# INTERACTIONS BETWEEN ACETYLCHOLINESTERASE AND TETRA-N-ALKYLAMMONIUM IONS

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Abstract—A study of the mechanism of interaction of acetylcholinesterase with some simple tetra-*N*-alkyammonium ions has been made. Kinetic schemes have been proposed which are consistent with the experimental results observed in the enzyme-tetra-*N*-alkylammonium system in the presence of substrate and in the presence of an organophosphorus inhibitor.

Some years ago it was postulated that the products of hydrolysis in acyl hydrolase systems were formed in two consecutive reactions, with an acylated enzyme intermediate being formed from the enzyme-substrate (Michaelis) complex [1, 2]. The hydrolysis of acetylcholine by acetylcholinesterase (AChE) is one such system. It has been previously shown that the rate of substrate hydrolysis in such a system may be modified by reversible inhibitors by two mechanisms: complex formation between the free enzyme and inhibitor, thereby reducing the concn of Michaelis complex, or complex formation between the acylated enzyme and inhibitor, thereby modifying the rate of de-acylation [3]. A third alternative, complex formation between the Michaelis complex and inhibitor, has been shown not to occur [3]. Reversible inhibitors, or modifiers, of the AChE-substrate system generally contain a quaternary nitrogen atom [3].

In order to gain a better understanding of the mechanisms by which quaternary nitrogen compounds interact with AChE, an investigation of the effects upon substrate hydrolysis of a series of simple tetra-N-alkylammonium salts ( $R_4N^+$ ) has been carried out, where R = methyl (Me), ethyl (Et), n-propyl ( $Pr^n$ ), n-butyl ( $Bu^n$ ), and n-pentyl ( $Pent^n$ ). A mechanistic description of the system has been developed which is kinetically rigorous and somewhat simpler than that previously proposed [3].

A kinetic analysis of the system where two different modifiers are present has been proposed and shown to apply to experimentally measured data. A limited number of experiments have also been performed on the AChE-modifier-organophosphorus (irreversible) inhibitor system in order to demonstrate that the AChE-modifier interactions proposed earlier still apply under these conditions.

#### **EXPERIMENTAL**

# Materials

AChE from bovine erythrocytes (EC 3.1.1.7) was obtained from Sigma Chemicals. Acetylcholine iodide (AChI) was obtained from BDH Biochemi-

cals. Me<sub>4</sub>N<sup>+</sup>I<sup>-</sup>, Et<sub>4</sub>N<sup>+</sup>I<sup>-</sup>, Pr<sup>2</sup>N<sup>+</sup>I<sup>-</sup> and Bu<sup>2</sup>N<sup>+</sup>I<sup>-</sup> were obtained from BDH Chemicals and were recrystallised from ethanol/water or ethyl acetate/ethanol before use. Pent<sup>2</sup>N<sup>+</sup>Br<sup>-</sup> was prepared at CDE (Porton, U.K.) [4].

AChE activities were measured titrimetrically using a Radiometer automatic titrator (pH meter 26, titrator 11, titrigraph SBR 2c, syringe burette SBU 1a) fitted with twin syringes containing NaOH (0.01 N) and AChI (0.01 M).

## Methods

Single-modifier experiments. To a solution of NaCl (0.1 M, 10 ml) containing AChE (1 unit\*) in a radio meter cell at 37° and at pH 7.4 was added AChI solution. The surface of the solution was bathed with N<sub>2</sub> to prevent CO<sub>2</sub> uptake. Five concns of AChI were used, ranging from  $4.0 \times 10^{-4}$  to  $0.6 \times 10^{-4}$  M. The rate of AChI hydrolysis was monitored on the radiometer by pH-statting at 7.4, and was constant because the AChI concn was maintained by means of the twin syringes. Hydrolysis rates at five different AChI concns were then measured at six different concns of tetra-N-alkylammonium ions. Eadie plots [5] of the rate of substrate hydrolysis (v) against v/[S] (where [S] = substrate concn) were constructed for each different modifier concn. They yielded gradients of -G (defined later in this report) and intercepts of  $V_m$ .

Figs 1 and 2 show examples of these plots, and Table 1 shows the results obtained.

Mixed-modifier experiments. These experiments were performed and the results were treated as before, except that two different modifiers were present in solution. The results are shown in Table 2.

Organophosphorus inhibitions. AChE solution (10 units/ml in 0.1 M NaCl and  $5 \times 10^{-3}$  M phosphate buffer at pH 7.4—0.5 ml) at 37° was made  $10^{-4}$  M with respect to O-ethyl S-n-propyl methyl phosphonothiolate. Aliquots of 0.08 ml were removed at various times and diluted into 0.1 M NaCl solution (10 ml), and the remaining AChE activity was assayed by making the solution  $5 \times 10^{-4}$  M with AChI and pH-statting at 7.4 on the radiometer: continuing inhibition by the organophosphate was

<sup>\* 1</sup> unit will hydrolyse 1  $\mu$ mole of acetylcholine to choline and acetate per minute at pH 8 and 37°.

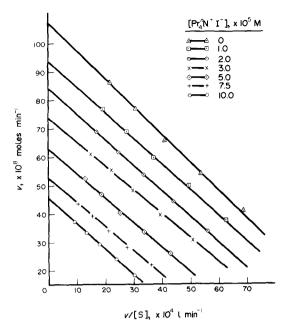


Fig. 1. Plots of v against v/[S] at different concns of AChI and Pr $2N^+I^-$ .

insignificant following this degree of dilution. The inhibition was repeated at different modifier concns. Apparent first-order rate coefficients were calculated as shown in the kinetic analysis. The results are shown in Table 3, and are the means of at least two experiments.

## Kinetic analysis

Single modifier. The system to be analysed is shown below; E is free enzyme, S is substrate, and M is the reversible inhibitor or modifier. ES and EM are the complexes formed between the enzyme and substrate and the enzyme and modifier respectively. EA is acylated enzyme and EAM the complex formed from EA and M.

$$E + S \stackrel{k_1}{\rightleftharpoons} ES \stackrel{k_2}{\rightarrow} EA \stackrel{k_3}{\rightarrow} E$$

$$+ \qquad \qquad + \qquad \qquad$$

 $[E_0] = [E] + [ES] + [EA] + [EM] + [EAM]$  (1) where  $[E_0]$  represents the initial enzyme concn.

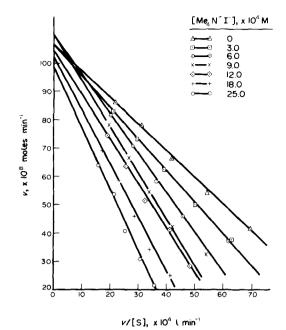


Fig. 2. Plots of v against v/[S] at different concns of AChI and Me<sub>4</sub>N<sup>+</sup>I<sup>-</sup>.

If the rate of acid production = v then  $v = k_3[EA] + k_6[EAM]$ , i.e.:

$$v = [EA](k_3 + k_6K_5[M]).$$
 (2)

Application of steady-state theory to [EA] and substitution into equations (1) and (2) leads to:

$$v = \frac{[E_0]k_2K_1[S](k_3 + k_6K_5[M])}{(1 + K_1[S] + K_4[M])(k_3 + k_6K_5[M]) + k_2K_1[S](1 + K_5[M])}.$$

This can be rearranged to the following equation:

$$v = \frac{-v}{[S]}G + V_m \tag{3}$$

where

$$G = \frac{(1 + K_4[M])(k_3 + k_6 K_5[M])}{K_1(k_2 + k_3) + (K_1 K_5[M])(k_2 + k_6)}$$
(4)

and

$$V_m = \frac{[E_0]k_2(k_3 + k_6K_5[M])}{(k_2 + k_3) + K_5[M](k_2 + k_6)}.$$
 (5)

Dividing equation (4) by equation (5) leads to:

$$\frac{G}{V_m} = \frac{1 + K_4[M]}{[E_0]k_2K_1}$$

Table 1.  $V_m$  and G values resulting from Eadie plots at different modifier concus

[Me <sub>4</sub> N <sup>+</sup> ] (×10 <sup>4</sup> M)	$V_m$ (×10 <sup>8</sup> moles min <sup>-1</sup> )	G (×10 <sup>4</sup> M)	[Et <sub>4</sub> N <sup>+</sup> ] (×10 <sup>3</sup> M)	$V_m$ (×10 <sup>8</sup> moles min <sup>-1</sup> )	G (×10 <sup>4</sup> M)	[Pr <sub>4</sub> <sup>n</sup> N <sup>+</sup> ] (×10 <sup>5</sup> M)	$V_m$ (×10 <sup>8</sup> moles min <sup>-1</sup> )	<i>G</i> (×10⁴ M)	[Bu <sub>4</sub> "N <sup>+</sup> ] (×10 <sup>5</sup> M)	$V_m$ (×10 <sup>8</sup> moles min <sup>-1</sup> )	<i>G</i> (×10⁴ M)	[Pent <sub>4</sub> <sup>n</sup> N <sup>+</sup> ] (×10 <sup>5</sup> M)	$V_m$ (×10 <sup>8</sup> moles min <sup>-1</sup> )	<i>G</i> (×10⁴ M)
0	107	0.95	0	106	0.91	0	108.0	0.98	0	104.0	0.91	0	108.0	0.98
3.0	106	1.12	0.5	108	1.58	1	93.7	0.90	0.5	92.7	0.88	3.0	97.6	1.12
6.0	110	1.40	1.0	117	2.36	2	84.2	0.90	1.0	83.9	0.91	5.0	91.3	1.14
9.0	110	1.60	1.5	114	3.10	3	73.7	0.85	2.0	73.7	1.10	8.0	82.2	1.18
12.0	104	1.58	2.0	116	3.71	5	63.0	0.86	3.0	58.9	0.89	12.0	82.0	1.52
18.0	102	1.90	3.0	106	4.50	7.5	52.9	0.88	4.0	50.9	0.87	15.0	74.2	1.59
25.0	98.2	2.12	5.0	96.3	6.18	10	42.8	0.87	6.0	41.7	0.93	20.0	71.3	1.72

Table 2. V<sub>m</sub> and G values resulting from Eadie plots at different mixed-modifier concus

$G \times 10^4 M$	0.98	1.00	1.01	1.00	1.00	96.0	1.09
$V_m$ $(\times 10^8 \text{ moles})$ min <sup>-1</sup>	94.2	82.0	73.9	61.0	52.2	45.0	37.1
[Pr <sub>1</sub> N <sup>+</sup> ] (×10 <sup>5</sup> M)	0	0.5	1.0	2.0	3.0	4.0	6.0
$\begin{bmatrix} Bu_t^2N^+ \\ (\times 10^5  M) \end{bmatrix}$	0	0.25	0.5	1.0	1.5	2.0	3.0
$G \times 10^4 \mathrm{M})$	1.09	1.40	1.45	2.47	2.78	3.78	5.30
$V_m$ (×108 moles min <sup>-1</sup> )	97.4	93.1	84.2	88.5	82.0	0.98	83.1
$\begin{bmatrix} Et_4N^+ \end{bmatrix} \\ (\times 10^3  \mathrm{M})$	0	0.25	0.5	1.0	1.5	2.0	3.0
$\frac{[\mathbf{B}\mathbf{u}_4^4\mathbf{N}^+]}{(\times 10^5\mathrm{M})}$	0	0.25	0.5	1.0	1.5	2.0	3.0
$G \times 10^4 \mathrm{M})$	0.99	1.10	1.19	1.21	1.54	1.51	1.56
$V_m \times 10^8 \text{ moles}$ min <sup>-1</sup> )	99.0	92.4	87.0	76.0	72.5	64.3	55.2
$[\mathrm{Me_4N^+}] \\ (\times 10^4  \mathrm{M})$	0	1.0	2.0	4.0	0.9	8.0	12.0
$\begin{array}{c} [\mathrm{Bu}_4^*\mathrm{N}^+] \\ (\times 10^5\mathrm{M}) \end{array}$	0	0.25	0.5	1.0	1.5	2.0	3.0

Table 3. Comparison of observed and calculated  $k_i$  values in the presence of modifiers

Modifier	Concn (M)	$K_4[M]$	$k_i$ (obs.) $(\times 10^3  \mathrm{sec}^{-1})$	$k_i$ (calc.) $(\times 10^3 \mathrm{sec}^{-1})$
  - 	Control		$3.30 \pm 0.15$ (N = 4)	
$Me_4N^+$	$1.79\times10^{-3}$	1	$1.50 \pm 0.05$	1.65
	$7.14 \times 10^{-3}$	4	$0.61 \pm 0.03$ (N = 2)	99.0
Bu7N⁺	$0.37\times10^{-4}$	-	$1.83 \pm 0.05$ (N = 2)	1.65
	$1.48 \times 10^{-4}$	4	$0.84 \pm 0.03$ (N = 2)	99.0
SAD 128	$0.45\times10^{-4}$	-	$1.74 \pm 0.06$	1.65
	$1.82 \times 10^{-4}$	4	$0.84 \pm 0.02$ $(N = 2)$	99.0

N = number of experiments.

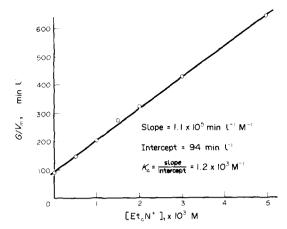


Fig. 3. Plot of  $G/V_m$  against  $[Et_4N^+]$ .

so that a plot of  $G/V_m$  against [M] will give a line of gradient  $K_4/([E_0]k_2K_1)$ , and an intercept (on the ordinate) of  $1/([E_0]k_2K_1)$ . Hence  $K_4$  can be calculated. Fig. 3 shows a typical example of this plot. When [M] = 0,  $V_m = V_0$ , i.e.:

$$V_0 = \frac{[E_0]k_2k_3}{k_2 + k_3}. (6)$$

Dividing equation (5) by equation (6) gives:

$$\frac{V_m}{V_0} = V_r = \frac{1 + \frac{k_6}{k_3} K_5[M]}{1 + K_5[M] \frac{(k_2 + k_6)}{(k_2 + k_3)}}.$$

Rearrangement gives:

$$\frac{1-V_{\rm r}}{[{\rm M}]} = V_{\rm r} K_5 \frac{(k_2+k_6)}{(k_2+k_3)} - \frac{k_6}{k_3} K_5.$$

Thus a plot of  $(1 - V_r)/[M]$  against  $V_r$  will give a line of gradient  $K_5$   $[k_2 + k_6)/(k_2 + k_3)]$  and an intercept (on the abscissa) of  $(k_6/k_3)[(k_2 + k_3)/(k_2 + k_6)]$ . Fig. 4 shows a typical example of this plot, and Table 4 lists the various parameters for this series of modifiers. Plots of  $(1 - V_r)/[M]$  against  $V_r$  were not possible for  $Me_4N^+$  and  $Et_4N^+$  because  $V_m$  was constant.

Mixed modifiers. The system to be analysed is shown below:  $M_1$  and  $M_2$  are different modifiers. Other symbols have been defined previously.

$$EM_{1} \qquad EAM_{1} \xrightarrow{K_{6 (1)}} EM_{1}$$

$$\downarrow \uparrow K_{4 (1)} \qquad \downarrow \uparrow K_{5 (1)}$$

$$M_{1} \qquad M_{1}$$

$$+ E + S \stackrel{K_{1}}{\rightleftharpoons} ES \stackrel{k_{2}}{\Rightarrow} EA \stackrel{k_{3}}{\Rightarrow} E$$

$$+ \qquad + \qquad M_{2} \qquad M_{2}$$

$$\downarrow \uparrow K_{4 (2)} \qquad \downarrow \uparrow K_{5 (2)}$$

$$EM_{2} \qquad EAM_{2} \xrightarrow{k_{6} \mathcal{P}} EM_{2}$$

 $K_1$  has been defined previously.

$$K_{4(1)} = \frac{[EM_1]}{[E][M_1]}; K_{4(2)} = \frac{[EM_2]}{[E][M_2]};$$

$$K_{5(1)} = \frac{[EAM_1]}{[EA][M_1]}; K_{5(2)} = \frac{[EAM_2]}{[EA][M_2]}$$

$$[E_0] = [E] + [EM_1] + [EM_2] + [ES] + [EAM_1] + [EAM_2] \quad (7)$$

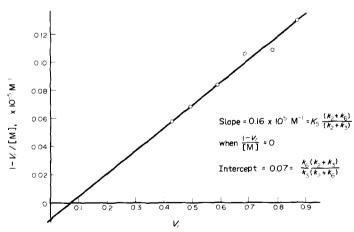


Fig. 4. Plot of  $(1 - V_r)/[M]$  against  $V_r$  for  $Pr_4^n N^-$ .

Table 4. Parameters calculated graphically

Parameter	Me <sub>4</sub> N <sup>+</sup>	Et <sub>4</sub> N-	Pr4 N+	Bu <sub>4</sub> N <sup>+</sup>	Pent <sup>n</sup> N <sup>+</sup>
$K_4$ (M <sup>-1</sup> )	$6.5 \times 10^{2}$	$1.2 \times 10^{3}$	$1.2 \times 10^{4}$	$2.7 \times 10^{4}$	$8.5 \times 10^{3}$
$K_5 \frac{(k_2 + k_6)}{(k_2 + k_3)} (M^{-1})$			$1.6 \times 10^4$	$2.5 \times 10^4$	$6.8 \times 10^{3}$
$\frac{k_6(k_2+k_3)}{k_3(k_2+k_6)}$	_	_	0.07	0	0.41

$$v = k_3 [EA] + k_{6(1)} [EAM_1] + k_{6(2)} [EAM_2], i.e.$$
:

$$v = [EA](k_3 + k_{6(1)}K_{5(1)}[M_1] + k_{6(2)}K_{5(2)}[M_2]).$$
(8)

Application of steady-state theory to [EA] and substitution into equations (7) and (8) leads to:

$$v = \frac{k_{2}[E_{0}](k_{3} + k_{6}_{(1)}K_{5}_{(1)}[M_{1}] + k_{6}_{(2)}K_{5}_{(2)}[M_{2}]}{(1 + K_{4}_{(1)}[M_{1}] + K_{4}_{(2)}[M_{2}])(k_{3} + k_{6}_{(1)}K_{5}_{(1)}[M_{1}] + k_{6}_{(2)}K_{5}_{(2)}[M_{2}])}{+(1 + K_{5}_{(1)}[M_{1}] + K_{5}_{(2)}[M_{2}])k_{2} + (k_{3} + k_{6}_{(1)}K_{5}_{(1)}[M_{1}] + k_{6}_{(2)}K_{5}_{(2)}[M_{2}])}}{K_{1}[S]}$$

This can be rearranged to the following equation:

$$v = \frac{-v}{|S|}G + V_m \tag{9}$$

where

$$G = \frac{(1 + K_{4}_{(1)}[M_1] + K_{4}_{(2)}[M_2]) (k_3 + k_{6}_{(1)}K_{5}_{(1)}[M_1] + k_{6}_{(2)}K_{5}_{(2)}[M_2])}{K_1 \left[k_2(1 + K_{5}_{(1)}[M_1] + K_{5}_{(2)}[M_2]) + k_3 + k_{6}_{(1)}K_{5}_{(1)}[M_1] + k_{6}_{(2)}K_{5}_{(2)}[M_2]\right]}$$
(10)

and

$$V_{m} = \frac{k_{2}[E_{0}](k_{3} + k_{6}_{(1)}K_{5}_{(1)}[M_{1}] + k_{6}_{(2)}K_{5}_{(2)}[M_{2}])}{k_{2}(1 + K_{5}_{(1)}[M_{1}] + K_{5}_{(2)}[M_{2}]) + k_{3} + k_{6}_{(1)}K_{5}_{(2)}[M_{1}] + k_{6}_{(2)}K_{5}_{(2)}[M_{2}]}.$$
(11)

Dividing equation (10) by equation (11) leads to:

$$\frac{G}{V_m} = \frac{1 + K_{4 (1)}[M_1] + K_{4 (2)}[M_2]}{[E_0]k_2K_1} 
= \frac{1}{[E_0]k_2K_1} + \frac{[M_1]}{[E_0]k_2K_1} (K_{4 (1)} + K_{4 (2)} \cdot C)$$

where  $C = [M_2]/[M_1]$ , so that a plot of  $G/V_m$  against  $[M_1]$  will give a gradient of  $[(K_{4(1)} + K_{4(2)} \cdot C)]/[M_1]$ 

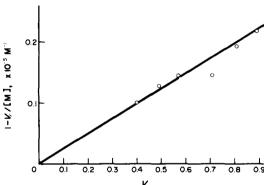


Fig. 5. Plot of  $(1 - V_r)/[M]$  against  $V_r$  for  $Bu_4^n N^+$ 

( $[E_0]k_2K_1$ ) and an intercept (on the ordinate) of 1/

Hence  $(K_{4(1)} + K_{4(2)} \cdot C)$  can be calculated and compared with the  $K_4$  values obtained from the single-modifier experiments. A comparison of these values is shown in Table 5.

 $V_{\rm r}$  is calculated as in the single-modifier analysis,

$$V_{r} = \frac{1 + \frac{k_{6 (1)}}{k_{3}} K_{5 (1)} [M_{1}] + \frac{k_{6 (2)}}{k_{3}} K_{5 (2)} [M_{2}]}{1 + K_{5 (1)} [M_{1}] \left(\frac{k_{2} + K_{5 (1)}}{k_{2} + k_{3}}\right) + K_{5 (2)} [M_{2}] \left(\frac{k_{2} + k_{6 (2)}}{k_{2} + k_{3}}\right). \quad (12)$$

Rearrangement of equation (12) gives:

$$\begin{split} \frac{1 - V_r}{[\mathbf{M}_1]} &= V_r \bigg[ K_{5 \ (1)} \bigg( \frac{k_2 + k_6 \ (1)}{k_2 + k_3} \bigg) \\ &\quad + C \cdot K_{5 \ (2)} \bigg( \frac{k_2 + k_6 \ (2)}{k_2 + k_3} \bigg) \, \bigg] \\ &\quad - \bigg( \frac{k_6 \ (1)}{k_3} K_{5 \ (1)} + C \cdot \frac{k_6 \ (2)}{k_3} K_{5 \ (2)} \bigg) \end{split}$$

Table 5. Comparison of observed parameters from mixed-modifier experiments with those calculated from the single-modifier experiments

Modifier	$K_{4(1)} + C \cdot I$	V	<b>7,</b> †	Gradient†		
mixture and ratio	Obs.	Calc.	Obs.	Calc.	Obs.	Calc.
Bu2N+: Me4N+ $C = 40$	$6.5 \times 10^{4}$	5.3 × 10 <sup>4</sup>	0.16	0.19	$3.6 \times 10^{4}$	$3.1 \times 10^{4}$
$Bu^{2}N^{+}:Et_{4}N^{+}$ $C=100$	$1.6 \times 10^{5}$	$1.5 \times 10^{5}$	0.73	0.74	$10.0 \times 10^4$	9.8 × 10 <sup>4</sup>
$Bu_4^n N^+: Pr_4^n N^+$ $C = 2$	$5.7 \times 10^{4}$	$5.1 \times 10^{4}$	0.06	0.04	$5.8 \times 10^{4}$	5.7 × 10 <sup>4</sup>

<sup>\*</sup> Observed values are from plots of  $G/V_m$  against  $[M_1]$ . † Observed values are from plots of  $(1 - V_r)/[M_1]$  against  $V_r$ .

so that a plot of  $(1 - V_r)/[M_1]$  against  $V_r$  will give a gradient of  $K_{5 (1)}[(k_2 + k_{6 (1)})/(k_2 + k_3)] + C \cdot K_{5 (2)}[(k_2 + k_{6 (2)})/(k_2 + k_3)]$  and an intercept (on the abscissa) of  $(k_{6 (1)}/k_3)K_{5 (1)} + C \cdot (k_{6 (2)}/k_3)K_{5 (2)}$ . Hence these values can be calculated and compared with the values obtained from the single-modifier experiments (see Table 4).

Organophosphorus inhibitions. The system to be analysed is shown below; E and M are enzyme and modifier respectively, I is the organophosphorus inhibitor, and EI is phosphonylated enzyme.

$$E + I \xrightarrow{k_i} EI + M$$

$$K_4 \downarrow \uparrow$$

$$EM$$

 $K_4$  has been defined previously, and  $k_i$  is the rate coefficient of inhibition of E by I. Complex formation between E and I, analogous to that between E and S, does occur, but is kinetically unimportant at the inhibitor concn used. It has been shown [6] that for systems of this type:

$$\ln \frac{[E_0] - [EI]}{[E_0]} = \frac{k_i[I]}{(1 + K_4[M])}.t.$$

 $[I] \gg [E_0]$ , and is considered to remain constant. Plots of  $\ln ([E_0] - [EI])/[E_0]$  against t produced a line of gradient  $-k_i/(1 + K_4[M])$ . Values of [M] were chosen (from the single-modifier results) that would reduce  $k_i$  by either a factor of 2 or a factor of 5. Three modifiers,  $Me_4N^+$ ,  $Bu_4^nN^+$  and SAD 128,\* were treated in this way.

#### DISCUSSION

The kinetic schemes developed for single and mixed modifiers have been shown to apply to the experimentally observed results. Furthermore, the parameters obtained from the analysis of the single-modifier experiments have been shown to be in agreement with those obtained from the analysis of the mixed-modifier experiments and those from the analysis of the AChE-modifier-organophosphorus inhibition system.

Reversible enzyme inhibitors can modify the rate of substrate hydrolysis by forming an unreactive complex with free enzyme, or by forming a complex with acylated enzyme which does not deacylate, or does so at a slower rate than uncomplexed acylated enzyme [3]. Examination of Table 4 shows that plots of  $(1 - V_r)/[M]$  against  $V_r$  were not possible for the modifiers  $Me_4N^+$  and  $Et_4N^+$ . This was because  $V_m$  values were constant (Table 1) over the experimental concns used, and indicates that either  $K_5 = 0$ , or that  $k_6 = k_3$ . It is not possible to unambiguously determine which possibility is correct, but since  $K_4 \neq 0$ ,  $K_5$  is unlikely to be zero and therefore  $k_3 = k_6$  is the

CMe<sub>3</sub>·2Cl<sup>-</sup>, and has been used in the treatment of nerve agent poisoning in animals. It has been treated in the same way as other modifiers in the single experiments. The results will be given in more detail in a later publication.

most likely explanation. These findings indicate that Me<sub>4</sub>N<sup>+</sup> and Et<sub>4</sub>N<sup>+</sup> inhibit the hydrolysis of substrate purely by competing with substrate for the active site on the free enzyme, and differ from previous findings [7], which showed that de-acylation occurred at slightly reduced rates with these two modifiers. Values of V, could be calculated for the other three modifiers (Pr4N+, Bu4N+ and Pent4N+) in the homologous series, indicating that inhibition of substrate hydrolysis occurs both because the rate of de-acylation is reduced, and because the concn of free enzyme is reduced due to complex formation  $(K_4)$ values were also able to be calculated). Examination of the  $K_4$  values (Table 4) of these latter three compounds shows that they have a greater affinity for free AChE than either Me<sub>4</sub>N<sup>+</sup> or Et<sub>4</sub>N<sup>-</sup>. Table 4 shows that the parameter  $(k_6/k_3)/[(k_2 + k_3)/(k_2 + k_3)]$  $k_6$ )] is zero for Bu<sub>4</sub>N<sup>+</sup>, i.e. the graph (Fig. 5) for this compound gives a line which intersects the origin, indicating that  $k_6 = 0$ , i.e. no de-acylation of the acylated enzyme-modifier complex occurs. Previous workers found that this complex de-acylated very slowly [7].

In order to obtain any more information from the calculated parameters certain assumptions have to be made. Previous workers [2] have shown that, in the hydrolysis of acetylcholine by electric eel AChE, the rate coefficient of breakdown of the Michaelis complex is between 10<sup>2</sup> and 10<sup>3</sup> times greater than that of the de-acylation of acylated enzyme, i.e. that  $k_2 \gg k_3$ . If this is also the case for bovine erythrocyte AChE, the parameters (Table 4) derived from the gradients and intercepts when  $(1 - V_r)/[M]$ ) is plotted against  $V_r$  can be simplified to  $K_5$  and the ratio  $k_6/k_3$  respectively. Consequently it can be seen that the ratio  $k_6/k_3$  is very small for the modifier PraN+, indicating that de-acylation of the acylated enzyme-modifier complex (EAM) is slow compared to that of the free acylated enzyme (EA). The ratio  $k_6/k_3$  is 0.41 for Pent<sup>n</sup> N<sup>+</sup>, indicating that the rate of de-acylation of EAM is significant compared to that of EA. The simplified parameter leading to  $K_5$ indicates that the three modifiers have similar affinities for free AChE and acylated AChE, and that the values of  $K_5$  are 3-4 times higher than those found by previous workers [7].

Conventional theories of the reversible inhibition of enzymes in the presence of substrate state that compounds which effect only  $K_m$  values (obtained from the slopes in Eadie plots) are termed competitive inhibitors, and those which effect only  $V_m$ values (obtained from the intercepts in Eadie plots) are termed non-competitive inhibitors [8]. Examination of Table 1 shows that the values of G (obtained from the slopes in Eadie plots) for the modifiers Pr<sub>4</sub><sup>n</sup>N<sup>+</sup> and Bu<sub>4</sub><sup>n</sup>N<sup>+</sup> are constant, and so the two modifiers can conventionally be termed noncompetitive inhibitors. However, examination of Table 4 shows that  $K_4$  values are able to be calculated for these two modifiers, indicating that they also form a complex with free AChE, in competition with substrate. It has also been shown that the modifier Bu<sup>n</sup>N<sup>+</sup> competes with an organophosphorus inhibitor for the active site on AChE (Table 3) thereby reducing the expected rate of phosphonylation, and again indicating that the action of this inhibitor is

<sup>\*</sup> SAD 128 is Me<sub>3</sub>C- \(\bigcap \text{N^+-CH<sub>2</sub>OCH<sub>2</sub>-\text{N}}\) -

not purely non-competitive. These findings indicate that conventional theories of reversible enzyme inhibition are in need of modification where enzyme-substrate interactions produce covalent enzyme intermediates.

O-Ethyl S-n-propyl methyl phosphonothiolate, the irreversible inhibitor used to phosphonylate AChE in the presence of various modifiers (Table 3), exists in two optically active forms whose rate coefficients of inhibition of AChE vary by a factor of 15 [9]. Phosphonylation of AChE is therefore caused essentially by the more active enantiomer in this case. From the results shown in Table 3 it is therefore not possible to gain any information on the stereochemical aspects of phosphonylation of AChE in the presence of modifiers, although it is expected that the presence of a modifier will reduce the observed rate of phosphonylation by the same degree irrespective of the configuration of the inhibitor.

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