

THE DUAL NATURE OF THE REACTION BETWEEN PORCINE ELASTASE
AND HUMAN PLASMA α_1 PROTEINASE INHIBITOR*S. Satoh[†], T. Kurecki, L.F. Kress and M. Laskowski, Sr.Laboratory of Enzymology, Roswell Park Memorial Institute
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Received November 13, 1978

SUMMARY: Only a portion of human plasma α_1 proteinase inhibitor (α_1 PI) forms a 1:1 complex with porcine elastase; the other portion is inactivated via proteolysis. High temperature (37°) and high salt (2 M) enhance complex formation. The complex is unstable, but no significant liberation of active elastase could be demonstrated. Probably the same two major products of ~50,000 and ~4,000 daltons are formed from α_1 PI via proteolysis and via disintegration of the complex. Iodination of α_1 PI or oxidation with chloramine-T prevents complex formation with elastase but not with trypsin. Iodinated elastase, however, forms a complex with α_1 PI.

Recently, it was reported that about one-half of α_1 PI formed a 1:1 complex with porcine elastase. The other half was hydrolyzed to form one large and one small fragment (1). Independently, several laboratories showed that α_1 PI was susceptible to limited proteolysis by a variety of proteinases (2-5), and that the inactivating cleavage occurred in the same region as that observed during inactivation by trypsin (6,7). It was decided to extend our studies of α_1 PI (6) to the pair, α_1 PI-elastase, hoping to find conditions that suppress or enhance the pathway leading to complex formation.

EXPERIMENTAL PROCEDURES

α_1 PI was prepared as described (6). Porcine trypsin was from Novo. Elastase was prepared according to Shotton (8) and further purified from trypsin and chymotrypsin by passing it through an affinity column containing immobilized soybean trypsin inhibitor and a second column containing turkey ovomucoid to remove elastase II (9,10). The final elastase preparation had no detectable tryp-

* Supported by grants HL15892, HL22996 and GM24950 from the National Institutes of Health, BMS73-06750 from the National Science Foundation, and a contract from DOE EY76S023225.

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sin, chymotrypsin, or elastase II activity and was devoid of any contaminant capable of direct proteolysis of α_1 PI. Molarity was calculated using $E_{1\%}^{1\text{cm}, 280\text{ nm}} = 20.2$ (elastase), 5.0 (α_1 PI), and 14.9 (trypsin); the latter was corrected for active trypsin (11). For titration curves elastase solution containing a specified amount in 0.2 ml of phosphate buffer, pH 6.8, was mixed with 0.8 ml containing variable amounts of α_1 PI in the same buffer. The mixture was preincubated 60 min at 1°, 10 min at 25° or 37°. An aliquot (0.5 ml) was withdrawn to the cuvette containing substrate and the remaining elastase activity determined by the method of Visser and Blout (12) using a Cary Model 16 Spectrophotometer at 25°. Trypsin was determined according to Schwert and Takenaka (13). In the experiments using gels α_1 PI and elastase were mixed in predetermined ratio and temperature in a total volume of 10 μ l. To stop the reaction quickly 2 μ l of a mixture containing 10% sodium dodecyl sulfate (SDS) and 10% 2-mercaptoethanol (kept in a boiling water bath) were added, and the tube was immediately transferred to a boiling water bath for 5 min. After cooling 2 μ l of fluorescamine solution (5 mg/ml acetone) were added, followed by 1 μ l of 50% sucrose solution. One μ l of this mixture was applied to an appropriate channel of the gradient gel slab (3).

Elastase- α_1 PI complex was prepared by gel filtration of the reaction mixture (α_1 PI/elastase = 2) through a column of Biogel P-150 (1.5 x 90 cm). The first peak contained a polymer of the ~50,000 dalton fragment. The second peak contained crude complex. Similar distribution has been observed with α_1 PI-chymotrypsin complex (14). Crude complex was further purified on DEAE Biogel A10 with a gradient of 0 to 0.15 M NaCl in 0.01 M sodium phosphate, pH 7.5. The peak containing the complex was concentrated with Ficoll 400. The complex was identified by gel electrophoresis with and without SDS (not shown).

Iodination was performed as described (15).

RESULTS AND DISCUSSION

Fig. 1 shows the difference in shape between the titration curves of trypsin and elastase with α_1 PI. Titration of trypsin is a typical straight line representing uni-unimolecular reaction. From the curving near the equivalence point the dissociation constant can be calculated (16). With elastase the intercept and the convex shape of the titration curves depend on temperature and salt concentration. If the inactivation curve results from two independent reactions, one leading to complex formation, the other to direct proteolysis, the convex shape and variability of the intercept would be expected. Lowering of temperature increases both the convexity and the value of the intercept, suggesting that the rate of complex formation is affected more than the rate of

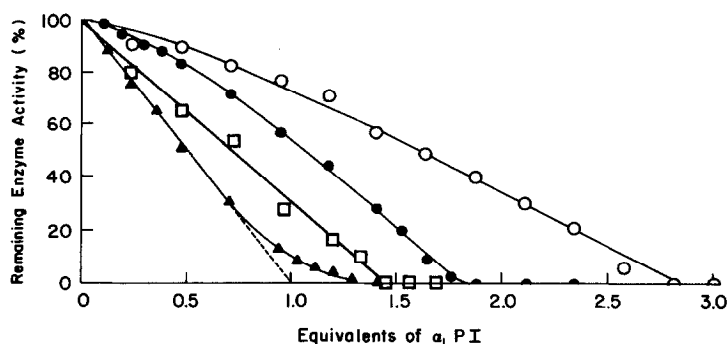


Fig. 1. Titration curves of trypsin (▲—▲) at 25°, and of elastase (○—○) at 25° and (◻—◻) at 37°. Trypsin 6.66×10^{-10} moles was titrated in 0.05 M Tris-HCl, pH 8.0 containing 1 mM CaCl_2 . Elastase (6.72×10^{-10} moles) titrations were in 0.05 M phosphate buffer, pH 6.8, except that the 37° (◻—◻) titration was made in the presence of 2 M NaCl, and corrected for the decrease in elastase activity in NaCl.

proteolysis. Under the most favorable conditions (37° and 2 M NaCl) only 1/3 of $\alpha_1\text{PI}$ is directly digested, whereas 2/3 forms the complex. This suggests that hydrophobic forces enhance complex formation with elastase. A similar conclusion was reached from experiments with dioxane (17).

The dual nature of elastase and $\alpha_1\text{PI}$ reaction is also shown by gradient gel electrophoresis (3). The experiments were performed using an $\alpha_1\text{PI}$ /elastase ratio of 1.6, at two different temperatures, 37° (Fig. 2A) and 25° (Fig. 2B). At 37° (without NaCl) only a trace of free elastase should be present, and indeed no band of free elastase is visible (Fig. 2A). The $\alpha_1\text{PI}$ band (54,000 daltons) is visible for about 30 sec. Only one type of complex (80,000 daltons) corresponding to 1:1 combination is immediately formed and is already visible at 10 seconds but disappears after 30 min. Two fragments of $\alpha_1\text{PI}$, ~50,000 and ~4,000 daltons are formed simultaneously, become visible at 30 sec, and persist throughout the experiment, suggesting that they are relatively stable to further digestion by elastase. At 25° (Fig. 2B) with the

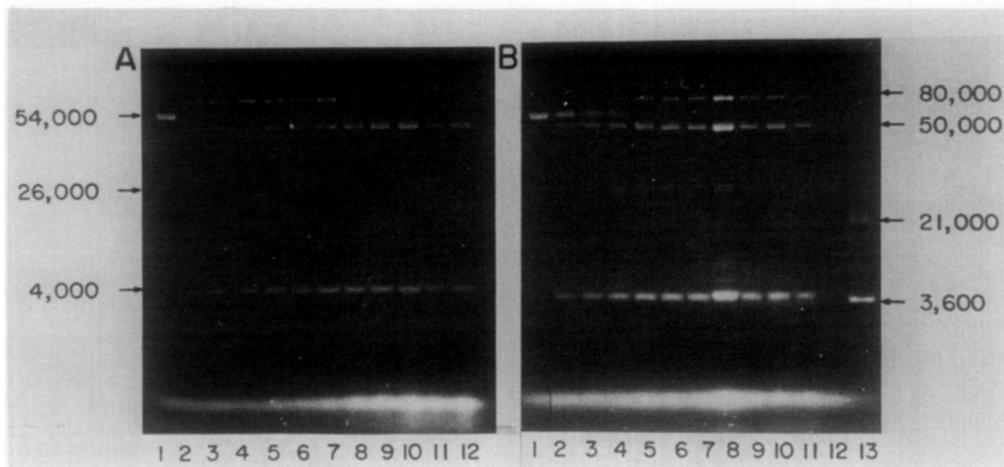


Fig. 2A. Progress of reaction between elastase and α_1 PI at 37° . Electrophoresis was run overnight at 100 volts (constant voltage); channel 1) markers: α_1 PI and elastase 2) 10 sec incubation 3) 30 sec, 4) 1 min, 5) 5 min, 6) 10 min, 7) 30 min, 8) 1 h, 9) 2 h, 10) 5 h, 11) 15 h, 12) 24 h. *2B.* Similar experiment at $\sim 1^\circ$. Channel 1) markers: α_1 PI and elastase 2) 30 sec incubation 3) 1 min, 4) 5 min, 5) 10 min, 6) 30 min, 7) 1 h, 8) 2 h, 9) 5 h, 10) 15 h, 11) 24 h, 12) elastase only, 13) markers: soybean trypsin inhibitor (Kunitz) and B chain of insulin.

ratio 1.6 about 50% of the elastase should be free (see Fig. 1), and indeed free elastase is seen throughout the experiment (Fig. 2B). Intact α_1 PI persists longer than in Fig. 2A. The complex (80,000 daltons) forms more slowly but remains visible for 24 hours. Neither at 37° nor at 1° are bands observed in the regions between 80,000 and 54,000 daltons. This contrasts with the α_1 PI-trypsin reaction (6,7) and earlier reports on α_1 PI-elastase reaction (1,18). An increase in intensity of the bands in the range of a few hundred daltons is apparent in Fig. 2. This is probably due to side reactions, one of which is instability during SDS 2-mercaptoethanol treatment, because a weak band is seen even with markers. Additional experiments of this type in which the α_1 PI/elastase ratio was 1 or 2, and temperature was either 1° or 37° were performed (data not shown). The conclusion was the same;

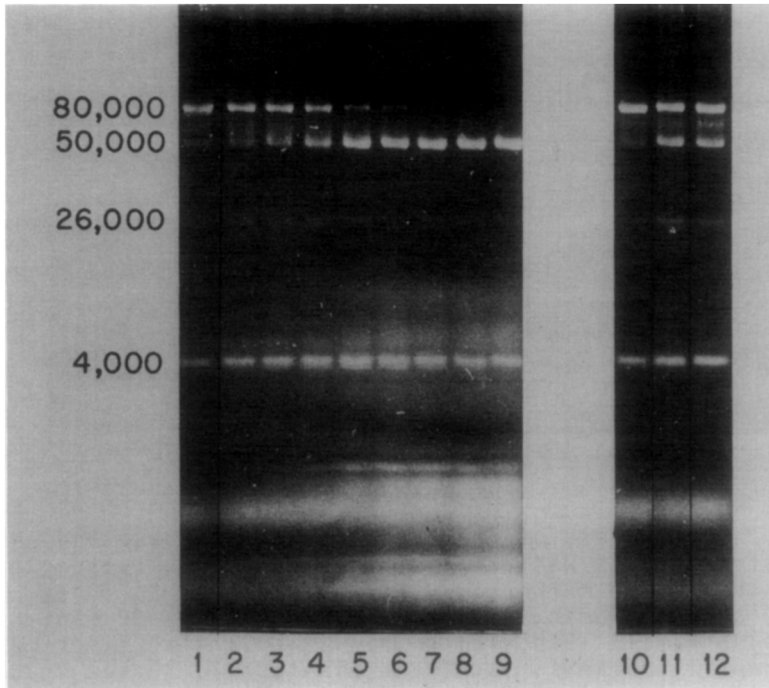


Fig. 3. Stability of the isolated α_1 PI-elastase complex in the presence and absence of free elastase at 37°. Complex (2.67 A_{280} units) and 5 μ g elastase were incubated in 50 μ l of 0.05 M sodium phosphate, pH 7.0. Electrophoresis as in *Fig. 2*. Channel 1) 10 sec incubation, 2) 30 sec, 3) 1 min, 4) 5 min, 5) 10 min, 6) 30 min, 7) 1 h, 8) 2 h, 9) 5 h; 10-12) controls incubated at 37° without added elastase: 10) 30 min, 11) 1 h, 12) 5 h.

namely, that 2 independent reactions occur. *Fig. 3* shows an experiment in which isolated complex was treated with elastase and incubated at 37°. The results confirm that $\sim 50,000$ and $\sim 4,000$ dalton major bands are formed from the complex, and that the degradation reactions are accelerated by added elastase.

Experiments involving radioiodination of either α_1 PI or elastase were then performed. The method of iodination requires simultaneous oxidation (15). Labeling of α_1 PI decreased its activity against trypsin by only 20%, but abolished its ability to form a complex with elastase. Formation of α_1 PI fragments of $\sim 50,000$ and $\sim 4,000$ daltons by elastase was not affected (data not

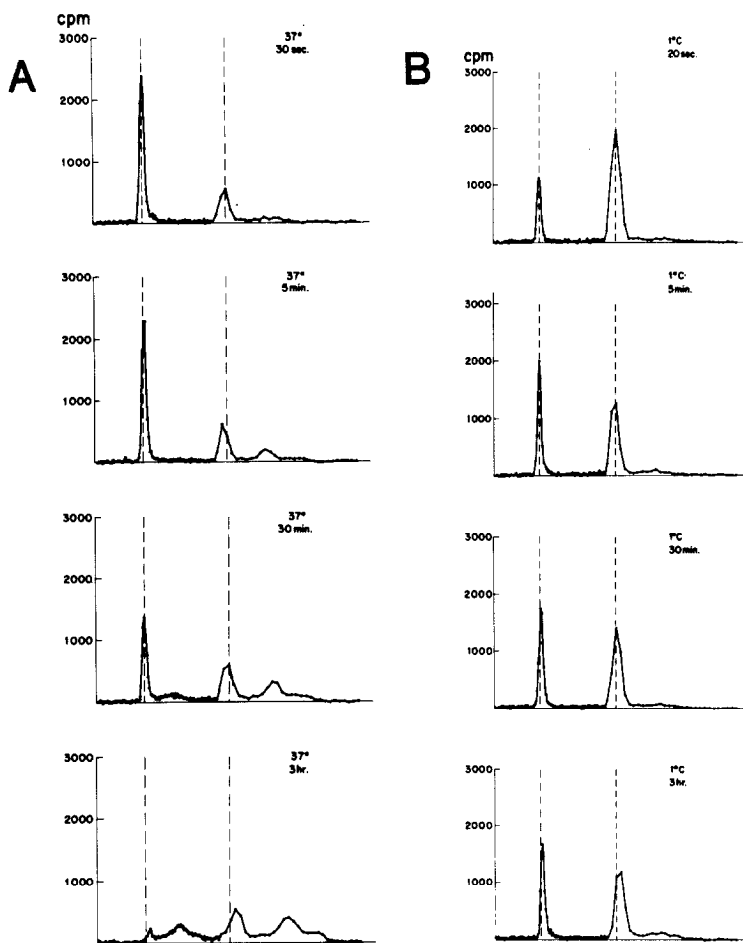


Fig. 4A. Reaction between ^{125}I -elastase and native $\alpha_1\text{PI}$ at 37° at $\alpha_1\text{PI}$ /elastase ratio = 2. At indicated time intervals the aliquots were withdrawn, terminated and electrophoresed for 5 h at 10 mA per tube (constant current). The gels were sliced into 0.5 mm fragments with a Gilson automatic gel slicer. The fragments were placed into scintillation vials and radioactivity was counted using a Packard Auto-Gamma Spectrometer. Two dotted lines were drawn at 80,000 and 26,000 daltons (the peak of complex at 30 sec and the peak of elastase). *4B.* Similar experiment to that in *4A* except that the temperature of the reaction was $\sim 1^\circ$.

shown). Oxidation of $\alpha_1\text{PI}$ with chloramine-T without iodination was sufficient to produce the same effects. Since only methionine is oxidized under the conditions used (19), this suggests that methionine is at the reactive center of $\alpha_1\text{PI}$ for elastase, and agrees with a recent independent conclusion (20). The reason for the different behavior of oxidized or iodinated $\alpha_1\text{PI}$ vs trypsin

remains temporarily unresolved.

Iodination of elastase, however, did not prevent complex formation. At 37° complex formed rapidly and disappeared within 3 hours (Fig. 4A). The elastase peak remained almost constant, but the counts increased in the regions between 80,000 and 26,000 daltons. This was further evidence that only a single α_1 PI-elastase complex is formed under the conditions used. The formation of products in both regions is due to direct digestion of the complex, because a control experiment with elastase alone showed only ~5% loss of activity during the 3-hour incubation. At 1° (Fig. 4B) complex forms slowly, never reaches the same height as at 37°, but lasts much longer. The amount of free elastase first decreases, and then stays constant. There is little radioactive material in the region from 80,000 to 26,000, but some accumulation of radioactive material in the region below 26,000 daltons is detectable. Neither at 37° nor at 1° is there any accumulation of counts in the regions corresponding to 50,000 and 4,000 daltons. The disintegration of complex (Fig. 3) proceeds by either cleaving the α_1 PI moiety into 50,000 and 4,000 dalton fragments, or by cleaving the elastase moiety at several sites (Figs. 3 and 4). Both processes apparently occur simultaneously.

ACKNOWLEDGMENTS

This work is dedicated to the memory of Mr. Lubomyr Stasiuk, a long time friend and colleague.

We are grateful to Dr. N. Tanigaki of the Immunology Research Department for his help in the radioiodination experiments, and to Dr. K. Oda (present address College of Agriculture, Laboratory of Applied Microbiology, University of Osaka Prefecture, Sakai, Osaka, Japan) who in 1977 made the first observation that temperature affects complex formation between elastase and α_1 PI.

REFERENCES

1. Baumstark, J.S., Lee, C.T. and Luby, R.J. (1977) *Biochim. Biophys. Acta* 482, 400-411.
2. Johnson, D. and Travis, J. (1977) *Biochem. J.* 163, 639-641.

3. Kurecki, T., Laskowski, M., Sr. and Kress, L.F. (1978) *J. Biol. Chem.* **253** (in press).
4. Kress, L.F. and Paroski, E.A. (1978) *Biochem. Biophys. Res. Commun.* **83**, 649-656.
5. Oda, K., Morihara, K. and Tsuzuki, H. (1978) *Infec. Immun.* (in press).
6. Oda, K., Laskowski, M., Sr., Kress, L.F. and Kowalski, D. (1977) *Biochem. Biophys. Res. Commun.* **76**, 1062-1070.
7. Moroi, M. and Yamasaki, M. (1974) *Biochim. Biophys. Acta* **359**, 130-141.
8. Shotton, D.M. (1970) *Methods in Enzymology* **19**, 113-140.
9. Ardelt, W. (1974) *Biochim. Biophys. Acta* **341**, 318-326.
10. Gertler, A., Weiss, Y. and Burstein, Y. (1977) *Biochemistry* **16**, 2709-2715.
11. Chase, T. and Shaw, E. (1967) *Biochem. Biophys. Res. Commun.* **29**, 508-514.
12. Visser, L. and Blout, E.R. (1972) *Biochim. Biophys. Acta* **268**, 257-260.
13. Schwert, G.W. and Takenaka, Y. (1955) *Biochim. Biophys. Acta* **16**, 570-575.
14. Bloom, J.W. and Hunter, M.J. (1978) *J. Biol. Chem.* **253**, 547-559.
15. McConahey, P.J. and Dixon, F.J. (1966) *Int. Arch. Allergy* **29**, 185-189.
16. Green, N.M. and Work, E. (1953) *Biochem. J.* **54**, 347-352.
17. Cohen, A.B. (1975) *Biochim. Biophys. Acta* **391**, 193-200.
18. Lo, T.N., Cohen, A.B. and James, H.L. (1976) *Biochim. Biophys. Acta* **453**, 344-356.
19. Shechter, Y., Burstein, Y. and Patchornik, A. (1975) *Biochemistry* **14**, 4497-4503.
20. Johnson, D. and Travis, J. (1978) *J. Biol. Chem.* **253**, 7142-7144.