



Structure dependence in the solvolysis kinetics of amino acid esters

John Haseltine*, Jason W. Runyon†

Department of Chemistry and Biochemistry, Kennesaw State University, Kennesaw, GA 30144, USA

ARTICLE INFO

Article history:

Received 19 March 2010

Accepted 14 April 2010

Available online 18 April 2010

Keywords:

Amino acid esters

Acyl transfer

Crowded amides

Amide polarity

ABSTRACT

To better understand acyl transfer reactions of oligopeptides, seventeen *N*-acyl amino acid esters were solvolyzed in mildly basic methanol-*d*₄. All show pseudo-first-order kinetics by ¹H NMR. The rate constant varies up to 400-fold with the identity of the amino acid and up to 6200-fold with the identity of the *N*-acyl group. The impact of the *N*-acyl group on the rate constant is discussed in terms of crowding, amide conformation, and amide C=O bond character.

© 2010 Elsevier Ltd. All rights reserved.

1. Introduction

We reported previously that under mildly basic solvolysis conditions, some closely related amino acid and oligopeptide esters have strikingly varied half-lives (Eqn. 1 and Table 1).¹ Each compound suffers a clean acyl transfer, yielding the corresponding methyl ester plus ethanol. The hypothesis under consideration was that oligopeptides of different lengths would show different degrees of acyl transfer reactivity. The data show an impact of structural differences that are four to seven bonds distant from the ester group. One would like to understand this impact well given that it is similar to rate/length dependences seen in the proteolysis of oligopeptides. Rate/length dependences in proteolysis are usually rationalized in terms of enzyme/substrate interactions. We have suggested, however, that substrate structure itself plays a larger and more consequential role than formerly suspected.¹

Table 1

Half-life values for ethyl esters undergoing solvolysis to their respective methyl esters in 2.06 M *i*-Pr₂NEt/MeOH at ambient temperature (from Ref. 1)

Compound	<i>t</i> _{1/2} (d)
Piv-Pro-OEt	1700
Piv-Pro-Pro-OEt	220
Ac-Pro-Pro-OEt	30
Piv-Sar-Pro-Pro-OEt	15
Ac-Pro-OEt	2.4

Piv = pivaloyl, Pro = prolyl, Ac = acetyl, Sar = sarcosyl.

* Corresponding author. Tel.: +1 770 499 3426; fax +1 770 423 6744.

E-mail address: jhaselti@kennesaw.edu (J. Haseltine).

† Present address: Department of Chemistry, University of Alabama, Tuscaloosa, AL 35487, USA.



As a next step in our project, we proceeded to inspect the simplest esters more closely. This Letter compares *N*-acyl amino acid esters undergoing acyl transfer. Even in such small compounds, the variation of reactivity with structure is pronounced and intriguing.

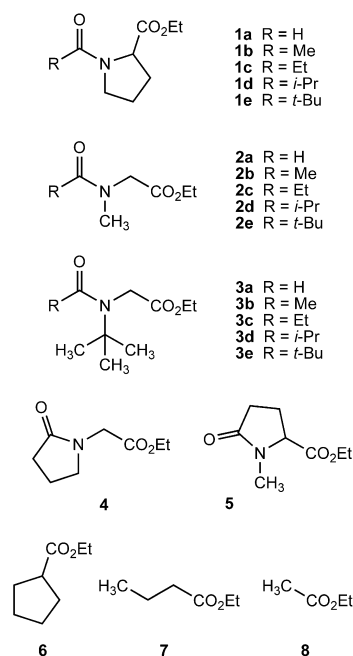


Figure 1. *N*-Acyl amino acid esters subjected to base-promoted solvolysis.

2. Results and discussion

Each of the esters in Figure 1 converts cleanly to its respective methyl ester plus ethanol under mild conditions (0.1 M ester and 1.03 M *i*-Pr₂NEt in methanol-*d*₄, 21.5 ± 1 °C).^{2–6} The reactions were monitored directly by ¹H NMR. The pseudo-first-order rate constant was calculated from successive integrations of the diminishing –CO₂CH₂CH₃ signal of the reactant or the growing DOCH₂CH₃ quartet.⁷ The results are listed in Table 2.

Series 1–3 differ in their absolute reactivity and in their rate constant patterns. Comparisons with control compounds show that the amide group can boost ester reactivity significantly. The rate constants for **1a** and **2a** are about 100-fold and 85-fold greater than those for the similar simple esters ethyl cyclopentanecarboxylate (**6**) and ethyl butanoate (**7**). The rate constant varies significantly with the identity of the *N*-acyl group, and most strongly in series 1.

The acyl transfer mechanism probably involves a base-assisted pyramidalization of the ester group by the solvent.^{8,9} Neighboring-group participations are unlikely. Deuteration next to the amide carbonyl was not evident in any of the kinetics trials. Also, when the base was omitted from the reaction, the rate constants were greatly reduced. This control experiment was run for all amide-esters except **1e** and **3e**. Conventional neighboring-group participations should show at least one of two consequences. They should show deuteration if amide enolization is occurring (not expected under our mild conditions). Alternatively, if the neutral amide group were the nucleophile, *k* should be largely unaffected by omitting the base. For most compounds in the study, deuteration next to the ester carbonyl was also either not evident or its rate constant was less than one-half that for acyl transfer. Enolization of the ester group and/or ketene formation is therefore not likely a part of the acyl transfer mechanism for those compounds.¹⁰ As for the remaining compounds, kinetic isotope data suggest that the ester group is not deprotonated as a part of acyl transfer events.

One possible explanation for the variation of *k* within each ester series is that the ester group is directly crowded to different ex-

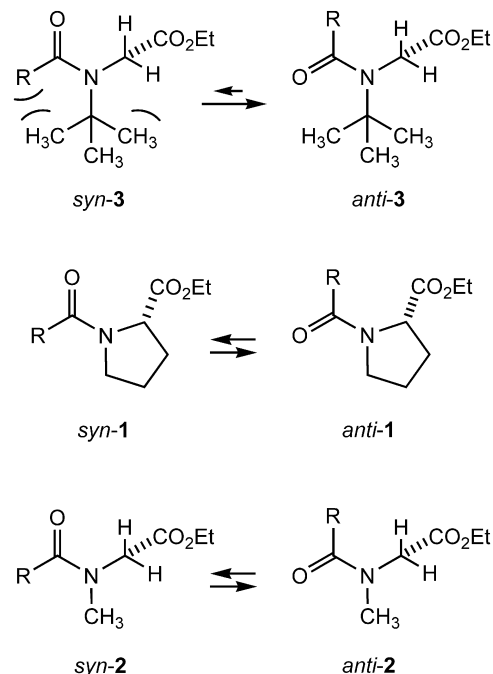


Figure 2. Amide conformation in series 1–3 according to NMR data and ab initio calculations.

tents by the different *N*-acyl groups. This explanation is most credible for series 3 since the bulky *N*-*t*-butyl group should generally favor conformations that have the *N*-acyl R group and the ester group close to each other (Fig. 2). The largest R groups might then interfere the most with ester pyramidalization and sponsor the lowest rate constants.

The explanation does not apply as neatly to series 1 and 2, however, since those series do not favor an amide conformation in which the *N*-acyl R group and the ester group are close to each other. Proton NMR integrations and carbon chemical shifts indicate the alternative amide conformation to be slightly favored for most of series 1 and 2 (*syn*-1 and *syn*-2 in Fig. 2). Therefore, for mutual crowding of the *N*-acyl and ester groups to be the main source of variation in *k*, the anti conformation would need to be the more reactive conformation for most of each series, and substantially more reactive than the *syn* conformation. The possibility is not ruled out. Note that ester **3a** has a smaller rate constant than **3b–d**. This is interesting because esters **1a** and **2a**, also bearing the smallest *N*-acyl group, each show the largest rate constants in their series. The predominant amide conformation in **3a** is evidently *syn*-**3a**, with NOE measurements supporting a prediction from ab initio modeling (gas phase; 6–31G* level). Therefore, the anti conformation may indeed have more of an activating effect on ester reactivity.

To test the importance of amide conformation more plainly, esters **4** and **5** were prepared and solvolyzed. The lactam rings in **4** and **5** lock the amide linkage into *syn* and *anti* conformations, respectively. The rate constants are shown in Table 2. Ester **5** is slightly more reactive than **4**, despite having more branching next to its ester carbonyl. The impact of this branching should be mild. The rate constants for esters **6–8** imply that each alkyl substituent next to the ester carbonyl in that control series lowers *k* by a factor of about 3. Thus the data are consistent with the anti conformation being more activating than *syn*, if only slightly so.

A different explanation for the variation of *k* in each ester series involves the activating effect of the amide group. Electron-withdrawing groups in esters are known to facilitate acyl transfer.

Table 2
Rate constants for ester solvolysis, IR frequencies of the amide carbonyl, and NMR chemical shifts of the amide carbonyl

Compound	<i>k</i> (10 ^{–5} s ^{–1})	ν _{C=O} ^a (cm ^{–1})	δ _{C=O} ^{b,c} (ppm)
1a	3.3	1663	160.8, 161.7
1b	0.099	1644	169.5, 169.7
1c	0.044	1644	172.5, 172.7
1d	0.034	1645	176.0, 176.5
1e	0.00053	1622	176.9
2a	8.1	1668	163.0, 163.2
2b	2.6	1646	171.2, 171.5
2c	1.7	1648	174.3, 174.6
2d	1.3	1646	177.5, 177.6
2e	0.22	1631	178.1
3a	0.10	1654	161.8
3b	0.32	1651	171.8
3c	0.22	1652	174.6
3d	0.22	1640	178.6
3e	0.0045	1628	178.7
4	2.1	1682	175.7
5	3.6	1688	175.2
6^d	0.031	—	—
7^e	0.094	—	—
8^f	0.27	—	—

^a Obtained for neat compounds.

^b Obtained for CDCl₃ solutions.

^c Values are given for both amide conformations, if observed.

^d Ethyl cyclopentanecarboxylate.

^e Ethyl butanoate.

^f Ethyl acetate.

Download English Version:

<https://daneshyari.com/en/article/5271629>

Download Persian Version:

<https://daneshyari.com/article/5271629>

[Daneshyari.com](https://daneshyari.com)