Table	2.	Main	interatomic	bond	distances	(Å)	and	
angles (°) with e.s.d.'s in parentheses								

PO₄ tetrahedron								
P	0(1)		O(2)	0(3)		0(4)		
$\frac{1}{0}$	1.493 (1	۱ I	12.3 (1)	106.5 (n 1	15.6 (1)		
$\tilde{O}(2)$	2.534 (2	ς i	.557 (2)	106.4 ($\frac{1}{10}$ $\frac{1}{10}$	1.0 (1)		
	2,450 (2	(501 (2)	1.566 ($\frac{1}{2}$	/4·9 (1)		
	2,524 (2		AD5 (2)	1.500 (2 1	502 (1)		
0(4)	2.334 (2) 2	•425 (2)	2.527 (2) <u>1</u>	<u>502 (1)</u>		
	F	P—P	4.7195 (1	7)				
C₄H ₁₂ NO tetrahedral group								
C(2)	C(1)		Č(3)	C(4)		N		
C (1)	1.515 (3) 1	08·9 (2)	111.6 (2) 1()8·0 (2)		
C(3)	2.459 (3	วั่า	·508 (3)	112.3	2) 10	18.1(2)		
Č(4)	2.502 (3	ý <u>ž</u>	·507 (3)	1.510 (3) 10	$\frac{1}{17.7}$ (2)		
Ň	2.435 (3	$\hat{1}$.429 (3)	2.425 (3 1 1.	494 (2)		
• •	(.	, -	, (5)	2 125 (<u>+)+(2)</u> .		
C(3)—O	1.419 (3)		C(2)C(3	i)—O 11	l0·6 (2)		
Hydrogen bo	onds							
O(N)—H…O) ()(N)—H	H H…	0 0(N)	—н—о	0(N)-0		
O(2)—H(2)…C	0(4)	0.81 (3)	1.78	(3)	75 (3)	2.586 (2)		
O(3)-H(3)-C)	0.65 (2)	1.98	(2) 1	77 (3)	2.635 (2)		
O-H…O(1)		0.75 (2)	1.93	(2) 1	73 (2)	2.673 (2)		
O(W) - H(W)	····O(4)	0.76 (2)	2.12	(2) 1	72 (3)	2.868 (2)		
O(W) - H(2W)	••• O (1)	0.74 (2)	2.07	(2) 1	67 (2)	2.797 (2)		
N—Ĥ(1N)…Ó	(4)		2.08	(2) 1	62 (2)	2.905 (2)		
	()	0.85(2)			(-)	2,00 (2)		
N—H(1N)…O	(2)	(-)	2.53	(2) 1	27 (2)	3.120 (2)		
N-H(2N)O	ά		2.08	(3) 1	42 (2)	2.865 (2)		
	(-)	0.92(3)		()	(_)	2 005 (2)		
NH(2N)O			2.53	(3) 1	21 (1)	2.776 (2)		
N-H(3N)O	(W)	0.89 (3)	1.89	(3) 1	76 (2)	2.773(2)		
	` '				- (-)			
	H (1И	V)—O(W)—H(2 <i>W</i>) 105	(3)			

of tris(ethylenediammonium) bis(*cyclo*-triphosphate), [NH₃(CH₂)₂NH₃]₃(P₃O₉)₂ (Averbuch-Pouchot, Durif & Guitel, 1989). One quasi-linear hydrogen bond joins N to the water molecule, another one joins O to the PO₄ tetrahedron.

So the succession of $C_4H_{12}NO^+$ and O(W) ribbons and $H_2PO_4^-$ chains is stabilized by a threedimensional network of hydrogen bonds and geometrical details of this hydrogen-bond scheme are listed in Table 2. Some characteristic features should be noted: there is no direct bonding between the different organic groups in a ribbon and no bonding at all between the ribbons or chains related by centrosymmetry.

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Structure of N-(3,6-Dimethoxy-17-methyl-4,5-epoxy-6,14 α -etheno-7 α -isomorphinanylcarbonyl)-L-phenylalanine Ethyl Ester Hydrochloride

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Abstract. $C_{33}H_{39}N_2O_6^+.Cl^-$, $M_r = 595 \cdot 13$, orthorhombic, $P2_12_12_1$, a = 10.678 (1), b = 11.365 (1), c = 24.664 (2) Å, V = 2993.1 (5) Å³, Z = 4, $D_x = 1.321$ g cm⁻³, λ (Mo K α) = 0.71073 Å, $\mu = 1.7$ cm⁻¹, F(000) = 1264, T = 295 K, R = 0.052 for 2450 observed reflections with $I > 2.5\sigma(I)$. The phenylalanine residue is found to have no significant effect on the geometry of the ethenoisomorphinan skeleton. The amide H is intramolecularly hydrogen 0108-2701/90/122384-04\$03.00

bonded to the carbonyl O of the peptide moiety and not to the methoxy ether O. This results in an orientation of the peptide side chain more or less parallel to the piperidine moiety.

Introducton. The interaction of opioids with the opioid receptor has still not been clarified, inspite of several valuable receptor models (Kolb, 1987; and references cited therein). Conformational analysis in © 1990 International Union of Crystallography

combination with pharmacological results is a helpful tool in the search for new information. Therefore, we synthesized new morphinan peptides [(1)-(4), Beverman, Lie, Maat & Noordam-Weissdorf, 1982; (5)-(7), Cappon, Lie & Maat, 1990] based on the similarities between morphinans and endogenous opioid peptides, such as enkephalin, and the message-address concept (Chavkin & Goldstein. 1981), which suggests that the N-terminal amino acids in opioid peptides are responsible for the pharmacological stimulance and that some C-terminal amino acids are of importance for the receptor affinity. In these morphinan peptides we have coupled an ethenoisomorphinan-7-carboxylic acid as the message moiety to several enkephalin residues as the address moiety. Some of these compounds appear to



(1*a,b*) $R_2 = Leu - OEt$ (2*a,b*) $R_2 = Phe - Leu - OEt$ (3*a,b*) $R_2 = Gly - Phe - Leu - OEt$ (4*a,b*) $R_2 = Phe - Leucinol$ $(5a,b) R_2 = Phe - OEt$ $(6a,b) R_2 = D - Phe - OEt$ $(7a,b) R_2 = Gly - Phe = OEt$ $a R_1 = OCH_3 b R_1 = OH$

be potent analgesics with high receptor affinity (Smith, Medzihradsky & Woods, 1986). For a detailed interpretation of the structure-activity relationships we have carried out conformational analyses on parts of the molecules using molecular mechanics (Cappon, Lie & Maat, 1990). The X-ray analysis of the title compound (5*a*) is the first crystal structure reported in the field of morphinan peptides. Points of interest were the conformation of the skeleton, the influence of the peptide substituent on the geometry and the orientation of this substituent. Intramolecular hydrogen bonding is thought to play an important role in the orientation.

Experimental. The title compound has been synthesized from the Diels–Alder adduct of thebaine with ethyl acrylate and the ethyl ester of L-phenylalanine, and was crystallized as its hydrochloride salt from ethanol.



Data were collected on an Enraf–Nonius CAD-4F diffractometer for a colourless transparent crystal $(0.22 \times 0.30 \times 0.42 \text{ mm})$ mounted on a glass fibre. Unit-cell parameters and their e.s.d.'s were derived

from a least-squares treatment of 25 SET4 reflections $(10 < \theta < 14^{\circ})$. Intensity data for 4660 reflections $(h - 13:0, k - 14:0, l : 0:32; \theta < 27.5^{\circ};$ Zr-filtered Mo K α radiation) were collected in the $\omega/2\theta$ scan mode with $\Delta \omega = (0.60 + 0.35 \tan \theta)^{\circ}$. Three reference reflections ($0\overline{2}4$, $\overline{1}\overline{2}0$ and $\overline{1}01$) indicated no decay during 70 h of X-ray exposure. The intensity data were corrected for Lp but not for absorption. The variance $\sigma^2(I)$ was calculated based on counting statistics plus an instability constant term $(0.023I)^2$ as derived from the excess variance in the reference reflections (McCandlish, Stout & Andrews, 1975). The space group was derived from the observed systematic absences. The structure was solved by direct methods (SHELXS86; Sheldrick, 1986) and refined on F by full-matrix least squares with SHELX76 (Sheldrick, 1976). The two H atoms bonded to N atoms were located in a difference Fourier map and their positions refined. All other H atoms were introduced at calculated positions (C-H = 0.95 Å) and refined with fixed geometry with respect to their carrier atoms with two common isotropic thermal parameters. Refinement with weights, $w^{-1} = \sigma^2(F) + 0.0005F^2$, converged at R = $0.052 \ [wR = 0.057; S = 1.11; 399 \text{ parameters}; 2450$ reflections, $(\Delta/\sigma)_{\text{max}} = 0.5$]. A final difference Fourier map did not show residual peaks outside -0.30 and 0.36 e Å⁻³. Scattering factors from Cromer & Mann (1968) and anomalous-dispersion terms from Cromer & Liberman (1970) were used. Final parameters are given in Table 1.* The programs PLATON and PLUTON (Spek, 1982) were used for the calculation of geometrical data and the plot respectively. All calculations were made on a microVAX-II.

Discussion. Fig. 1 shows the molecule with adopted numbering. Data on the geometry have been assembled in Table 2. The Cl anion is H bonded to the protonated N(17) $[N(17)-H(17) 0.97 (5), H(17)\cdots Cl 2.12 (5) \text{ Å}, N(17)-H(17)\cdots Cl 154 (4)^{\circ}]$. The conformation of the ethenoisomorphinan skeleton is very similar to that observed in the X-ray structure of 3-methoxyetorphine (van den Hende & Nelson, 1967).



* Lists of structure factors, anisotropic thermal parameters and H-atom parameters have been deposited with the British Library Document Supply Centre as Supplementary Publication No. SUP 53086 (21 pp.). Copies may be obtained through The Technical Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

Table 1. Final coordinates and equivalent isotropic Table 2. Bond distances (Å) and bond angles (°) for thermal parameters and their e.s.d.'s in parentheses

non-H atoms

	$U_{\rm eq} = (1/3) \sum_i \sum_j U_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j.$				O(1)C(3)	1.363 (7)	C(7)—C(8) 1	553 (6)
			· · ·	XI (\$ 2)	O(1)C(22)	1.435 (8)	C(7) - C(20) 1	506 (6)
	x	у	Z	$U_{eq}(A^{-})$	O(2) - C(4)	1.376 (6)	C(8) - C(14) 1	555 (6)
Cl	0.2833 (1)	0.2333(1)	-0.06108(7)	0.0607 (5)	O(2) = O(3)	1.403 (0)	C(9) = C(10) 1	540 (7)
O(1)	0.0181 (4)	0.7029 (4)	0.2413(1)	0.055(2)	O(3) - C(0)	1.414 (7)	C(9) = C(14) 1	517 (7)
0(2)	-0.0281 (3)	0.5807(3)	0.1430 (1)	0.037(1)	O(3) - C(23) O(4) - C(20)	1.236 (6)	C(1) - C(12) = 1	365 (7)
0(3)	-0.1554(3)	0.6003(3)	-0.0810(1)	0.040(1)	O(5) - C(32)	1.199 (8)	C(12) - C(13) = 1	488 (6)
0(4)	-0.3973 (4)	0.4488 (5)	-0.0888(2)	0.045(1)	O(6) - C(32)	1.313 (8)	C(13)-C(14) 1	551 (6)
0(6)	-0.4380 (4)	0.5539 (4)	-0.1624(2)	0.067(2)	O(6)-C(33)	1.474 (8)	C(13)-C(15) 1	530 (6)
N(1)	-0.1706(4)	0.5330(4)	-0.0775(2)	0.038(1)	N(1)-C(20)	1.336 (6)	C(14)—C(18) 1	508 (6)
N(17)	0-3469 (3)	0.3762 (3)	0.0383 (2)	0.031 (1)	N(1)-C(24)	1.457 (7)	C(15)—C(16) 1	500 (7)
C(1)	0.3359 (5)	0.6361 (5)	0.1904 (2)	0.046 (2)	N(17)—C(9)	1.516 (6)	C(18)—C(19) 1	315 (6)
C(2)	0.2413 (5)	0.6725 (5)	0.2249 (2)	0.048 (2)	N(17)—C(16)	1.499 (6)	C(24)—C(25) 1	521 (7)
C(3)	0.1167 (5)	0.6578 (5)	0.2130 (2)	0.041 (2)	N(17) - C(21)	1.493 (5)	C(24) - C(32) = 1	504 (8)
C(4)	0.0881 (5)	0.5937 (4)	0.1670 (2)	0.034 (2)	C(1) - C(2)	1.384 (7)	C(25) - C(26) = 1	504 (8)
C(5)	-0.0112(4)	0.5268 (4)	0.0903 (2)	0.030 (1)	$C(1) \rightarrow C(1)$	1.412 (7)	C(26) - C(21) = 1	300 (0)
C(6)	-0.0351 (4)	0.6125 (4)	0.0430(2)	0.031(1)	C(2) - C(3)	1.382 (7)	C(20) - C(31) = 1	37 (1)
C(7)	-0.0121(4)	0.5311 (4)	-0.0083(2) -0.0155(2)	0.029(1)	C(4) - C(12)	1.390 (7)	C(28) - C(29) = 1	38 (1)
	0.3352 (4)	0.5087 (4)	0.0435(2)	0.030(1)	C(5) - C(6)	1.539 (6)	C(29) - C(30) = 1	37 (1)
CUM	0.3956 (4)	0.5524 (5)	0.0965(2)	0.035(2)	C(5)-C(13)	1.556 (6)	C(30)—C(31) 1	39 (1)
C(11)	0-3060 (5)	0.5720 (4)	0.1431(2)	0.033 (2)	C(6)-C(7)	1.587 (6)	C(33)—C(34) 1	43 (1)
C(12)	0.1827 (4)	0.5447 (4)	0.1355 (2)	0.030 (2)	C(6)-C(19)	1.497 (6)		
C(13)	0.1281 (4)	0.4874 (4)	0.0867 (2)	0.027 (1)				
C(14)	0.1973 (4)	0-5436 (4)	0.0378 (2)	0.027 (1)	$C(3) \rightarrow O(1) \rightarrow C(22)$	116.7 (5)	C(4) - C(12) - C(13)	110-0 (4)
C(15)	0.1425 (4)	0.3535 (4)	0.0849 (2)	0.031 (1)	C(4) - O(2) - C(5)	108.1 (3)	C(11) - C(12) - C(13)	126.0 (4)
C(16)	0.2767 (5)	0.3147(4)	0.0827 (2)	0.037(2)	C(0) = O(3) = C(23)	118.3 (4)	C(5) - C(13) - C(12)	101-7 (3)
C(18)	0.0606(4)	0.7097 (4)	0.0409(2)	0.030 (1)	C(32) = O(0) = C(33) C(20) = N(1) = C(24)	123.8 (4)	C(5) - C(13) - C(15)	112.4 (4)
C(19)	-0.0675 (4)	0.5853 (4)	-0.0586(2)	0.031(1) 0.032(2)	C(9) - N(17) - C(16)	$111 \cdot 1$ (3)	C(12) - C(13) - C(14)	105.2 (4)
C(20)	0.4798 (4)	0.3359 (5)	0.0350(2)	0.032(2)	C(9) - N(17) - C(21)	112.8 (3)	C(12) - C(13) - C(15)	114.8 (4)
C(22)	0.0465 (8)	0.7922 (6)	0.2806 (3)	0.074(3)	C(16)-N(17)-C(21) 111.9 (4)	C(14)-C(13)-C(15)	109-8 (4)
C(23)	-0.2590 (5)	0.6004 (6)	0.0611 (3)	0.060 (2)	C(2) - C(1) - C(11)	119.8 (5)	C(8)C(14)-C(9)	116-9 (4)
C(24)	-0.2362 (5)	0-5703 (5)	-0.1264 (2)	0.041 (2)	C(1)-C(2)-C(3)	122.6 (5)	C(8)-C(14)-C(13)	109-3 (3)
C(25)	-0.1670 (6)	0.5429 (5)	-0.1788 (2)	0.051 (2)	O(1) - C(3) - C(2)	126.3 (5)	C(8) - C(14) - C(18)	104.0 (4)
C(26)	-0.1265 (6)	0.4175 (5)	-0.1872 (2)	0.051 (2)	O(1) - C(3) - C(4)	116.7 (5)	C(9) - C(14) - C(13)	106-3 (4)
C(27)	-0.0122 (6)	0.3764 (6)	-0.1667 (2)	0.057 (2)	C(2) - C(3) - C(4) C(2) - C(4) - C(3)	11/0(5)	C(9) - C(14) - C(18)	112.4 (4)
C(28)	0.0299 (7)	0.2640 (7)	-0.1/58(3)	0.076(3)	O(2) - C(4) - C(3)	120.8 (3)	C(13) - C(13) - C(16)	107.7 (4)
C(29)	-0.1536(9)	0.1397(7)	-0.2075(3) -0.2285(3)	0.085(3)	C(3) - C(4) - C(12)	120.6 (5)	N(17) - C(16) - C(15)	111.5 (4)
C(30)	-0.1966(7)	0.3308 (6)	-0.2184(3)	0.068 (3)	O(2) - C(5) - C(6)	113.3 (4)	C(14) - C(18) - C(19)	114.8 (4)
C(32)	-0.3645 (6)	0.5161 (5)	-0.1234(3)	0.047(2)	O(2) - C(5) - C(13)	106.8 (3)	C(6)-C(19)-C(18)	116-1 (4)
C(33)	-0.5698 (6)	0.5155(7)	-0.1632(3)	0.079(3)	C(6) - C(5) - C(13)	107.3 (3)	O(4)-C(20)-N(1)	121.9 (4)
C(34)	-0.6501 (8)	0.6034 (9)	-0.1410 (4)	0.111 (4)	O(3)-C(6)-C(5)	115.5 (4)	O(4)-C(20)-C(7)	122.6 (4)
. ,		.,		.,	O(3)-C(6)-C(7)	112.4 (3)	N(1)-C(20)-C(7)	115-5 (4)
					O(3) - C(6) - C(19)	107.6 (4)	N(1)-C(24)-C(25)	114.2 (4)
					C(5) - C(6) - C(7)	102-1 (3)	N(1) - C(24) - C(32)	106-1 (5)
	0.0	~			C(3) - C(6) - C(19)	110.7 (4)	C(23) - C(24) - C(32)	115.0 (5)
	- AP	()	0		C(7) - C(8) - C(19)	110-1 (3)	C(24) - C(25) - C(20)	121.1 (5)
	C21	\sum_{α}	Q L C29		C(0) - C(7) - C(3)	111.0 (4)	C(25) - C(26) - C(21)	121.6 (6)
	NIZ-	//	(E)		C(8) - C(7) - C(20)	$111 \cdot 1 (4)$	C(27) - C(26) - C(31)	117.2 (6)
			C28 The West	Ð	C(7) - C(8) - C(14)	108.0 (3)	C(26) - C(27) - C(28)	122.4 (6)
	Store Falls		al M	C 30	N(17)-C(9)-C(10)	110.9 (4)	C(27)-C(28)-C(29)	118.7 (7)
	C10 C	Ω.	C27)	N(17)-C(9)-C(14)	109.2 (3)	C(28)-C(29)-C(30)	120.9 (7)
		Lie Co	(26) 11 (3)	[≮] ≻	C(10)-C(9)-C(14)	113.4 (4)	C(29)-C(30)-C(31)	120-1 (7)
	CII CIS			~	C(9)-C(10)-C(11)	115.3 (4)	C(26) - C(31) - C(30)	120.7 (7)
	Carlo Altin	All CIO LA C7	_04	Ð	C(1) - C(11) - C(10)	124-1 (5)	O(5) - C(32) - O(6)	123.7 (6)
					$C(1) \rightarrow C(1) \rightarrow C(12)$	110.0 (3)	O(5) - C(32) - C(24)	124.2 (6)
	C12	C17 C20	C24		C(4) = C(1) = C(1)	122.2 (4)	O(6) - C(32) - C(24)	111.2 (3)
	<u>°Щ Щ бо</u>	506	N. M.			122 2 (4)	O(0) O(33) O(34)	
		^{co} M		05				
		1 03	05 132					
	III C3 02	AR "	1	R.J.	ceivable int	amalagular	hydrogen bonds	around

Fig. 1. View of the molecule with adopted numbering.

This means that the influence of the phenylalanine substituent on the skeleton geometry is negligible. The conformation within the phenylalanine ethyl ester is rather flexible (Wei, Doherty & Einstein, 1972; Yamashita, Kato, Yamane & Ashida, 1982; Ishida, Tanabe & Inoue, 1983). Of the two conceivable intramolecular hydrogen bonds around the peptide bond, the one between the amide H and the carbonyl O of the phenylalanine ethyl ester residue [N(1)-H(10) 0.94(5), H(10)-O(5) 2.13(5) Å,N(1)-H(10)...O(5) 111 (4)°] is formed. This is of interest because formation of the other one, namely between the amide H and the 6-methoxy O achievable by rotation about C(7)—C(20), would dramatically alter the relative orientation of the phenylalanine and ethenoisomorphinan moieties. The present conformation is probably favoured for packing reasons. No intermolecular H bonds are found.

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Structure and Absolute Configuration of Two *ent*-Atisane Diterpenes from *Euphorbia fidjiana*

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Abstract. ent-(13S)-Hydroxyatis-16-en-3,14-dione, $C_{20}H_{28}O_3$, $M_r = 316.4$, orthorhombic, $P2_12_12_1$, a =7.335 (2), b = 12.539 (1), c = 18.325 (1) Å, V = 1685.4 Å^3 , Z = 4, $D_x = 1.247 \text{ g cm}^{-3}$, $\lambda(Mo K\alpha) =$ $0.71069 \text{ Å}, \ \mu = 0.80 \text{ cm}^{-1}, \ F(000) = 688, \ R = 0.042$ for 1626 observed data $[I > 2.5\sigma(I)]$ at room tement-16\alpha-Hydroxy-3-oxoatisan-17-yl 4perature. bromobenzoate, $C_{27}H_{35}BrO_4$, $M_r = 503.5$, orthorhombic, $P2_12_12_1$, a = 7.029 (2), b = 11.642 (3), c = $V = 2342.5 \text{ Å}^3$ 28.626 (6) Å, 1.427 g cm⁻³, Z = 4, $D_{\rm r} =$ $\mu =$ $\lambda(\text{Mo } K\alpha) = 0.71069 \text{ Å},$ 17.8 cm^{-1} , F(000) = 1056, R = 0.034 for 1801 observed data $[I > 3\sigma(I)]$ at 170 K. These compounds are new diterpenoids whose stereochemistry has been determined.

Introduction. The Euphorbiaceae family is known as a source of antileukaemic and cocarcinogenic compounds (Kupchan, Uchida, Branfman, Dailey & Fei, 1976). Preparations of *Euphorbia fidjiana*, a Fijian plant known locally as vasa damu, have been used locally for the treatment of a variety of ailments. Extracts of the heartwood of *E. fidjiana* have yielded 15 diterpenoid compounds with atisane or seco-

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atisane skeletons (Lal, Cambie, Rutledge & Woodgate, 1990). The structure of two of these, *ent*-(13S)-hydroxyatis-16-en-3,14-dione (I) and *ent*- 16α ,17-dihydroxyatisan-3-one as the *p*-bromobenzoate (II), along with the NMR spectra establish the stereochemistry of the whole series. In addition the absolute configuration of (II) establishes the absolute stereochemistry of all of the compounds.

Experimental. Suitable single crystals were obtained from slow evaporation of hexane solutions. Crystal samples of dimensions $0.4 \times 0.3 \times 0.2$ mm (I) and 0.5 $\times 0.2 \times 0.2$ mm (II) were mounted in random orientations on an Enraf-Nonius CAD-4 diffractometer. The lattice parameters and orientation matrix were obtained from 25 reflections in the range 22 < $2\theta < 26^{\circ}$. The Laue symmetry is *mmm* and the space groups were uniquely determined by the systematic absences. The intensity data were collected using Zr-filtered Mo K α radiation (II) using the ω -2 θ scan technique. Three reflections monitored every hour showed no non-statistical variations in intensity. Unique data sets were collected for each structure. The number of reflections surveyed was 2352 in the range $2 < 2\theta < 56^{\circ}$ with *h*, *k*, *l* ranges 10, 17, 25 (I) and 2480 in the range $2 < 2\theta < 50^{\circ}$ with h, k, l ranges © 1990 International Union of Crystallography

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