SOME REACTIONS OF PHENOLIC MALEAMIC ACIDS RELEVANT TO CARBOXYPEPTIDASE CATALYSIS

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Summary: In neutral aqueous acetonitrile, N-o-hydroxybenzyl-2,3-dimethylmaleamic acid is rapidly converted to N-o-hydroxybenzyl-2,3-dimethylmaleimide, while N-propyl-N-o-hydroxybenzyl-2,3-dimethylmaleamic acid is converted to dimethylmaleic anhydride; both reactions are catalyzed by the phenolic groups.

The enzyme carboxypeptidase A cleaves peptides with the assistance of three catalytic groups: the carboxylate ion of a glutamate residue, the phenolic hydroxyl of a tyrosine, and a zinc ion. For this reason we have been investigating molecules in which a carboxylate ion and a phenol could cooperate in cleaving an intramolecular amide linkage. A few years ago a report from our laboratory¹ described the change which occurred when one such species, compound (1), was allowed to react in 1 \underline{M} H₂O/CH₂CN under acidic and neutral conditions. Under acidic conditions the well-precedented² conversion of maleamic acid (1) to the maleic anhydride (2) and amine was observed, but at neutrality a new process took over in which the anhydride was not produced. This process was promoted by the phenolic group, and was proposed¹ to be a hydrolysis of the amide group by a general base mechanism catalyzed by carboxylate and phenol. On further investigation, we find that the products of this new process were mis-identified, and the reaction at neutrality is actually a closure of the maleamic acid ($\underline{1}$) to the This closure³ can be blocked by N-propylation in compound (4); maleimide (3). under both neutral and acidic conditions (4) reacts to form the anhydride (2). At neutrality, the maleimide formation from $(\underline{1})$ and the anhydride formation from (4) are catalyzed by the phenol groups.

Maleamic acids (<u>1</u>) and (<u>4</u>) were obtained, as the amine salts, by reaction of two equiv. of the appropriate amine⁴ with dimethylmaleic anhydride. An authentic sample of the maleimide (<u>3</u>), (m.p. 79-80[°], M + 1 = 232, ¹H NMR in CDCl₃ δ 1.98 (6H, s), 4.63 (2H, s), 7.50 to 6.73 (4H, m); UV λ_{max} in H₂O, 270 nm (ε 8,000) 310 nm (ε 500); IR 1700 cm⁻¹) was obtained from (<u>1</u>) by heating^{3(c)} (150[°], 3 hrs). Kinetic studies at 25.0[°] were performed, as described¹, in 1 M

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 H_2O/CH_3CN with 5 x 10⁻³ <u>M</u> HOAc and 5 x 10⁻⁴ <u>M</u> KOAc (10:1, acidic), 5 x 10⁻⁴ <u>M</u> HOAc and 5 x 10⁻³ <u>M</u> KOAc (1:10, neutral), and 5 x 10⁻³ <u>M</u> KOAc with no added HOAc (referred to as 1:50, neutral). The absorbance was followed at 250 nm, which is λ_{max} for dimethylmaleic anhydride. In all cases the kinetic runs gave good first-order plots for at least two half lives.

In the acidic buffer, compound $(\underline{4})$ underwent ring closure to form the anhydride $(\underline{2})$ (increase in absorbance) with $k = 12.0 \pm 0.5 \times 10^{-4} \text{ s}^{-1}$; this decreased by only a factor of 5, to $k = 2.5 \pm 0.5 \times 10^{-4} \text{ s}^{-1}$, in the neutral (1:10 and 1:50) buffers, in which closure to form the anhydride was followed by a slower decrease in absorbance as the anhydride hydrolyzed. Since we had found¹ that in maleamic acids without a phenolic catalytic group the rates of ring closure decrease by ca. 10^{3} in proceeding from the 10:1 to the 1:50 buffer system, the phenolic group in $(\underline{4})$ is contributing more than a factor of 10^{2} to the ring-closure process at neutrality. This catalysis is presumably due to protonation of the leaving group nitrogen by the phenol, as shown.

In the acidic buffer, compound $(\underline{1})$ also underwent closure to dimethylmaleic anhydride $(\underline{2})$ with an increase in absorbance at 250 nm, as we had reported.¹ We find that this process has $k = 20.0 \pm 0.5 \times 10^{-4} s^{-1}$ (rept'd.¹ 45.5 $\times 10^{-4} s^{-1}$). The product $(\underline{2})$ was further identified by high performance liquid chromatography (HPLC). When the reaction mixture was allowed to stand, this first anhydride product $(\underline{2})$ disappeared and was replaced by the maleimide $(\underline{3})$, identified by HPLC and UV (broad maximum at 310 nm). In the neutral buffers the anhydride $(\underline{2})$ was not formed, even as a transient intermediate. (We have reported¹ that $(\underline{2})$ could not be trapped with amines present in the reaction mixture which successfully trap authentic $(\underline{2})$ under the same conditions, and this has been confirmed.⁵) Instead, it underwent a new process with $k = 1.7 \times 10^{-4} s^{-1}$, in agreement with our previous report¹ of $k = 1.9 \pm 0.1 \times 10^{-4} s^{-1}$, with a steady decrease in absorbance at 250 nm. The product of this process was the maleimide $(\underline{3})$, identified by HPLC and UV spectra. It was formed quantitatively (HPLC) at the expected rate determined from the UV kinetic studies.

As reported¹, this new reaction of (<u>1</u>) at neutrality is ca. 70 times as rapid as is the disappearance of an analog of (<u>1</u>) without the phenolic group under the same conditions. Thus at neutrality the phenol group catalyzes maleimide formation. The most likely mechanism is that shown, involving the kinetically equivalent carboxylic acid/phenoxide state (<u>5</u>) of the substrate. Less likely alternatives would involve reversible formation of a tetrahedral intermediate, with phenol catalysis in a subsequent rate-determining step. Curiously, in acid the closure of (<u>1</u>) to the anhydride is <u>faster</u> than is the closure of (<u>4</u>), but at neutrality (<u>4</u>) continues to form anhydride while (<u>1</u>) substitutes a new process, closure to the maleimide; however, the reaction of (<u>1</u>) is now <u>slower</u> than the reaction of (<u>4</u>), so the anhydride formation from (<u>1</u>), no longer the major process, must have a greatly depressed rate compared with that of $(\underline{4})$.

We have proposed⁶ that the cleavage of peptide substrates by carboxypeptidase A occurs by a general base mechanism, while Makinen⁷ has demonstrated that cleavage of an ester substrate occurs by a nucleophilic mechanism, with formation of an intermediate anhydride. It is now clear that the chemistry of (<u>1</u>) is not related to the general base mechanism for catalysis by carboxypeptidase A, as had been suggested¹. Furthermore, compound (<u>4</u>) continues to use the nucleophilic mechanism, with phenol catalysis, even at neutrality. Thus other models must be sought for the mechanism we have proposed⁶ for the cleavage of amides by a carboxylate and a phenol group (and a zinc atom) in carboxypeptidase catalysis.⁸

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