Radiocarbon dating of the Shroud of Turin

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Very small samples from the Shroud of Turin have been dated by accelerator mass spectrometry in laboratories at Arizona, Oxford and Zurich. As controls, three samples whose ages had been determined independently were also dated. The results provide conclusive evidence that the linen of the Shroud of Turin is mediaeval.

THE Shroud of Turin, which many people believe was used to wrap Christ's body, bears detailed front and back images of a man who appears to have suffered whipping and crucifixion. It was first displayed at Lirey in France in the 1350s and subsequently passed into the hands of the Dukes of Savoy. After many journeys the shroud was finally brought to Turin in 1578 where, in 1694, it was placed in the Royal Chapel of Turin Cathedral in a specially designed shrine.

Photography of the shroud by Secondo Pia in 1898 indicated that the image resembled a photographic 'negative' and represents the first modern study. Subsequently the shroud was made available for scientific examination, first in 1969 and 1973 by a committee appointed by Cardinal Michele Pellegrino¹ and then again in 1978 by the Shroud of Turin Research Project (STURP)². Even for the first investigation, there was a possibility of using radiocarbon dating to determine the age of the linen from which the shroud was woven. The size of the sample then required, however, was ~500 cm², which would clearly have resulted in an unacceptable amount of damage, and it was not until the development in the 1970s of small gas-counters and accelerator-mass-spectrometry techniques (AMS), requiring samples of only a few square centimetres, that radiocarbon dating of the shroud became a real possibility.

To confirm the feasibility of dating the shroud by these methods an intercomparison, involving four AMS and two small gas-counter radiocarbon laboratories and the dating of three known-age textile samples, was coordinated by the British Museum in 1983. The results of this intercomparison are reported and discussed by Burleigh *et al.*³.

Following this intercomparison, a meeting was held in Turin in September-October 1986 at which seven radiocarbon laboratories (five AMS and two small gas-counter) recommended a protocol for dating the shroud. In October 1987, the offers from three AMS laboratories (Arizona, Oxford and Zurich) were selected by the Archbishop of Turin, Pontifical Custodian of the shroud, acting on instructions from the Holy See, owner of the shroud. At the same time, the British Museum was invited to help in the certification of the samples provided and in the statistical analysis of the results. The procedures for taking the samples and treating the results were discussed by representatives of the three chosen laboratories at a meeting at the British Museum in January 1988 and their recommendations⁴ were subsequently approved by the Archbishop of Turin.

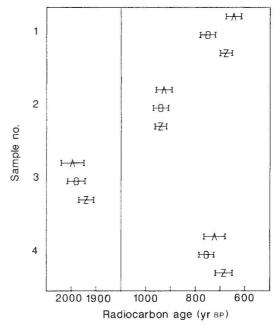


Fig. 1 Mean radiocarbon dates, with $\pm 1\sigma$ errors, of the Shroud of Turin and control samples, as supplied by the three laboratories (A, Arizona; O, Oxford; Z, Zurich) (See also Table 2.) The shroud is sample 1, and the three controls are samples 2-4. Note the break in age scale. Ages are given in yr BP (years before 1950). The age of the shroud is obtained as AD 1260-1390, with at least 95% confidence.

Removal of samples from the shroud

The sampling of the shroud took place in the Sacristy at Turin Cathedral on the morning of 21 April 1988. Among those present when the sample was cut from the shroud were Cardinal Anastasio Ballestrero (Archbishop of Turin), Professor L. Gonella (Department of Physics, Turin Polytechnic and the Archbishop's scientific adviser), two textile experts (Professor F. Testore of Department of Materials Science, Turin Polytechnic and G. Vial of Musée des Tissues and Centre International d'Étude des Textiles Anciens in Lyon), Dr M. S. Tite of the British Museum, representatives of the three radiocarbon-dating laboratories (Professor P. E. Damon, Professor D. J. Donahue, Professor E. T. Hall, Dr R. E. M. Hedges and Professor W. Woelfli) and G. Riggi, who removed the sample from the shroud.

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The shroud was separated from the backing cloth along its bottom left-hand edge and a strip (~10 mm×70 mm) was cut from just above the place where a sample was previously removed in 1973 for examination. The strip came from a single site on the main body of the shroud away from any patches or charred areas. Three samples, each ~50 mg in weight, were prepared from this strip. The samples were then taken to the adjacent Sala Capitolare where they were wrapped in aluminium foil and subsequently sealed inside numbered stainless-steel containers by the Archbishop of Turin and Dr Tite. Samples weighing 50 mg from two of the three controls were similarly packaged. The three containers containing the shroud (to be referred to as sample 1) and two control samples (samples 2 and 3) were then handed to representatives of each of the three laboratories together with a sample of the third control (sample 4), which was in the form of threads. All these operations, except for the wrapping of the samples in foil and their placing in containers, were fully documented by video film and photography.

The laboratories were not told which container held the shroud sample. Because the distinctive three-to-one herringbone twill weave of the shroud could not be matched in the controls, however, it was possible for a laboratory to identify the shroud sample. If the samples had been unravelled or shredded rather than being given to the laboratories as whole pieces of cloth, then it would have been much more difficult, but not impossible, to distinguish the shroud sample from the controls. (With unravelled or shredded samples, pretreatment cleaning would have been more difficult and wasteful.) Because the shroud had been exposed to a wide range of potential sources of contamination and because of the uniqueness of the samples available, it was decided to abandon blind-test procedures in the interests of effective sample pretreatment. But the three laboratories undertook not to compare results until after they had been transmitted to the British Museum. Also, at two laboratories (Oxford and Zurich), after combustion to gas, the samples were

recoded so that the staff making the measurements did not know the identity of the samples.

Controls

The three control samples, the approximate ages of which were made known to the laboratories, are listed below. Two were in the form of whole pieces of cloth (samples 2 and 3) and one was in the form of threads (sample 4).

Sample 2. Linen (sample Q1.T/32) from a tomb excavated at Qasr Ibrîm in Nubia by Professor J. M. Plumley for the Egypt Exploration Society in 1964. On the basis of the Islamic embroidered pattern and Christian ink inscription, this linen could be dated to the eleventh to twelfth centuries AD.

Sample 3. Linen from the collection of the Department of Egyptian Antiquities at the British Museum, associated with an early second century AD mummy of Cleopatra from Thebes (EA6707). This linen was dated in the British Museum Research Laboratory using liquid scintillation counting, giving a radiocarbon age of $2,010\pm80$ yr BP (BM-2558). This corresponds to a calendar age, rounded to the nearest 5 years, of 110 cal BC-AD 75 cal at the 68 per cent confidence level⁵ (where cal denotes calibrated radiocarbon dates).

Sample 4. Threads removed from the cope of St Louis d'Anjou which is held in a chapel in the Basilica of Saint-Maximin, Var, France. On the basis of the stylistic details and the historical evidence the cope could be dated at ~AD 1290-1310 (reign of King Phillipe IV).

Measurement procedures

Because it was not known to what degree dirt, smoke or other contaminants might affect the linen samples, all three laboratories subdivided the samples, and subjected the pieces to several different mechanical and chemical cleaning procedures.

All laboratories examined the textile samples microscopically to identify and remove any foreign material. The Oxford group

			Tal	ble 1 Basic	data (individ	ual measure	ments)		
	Sam	ple 1	Sam	ple 2	Sam	ple 3	Sam	ple 4	Pretreatment and replication codes
Arizona	AA-3367		AA-3368		AA-3369		AA-3370		
	A1.1b* A1.2b A1.3a A1.4a	591 ± 30 690 ± 35 606 ± 41 701 ± 33	A2.1b A2.2a A2.3a (1) A2.4a (2) A2.5b	922 ± 48 986 ± 56 829 ± 50 996 ± 38 894 ± 37	A3.1b A3.2a (1) A3.3a A3.4a(2) A3.5b	1,838 ± 47 2,041 ± 43 1,960 ± 55 1,983 ± 37 2,137 ± 46	A4.1b A4.2a A4.3a (1) A4.4a (2) A4.5b	724 ± 42 778 ± 88 764 ± 45 602 ± 38 825 ± 44	a, method a b, method b (), same subsample
δ ¹³ C (%)		-25.0		-23.0		-23.6		-25.0	
Oxford	2575		2574		2576		2589		
	O1.1u O1.2b O1.1b	795 ± 65 730 ± 45 745 ± 55	O2.1u O2.1b O2.2b†	980 ± 55 915 ± 55 925 ± 45	O3.1u O3.1b O3.2b	$1,955 \pm 70$ $1,975 \pm 55$ $1,990 \pm 50$	O4.2u O4.2b (1) O4.2b (2)	785 ± 50 710 ± 40 790 ± 45	u, unbleached b, bleached (), same pretreatment run combination
δ^{13} C‡ (%)		-27.0		-27.0		-27.0		-27.0	run comomation
Zurich	ETH Z1.1u Z1.1w Z1.1s	733±61 722±56 635±57	ETH- Z2.1u Z2.1w Z2.1s	-3884 890 ± 59 1,036 ± 63 923 ± 47	ETH- Z3.1u Z3.2w Z3.2s	3885§ 1,984±50 1,886±48 1,954±50	ETH- Z4.1u Z4.1w Z4.1s	-3882 739 ± 63 676 ± 60 760 ± 66	u, ultrasonic only w, weak
	Z1.2w Z1.2s	639 ± 45 679 ± 51	Z2.2w Z2.2s	980 ± 50 904 ± 46			Z4.2w Z4.2s	646 ± 49 660 ± 46	s, strong
δ ¹³ C (%)		-25.1		-23.6		-22.0		-25.5	

In years BP, corrected for δ^{13} C fractionation; errors at 1σ level; see text for pretreatment details.

^{*} The identification code for each measurement shows, in order, the laboratory, sample, measurement run, pretreatment and any replication involved.

† One anomalous replicate (of 6) obtained for independent measurement O2.2b; if rejected it reduces date by 40 yr; final date quoted actually reduced by 20 yr.

[‡] Measured for samples 1 and 3; assumed for samples 2 and 4.

[§] The loose weave of sample Z3.1 led to its disintegration during strong and weak chemical treatments. Z3.2 was centrifuged to avoid the same loss of material.

Average of separate determinations by AMS.

cleaned the samples using a vacuum pipette, followed by cleaning in petroleum ether (40 °C for 1 h) to remove lipids and candlewax, for example. Zurich precleaned the samples in an ultrasonic bath. After these initial cleaning procedures, each laboratory split the samples for further treatment.

The Arizona group split each sample into four subsamples. One pair of subsamples from each textile was treated with dilute HCl, dilute NaOH and again in acid, with rinsing in between (method a). The second pair of subsamples was treated with a commercial detergent (1.5% SDS), distilled water, 0.1% HCl and another detergent (1.5% triton X-100); they were then submitted to a Soxhlet extraction with ethanol for 60 min and washed with distilled water at 70 °C in an ultrasonic bath (method b).

The Oxford group divided the precleaned samples into three. Each subsample was treated with 1M HCl (80 °C for 2 h), 1M NaOH (80 °C for 2 h) and again in acid, with rinsing in between. Two of the three samples were then bleached in NaOCl (2.5% at pH-3 for 30 min).

The Zurich group first split each ultrasonically cleaned sample in half, with the treatment of the second set of samples being deferred until the radiocarbon measurements on the first set had been completed. The first set of samples was further subdivided into three portions. One-third received no further treatment, one-third was submitted to a weak treatment with 0.5% HCl (room temperature), 0.25% NaOH (room temperature) and again in acid, with rinsing in between. The final third was given a strong treatment, using the same procedure except that hot (80 °C) 5% HCl and 2.5% NaOH were used. After the first set of measurements revealed no evidence of contamination, the second set was split into two portions, to which the weak and strong chemical treatments were applied.

All of the groups combusted the cleaned textile subsample with copper oxide in sealed tubes, then converted the resulting CO_2 to graphite targets. Arizona and Oxford converted CO_2 to CO in the presence of zinc, followed by iron-catalysed reduction to graphite, as described in Slota *et al.*⁶. Zurich used cobalt-catalysed reduction in the presence of hydrogen, as described by Vogel *et al.*^{7,8}.

Each laboratory measured the graphite targets made from the textile samples, together with appropriate standards and blanks, as a group (a run). Each laboratory performed between three and five independent measurements for each textile sample which were carried out over a time period of about one month. The results of these independent measurements (Table 1) in each case represent the average of several replicate measurements made during each run (samples are measured sequentially, the sequence being repeated several times). The specific measurement procedures for each laboratory are given by Linick et al.⁹ for Arizona, by Gillespie et al.¹⁰ for Oxford and by Suter et al.¹¹ for Zurich. Arizona and Oxford measured ¹⁴C/¹³C ratios by AMS and determined the ¹³C/¹²C ratios using conventional mass spectrometry. Zurich determined both ¹⁴C/¹²C and ¹³C/¹²C quasi-simultaneously using AMS only.

The conventional radiocarbon ages were all calculated using the procedures suggested by Stuiver and Polach¹², with normalization to $\delta^{13}C = -25\%$, and were accordingly reported in yr BP (years before 1950). The errors, which are quoted in Table 1 at the 1σ level (σ is standard deviation), include the statistical (counting) error, the scatter of results for standards and blanks, and the uncertainty in the $\delta^{13}C$ determination (Arizona includes the $\delta^{13}C$ error at a later stage, when combining subsample results; Oxford errors below 40 yr are rounded up to 40).

Results

On completion of their measurements, the laboratories forwarded their results to the British Museum Research Laboratory for statistical analysis. The individual results as supplied by the laboratories are given in Table 1. Each date represents a unique combination of pretreatment and measurement run and applies

Table 2 Summary of mean radiocarbon dates and assessment of interlaboratory scatter

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Sample	1	2	3	4			
Arizona	646 ± 31	927 ± 32	$1,995 \pm 46$	722 ± 43			
Oxford	750 ± 30	940 ± 30	$1,980 \pm 35$	755 ± 30			
Zurich	676 ± 24	941 ± 23	$1,940\pm30$	685 ± 34			
Unweighted mean*	691 ± 31	936±5	$1,972 \pm 16$	721 ± 20			
Weighted mean†	689 ± 16	937 ± 16	$1,964 \pm 20$	724 ± 20			
χ^2 value (2 d.f.)	6.4	0.1	1.3	2.4			
Significance‡ level (%)	5	90	50	30			

Dates are in yr BP. d.f., degrees of freedom.

- * Standard errors based on scatter.
- † Standard errors based on combined quoted errors.
- ‡ The probability of obtaining, by chance, a scatter among the three dates as high as that observed, under the assumption that the quoted errors reflect all sources of random variation.

to a separate subsample, except where indicated by the identification code. From these data it can be seen that, for each laboratory, there are no significant differences between the results obtained with the different cleaning procedures that each used.

The mean radiocarbon dates and associated uncertainties for the four samples, as supplied by each of the three laboratories, are listed in Table 2 and shown in Fig. 1. Also included in Table 2 are the overall unweighted and weighted means, the weights being proportional to the inverse squared errors as quoted by the laboratories. The underlying principle of the statistical analysis has been to assume that, unless there is strong evidence otherwise, the quoted errors fully reflect all sources of error and that weighted means are therefore appropriate. An initial inspection of Table 2 shows that the agreement among the three laboratories for samples 2, 3 and 4 is exceptionally good. The spread of the measurements for sample 1 is somewhat greater than would be expected from the errors quoted.

More quantitatively, to establish whether the scatter among the three laboratory means was consistent with their quoted errors, a χ^2 test was applied to the dates for each sample, in accordance with the recommended procedure of Ward and Wilson¹³. The results of this test, given in Table 2, show that it is unlikely that the errors quoted by the laboratories for sample 1 fully reflect the overall scatter. The errors might still reflect the uncertainties in the three dates relative to one another, but in the absence of direct evidence on this, it was decided to give the three dates for sample 1 equal weight in determining the final mean, and to estimate the uncertainty in that mean from the scatter of results.

As shown in Table 2, the unweighted mean of the radiocarbon age of sample 1 and its uncertainty are $691 \pm 31 \text{ yr BP}$. The confidence limits for sample 1 were obtained by multiplying the uncertainty by t_d , the value of a Student's t distribution with d degrees of freedom at the appropriate probability level. The value of d, which lies between the inter- and intra-laboratory degrees of freedom-that is, between 2 and 9-was estimated at 5 on the basis of an analysis of variance on the 12 individual measurements supplied by the laboratories14. Individual measurements from a particular laboratory were weighted according to their inverse squared errors, but the contributions from different laboratories were weighted equally, thus ensuring consistency with Table 2. Thus for sample 1, where the error has been estimated from the scatter, ~68% and 95% confidence limits for the true radiocarbon date were found from the 1.1σ and 2.6σ errors about the unweighted mean respectively, the multiplying factors being obtained from standard tables of the t_5 distribution. However, for samples 2, 3 and 4, the limits were obtained in the usual way from 1σ and 2σ quoted errors about the weighted means, assuming normality.

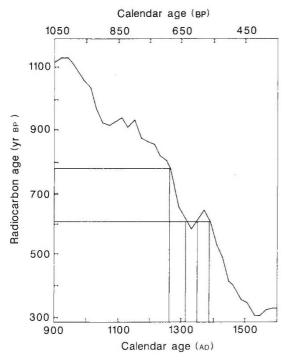


Fig. 2 Calibration of the overall mean radiocarbon date for sample 1 (the Shroud of Turin) using the 'intercept' method. (See also Table 3.) Calibration is necessary because of natural variations in atmospheric ¹⁴C. The calibration curve for the relevant period is that of Stuiver and Pearson⁵, a portion of which is illustrated. The uncertainty in the calibration curve has been combined with the error in the mean radiocarbon date, giving the 95% confidence limits on the radiocarbon scale; the error envelope on the curve has therefore been omitted from the diagram. The stippled areas show how the 95% confidence limits are transformed from the radiocarbon to the calendar scale.

The calendar-age ranges which correspond to the radiocarbon confidence limits are shown in Table 3. These were determined from the high-precision curve of Stuiver and Pearson⁵ based on dendrochronological dating. Method A (the intercept method) in revision 2.0 of the University of Washington Quaternary Isotope Laboratory Radiocarbon Calibration Program¹⁵ was used. In this method, the error in the calibration curve is first incorporated into the radiocarbon error, thus widening the limits on the radiocarbon scale slightly; calendar ages are then found that correspond to these limits, without transforming the complete probability distribution of radiocarbon dates. No additional uncertainty has been added to take account of the short growth period of textile samples. There is little published guidance on this, although it has been suggested that 15 years should be added in quadrature to the overall uncertainty in the radiocarbon date for samples of growth period less than one year, such as linen. In general, such additional uncertainty would widen the 95% calendar limits by ~2-4 years at either end, except for sample 3 where the 9 cal BC limit would be changed to 34 cal BC.

The 95% limits for the shroud are also illustrated in Fig. 2, where it is apparent that the calibration of the radiocarbon date for sample 1 gives a double range. The correct transformation of probability distributions from the radiocarbon to the calendar scale is still subject to debate, there being two different methods of dealing with multiple intercepts. However, both methods agree that the major probability peak lies in the earlier of the two ranges, in the 68% range at the end of the thirteenth century. Sample 4 has a very narrow calendar range: this is due to the steep slope in the calibration curve at this point, and is an occasional instance of calibration reducing rather than increas-

Table 3 Calibrated date ranges at the 68% and 95% confidence levels

Sample	Mean date (yr BP)		Calendar date ranges	
1*	691 ± 31	68%	AD 1273-1288	
		95%	AD 1262-1312, 1353-1384 cal	
2†	937 ± 16	68%	AD 1032-1048, 1089-1119,	
			1142-1154 cal	
		95%	AD 1026-1160 cal	
3†	$1,964 \pm 20 \ddagger$	68%	AD 11-64 cal	
		95%	9 cal BC-AD 78 cal	
4†	724 ± 20	68%	AD 1268-1278 cal	
		95%	AD 1263-1283 cal	

^{*} Confidence limits on the radiocarbon scale found from the standard error on the unweighted mean, assuming a t_5 distribution (multiplying factors 1.1 and 2.6 for 68% and 95% limits respectively). Standard error estimated from scatter.

ing a confidence range. Sample 3 compares well with the date obtained by conventional radiocarbon dating; there is no evidence for a difference between the two results. The dates for samples 2 and 4 agree with the historical evidence, which places them in the eleventh to twelfth centuries and late thirteenth/early fourteenth centuries AD respectively.

The results, together with the statistical assessment of the data prepared in the British Museum, were forwarded to Professor Bray of the Istituto di Metrologia 'G. Colonetti', Turin, for his comments. He confirmed that the results of the three laboratories were mutually compatible, and that, on the evidence submitted, none of the mean results was questionable.

Conclusions

The results of radiocarbon measurements at Arizona, Oxford and Zurich yield a calibrated calendar age range with at least 95% confidence for the linen of the Shroud of Turin of AD 1260-1390 (rounded down/up to nearest 10 yr). These results therefore provide conclusive evidence that the linen of the Shroud of Turin is mediaeval.

The results of radiocarbon measurements from the three laboratories on four textile samples, a total of twelve data sets, show that none of the measurements differs from its appropriate mean value by more than two standard deviations. The results for the three control samples agree well with previous radiocarbon measurements and/or historical dates.

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[†] Confidence limits on the radiocarbon scale found from the standard error on the weighted mean, assuming a normal distribution (multiplying factors 1 and 2 for 68% and 95% limits respectively). Standard error computed from quoted errors.

[‡] Date by conventional radiocarbon dating at the British Museum: $2010\pm80\,\mathrm{yr}$ BP (BM-2558).

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Three-dimensional structure of aspartyl protease from human immunodeficiency virus HIV-1

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The crystal structure of the protease of the human immunodeficiency virus type 1 (HIV-1), which releases structural proteins and enzymes from viral polyprotein products, has been determined to 3 Å resolution. Large regions of the protease dimer, including the active site, have structural homology to the family of microbial aspartyl proteases. The structure suggests a mechanism for the autoproteolytic release of protease and a role in the control of virus maturation.

THE HIV-1 retrovirus has been shown to be the causative agent in AIDS^{1,2}. In the course of viral replication, many retroviral structural proteins and enzymes are initially translated as polyproteins which undergo enzymatic cleavage to generate the functional proteins found in mature virions3. Genetic and biochemical studies have demonstrated that a virally encoded protease is responsible for the release of structural proteins from the gag gene product, and for the release of protease (implying autoproteolysis), reverse transcriptase and integrase from the gag-pol fusion protein⁴⁻⁸. Analysis of the sequences of retroviral proteases led to the suggestion that these enzymes were members of the aspartyl protease family, on the basis of the observed conservation of a characteristic Asp-Thr-Gly active-site sequence9. HIV-1 protease was found to be inhibited in vitro by pepstatin¹⁰⁻¹², a general inhibitor of aspartyl proteases, whose action operationally defines members of this class of enzymes¹³. Furthermore, single amino-acid substitutions of the HIV-1 protease sequence in which the highly conserved Asp 25 was converted to Asn^{12,14}, Thr¹⁰ or Ala¹⁵ resulted in the elimination of protease activity. Upon transfection into mammalian cells, a provirus containing one of these mutations (Asn 25) was shown to generate non-infectious virions containing unprocessed polyprotein¹⁴. Consequently, we have focused on HIV-1 protease as a therapeutic target, and have now determined the structure of this enzyme to accelerate the rational design of protease inhibitors as potential agents in the control of AIDS.

HIV-1 protease, with only 99 amino-acid residues 16-19, is the smallest of the retroviral proteases, and is much smaller than the microbial and mammalian aspartyl proteases, each of which contains approximately 325 residues. These pepsin-like enzymes contain homologous N- and C-terminal domains that may have evolved from a primordial gene duplication and fusion event^{20,21}. The domains, which are related by an approximate twofold axis of symmetry, each contribute one Asp-Thr-Gly triad to the pseudo-symmetric active site. We show here that in HIV-1 protease, a strict crystallographic twofold relationship exists, in which each monomer contributes one Asp-Thr-Gly sequence to a truly symmetric active site. The structure of this active site, including the interaction of the catalytic aspartic acid residues across the twofold axis, closely resembles the active site of the pepsin-like aspartyl proteases.

Structure of HIV-1 protease

The isolation and crystallization of the HIV-1 protease have been described elsewhere²². Details of the structure determination are given in Table 1 legend. C_{α} coordinates will be deposited with the Brookhaven Protein Data Bank²³.

The tertiary structure of the HIV-1 protease monomer contains exclusively β -sheet, turn and extended polypeptide structural elements. Schematic descriptions of the protease structure are presented in Fig. 1. In the 3.0 Å multiple isomorphous replacement (MIR) electron density map from which this structure was derived, strong, continuous electron density is observed for all but the first five residues at the N terminal (Pro 1 to Leu 5). Chemical sequence analysis of washed crystals, however, confirmed the presence of the correct N-terminal sequence, beginning with Pro 1 (J. Rodkey, unpublished results). We conclude that the first five residues are present in our crystallized protein, but that they are disordered crystallographically, probably because of an inherent flexibility (see below). It should be noted that our current model leaves no regions of electron density in the MIR map unassigned.

In this crystal structure, the two monomer components of each HIV-1 protease dimer are related by a crystallographic twofold axis of symmetry. The overall shape of the dimer is very asymmetric; it can be approximated by an oblate ellipsoid, roughly 55 by 35 by 25 Å. To simplify the description of the dimer, the two monomers will arbitrarily be denoted A and B, even though they are crystallographically equivalent (Fig. 2). The dimer is stabilized by two sets of antiparallel interactions,