## Thymosin  $\alpha_{11}$ : A peptide related to thymosin  $\alpha_1$  isolated from calf thymosin fraction 5

(thymic peptides/amino acid sequences/Candida albicans infection/thymosin  $\alpha_1$  analogues)

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ABSTRACT Two peptides related to thymosin  $\alpha_1$  have been isolated from preparations of calf thymosin fraction 5. One, lacking four amino acid residues at the COOH terminus, is designated des-(25-28)-thymosin  $\alpha_1$ . The other, named thymosin  $\alpha_{11}$ , contains seven additional amino acid residues at the COOH terminus. The sequence of this peptide is: AcSer-Asp-Ala-Ala-Val-Asp-Thr-Ser-Ser-Glu-Ile-Thr-Thr-Lys-Asp-Leu-Lys-Glu-Lys-Lys-Glu-Val-Val-Glu-Glu-Ala-Glu-Asn-Gly-Arg-Glu-Ala-Pro-Ala-AsnOH. Thymosin  $\alpha_{11}$ , in doses of <300 ng per mouse, protects susceptible inbred murine strains against opportunistic infections with Candida albicans. It is  $\approx$  30 times as potent as thymosin fraction 5 and approximately equal in potency to thymosin  $\alpha_1$ .

A preparation from calf thymus designated thymosin fraction  $5(1, 2)$  has been shown to enhance parameters of immune function in several in vivo and in vitro systems (for reviews, see refs. 3 and 4). Analytical isoelectric focusing (IEF) and gel filtration chromatography of thymosin fraction 5 indicated the presence of as many as 30 distinct peptides, ranging in isoelectric points from pH 4.0 to pH 7.0 and in molecular weights from 1,000 to 15,000 (5, 6). The first peptide to be isolated from thymosin fraction 5 was thymosin  $\alpha_1$ ; sequence analysis showed this peptide to contain 28 amino acid residues, corresponding to a  $M_r$ of 3,107 (5). The sequence was confirmed by chemical synthesis (7). Thymosin  $\alpha_1$  was reported to exhibit many, but not all, of the biological activities of thymosin fraction 5 (6).

The biosynthesis of thymosin  $\alpha_1$  appears to involve the formation of a larger precursor polypeptide,  $M_r = 16,000$ , based on experiments in which calf thymus mRNA was translated in vitro (8, 9). The identity of proteinases involved in the processing of this putative precursor remains unknown.

We report here the isolation from calf thymus fraction 5 of two peptides structurally related to thymosin  $\alpha_1$ . One is a shorter peptide derived from residues 1-24 of thymosin  $\alpha_1$ , designated des-(25-28)-thymosin  $\alpha_1$ . The other, named thymosin  $\alpha_{11}$ , is a longer peptide composed of 35 amino acid residues, the first 28 of which are identical to thymosin  $\alpha_1$ . The sequence of the COOH-terminal extension in thymosin  $\alpha_{11}$  is Gly-Arg-Glu-Ala-Pro-Ala-AsnOH. We also report that thymosin  $\alpha_{11}$  can enhance resistance to intravenous challenge with Candida albicans to a degree approximately equal to that observed with thymosin  $\alpha_1$ .

## EXPERIMENTAL PROCEDURES

Materials. All chemicals and solvents employed were chromatography grade. Trypsin (TPCK-treated), Staphylococcus aureus V-8 protease, carboxypeptidase Y, and Polybrene were from Worthington, Miles, Sigma, and Pierce, respectively.

Thymosin fraction 5 (batches 577 and 677), prepared by the Roche Biopolymer Department, was generously provided by Courtney McGregor. The pentapeptide Glu-Ala-Pro-Ala-AsnOH was synthesized by the solid-phase method (10).

Methods. Digestion with proteinases was carried out as described in the tables. Separation of peptides by HPLC was performed with an Ultrasphere ODS C18 column (5  $\mu$ m, 4.6  $\times$  250 mm, Altex, Berkeley, CA) with a fluorescamine detection system as described by Stein and Moschera (11). Amino acid analyses and the manual Edman degradation procedures were carried out as described (12), except that for the latter, Polybrene (1 mg) was added to the peptide solutions. Automated Edman degradation was by the method of Hewick et al. (13).

Preliminary separation of the components of thymosin fraction 5 was by preparative IEF (14, 15). After electrofocusing for <sup>17</sup> hr at <sup>a</sup> maximal current of <sup>20</sup> mA and <sup>a</sup> maximal voltage of 1.1 kV, the gel bed was divided into 30 sections with a stainless steel grid and the peptides in each fraction were eluted with 5 ml of water. The pH of each eluate was determined with <sup>a</sup> Radiometer PHM <sup>83</sup> Autocal pH meter.

For analysis of IEF fractions by HPLC, aliquots were lyophilized and dissolved in <sup>a</sup> small volume of buffer A (0.2 M pyridine/1.0 M HCOOH). Elution was with buffer A and <sup>a</sup> linear gradient of 1-propanol.

Fungus. The isolate of C. albicans, originally obtained from <sup>a</sup> human patient, was maintained on Sabouraud's dextrose agar slants at room temperature. For inoculation into mice, the yeast cells were grown in Sabouraud's dextrose broth on <sup>a</sup> shaker at 37°C for 18-20 hr.

Mouse Protection Assay. The inbred strain of mice  $(C_3H/HeJ)$ was purchased from The Jackson Laboratory. The mice were inoculated daily intraperitoneally, each with 0.5 ml of thymosin fraction 5 or one of the purified thymosins therefrom, beginning 2 days before challenge with C. albicans and continuing daily until the termination of the experiment. Each mouse was challenged intravenously with  $4 \times 10^4$  viable cells of C. albicans and the numbers of fungus cells in the left kidneys were determined quantitatively at specified times after infection by grinding the tissue in sterile sand and plating increasing dilutions of the suspensions on Sabouraud's agar. The numbers of colonies of C. albicans were counted after incubation of the plates for  $72$  hr at  $37^{\circ}$ C  $(15)$ .

## RESULTS

**Isolation of Peptides by HPLC.** Thymosin  $\alpha_1$  was found to be present in fractions 4-7 from the IEF separations, corresponding to <sup>a</sup> pH range of 3.72-3.97. In addition to thymosin  $\alpha_1$  (peak b, Fig. 1), two other peptides were recovered that

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Abbreviation: IEF, isoelectric focusing.



FIG. 1. Separation by HPLC of peptides in the IEF fractions. Calf thymosin fraction 5 (2.0 g, batch 577) was fractionated by IEF. Each of the 30 fractions obtained was analyzed by HPLC as described (14, 15). The results are shown for IEF fractions 6 and 7 by using a linear gradient of 1-propanol in buffer A as shown  $(--)$ . Peaks a, b, and c were identified as des-(25-28)-thymosin  $\alpha_1$ , thymosin  $\alpha_1$ , and thymosin  $\alpha_{11}$ , respectively, as described in the text. Some additional thymosin  $\alpha_1$  was recovered from IEF fractions 4 and 5 and a small quantity of thymosin  $\alpha_{11}$  was also present in IEF fraction 5 (data not shown).

emerged slightly before and slightly later than thymosin  $\alpha_1$  (peaks a and c, Fig. 1). The recoveries from 2.0 g of thymosin fraction 5 (batch 577) were 1.0 mg in peak a, 3.2 mg in peak b, and 4.2 mg in peak c, respectively. From its amino acid composition (Table 1) and analysis of tryptic fragments (data not shown), the peptide in peak b was identified as thymosin  $\alpha_1$ .

Table 1. Amino acid composition of peptides isolated from thymosin fraction 5

| Residue*   | Des-(24–25)-<br>thymosin $\alpha_1$ | Thymosin $\alpha_1$     | Thymosin $\alpha_{11}$ |  |
|------------|-------------------------------------|-------------------------|------------------------|--|
| Asp        | 4.0(3)                              | 4.7(4)                  | 6.1(5)                 |  |
| Thr        | 3.4(3)                              | 2.6(3)                  | 2.9(3)                 |  |
| Ser        | 3.8(3)                              | 2.8(3)                  | 3.1(3)                 |  |
| Glu        | 3.5(4)                              | 6.2(6)                  | 7.9(7)                 |  |
| Gly        | (0)<br>0                            | (0)<br>0                | 2.1(1)                 |  |
| Ala        | 2.6(2)                              | 3.1(3)                  | 5.5(5)                 |  |
| Val        | 3.0(3)                              | 2.4(3)                  | 2.5(3)                 |  |
| <b>Ile</b> | 1.0(1)                              | 1.0(1)                  | 1.0(1)                 |  |
| Leu        | 1.0(1)                              | 1.0(1)                  | 1.1(1)                 |  |
| Lys        | 4.0(4)                              | 2.9(4)                  | 4.0(4)                 |  |
| Arg        | (0)<br>0                            | (0)<br>0                | 1.2(1)                 |  |
| Pro        | (0)<br>0                            | $\left( 0 \right)$<br>0 | 1.0(1)                 |  |

The results are expressed as ratios to the value for isoleucine, which was taken as 1.0. The values in parentheses are those predicted from the amino acid sequences.

\* Met, Tyr, Phe, and His were absent or present in only trace quantities.

Identification of the Peptide in Peak a. This peptide was characterized as a fragment derived from residues 1-24 of thymosin  $\alpha_1$ , based on its amino acid composition (Table 1), the absence of a tryptic peptide corresponding to residues 21-28, and the presence of a new tryptic peptide corresponding to residues 21-24 (data not shown).

Sequence Analysis of Thymosin  $\alpha_{11}$  Purified from Peak C. The fractions in peak c were combined and rechromatographed as described in the legend to Fig. 1. The major peptide recovered (designated thymosin  $\alpha_{11}$ , Table 1) was digested with trypsin (Fig. 2). Six fragments were recovered and identified by their amino acid composition (Table 2). Peptides T6, T3, and T1 were identical to peptides derived from residues 1-14, 15- 17, and 18-20, respectively, of thymosin  $\alpha_1$ . Peptides T4 and T5 were similar in amino acid composition, differing only in the presence of a trace of lysine in peptide T4. Their composition indicated that they corresponded to residues 20-28 of thymosin  $\alpha_1$ , plus glycine and arginine. Automated sequence analysis established the structure of peptide T5 (Fig. 3). Peptide T4, which was identical to peptide T5 in amino acid composition, was found to contain aspartic acid, instead of asparagine, at position 28. The tryptic digests contained an additional peptide (T2) that was not present in tryptic digests of thymosin  $\alpha_1$ . This peptide contained no lysine or arginine, and must therefore have arisen from the COOH terminus of thymosin  $\alpha_{11}$ .

Manual Edman degradation of tryptic peptide T2 yielded the sequence Glu-Ala-Pro-Ala-AsnOH. Asparagine was recovered as the free amino acid after the fourth step of the Edman procedure. The structure of peptide T2 was confirmed by its cochromatography with the synthetic pentapeptide. Localization of peptide T2 at the COOH terminus of thymosin  $\alpha_{11}$ was confirmed by digestion of the latter with carboxypeptidase Y, which released  $\approx$  1 equivalent of asparagine, followed by alanine (2 equivalents) and proline (1 equivalent). The location of arginine at position 30 was confirmed by the isolation of a major fragment containing arginine after digestion of thymosin



FIG. 2. Peptides recovered from the tryptic digest of thymosin  $\alpha_{11}$ . The peptides in peak c from IEF fractions 6 and 7 (Fig. 1) were combined, lyophilized, and purified by rechromatography as described in the legend to Fig. 1. An aliquot (600  $\mu$ g) was digested with 42.9  $\mu$ g of TPCK-treated trypsin in 100  $\mu$ l of 0.4 M pyridine (pH 7.5). After 15 hr at 25°C the reaction mixture was lyophilized and the tryptic peptides were separated by HPLC using <sup>a</sup> gradient of acetonitrile in buffer A as shown (---). Fractions (0.65 ml) were collected every minute. At 6-sec intervals,  $5-\mu l$  samples were diverted to the fluorescamine detector.





Calculations were based on assigning a value of 1.0 for the residue shown in italics. ND, not determined.

mmol recovered from a digest of 200 nmol of thymosin  $\alpha_{11}$ . <sup>t</sup> nmol recovered from a digest of 8.7 nmol of thymosin  $\alpha_{11}$ .

 $\alpha_{11}$  with S. aureus V8 protease (peptide S7, Table 2). The amino acid composition of this peptide corresponded to that predicted for residues 26-31 of thymosin  $\alpha_{11}$ , including the last four residues of thymosin  $\alpha_1$ , plus the first three amino acid residues, glycine, arginine, and glutamic acid, found in the COOH-terminal extension of thymosin  $\alpha_{11}$ . Smaller quantities of two other fragments, whose amino acid compositions corresponded to residues 19-35 (peptide S3) and 25-35 (peptide S2) of thymosin  $\alpha_{11}$ , were also isolated from the S. aureus protease digests (Table 2 and Fig. 3). The results establish thymosin  $\alpha_{11}$ as containing the thymosin  $\alpha_1$  sequence plus seven additional amino acids at the COOH terminus.

Mouse Protection Assay. Inbred strains of mice vary in their susceptibility to infection with C. albicans (16). Thus, mice of such strains as C3H/HeJ or CBA/CaJ were found to be highly susceptible to infection, whereas mice of such strains as C57BL/ 1OSNJ or C57BL/KsJ were highly resistant to challenge. Because resistance to infection with C. albicans is associated with cell-mediated processes (17), and therefore with T lymphocytes, thymic hormones should have an effect on the host response. Thymosin fraction 5 and some peptides derived therefrom have been found to enhance maturation and replication of T lymphocytes (4) and, accordingly, should influence the resistance of a susceptible murine strain, such as C3H/HeJ, to infection with C. albicans.

Thymosin fraction 5, thymosin  $\alpha_1$ , or thymosin  $\alpha_{11}$  was injected daily in graded doses into three different groups of mice, beginning 2 days before intravenous challenge with  $4 \times 10^4$  cells of C. albicans. In comparison with control mice, at optimal doses all three preparations provided protection (Table 3). The polypeptides, thymosin  $\alpha_1$ , and thymosin  $\alpha_{11}$ , were approximately equal in potency, being most active in daily doses of 160-320 ng per mouse. Because the optimal dose for thymosin fraction 5 was  $5-10 \mu$ g, the peptides were therefore about 30 times more potent than fraction 5 in their ability to induce resistance to infection with C. albicans.

## DISCUSSION

The isolation from thymosin fraction 5 of two peptides structurally related to thymosin  $\alpha_1$ , one lacking four amino acids from the COOH terminus and the other containing seven additional amino acids, supports our previous suggestion that thymosin  $\alpha_1$ is not the native form of the thymic hormone and that peptides present in thymosin fraction 5 may be proteolytic fragments of larger peptides (18). We have previously identified thymosin  $\beta_8$ , also isolated from thymosin fraction 5, as a modified form of the native peptide, thymosin  $\beta_9$ , the latter containing two additional amino acid residues at the COOH terminus (14).

Both thymosin  $\alpha_1$  and thymosin  $\alpha_{11}$  contain COOH-terminal asparagine residues, suggesting the existence of a proteinase with some selectivity for asparaginyl bonds. We have identified a lysosomal proteinase in rat liver that catalyzes the hydrolysis of an asparaginyl-valine bond in rabbit liver fructose-1,6-bisphosphatase (19). An internal asparaginyl-valine bond in cathepsin B also appears to be the target of a proteinase in rat liver, possibly cathepsin B itself (20).

Both thymosin  $\alpha_1$  and thymosin  $\alpha_{11}$  enhanced the resistance of susceptible C3H/HeJ mice to challenge with C. albicans, when administered at a daily optimal dose of 160-320 ng per mouse. Increasing or decreasing the daily dosage caused a decline in the degree of host resistance.

Thymosin fraction 5, or its individual components, has been reported to enhance resistance to infection with a variety of microorganisms under a variety of host conditions. Thus, thymosin fraction 5 enhanced the cell-mediated resistance of moderately protein-malnourished mice to Listeria monocytogenes



FIG. 3. The amino acid sequence of thymosin  $\alpha_{11}$ . The sequence shown is based on the composition of the tryptic peptides (T1-T6) and of peptides generated by digestion with S. aureus V8 protease (S2, S3, and S7). The tryptic peptide lacking lysine and arginine, presumably derived from the COOH terminus, was analyzed by Edman degradation  $\leftrightarrow$ ). For peptide T5, amino acids released from thymosin  $\alpha_{11}$  by carboxypeptidase Y are indicated ( $\leftarrow$ ). Des-(25-28)-thymosin  $\alpha_1$  and thymosin  $\alpha_1$  would be derived from residues 1-24 and residues 1-28 of thymosin  $\alpha_{11}$ , respectively.

Table 3. Effect of thymosin fraction 5 and thymic peptides on the growth of C. albicans in C3H/HeJ mice

| Thymosin fraction 5   |                            | Thymosin $\alpha_1$   |                            | Thymosin $\alpha_{11}$ |                            |
|-----------------------|----------------------------|-----------------------|----------------------------|------------------------|----------------------------|
| Dose, ng<br>per mouse | C. albicans<br>cell count* | Dose, ng<br>per mouse | C. albicans<br>cell count* | Dose, ng<br>per mouse  | C. albicans<br>cell count* |
| <b>None</b>           | 5,100                      |                       |                            |                        |                            |
| 2,560                 | 8.500                      | 80                    | 5,870                      | 80                     | 4,200                      |
| 5,120                 | 440                        | 160                   | 190                        | 160                    | 510                        |
| 10,240                | 320                        | 320                   | 780                        | 320                    | 320                        |
| 20,480                | 1.600                      | 640                   | 1,410                      | 640                    | 1,260                      |

Mice were treated daily with the indicated doses of thymosin fraction 5, thymosin  $\alpha_1$ , or thymosin  $\alpha_{11}$ 

and challenged with  $4 \times 10^4$  cells of C. albicans 2 days after the start of treatment (see Methods).

\* Three mice from each set were sacrificed on days 7, 14, and 21 after infection and the values represent the average number of organisms in the left kidneys of the nine mice in each set.

(21) and of alloxan-diabetic mice to C. albicans (22). Also, administration of thymosin  $\alpha_1$  at optimal dose and schedule protected  $CD2F_1$  mice infected with lethal doses of C. albicans (23) and of ddY or C57BL/6 mice infected with C. albicans, Listeria monocytogenes, or Serratia marcescens after immunosuppression with 5-fluorouracil (24).

It is highly important that the individual components of thymosin fraction 5 be separated, purified, and characterized. This concept is based on the observation that the action of thymosin fraction 5 is often quite variable. For example, when the effect of thymosin fraction 5 on human peripheral blood lymphocytes was determined *in vitro* in an assay employing an allogeneic "mixed lymphocyte reaction," the outcome of the response was unpredictable (25). This result was attributed to an ability of thymosin fraction 5 to stimulate both helper and suppressor T lymphocytes. In addition, administration of thymosin fraction 5 to some susceptible murine strains-i.e., C3H/HeJ-enhanced cell-mediated immunity, whereas administration to resistant murine strains-i.e., C57BL/10SNJ-decreased cellmediated immunity (16, 26).

Thymosin  $\alpha_{11}$  may itself represent a proteolytically modified fragment of the native thymic peptide; if so, it is a larger fragment than thymosin  $\alpha_1$ , closer in structure to the native peptide and retaining biological activity. The isolation of the native peptide and evaluation of its chemical and biological properties will require extraction conditions that preclude any possibility of proteolysis.

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- 1. Goldstein, A. L., Guha, A., Zatz, M. M., Hardy, M. A. & White, A. (1972) Proc. Natl. Acad. Sci. USA 69, 1800-1803.
- 2. Hooper, J. A., McDaniel, M. C., Thurman, G. B., Cohen, G. H., Schulof, R. S. & Goldstein, A. L. (1975) Ann. N.Y. Acad. Sci. 249, 125-144.
- 3. White, A. (1980) in Biochemical Action of Hormones, ed. Litwack, G. (Academic, New York), pp. 1-46.
- 4. Goldstein, A. L., Low, T. L. K., Thurman, G. B., Zatz, M. M., Hall, N., Chenk, J., Hu, S.-K., Naylor, P. B. & McClure, J. E. (1981) Recent Prog. Horm. Res. 37, 369-415.
- 5. Goldstein, A. L., Low, T. L. K., McAdoo, M., McClure, J., Thurman, G. B., Rossio, J., Lai, C.-Y., Chang, D., Wang, S.-S., Harvey, C., Ramel, A. H. & Meienhofer, J. (1977) Proc. Natl. Acad. Sci. USA 74, 725-729.
- 6. Low, T. L. K., Thurman, G. B., McAdoo, M., McClure, J., Rossio, J. L., Naylor, P. H. & Goldstein, A. L. (1979) J. Biol. Chem. 254, 981-986.
- 7. Wang, S.-S., Kulesha, I. D. & Winter, D. P. (1979) J. Am. Chem. Soc. 101, 253-254.
- 8. Freire, M., Crivellaro, O., Isaacs, C., Moschera, J. & Horecker, B. L. (1978) Proc. Natl. Acad. Sci. USA 75, 6007-6011.
- 9. Freire, M., Hannappel, E., Rey, M., Freire, J. M., Kido, H. & Horecker, B. L. (1981) Proc. Natl. Acad. Sci. USA 78, 192-195.
- 10. Wong, T. W. & Merrifield, R. B. (1980) Biochemistry 19, 3233- 3238.
- 11. Stein, S. & Moschera, J. (1981) Methods Enzymol. 79, 7-16.
- 12. Erickson-Viitanen, S., Ruggieri, S., Natalini, P. & Horecker, B. L. (1983) Arch. Biochem. Biophys. 221, 570-576.
- 13. Hewick, R. M., Hunkapiller, M. W., Hood, L. E. & Dreyer, W. J. (1981) J. Biol. Chem. 256, 7990-7997.
- 14. Hannappel, E., Davoust, S. & Horecker, B. L. (1982) Proc. Natl. Acad. Sci. USA 79, 1708-1711.
- 15. Hannappel, E., Davoust, S. & Horecker, B. L. (1982) in Protides of the Biological Fluids, ed. Peeters, H. (Pergamon, Oxford), pp. 739-742.
- 16. Salvin, S. B. & Neta, R. (1983) Cell. Immunol. 75, 160–172.<br>17. Salvin, S. B. & Neta, R. (1982) in The Reticuloendothelial Su.
- Salvin, S. B. & Neta, R. (1982) in The Reticuloendothelial System, eds. Ross, N. R. & Siegel, B. V. (Plenum, New York), Vol. 4, pp. 145-186.
- 18. Hannappel, E., Davoust, S. & Horecker, B. L. (1982) Biochem. Biophys. Res. Commun. 104, 266-271.
- 19. Lazo, P. S., Tsolas, O., Sun, S. C., Pontremoli, S. & Horecker, B. L. (1978) Arch. Biochem. Biophys. 188, 308-314.
- 20. Katunuma, N. & Kominami, E. (1983) in Current Topics in Cellular Regulation, eds. Horecker, B. L. & Stadtman, E. R. (Academic, New York), Vol. 22, pp. 71-101.
- 21. Petro, T. M., Chien, G. & Watson, R. R. (1982) Infect. Immun. 37, 601-608.
- 22. Salvin, S. B. & Tanner, E. P. (1983) Clin. Exp. Immunol. 54, in press.
- 23. Bistoni, F., Marconi, P., Frati, L., Bonmassar, E. & Garaci, E. (1982) Infect. Immun. 36, 609-614.
- 24. Ishitsuka, H., Umeda, Y., Nakamura, J. & Yagi, Y. (1983) Cancer Immunol. Immunother. 14, 145-150.
- 25. Kaufman, D. B. (1980) Clin. Exp. Immunol. 39, 722-727.
- 26. Neta, R. & Salvin, S. B. (1983) Cell. Immunol. 75, 173-180.