THE RESOLUTION OF RACEMIC 2-ALKYL-1-AMINOCYCLOPROPANE-1-CARBOXYLIC ACIDS Jack E. Baldwin*, Robert M. Adlington, Bernard J. Rawlings and Richard H. Jones Dyson Perrins Laboratory, University of Oxford, South Parks Road, Oxford OX1 3QY, U.K.

<u>Abstract</u>: Practical procedures for the resolution of racemic modification of $(1\underline{R}, 2\underline{S})$ - and $(1\underline{S}, 2\underline{R})$ -1-amino-2-ethylcyclopropane-1-carboxylic acid <u>1a,b</u>,(1\underline{R}, 2\underline{S})- and (1\underline{S}, 2\underline{R})-1-amino-2-methylcyclopropane-1-carboxylic acid <u>2a,b</u>, and (1\underline{R}, 2\underline{R})- and (1\underline{S}, 2\underline{S})-1-amino-2-methylcyclopropane-1-carboxylic acid <u>3a,b</u> are described; the structures as <u>1a,2a</u>, and <u>3a</u> were confirmed by X-ray crystallographic methods.

In the preceding paper¹ we described a convenient synthesis of the racemic forms of 2alkylated-1-aminocyclopropane-1-carboxylic acids <u>1a,b</u>, <u>2a,b</u>, and <u>3a,b</u>. In order to develop our studies on ethylene biosynthesis^{2,3} we required reliable⁴ methods for the resolution of these amino-acids so that they may be tested as substrates or competitive inhibitors of the ethylene synthetase system found in apple tissue.

Initially, the racemates <u>1a,b</u>, <u>2a,b</u>, and <u>3a,b</u> were converted to their <u>N</u>-chloroacetyl derivatives and tested as substrates for Porcine Kidney Acylase I⁵. By following the reaction by N.M.R. spectroscopy it was shown that only <u>4b</u> was selectively hydrolysed by this enzyme.⁶ Standard work up of the crude product from <u>4a,b</u> gave (1<u>S</u>, <u>2S</u>)-<u>3b</u>, $[\alpha]_{D}^{20}$ +44^O (C = 1.0, H₂O), δ H(300MHz, D₂O) as described¹, and unreacted <u>4a</u>. Acidic hydrolysis (6M HCl, reflux, 6h.) of <u>4a</u> and purification by ion-exchange [Dowex 50W X 8(H)] gave (1<u>R</u>, <u>2R</u>)-<u>3a</u>, $[\alpha]_{D}^{20}$ -42^O (C = 1.4, H₂O), δ H(300MHz, D₂O) as described.¹ The structure of the chloroacetyl derivative as (1<u>R</u>, <u>2R</u>)-<u>4a</u> was confirmed by X-ray crystallographic analysis of the crystals from ethylacetate/hexane (figure 1).

Crystal Data: $C_{r}H_{10}NO_{3}Cl$, m = 200.0, Monoclinic, space group $P2_{1}$, a = 5.961(1) b = 10.066(4), c = 7.424(1) Å, β = 90.35(1)^O, U = 445.2 Å³, Z = 2, D_C = 1.43 cm⁻¹. 1568 independent reflections (0 < 20 < 55^O) gave 1461 observed reflections (I > 3 σ (I)). The structure was solved by Patterson and electron density methods. The Final R value is 0.030 (R_W 0.038). All data were measured on an Enraf-Nonius CAD4 diffractometer using graphite monochromated Mo-K_α radiation (λ = 0.71069 Å). Refinement was by full matrix least squares in all cases.⁷ The absolute configuration in <u>4a</u> was determined by including a polarity parameter⁸ in the refinement which converged to 0.97(13), thus showing that the molecule illustrated in Fig.1

to have the correct absolute configuration. This was confirmed by the careful measurement of 22 Friedel pairs followed by the application of Bijvoet's method.⁹

As the enzymic method failed for the resolution of the racemates $\underline{1a, b}$ and $\underline{2a, b}$ an alternative chemical method was sought. Thus the racemate $\underline{1a, b}$ was converted to its amine hydrochloride methyl ester (MeOH, HCl(g), 50°, 18h.) which was coupled with \underline{S} -(+)-2-hydroxy-2-phenylacetic acid [N-Ethoxycarbonyl-2-ethoxy-1,2 dihydroquinoline (EEDQ) (2 equivs.)^{1°}, triethylamine (1 equiv.), dichloromethane, 18h.] to the diastereoisomeric derivatives $\underline{5a, b}$ (75% from $\underline{1a, b}$), $\delta H(300 \text{MHz, } \text{C}^2 \text{HCl}_3)^{11}$ 3.62, 3.64 p.p.m. (2x3H, 2xs, 2xCO₂Me). Double recrystallisation (from ethyl acetate/hexane) gave the essentially pure diastereoisomer $\underline{5a}$, m.p. 153-4°, $[\alpha]_D^{2°}$ +113° (C = 1.5, EtOH), $\delta H(300 \text{MHz, } \text{C}^2 \text{HCl}_3)$ 0.94-1.00(1H, m, 3-H), 0.98(3H, t, J 8Hz, 2-CH₂CH₃), 1.10-1.25(1H, m, 2-CH₂CH₃), 1.31(3H, t, J 7Hz, 0CH₂CH₃), 1.44-1.55(1H, m, 2-CH₂CH₃), 1.70-1.78(2H, m, 2,3-H), 3.64(3H, s, CO₂Me), 4.19-4.27(2H, <u>ca</u> dq, 0CH₂CH₃), 6.00(1H, s, CHPh), 6.66(1H, s, NH), and 7.36-7.53(5H, m, Ph-H), m/e (ammonium chemical ionisation) 336(MH⁺, 100%), which upon acid hydrolysis and ion-exchange gave¹² 1<u>R</u>, 2<u>S-1a</u> $[\alpha]_D^{2°}$ *65° (C = 1.3, H₂O), $\delta H(300 \text{MHz, D}_2O)$ as described.¹ The structure as <u>5a</u> was confirmed by X-ray crystallographic methods (Figure 2).

Crystal Data: $C_{18}H_{23}NO_6$, m = 349.4, Orthorhombic, space group $P2_12_12_1$, a = 9.895(2), b = 10.381(3), c = 18.405(5) Å₁U = 1890.6 Å³, Z = 4, D_c = 1.23g cm⁻³. 1904 independent reflections (0 < 20 < 50°) gave 1448 observed reflections (I > 2 σ (I)). The structure was solved by direct methods¹³ and the final R value is 0.034(R_W 0.041).

By similar means, the amine hydrochloride methyl ester from <u>1a,b</u> was coupled with <u>R</u>-(-)-2-hydroxy-2-phenylacetic acid to yield <u>6a,b</u> (79% from <u>1a,b</u>). Double recrystallisation gave the essentially pure diastereoisomer <u>6b</u> m.p. 154-5°, $[\alpha]_D^{2^\circ}$ -109° (C = 1.8, EtOH), which upon acidic hydrolysis and ion-exchange gave (1<u>S</u>, 2<u>R</u>)-<u>1b</u>, $[\alpha]_D^{2^\circ}$ -67° (C = 1.5, H₂O), δ H(300MHz, D₂O) as described.¹

An identical operation was used to resolve the racemate <u>2a,b</u>. Thus <u>2a,b</u> was converted to its amine hydrochloride methyl ester form which was coupled, as before, with <u>S</u>-(+)-2hydroxy-2-phenylacetic acid to the diastereoisomeric derivatives <u>7a,b</u> (85% from <u>2a,b</u>) $\delta H(300MHz, C^2HCl_s)$ 1.06(3H, d, <u>J</u> 6Hz, 2-Me) 1.155(3H, d, <u>J</u> 6Hz, 2-Me), 3.62, 3.64(2x3H, 2xs, 2xCO_2Me). Double recrystallisation (from ethyl acetate/hexane) gave the essentially pure diastereoisomer isomer <u>7a</u>, m.p. 139-40°, $[\alpha]_D^{2\circ}$ +106° (C = 3.1, EtOH), $\delta H(300MHz, C^2HCl_s)$ 0.88(1H, dd, <u>J</u> 5,7Hz, 3-H), 1.06(3H, d, <u>J</u> 6Hz, 2-Me), 1.31(3H, t, <u>J</u> 7Hz, OCH₂CH₃), 1.69-1.83(2H, m, 2,3-H), 3.64(3H, s, CO_2Me), 4.18-4.26(2H, <u>ca</u> dq, OCH₂CH₃), 6.00(1H, s, <u>CHPh</u>), 6.70(1H, s, NH) and 7.36-7.39, 7.49-7.52 (5H, m, Ph-H), m/e (ammonium desorption chemical ionisation) 350(MH⁺, 100%), which upon acidic hydrolysis and ion-exchange gave (1<u>R</u>, 2<u>S</u>)-<u>2a</u>, $[\alpha]_D^{2\circ}$ +73.5 (C = 0.4, H₂O), $\delta H(300MHz, D_2O)$ as described.¹ The structure as <u>7a</u> was confirmed by X-ray crystallographic methods (Figure 3).

Crystal Data: $C_{17}H_{21}NO_6$, m = 335.4, orthorhombic, space group $P2_12_12_1$ a = 9.765(6) b = 10.140(2), c = 18.482(4) Å, U = 1830.0 Å³, Z = 4, D_c = 1.22g cm⁻³. 1845 independent reflections (0 < 20 < 50^o) gave 1225 observed reflections (I > 2 σ (I)). The structure was solved by direct methods¹³. The current R value is 0.050(R_W = 0.060).

When the amine hydrochloride methyl ester from 2a,b was coupled with \underline{R} -(-)-2-hydroxy-2phenylacetic acid, a mixture of $\underline{8a}, \underline{b}$ (89% from $\underline{2a}, \underline{b}$) was formed. Double recrystallisation gave the essentially pure diasteroisomer $\underline{8b}$, m.p. 139-40°, $[\alpha]_D^{20}$ -104° (C = 3.0 in EtOH), which upon acidic hydrolysis and ion-exchange gave (1<u>S</u>, 2<u>R</u>)-2<u>b</u>, $[\alpha]_D^{20}$ -69° (C = 0.3, H₂O), $\delta H(300MHz, D_2O)$ as described.¹

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$$2 \underbrace{\underbrace{\overset{(0)}{\underset{Et}{\overset{1}{\underset{NH_{3}}{+}}}}_{Et}}^{2} \underbrace{\underbrace{\overset{(0)}{\underset{NH_{3}}{+}}}_{1}}_{NH_{3}+} \underbrace{\underbrace{1}_{\overset{(1)}{\underset{NH_{3}}{+}}}}_{1} \underbrace{\underbrace{1}_{\overset{(1)}{\underset{NH_{3}}{+}}}_{NH_{3}+}}$$

$$Me = \frac{1}{NH_3+} \frac{2b}{2b} \left(\frac{|S|}{2R} \right)$$

$$\begin{array}{c}
\text{Me} \\
2 \\
2 \\
+ NH_{3}
\end{array}$$

$$\begin{array}{c}
\text{Me} \\
\frac{3b}{(15, 25)}
\end{array}$$









References

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- R. M. Adlington, R. T. Aplin, J. E. Baldwin, B. J. Rawlings, and D. Osborne, <u>J.Chem.</u> Soc., Chem.Commun, 1982, 1086.
- 3. R. M. Adlington, J. E. Baldwin, and B. J. Rawlings, J.Chem.Soc., Chem.Commun., 1983, 290.
- 4. A report of the resolution of (±) allocoronamic acid (<u>1a,b</u>) into its (+) (1<u>S</u>, 2<u>R</u>) $[\alpha]_{D}^{23}$ +65.0° (C = 1.83, H₂O) and (-)(1<u>R</u>, 2<u>S</u>) $[\alpha]_{D}^{21}$ -68.4° (C =1.15, H₂O) forms [A. Ichihara et al, <u>Agric.Biol.Chem.</u>, 1977, <u>41</u>, 2497] in which the absolute configurations were assigned "by <u>application</u> of the sector rule in ORD" was subsequently reversed by the same author (A. Ichihara et al, <u>Tetrahedron Lett.</u>, 1979, 365). However conclusive proof of the correct absolute configuration by X-ray crystallographic methods was not reported.
- Obtained from Sigma Chemical Company, Poole, Dorset, BH17 7NH, England, order number A 3010, Acylase 1, grade 1, Activity 2,000-3,000 units per mg protein.
- 6. The chloroacetyl derivatives were incubated with Acylase 1 at pH 7-7.5, 37°, for 12h. Direct n.m.r. analysis (500MHz, D₂0) showed the presence of <u>3b</u> 6H 0.97(3H, d, <u>J</u> 6Hz, 2-Me) and unreacted <u>4a</u>, 6H 0.92(3H, d, <u>J</u> 6Hz, 2-Me) ratio <u>ca</u> 1:1. The chloroacetyl derivatives of <u>1a,b</u> and <u>2a,b</u> were not substrates for this enzyme. Hence in these cases, the C(2) alkyl group which is <u>cis</u>- to the amide function must seriously retard the enzymatic rate of amide hydrolysis. The selective enzymatic hydrolysis of <u>4b</u> over <u>4a</u> was expected as this entantiomer contains the <u>1S</u>-aminoacid configuration.
- 7. D. J. Watkin, and J. R. Carruthers, CRYSTAL Manual, Chemical Crystallography Laboratory, University of Oxford, 1981.
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- 10. EEDQ (2 equivs.) was used both to form the amide bond and to convert the free hydroxy function to an ethyl carbonate group.
- 11. N.M.R. shifts in deuteriochloroform are referenced to CHCl₃ = 7.27 p.p.m.
- 12. <u>S</u>-(+)-2-Hydroxy-2-phenylacetic acid was recovered <u>after</u> hydrolysis with identical ¹H n.m.r. spectrum and optical rotation $[\alpha]_D^{20}$ +146 (C = 2.5, EtOH) to that initially used $[\alpha]_D^{20}$ +150 (C = 1.5,EtOH) to form <u>5a,b</u>.
- 13. P. Main, S. J. Fiske, S. E. Hull, L. Lessinger, G. Germain, J. P. Declerq, and M. M. Woolfson, MULTAN 80, A system of computer programs for the solution of crystal structures from X-ray diffraction data, Department of Physics, University of York, England, 1980. The atomic co-ordinates for Figure 1,2,3 are available on request from the Director of the Cambridge Crystallographic Data Centre, University Chemical Laboratory, Lensfield Road, Cambridge CB2 1EW. Any request should be accompanied by the full literature citation for this communication.