Purification of ADAM 10 from bovine spleen as a TNF α convertase

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Abstract We have purified a protease with characteristics of TNF α convertase from bovine spleen membranes. Peptide sequencing of the purified protein identified it as ADAM 10 (Genbank accession no. Z21961). This metalloprotease cleaves a recombinant proTNF α substrate to mature TNF α , and can cleave a synthetic peptide substrate to yield the mature TNF α amino terminus in vitro. The enzyme is sensitive to a hydroxamate inhibitor of MMPs, but insensitive to phosphoramidon. In addition, cloned ADAM 10 mediates proTNF α processing in a processing-incompetent cell line.

Key words: Tissue necrosis factor α ; TNF α convertase; ADAM 10; Metalloprotease

1. Introduction

The pro-inflammatory cytokine TNF α [1-3] is synthesized as a 233 amino acid precursor. Conversion of this membranebound precursor to a secreted 17 kDa mature protein [4,5] is carried out by an unidentified protease, termed the TNF α convertase. Serine proteases are capable of releasing proTNFa from the membrane-bound form, although the resulting amino-terminus is not identical with that observed in vivo [6]. Many investigators believe TNFa convertase is a membrane-bound metalloprotease [7-9]. This activity is inhibited by EDTA and many prototypic MMP inhibitors but not by phosphoramidon, thiorphan, captopril and TIMP-1. This activity cleaves the proTNF α at the correct position, yielding an amino-terminal valine. We have purified one activity from bovine spleen with enzymatic characteristics consistent with TNF α convertase. In addition, we show that this protein, identified as ADAM 10, can reconstitute TNFa processing in a processing-incompetent cell line.

2. Materials and methods

2.1. In vitro assay for TNFa convertase activity

A protein-based assay for TNF α convertase activity was based on described procedures [7], using a ¹²⁵I-labelled peptide-tagged recombinant proTNF α polypeptide substrate (Flag-proTNF α). A membrane protein sample was mixed with 2 ng of ¹²⁵I-polypeptide substrate (approx. 50 000 cpm) in the presence of inhibitors (0.2 mM PMSF, 2 μ M pepstatin, 200 μ M leupeptin, 1 mM *N*-methoxysuccinyl-AAPV-chloromethyl ketone [AAPVCMK]). Following incubation at 37°C overnight, samples were fractionated by SDS-PAGE on a 16% gel. Polypeptides were visualized by autoradiography.

A peptide-based assay for TNF α convertase activity was carried out similarly to the procedure described in [7] using an amino-terminal dinitrophenylated peptide substrate (DNP-SPLAQAVRSSSR-NH₂). A membrane protein sample was mixed in 50 µl with 1 µg peptide substrate in the presence of inhibitors (see above). Following incubation at 37°C, the protein was precipitated by cold 5% trichloroacetic acid, 20% acetonitrile and the soluble peptide fraction was applied to a YMC 120 Å C-18 ODS-AQ column (4.6×100 mm). The column was eluted isocratically at 1 ml/min using 40% acetonitrile plus 0.06% trifluoroacetic acid. The elution of DNP-peptide was monitored at 360 nm using a Waters 625 LC system.

2.2. Purification from bovine spleen

Bovine spleens were purchased from Pel-Freez. The washed tissue was homogenized in a Waring blender, resuspended in lysis buffer (20 mM Tricine, pH 7.8, 8% sucrose) containing 0.1% PMSF, then homogenized using a Brinkmann Polytron tissue homogenizer. The cell debris was removed by two low-speed centrifugations and the membranes isolated by ultracentrifugation. The isolated membranes were washed in 10 mM HEPES, pH 7.5.

The membrane fraction was resuspended to 8 mg/ml in buffer A (20 mM Tris, pH 7.5, 1 mM MgSO₄, 10 mM NaCl, 10 μ M ZnSO₄) plus 2% Brij 35, and incubated at 4°C for 30 min. The membranes were collected by ultracentrifugation, then resuspended to 4 mg/ml in buffer A plus 2% Thesit (Boehringer Mannheim). The insoluble protein was removed by ultracentrifugation.

The detergent-solubilized fraction was applied to a chelating Sepharose (Pharmacia) column charged with nickel sulfate in buffer A with 0.3 M NaCl/0.1% octyl glucoside (all subsequent steps included 0.1% octyl glucoside). The 50 mM imidazole eluate was concentrated, then applied to a Sephacryl 300 (Pharmacia) sieving column equilibrated in buffer A. A retained fraction (approx. 60 kDa) was pooled, then applied to a wheat germ agglutinin column in 0.3 M NaCl and eluted with buffer A containing 0.5 M *N*-acetylglucosamine and 0.5 M NaCl. The eluate was dialyzed vs. 1 mM NaPO₄ pH 7, then applied to an hydroxyapatite (BioRad) column. An eluted fraction (40–50 mM NaPO₄) was then dialyzed vs. buffer A and passed successively over HS-20 and HQ-20 (Perseptive Biosystems) ion exchange columns. The unbound fraction from both columns contained the TNF α convertase activity.

The final protein fraction (approx. 100 μ g) was fractionated by SDS-PAGE on an 8% gel (Novex), then electrophoretically transferred to an Immobilon filter membrane in 10 mM CAPS buffer, pH 10, 10% methanol. After staining the membrane with 0.1% Amido Black in 40% methanol, 10% acetic acid, the major protein band was excised and used for microsequencing.

2.3. Cloning and in vivo activity of ADAM 10

Bovine ADAM 10 (Genbank accession no. Z21961) was cloned from 5 μ g poly(A)⁺ RNA (Clontech) using a GibcoBRL SuperScript Preamplification System and an oligo(dT) primer. The resulting cDNA was used directly for PCR using the listed primers:

CTTCCGGGTACCCGGAAGATGGTGTTGCTGAGAGTG (5' forward)

CGTTAAAAGCTTTTAACGTCTCATGTGTCCCATCTG (3' reverse).

PCR conditions were: 94°C, 30 s/60°C, 30 s/72°C, 2 min for 30 cycles. The PCR product was digested with *KpnI* and *HindIII* restriction enzymes, and was ligated into pCEP4 vector (Invitrogen)

Human 293EBNA cells (Invitrogen) were seeded into 96-well plates at 10⁴ cells/100 µl per well in DMEM containing 10% fetal bovine serum, and incubated at 37°C, 5% CO₂ for 20 h. Cells were transfected with 50 ng pSR α SPORT/proTNF α vector with or without 100 ng pCEP4/ADAM 10 using Lipofectin (Gibco-BRL). After 20 h, human TNF α was detected by ELISA using a human TNF α capture antibody (Pharmingen no. 18631D) and a biotinylated anti-TNF α (Pharmingen no. 18642D). The bound biotinylated antibody was detected using streptavidin-HRP (BioSource) in concert with ABTS sub-

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Fig. 1. Characterizing the TNF α convertase pool by SDS-polyacrylamide gel electrophoresis. TNF α convertase activity was purified from bovine spleen as described. Serial dilutions of final preparation were fractionated on an 8–16% polyacrylamide gel in SDS. Polypeptides were visualized by Coomassie Blue staining. Black arrows denote major protein species.

strate (KP Labs). Visible color development was quantitated at 405 nm vs. standards.

3. Results

After screening several bovine tissues for the ability to cleave ¹²⁵I-Flag-proTNFa to a 17 kDa protein (membrane fractions from kidney, spleen and lung contained detectable TNFα convertase activity, while heart and liver were inactive), we selected bovine spleen for purification of TNFa convertase. 9 g of bovine spleen membranes were fractionated as listed in Section 2. The final pool (Fig. 1) contained two major protein bands centered around 60 kDa. Fig. 2 shows the protein sequences of these bands are consistent with a known protein ADAM 10 [10], a metalloprotease with an ill-defined biological activity [11,12] (the lower band is a proteolyzed product starting at residue 285). ADAM 10 could cleave $^{125}\text{I-proTNF}\alpha$ to a 17 kDa protein (Fig. 3), and could cleave a peptide substrate between Ala and Val, consistent with the cleavage site in native TNFa. The peptide-cleaving activity was inhibited by EDTA (1 mM) and SCH43534, a peptide



Fig. 2. Amino acid sequence analysis of the TNF α convertase polypeptide. The polypeptides (black arrows in Fig. 1) were transferred to an Immobilon filter membrane, excised and sequenced. Figure shows the alignment of the obtained sequence information with that of the known metalloproteinase ADAM 10.



Fig. 3. ADAM 10 cleaves a proTNF α substrate. Aliquots of purified ADAM 10 were mixed with ¹²⁵I-Flag-proTNF α in a reaction containing 0.2 mM PMSF, 2 μ M pepstatin, 0.2 mM leupeptin and 1 mM AAPVCMK. Following incubation at 37°C overnight, samples were fractionated by SDS-PAGE on a 16% gel and radioactivity detected by autoradiography. Lanes: 1, no enzyme control; 2, ADAM 10. The upper arrow indicates the uncut proTNF α , about 21 kDa; the lower arrow denotes the 17 kDa processed TNF α band.

hydroxamate inhibitor of collagenase similar to BB2284 [9], but was not inhibited by phosphoramidon (300 μ M) and thiorphan (100 μ M).

To confirm this result, we sought to determine whether cloned bovine ADAM 10 could process proTNF α in intact cells. Screening numerous cell lines (including CHO, HeLa, IEC-6, C₂C₁₂, clone-9) showed that a human embryonic kidney cell line 293EBNA (as well as the parental cell line 293) was unable to process recombinant proTNF α , although recombinant growth hormone was effectively processed. We transfected pSR α SPORT/proTNF α with or without pCEP4/ ADAM 10, then measured TNF α production 20 h later. Fig. 4 shows that 293EBNA cannot process recombinant proTNF α (sample 1) unless co-transfected with pCEP4/ ADAM 10 (samples 2–5).



Fig. 4. ADAM 10 mediates the processing and release of proTNF α in vivo. The graph shows the amount of soluble TNF α produced with transfection of pSR α SPORT/proTNF α alone (sample 1) and after co-transfection with four independent pCEP4/ADAM 10 vector isolates (samples 2–5).

4. Discussion

We have purified an enzyme capable of proteolytically processing a recombinant proTNF α to native TNF α from bovine spleen. A homologous protein was recently purified from human Monomac 6 cells (data not shown). The protein was identified as ADAM 10 [10], a protein first purified from bovine brain based on its ability to cleave MBP [11,12]. The protein has since been found in many tissues, and is homologous to the *Drosophila kuz* protein, suggesting a role in neurogenesis [13]. We show that this protein can cleave proTNF α in vitro and in cultured human cells. This result, coupled with the numerous other proteolytic activities present in bovine tissue that conform to the inhibitor profile and cleavage site specificity established for TNF α convertase (data not shown) suggests that in vitro activity alone is not sufficient to establish a protease as TNF α convertase.

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References

- Lorenz, H.-M. Antoni, C., Valerius, T., Repp, R., Grunke, M., Schwerdtner, N., Nusslein, H., Woody, J., Kalden, J.R. and Manger, B. (1996) J. Immunol. 156, 1646–1653.
- [2] Elliott, M.J., Maini, R.N., Feldmann, M., Kalden, J.R., Antoni,

C., Smolen, J.S., Leeb, B., Breedveld, F.C., Macfarlane, J.D., Biji, H. and Woody, J.N. (1994) Lancet 344, 1105-1110.

- [3] Van Dullemen, Van Deventer, S.J., Hommes, D.W., Bijl, H.A., Jansen, J., Tytgat, G.N. and Woody, J. (1995) Gasteroenterology 109, 129–135.
- [4] Kriegler, M., Perez, C., DeFay, K., Albert, I. and Lu, S.D. (1988) Cell 53, 45–53.
- [5] Jue, D.-M., Sherry, B., Leudke, C., Manogue, K.R. and Cerami, A. 1990) Biochemistry 29, 8371–77.
- [6] Robache-Gallea, S., Morand, V., Bruneau, J.M., Schoot, B., Taggat, E., Realo, E., Chouaib, S. and Roman-Roman, S. (1995) J. Biol. Chem. 270, 23688–23692.
- [7] Mohler, K.M., Sleath, P.R., Fitzner, J.N., Cerretti, D.P., Alderson, M., Kerwar, S.S., Torrance, D.S., Otten-Evans, C., Greenstreet, T., Weerawarna, K., Kronhelm, S.R., Petersen, M., Gerhart, M., Kozlosky, C.J., March, C.J. and Black, R.A. (1995) Nature 370, 218–220.
- [8] McGeehan, G.M., Becherer, J.D., Bast, R.C., Boyer, C.M., Champion, B., Connolly, K.M., Conway, J.G., Furdon, P., Karp, S., Kidao, S., McElroy, A.B., Nichols, J., Pryzwansky, K.M., Schoenen, F., Sekut, L., Truesdale, A., Verghese, M., Warner, J. and Ways, J.P. (1994) Nature 370, 558-561.
- [9] Gearing, A.J., Beckett, P., Christodoulou, M., Churchill, M., Clements, J., Davidson, A.H., Drummond, A.H., Galloway, W.A., Gilbert, R., Gordon, J.L., Leber, T.M., Mangan, M., Miller, K., Nayee, P., Owen, K., Patel, S., Thomas, W., Wells, G., Wood, L.M. and Woolley, K. (1994) Nature 370, 555–557.
- [10] Wolfsberg, T.G., Primakoff, P., Myles, D.G. and White, J.M. (1995) J. Cell Biol. 131, 275–278.
- [11] Howard, L. and Glynn, P. (1995) Methods Enzymol. 248, 388– 395
- [12] Chantry, A., Gregson, N. and Glynn, P. (1992) Neurochem. Res. 17, 861–868.
- [13] Rooke, J., Pan, D., Xu, T. and Rubin, G.M. (1996) Science 273, 1227–1231.