

PURIFICATION OF CARP (*Cyprinus carpio*) KIDNEY CATHEPSIN C

Henneke Pangkey

Staf Pengajar pada Program Studi Budidaya Perairan,
Fakultas Perikanan dan Ilmu Kelautan. UNSRAT. Manado. 95115.

RINGKASAN

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Pemurnian enzim cathepsin C diperoleh melalui penggunaan kolom kromatografi S-Sepharose FF, Sephacryl S-200 HR, Concanavalin A-Agarose, Affi-gel 501 dan DEAE-Sephacel. Dari 160 g ginjal ikan mas diperoleh 0,33 mg enzim cathepsin C, dengan hasil pemurnian 150 kali. Berat molekul cathepsin C adalah 170 kDa yang terdiri dari 5 subunit, dengan berat molekul berkisar antara 10 sampai 20 kDa. Cathepsin C adalah oligomerik protein dan aktif sebagai eksopeptidase (pH 5) maupun endopeptidase (pH 7).

Kata Kunci: Pemurnian enzim, enzim Cathepsin C, ikan mas

INTRODUCTION

Cathepsin C [EC 3.4.14.1] is a lysosomal cathepsin^{6,10} and involves in intracellular protein degradation. Because of its specificity (the ability toward dipeptidyl derivatives), cathepsin C has other common names. It is also familiar to be designated as aminopeptidase, dipeptidyl aminopeptidase I, or dipeptidyl peptidase^{5,12}. Lately, cathepsin C was defined as a lysosomal exopeptidase which is capable of removing dipeptides sequentially from the amino terminus of a peptide chain⁶. The function of cathepsin C besides the important role in intracellular protein degradation, it appears also to operate in cell growth and neurominidase activation. In addition, cathepsin C is supposed to be involved in the function of the alimentary tract^{3,4,6}. So far, this enzyme has been purified in bluefly, frog, gastropods, squid, chicken, mouse, rabbit, lamb, ox and human⁵. The purpose of the present study is to purify cathepsin C from kidney carp *Cyprinus carpio*.

METHOD

Purification of cathepsin C

The purification method of cathepsin C was done following the method as described by Minotani¹³ with a slight modification. About 160 g of frozen carp kidney thawed at 4°C, minced, and homogenized with 5-fold of 50 mM sodium acetate buffer, pH 5.5, containing 1.0 mM DFP and 10 mM EDTA. The homogenized solution was centrifuged at 9,000 x g for 30 min to get a crude extract. The fraction precipitated with 20-70% saturation of ammonium sulfate was dissolved in a minimum volume in 50 mM sodium acetate buffer, pH 4.0 then dialyzed against the same buffer. After centrifugation at 12,000 x g for 30 min, the supernatant was applied to an S-Sepharose Fast Flow column (2.64x45 cm) equilibrated with the same buffer as in dialysis. The enzyme was eluted with a linear gradient of NaCl from 0 to 1.0 M in the same buffer. The active fractions

were pooled, concentrated by ultrafiltration (Amicon YM-10). The enzyme concentration was subjected to a Sephacryl S-200 column (1.5x100 cm) equilibrated with 50 mM sodium acetate buffer, pH 5.5, containing 0,2 M NaCl and 10 mM EDTA and eluted with the same buffer. The active fraction was collected and dialyzed against 50 mM potassium phosphate buffer, pH 6.0, containing 0,15 M NaCl and applied to a Concanavalin A-Agarose column (1.0x8.5 cm) equilibrated with the same buffer as in dialysis. It was washed with the same buffer followed by 0.1 M methyl α -D-mannopyranoside in the buffer. The active fractions were pooled and put on an Affi-Gel 501 column (1.0x5.0 cm) equilibrated with 50 mM sodium acetate buffer, pH 5.0, containing 0.15 M NaCl and eluted with the same buffer containing 10 mM cysteine. The active fractions were pooled and put on a DEAE-Sephacel column (0.5x7.7 cm) equilibrated with 50 mM sodium phosphate buffer, pH 7.5. The column was washed with the same buffer, then eluted with 50 mM sodium dihydrogen phosphate and followed by 0.3 M NaCl in 50 mM sodium dihydrogen phosphate.

RESULT AND DISCUSSION

On the present study carp cathepsin C could hydrolyze H-Gly-Phe- β nap and Z-Phe-Arg-MCA as well. The latter was found to be the excellent substrate for cathepsin B, L and S which are the endopeptidase. This finding was proved that cathepsin C was an endopeptidase too. Previous study described that cathepsin C could not digest Z-Phe-Arg-MCA⁷, however, bovine spleen cathepsin C can degrade Z-Phe-Arg-MCA efficiently¹¹. The purification step on column chromatographies could be seen in Fig. 1 – 5 and the enzyme was purified about 150 fold and produced 0.33 mg cathepsin C. Cathepsin C was not sufficiently separated from cathepsin S on S-Sepharose column chromatography because the main peak of cathepsin S was eluted just before cathepsin C. It was effectively separated from cathepsin S on Sephacryl S-200 gel filtration because the molecular mass of cathepsin C was larger (170 kDa) than cathepsin S¹⁴ (37 or 40 kDa). Cathepsin C was expected to contain carbohydrate. The enzyme was found to bind strongly to a Concanavalin A-agarose column. This property shares with cathepsin H (an exopeptidase)^{1,14} and cathepsin L (an endopeptidase)^{2,8-9}. The affinity chromatography, Affi Gel 501, also was effective in the purification of this enzyme. Finally, on the DEAE-Sephacel column chromatography, it was found that this column was very effective in purification of this enzyme. The column has the capacity to separate cathepsin C from the other protein. The purified cathepsin C was examined through the analytical polyacrylamide gel electrophoresis using a 16-18% gradient gel (Fig. 6). A single protein band was seen when a gel was stained for protein with silver stain and the protein band was reacted with anti-rat cathepsin C (Fig. 7).

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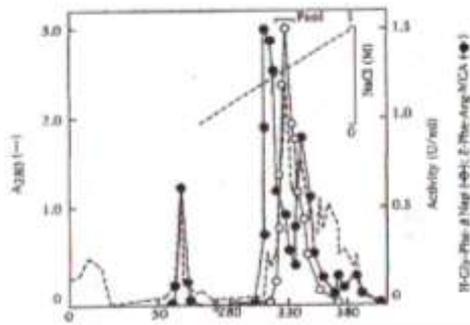


Fig.1 Chromatography of the ammonium sulfate fraction on S-Sepharose column

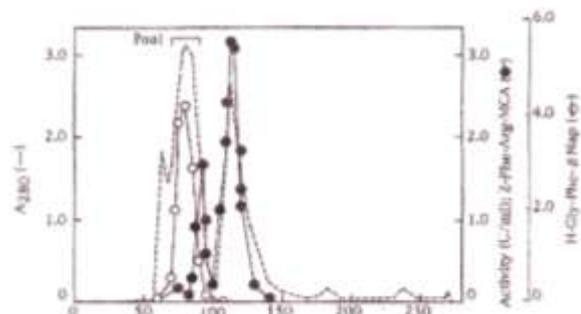


Fig.2 Chromatography of S-Sepharose fraction on a Sepharyl S-200 HR column

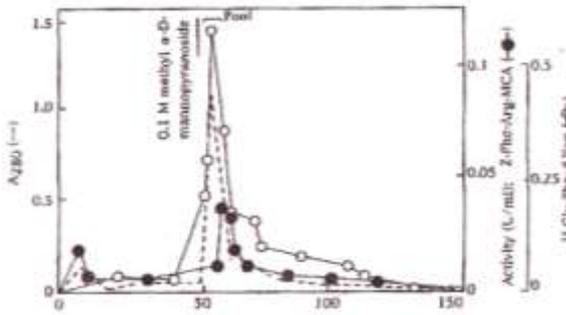


Fig.3 Affinity Chromatography of S-Sepharose HR fraction on a Concanavalin A-agarose column

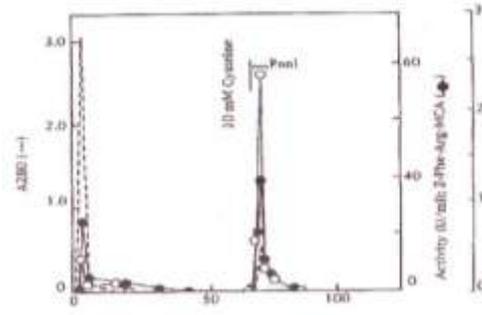


Fig.4 Affinity Chromatography of Con A-agarose fraction on Affi-Gel 501 column

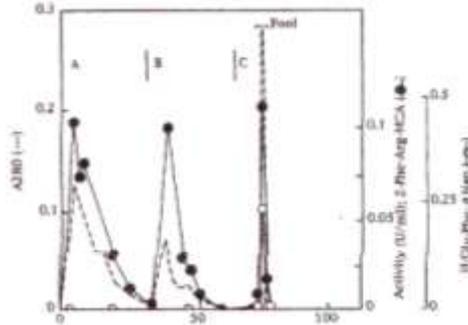


Fig.5 Affinity Chromatography of Affi-Gel 501 fraction on DEAE-Sephacel column



Fig.6 Analytical polyacrylamide gel electrophoresis and immunoblotting of the purified carp cathepsin C

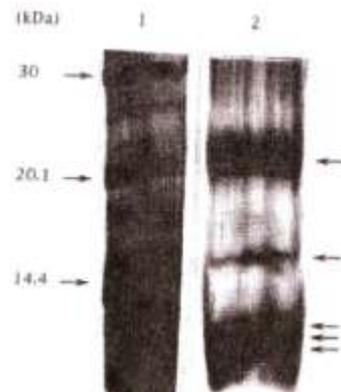


Fig.7 SDS-PAGE of carp cathepsin C on 16-18 gradient gel