# Isolation and partial characterization of follistatin: A single-chain $M_r$ 35,000 monomeric protein that inhibits the release of follicle-stimulating hormone

(ovary/pituitary/luteinizing hormone/gonadotropins/porcine follicular fluid)

NAOTO UENO, NICHOLAS LING\*, SHAO-YAO YING, FREDERICK ESCH, SHUNICHI SHIMASAKI, AND ROGER GUILLEMIN

The Salk Institute for Biological Studies, Laboratories for Neuroendocrinology, 10010 North Torrey Pines Road, La Jolla, CA 92037

Contributed by Roger Guillemin, August 6, 1987

A  $M_r$  35,000 protein with follicle-stimulating ABSTRACT hormone release-inhibitory activity was isolated from porcine ovarian follicular fluid by heparin-Sepharose affinity chromatography, gel filtration on Sephacryl S-200, and multiple steps of high-performance liquid chromatography. The isolated molecule is highly enriched in cysteines and is composed of a single polypeptide chain. In addition, it has no sequence homology with the previously characterized follicular fluid inhibins, which are heterodimeric proteins of  $M_r$  32,000 with follicle-stimulating hormone release-inhibiting activity. This protein specifically inhibits the basal secretion of follicle-stimulating hormone, but not that of luteinizing hormone, in the rat anterior pituitary monolayer culture system with a halfmaximal effective dose of 2.5-6.0 ng/ml. Another form of the molecule of  $M_r$  32,000 present in much lower concentration in follicular fluid was also isolated. It may differ from the M. 35,000 form in glycosylation or carboxyl-terminal truncation. We suggest that this compound be called "follistatin" to signify its structural difference from inhibin.

Secretion of pituitary follicle-stimulating hormone (FSH) is principally regulated by the hypothalamic hypophysiotropic gonadotropin-releasing hormone and gonadal steroids (1). But for many years the existence of a nonsteroidal, watersoluble factor from the gonad that feeds back on the pituitary specifically to suppress the secretion of FSH has been advocated by many researchers (2); the name "inhibin" was coined in 1932 to describe such a factor (3). However, the search for the elusive inhibin did not succeed until 1985, when two closely related glycoproteins of apparent  $M_r$  32,000, named inhibins A and B, were isolated from porcine ovarian follicular fluid (4-6). Subsequently, a similar  $M_r$  32,000 inhibin was also isolated from bovine ovarian follicular fluid (7, 8). Each inhibin was found to be composed of a common  $M_r$  18,000 glycosylated  $\alpha$  subunit linked by interchain disulfide bond(s) to either a  $\beta_A$  subunit of  $M_r$  14,700 to yield inhibin A or a  $\beta_{\rm B}$  subunit of  $M_{\rm r}$  14,000 to form inhibin B; the two  $\beta$ subunits were closely related but distinct from each other (4). The common  $\alpha$  subunit can exist also in a larger molecular form of  $M_r$  44,000 and, in combination with the  $\beta$  subunit of inhibin A, can form an inhibin of  $M_r$  56,000 (9).

Using partial amino acid sequence information, the messages encoding the  $\alpha$ ,  $\beta_A$ , and  $\beta_B$  subunits of porcine inhibin were cloned from an ovarian cDNA library (10) as were the  $\alpha$  and  $\beta_A$  subunits of bovine inhibin (11). DNA sequence analyses showed that all of the subunits were initially synthesized as large precursor proteins with the mature subunits residing at the carboxyl-terminal portion of the precursors. Surprisingly, the two  $\beta$  subunits, besides being homologous to each other, were found to be structurally related to a homodimeric protein, transforming growth factor  $\beta$ , which has totally different biological activities (10). Moreover, in addition to the inhibins, two FSH-releasing proteins of apparent  $M_r$  24,000, named activin A and activin AB, also were isolated from porcine follicular fluid (12–14). The former was characterized as a homodimer composed of two  $\beta$ subunits of inhibin A linked by disulfide bridge(s), while the latter is a heterodimer composed of the  $\beta$  subunits of inhibins A and B.

In our original work on the isolation of the two  $M_r$  32,000 heterodimeric inhibins from porcine ovarian follicular fluid, we had noted a side fraction migrating at the beginning of the first reversed-phase high-performance liquid chromatogram ahead of the activins and inhibins, which also could specifically suppress pituitary FSH secretion in the inhibin bioassay. In this paper we present the purification and isolation of this FSH release-inhibiting protein named "follistatin," present in porcine ovarian follicular fluid, and show that it is a glycoprotein of  $M_r$  35,000 composed of a single polypeptide chain.

# MATERIALS AND METHODS

**Bioassay.** Throughout the purification procedure, the FSH release-inhibiting activity was monitored by an *in vitro* bioassay using immature female rat anterior pituitary monolayer culture as in the isolation of the inhibins (4).

**Reversed-Phase HPLC.** The reversed-phase HPLC system used for the purification of the follistatin consisted of a Beckman series 322 gradient liquid chromatography system equipped with a Spectroflow 757 UV detector (Kratos, Ramsey, NJ), a Soltec 220 recorder (Soltec, Sun Valley, CA), and a Redirac 2112 fraction collector (LKB). Two reversedphase HPLC solvent systems were used. In the triethylammonium phosphate (Et<sub>3</sub>NHP<sub>i</sub>) system, solvent A consists of 0.25 M Et<sub>3</sub>NHP<sub>6</sub> (pH 3); solvent B is 80% (vol/vol) acetonitrile in solvent A. In the CF<sub>3</sub>COOH system, solvent A contains 1 ml of CF<sub>3</sub>COOH in 999 ml of water; solvent B is 1 ml of CF<sub>3</sub>COOH in 199 ml of water and 800 ml of acetonitrile. Other reversed-phase HPLC conditions are given in the legends to Figs. 1 and 2.

Anion-Exchange HPLC. The anion-exchange HPLC system used for the purification consisted of a Beckman series 332 gradient liquid chromatography system equipped with a Beckman 160 UV detector, a Soltec 1242 recorder, and a Redirac 2112 fraction collector. The solvent system used for the anion-exchange HPLC consisted of buffer A, which is

The publication costs of this article were defrayed in part by page charge payment. This article must therefore be hereby marked "*advertisement*" in accordance with 18 U.S.C. \$1734 solely to indicate this fact.

Abbreviations: FSH, follicle-stimulating hormone;  $Et_3NHP_i$ , triethylammonium phosphate.

<sup>\*</sup>To whom reprint requests should be addressed.

 $0.01~M~Na_2HPO_4$  (pH 6.5), and buffer B, which is 1 M NaCl in buffer A.

Initial Purification. The starting material for the purification of the follistatin was a side fraction derived from the purification of the porcine ovarian inhibins and activins (4, 12, 14). Briefly, porcine follicular fluid was processed batchwise through a heparin-Sepharose affinity chromatography column, and the adsorbed protein was further size-fractionated on a Sephacryl S-200 column in acidic medium. Fractions containing the  $M_r$  32,000 inhibins and FSH-releasing activities from the column (fractions 35–43 in figure 1a of ref. 4) were pooled and lyophilized. The lyophilized material ( $\approx 40$ mg) was dissolved in 40 ml of 0.2 M acetic acid and filtered through a Millex-HA 0.45- $\mu$ m filter (Millipore). The filtrate was applied directly to a 10 × 250 mm Vydac 5- $\mu$ m C<sub>4</sub> column (The Separation Group, Hesperia, CA). After all of the filtrate had been loaded, the column was washed with solvent A until the UV absorption reached baseline. The inhibins and activins were separated with a linear gradient of 27-36% acetonitrile in the Et<sub>3</sub>NHP<sub>i</sub> solvent system in 90 min at a flow of 3 ml/min. Three zones of FSH release-inhibiting activity corresponding to inhibins A and B and follistatin were detected, as were two zones of FSH-releasing activity as shown in Fig. 1*a*. Inhibins A and B, as well as the FSH-releasing substances (activin A and activin AB) have been purified to homogeneity and their primary structures characterized as reported (4, 10, 12, 14). The remaining FSH release-inhibiting zone migrating ahead of the activins and inhibins on the HPLC column was saved for the present work.

NaDodSO<sub>4</sub>/PAGE. The follistatin isolated from the final reversed-phase HPLC purification was analyzed in a 1-mm thick, 10% acrylamide gel by the method of Laemmli (15). The protein band was revealed by Coomassie blue reagent (Bio-Rad). The following molecular weight standards were



FIG. 1. (a) Reversed-phase HPLC purification of the FSH release-inhibiting as well as FSH-releasing proteins recovered from gel filtration. Three zones of FSH release-inhibiting activity (inhibins A, B, and follistatin) and two zones of FSH-releasing activity (activins A and AB), indicated by the solid bars, were recovered. (b) Reversed-phase HPLC purification of follistatin. Active fractions 3 and 4 in a were pooled and, after dilution with 0.2 M acetic acid, applied directly onto a Vydac C<sub>4</sub> column and eluted with the indicated gradient of acetonitrile in the CF<sub>3</sub>COOH system at 3 ml/min. (c) The active material (denoted by solid bar in b) was pooled and, after removal of the acetonitrile and adjustment to pH 6.5, was applied to a Spherogel-TSK DEAE-5PW column. The activity was separated by the indicated gradient of NaCl in sodium phosphate buffer.

used to calibrate the gel: bovine serum albumin ( $M_r$  67,000), ovalbumin ( $M_r$  43,000), and  $\alpha$  chymotrypsinogen ( $M_r$  25,700). The purified protein was analyzed under nonreducing and reducing conditions as before (4).

Amino Acid Analyses and Microsequencing. Amino acid analyses and amino-terminal sequence determination of follistatin were performed as described (16, 17).

Concanavalin A-Sepharose 4B Affinity Chromatography. Approximately 1  $\mu$ g of the purified  $M_r$  35,000 follistatin was iodinated with lactoperoxidase (Calbiochem), and the labeled protein was purified on a 0.7 × 50 cm Sephadex G-50 column with 0.01 M phosphate buffer (pH 7.5). The labeled follistatin was loaded onto a 1-ml concanavalin A-coupled Sepharose 4B column (Sigma) and washed successively with the loading buffer, 0.2 M and 1.0 M methyl  $\alpha$ -D-mannopyranoside in loading buffer, and 0.1 M acetic acid as described (18).

## RESULTS

For the purification of follistatin, fractions 3 and 4 in Fig. 1a, saved from the previous purification of the inhibins and

activins, were pooled, mixed with an equal volume of 0.2 M acetic acid, and applied directly onto a  $10 \times 250$  mm Vydac 5- $\mu$ m C<sub>4</sub> column as before (4). The active material was eluted from the column with a linear gradient of 21-30% acetonitrile in the CF<sub>3</sub>COOH system in 90 min at a flow of 3 ml/min as shown in Fig. 1b. Fractions with FSH release-inhibiting activity (fractions 19-23) detected by the bioassay were pooled and, after the acetonitrile was removed in a Speed-Vac concentrator (Savant) and the pH was adjusted to 6.5 with 0.1 M ammonium hydroxide, were pumped directly onto a 7.5  $\times$ 75 mm Spherogel-TSK 10-µm DEAE-5PW column (Toyo Soda, Tokyo). After loading, the column was washed with buffer A until the UV absorption reached baseline. The follistatin activity was separated with a linear gradient of 0-0.3 M NaCl in 90 min at a flow rate of 1 ml/min as shown in Fig. 1c. The active material eluted in fractions 19-22 was pooled and, after dilution to 4 times its original volume with 0.2 M acetic acid, was further purified on a 10  $\times$  250 mm Vydac 5- $\mu$ m C<sub>4</sub> column and eluted with a linear gradient of 21-30% acetonitrile in the CF<sub>3</sub>COOH solvent system in 90



FIG. 2. Reversed-phase HPLC purification of follistatin. (a) Active fractions (solid bar in Fig. 1c) were pooled and after dilution were applied directly onto a Vydac C<sub>4</sub> column and eluted with the indicated gradient of acetonitrile in the CF<sub>3</sub>COOH system at 1 ml/min. (b) The active material (denoted by solid bar I in a) was pooled, diluted, and chromatographed on a Vydac phenyl column with the indicated gradient of acetonitrile in the Et<sub>3</sub>NHP<sub>i</sub> system at 1 ml/min. (c) The material denoted by the solid bar in b was pooled and concentrated by chromatography on an Aquapore RP-300 column with the indicated gradient of acetonitrile in the CF<sub>3</sub>COOH system at 0.5 ml/min.

### Biochemistry: Ueno et al.

min at a flow of 1 ml/min as shown in Fig. 2a. The active material eluted in fractions 33–35 (peak I) was pooled and diluted to twice its original volume and rechromatographed on a 10  $\times$  250 mm Vydac 5- $\mu$ m phenyl column with a linear gradient of 18–27% acetonitrile in the Et<sub>3</sub>NHP<sub>i</sub> system in 90 min at a flow rate of 1 ml/min as shown in Fig. 2b. Finally, the active material in fractions 34–36 was pooled and, after dilution with 0.2 M acetic acid, was concentrated by chromatography on a 4.6  $\times$  250 mm Aquapore 10- $\mu$ m RP-300 column with a linear gradient of 20–80% acetonitrile in the CF<sub>3</sub>COOH system in 60 min at a flow rate of 0.5 ml/min as shown in Fig. 2c. Altogether,  $\approx$ 400  $\mu$ g of follistatin was isolated from 18 liters of follicular fluid.

The purified protein showed half-maximal inhibition of FSH release at a concentration of 2.5-6.0 ng/ml (0.07-0.17 nM). It had no effect on the secretion of luteinizing hormone, growth hormone, prolactin, or thyroid-stimulating hormone.

Amino acid analyses showed that follistatin, like inhibins A and B, is highly enriched in cysteines (Table 1). On NaDod-SO<sub>4</sub>/PAGE under nonreducing conditions, follistatin showed a single band migrating at an apparent  $M_r$  of 35,000 as shown in Fig. 3*a*. In contrast to inhibins A and B, follistatin exhibited a single band that migrated at an apparent  $M_r$  of 42,000 under reducing conditions (Fig. 3*b*), indicating that it is composed of a single polypeptide chain. Microsequence analyses of follistatin revealed its amino terminus to be Gly-Asn-Cys-Trp-Leu-Arg-Gln-Ala. In addition, 63% of the labeled follistatin was retained on the concanavalin A-Sepharose 4B affinity column, which could be displaced with 0.2 M methyl  $\alpha$ -D-mannopyranoside, indicating that follistatin is glycosylated.

Using similar reversed-phase HPLC conditions (data not shown), a  $M_r$  32,000 form (Fig. 4a) of follistatin with an identical amino-terminal sequence as that of the  $M_r$  35,000 follistatin was isolated from fractions 36 and 37 of Fig. 2a (peak II). Its amino acid composition is related to the  $M_r$  35,000 form (Table 1), and upon reduction, this protein migrated as a single band at an apparent  $M_r$  of 40,000 on NaDodSO<sub>4</sub>/PAGE (Fig. 4b).

Table 1. Amino acid composition of purified follistatin from porcine follicular fluid

Amino acid	<i>M</i> <sub>r</sub> 35,000 form	<i>M</i> <sub>r</sub> 32,000 form
Asx	$34.1 \pm 0.2$	$28.4 \pm 0.0$
Thr	$17.7 \pm 0.3$	$15.9 \pm 0.0$
Ser	$25.6 \pm 0.2$	$22.2 \pm 0.2$
Glx	$36.7 \pm 0.1$	$31.1 \pm 0.1$
Gly	$23.3 \pm 0.2$	$24.5 \pm 0.0$
Ala	$17.1 \pm 0.2$	$15.6 \pm 0.0$
Val	$15.5 \pm 0.3$	$14.0 \pm 0.0$
Met	$3.2 \pm 0.1$	$3.4 \pm 0.0$
lle	$10.1 \pm 0.1$	$7.5 \pm 0.1$
Leu	$20.9 \pm 0.2$	$19.1 \pm 0.0$
Tyr	$9.5 \pm 0.1$	$8.8 \pm 0.1$
Phe	$5.0 \pm 0.1$	$4.6 \pm 0.0$
His	$2.2 \pm 0.0$	$2.0 \pm 0.0$
Тгр	$5.8 \pm 0.1$	$5.6 \pm 0.1$
Lys	$26.2 \pm 0.0$	$25.2 \pm 0.1$
Arg	$12.7 \pm 0.1$	$13.7 \pm 0.1$
Cys*	$35.9 \pm 0.2$	$33.5 \pm 0.1$
Pro	$13.5 \pm 0.3$	$13.0 \pm 0.1$

Values represent means  $\pm$  SD of two analyses, normalized to a protein of 35,000 daltons for the  $M_r$  35,000 form of follistatin and a protein of 32,000 daltons for the  $M_r$  32,000 form.

\*Determined as cysteic acid after performic acid oxidation.



FIG. 3. NaDodSO<sub>4</sub>/PAGE analysis of the purified  $M_r$  35,000 follistatin under nonreducing conditions (*a*) and under reducing conditions (*b*). Positions of the molecular weight standards (shown × 10<sup>-3</sup>) are indicated at the left.

### DISCUSSION

Two novel proteins with FSH release-inhibiting activity were purified from porcine follicular fluid by using essentially the same procedures employed for the isolation of the inhibins and activins (4, 12, 14). The protein obtained in much higher yield has an apparent  $M_r$  of 35,000, whereas the other has an apparent  $M_r$  of 32,000. Both proteins were found to have an identical amino terminus and are composed of a single polypeptide chain. In view of the similar amino acid compositions and identical amino termini, the two forms of follistatin are probably related, and their difference in molecular size might be due to variation in glycosylation or carboxylterminal truncation.

Aside from the high content of cysteines, there is no relation in amino acid composition between follistatin and inhibins A and B. In addition, follistatin is a monomeric protein, whereas inhibins A and B and all higher molecular weight inhibins reported thus far are heterodimers (9, 19).

In a recent publication, Leversha *et al.* (20) reported the copurification of a  $M_r$  29,000–30,000 single-chain protein together with inhibin A from ovine follicular fluid. They demonstrated that this protein also has FSH release-inhibit-



FIG. 4. NaDodSO<sub>4</sub>/PAGE analysis of the purified  $M_r$  32,000 follistatin under nonreducing conditions (*a*) and under reducing conditions (*b*). Positions of the molecular weight standards (shown × 10<sup>-3</sup>) are indicated at the left.

ing activity and upon reduction migrated as a single band of apparent  $M_r$  36,000 on NaDodSO<sub>4</sub>/PAGE. However, no further purification and characterization of this protein was attempted. It is interesting to speculate that this single-chain protein with inhibin-like activity may be related to the follistatin isolated in our laboratory.

At present we do not know the physiological significance of follistatin. Preliminary biological studies showed that this molecule can specifically inhibit the spontaneous release of pituitary FSH with no effects on luteinizing hormone. Its relative potency is about one-third that of inhibin A (4). Undoubtedly, more physiological studies will be forthcoming to unravel the function of this new ovarian regulatory protein.

We thank F. Castillo, R. Schroeder, M. Regno, A. Becker, J. Czvik, G. Swanson, and P. Tan for technical assistance and D. Higgins for preparing the manuscript. This work was supported by National Institute of Child Health and Human Development contract N01-HD-6-2944, National Institutes of Health Program Project Grants HD-09690 and DK-18811, and a grant from the Robert J. and Helen C. Kleberg Foundation.

- Griffin, J. E. & Wilson, J. D. (1985) in Williams' Textbook of Endocrinology, eds. Wilson, J. D. & Foster, D. W. (Saunders, Philadelphia), 7th Ed., pp. 259-311.
- Franchimont, P., Verstraelen-Proyord, J., Hazee-Hagelstein, M. T., Renard, Ch., Demoulin, A., Bourguignon, J. P. & Hustin, J. (1979) Vitam. Horm. (N.Y.). 37, 243-302.
- 3. McCullagh, D. R. (1932) Science 76, 19-20.
- Ling, N., Ying, S.-Y., Ueno, N., Esch, F., Denoroy, L. & Guillemin, R. (1985) Proc. Natl. Acad. Sci. USA 82, 7217-7221.
- Miyamoto, K., Hasegawa, Y., Fukuda, M., Nomura, M., Igarashi, M., Kangawa, K. & Matsuo, H. (1985) Biochem. Biophys. Res. Commun. 129, 396-403.
- 6. Rivier, J., Spiess, J., McClintock, R., Vaughan, J. & Vale, W. (1985) Biochem. Biophys. Res. Commun. 133, 120-127.

- Fukuda, M., Miyamoto, K., Hasegawa, Y., Nomura, M., Igarashi, M., Kangawa, K. & Matsuo, H. (1986) Mol. Cell. Endocrinol. 44, 55-60.
- Robertson, D. M., deVos, F. L., Foulds, L. M., McLachlan, R. I., Burger, H. G., Morgan, F. J., Hearn, M. T. W. & deKretser, D. M. (1986) Mol. Cell. Endocrinol. 44, 271-277.
- 9. Robertson, D. M., Foulds, L. M., Leversha, L., Morgan, F. J., Hearn, M. T. W., Burger, H. G., Wettenhall, R. E. H. & deKretser, D. M. (1985) *Biochem. Biophys. Res. Commun.* 126, 220-226.
- Mason, A. J., Hayflick, J. S., Ling, N., Esch, F., Ueno, N., Ying, S.-Y., Guillemin, R., Niall, H. & Seeburg, P. (1985) *Nature (London)* 318, 659-663.
- Forage, R. G., Ring, J. M., Brown, R. W., McInerney, B. V., Cobon, G. S., Gregson, R. P., Robertson, D. M., Morgan, F. J., Hearn, M. T. W., Findlay, J. K., Wettenhall, R. E. H., Burger, H. G. & deKretser, D. M. (1986) Proc. Natl. Acad. Sci. USA 83, 3091-3095.
- 12. Ling, N., Ying, S.-Y., Ueno, N., Shimasaki, S., Esch, F., Hotta, M. & Guillemin, R. (1986) *Nature (London)* 321, 779-792.
- Vale, W., Rivier, J., Vaughan, J., McClintock, R., Corrigan, A., Woo, W., Karr, D. & Spiess, J. (1986) *Nature (London)* 321, 776-779.
- Ling, N., Ying, S.-Y., Ueno, N., Shimasaki, S., Esch, F., Hotta, M. & Guillemin, R. (1986) *Biochem. Biophys. Res. Commun.* 138, 1129–1137.
- 15. Laemmli, U. (1970) Nature (London) 227, 1677-1685.
- 16. Böhlen, P. & Schroeder, R. (1982) Anal. Biochem. 126, 144-152.
- 17. Esch, F. (1984) Anal. Biochem. 136, 39-47.
- Shibasaki, T., Ling, N. & Guillemin, R. (1980) Nature (London) 285, 416-417.
- Miyamoto, K., Hasegawa, Y., Fukuda, M. & Igarashi, M. (1986) Biochem. Biophys. Res. Commun. 136, 1103-1109.
- Leversha, L. J., Robertson, D. M., deVos, F. L., Morgan, F. J., Hearn, M. T. W., Wettenhall, R. E. H., Findlay, J. K., Burger, H. G. & deKretser, D. M. (1987) J. Endocrinol. 113, 213-221.