3(6)
C(17)	258(7)	213(8)	227(7)	3(6)	26(6)	-44(6)
C(18)	333(9)	164(8)	207(7)	16(6)	41(6)	64(6)
C(19)	502(10)	177(8)	340(8)	-71(7)	15(8)	13(7)
C(20)	355(8)	187(8)	221(7)	-19(6)	30(6)	-62(6)
C(21)	356(9)	283(9)	239(8)	-24(7)	79(7)	-139(7)
C(22)	701(13)	639(13)	554(11)	-180(10)	350(10)	-266(11)
C(23)	465(11)	926(16)	653(12)	-139(12)	-236(10)	253(12)
C(24)	328(8)	275(9)	231(7)	-46(7)	-3(7)	46(7)
C(25)	713(13)	542(14)	728(13)	-340(11)	146(11)	-5(11)
C(26)	682(12)	647(15)	547(11)	-108(10)	-309(10)	67(11)
C(27)	716(12)	432(12)	530(11)	-51(9)	30(10)	-201(9)