

ISOLATION OF A COMPLEX OF THE SUBCOMPONENTS OF THE ACTIVATED FIRST COMPONENT OF COMPLEMENT, C1f-C1s, FROM ACD-HUMAN PLASMA

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1. Introduction

The first component of complement, C1, is a Ca^{2+} -dependent complex of a molecular weight of about 10^6 daltons [1] and can be dissociated into three subcomponents, C1q, C1r and C1s, by treatment with EDTA [2]. It is well known that C1q has a binding activity to immune complexes [3], and the active form of C1f converts enzymatically C1s into active C1s [4] which is an estero-protease and represents the biological activities of the activated C1 [5].

However, little is known how these subcomponents are interacted to unite the C1 macromolecule.

This paper presents evidence indicating that C1f and C1s were isolated as a Ca^{2+} -dependent complex form by polyethylene glycol fractionation of ACD*-human plasma followed by chromatographies with IgG-Sepharose 6B and DEAE-Sephadex A-50. This result suggested that the inter-subcomponent bonds in the C1 macromolecule are not equivalent, and C1f and C1s have a high affinity via Ca^{2+} for each other but not for C1q. In addition, this isolation procedure seemed to be practical for rapid and concomitant purification of C1f and C1s from ACD-human plasma.

2. Materials and methods

ACD- and heparinized human bloods were

obtained from the blood bank of Japanese Red Cross.

2.1. Precipitation of C1 from ACD-human plasma with polyethylene glycol

One hundred ml of ACD-human plasma were mixed with 10 ml of 50% (W/V) of polyethylene glycol (# 4000) under stirring at 4°C. After 30 min, the precipitate was collected by centrifugation at 3 000 rpm for 15 min and extracted with 50 ml of 0.05 M NaCl. The insoluble materials which contained C1, fibrinogen and other macromolecular weight plasma proteins were then dissolved in 20 ml of 0.02 M Tris-HCl buffer, pH 8.5, containing 0.15 M NaCl and allowed to clot by the addition of 1 ml of 1 M CaCl_2 overnight at 4°C. The defibrinated fraction was used as the starting material for further purification of C1.

2.2. Determination of C1

C1 activity was determined by measuring either C1s esterase activity with ATTEE as substrate [5] or hemolytic activity with EAC4 cells [6]. A functional activity of C1f to activate C1s was determined by the method of Naff and Ratnoff [4], with slight modifications. In the most case, aliquots of 10 μl of fractions were incubated with C1s (about 2 μg) in 1 ml of 0.01 M Tris-HCl buffer, pH 8.5, containing 0.10 M NaCl and 5 mM EDTA-3Na at 37°C for 10 min, and the esterase activity of C1s thus activated was determined with AGLME [4] as substrate. The value was corrected for the hydrolysis of AGLME by the fractions alone. Human C1s was purified from heparinized plasma according to the method of Sakai and Stroud [7].

* Abbreviations: ATTEE, *N*- α -acetyl-L-tyrosine ethyl ester; AGLME, *N*- α -acetylglycyl-L-lysine methyl ester; ACD, acid citrated dextrose.

3. Results

When ACD-human plasma was fractionated with

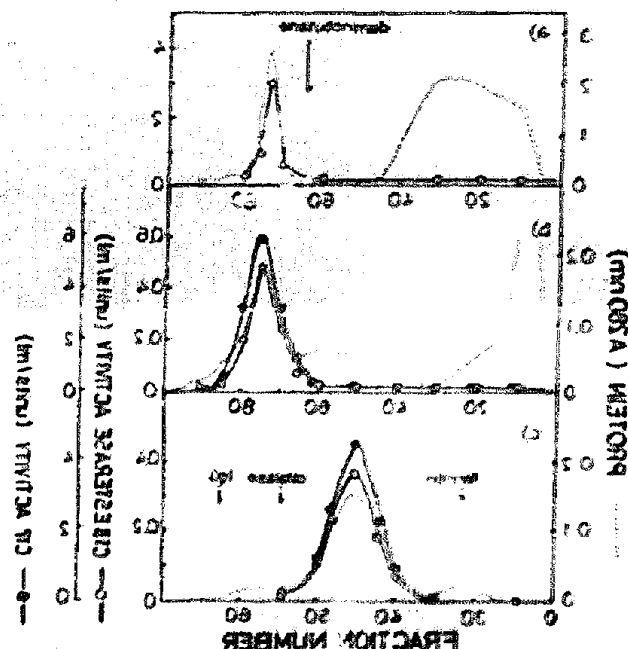


Fig. 1. Chromatographic profiles of the C fraction obtained from 200 ml of human plasma was diluted with equal volume of water and subjected to the affinity chromatography with a column (2.5 x 2 cm) of IgG-Sepharose as described by Bing [9]. The activity was eluted with 0.01 M Tris-HCl buffer, pH 8.2, containing 0.02 M NaCl and 2 mM CaCl₂. Fractions of each 2 ml were collected at room temperature. The amount of ATE hydrolyzed by C₁ were determined with the hydrolytic reagents [10], and one unit of esterase activity was defined as the activity capable of hydrolyzing 1 μ mol of the ester per min. (b) The C fraction eluted with the dialysis solution was dialyzed against 0.01 M Tris-HCl buffer, pH 8.2, containing 0.02 M NaCl and 2 mM CaCl₂, overnight and applied to a column (2.5 x 2 cm) of DEAE-Sepharose A-20 equilibrated with the dialysis buffer. The column was eluted by linear gradient increase of NaCl concentration with each 200 ml of 0.02 M and 0.30 M NaCl in the buffer. Fractions of each 4 ml were collected at 4°C. The activity of C₁ was expressed as the AUMU. The esterase activity of C₁ developed by the action of C₁. The peak fractions 70-80, were pooled, concentrated by ultrafiltration to 2 ml and gel filtered with a column (2 x 80 cm) of Sepharose 6B equilibrated with 0.01 M Tris-HCl buffer, pH 8.0, containing 0.12 M NaCl and 2 mM CaCl₂. The molecular weight was estimated according to the method of Andrews [15] with the standard proteins of known molecular weight (Boehringer's protein calibration kit II). Fractions of each 4 ml were collected at 4°C.

As shown in fig. 1c, the C₁ as measured by C₁ and C₂ activities was eluted from the calibrated column of Sepharose 6B as a single peak of the molecular weight of 3.2×10^5 daltons. This value was smaller than those of the C₁ macromolecule, 10^6 daltons [1], or C_{1d} 4×10^5 daltons [13], and was comparable to the sum of the molecular weights of C_{1f}, 1.8×10^5 daltons [14], and C_{1g}, 1.1×10^5 daltons [17].

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Fig. 2. Polyacrylamide gel electrophoresis. (a) The peak fraction (75 in fig. 1b). (b) The peak fraction after addition of EDTA-3Na to 2 mM. (c) The EDTA-treated fraction as in (b) after recalcification to final 10 mM. (d) Gel electrophoresis was performed with 2% polyacrylamide gels [11] and proteins were stained with Coomassie blue. (b) Electrophoresis of the peak fraction (42 in fig. 1c) in the presence of 0.2% sodium dodecyl sulfate with a 2% polyacrylamide gel [17].

daltons [15]. Gel electrophoresis of the peak fraction in the presence of sodium dodecyl sulfate exhibited two protein components of the molecular weights of 1.3×10^5 daltons and 1.1×10^5 daltons, respectively. The protein components corresponding to Clq itself or its non-covalent subunit of the molecular weight of 7×10^4 daltons [16] were not detected in the electrophoretogram (fig. 2d). In addition, EDTA-treatment of the peak fraction resulted in the dissociation into two protein components (fig. 2b), and Clf and Cls activities eluted from the unstained sectioned gels were found to be associated with the protein components in the β - and α -globulin regions, respectively. The dissociated two components were found to be complexed again with each other by the addition of Ca^{2+} (fig. 2c).

From these results, it was proposed that the Cl thus isolated was a Ca^{2+} -dependent complex of Clf and Cls.

4. Discussion

As presented above, the Cl thus isolated from ACD-human plasma was found to be lacking in Clq subcomponent and was proposed to be a Ca^{2+} -dependent complex of Clf and Cls.

As an additional information to support this conclusion, experiments in progress indicate that the purified Clf-Cls can not bind to the IgG-Sepharose 6B and needs the addition of Clq for its binding to the IgG-Sepharose 6B. In the present experiment, Clq was found to be recovered in the unadsorbed fraction of the DEAE-Sephadex A-50 chromatography. The fact that Clq and the Clf-Cls complex were separated from each other by DEAE-Sephadex A-50 chromatography even in the presence of Ca^{2+} may suggest the affinity of Clq for the Clf-Cls complex to be very weak.

The question on the origin of the Clf-Cls complex remains to be solved. However, the precursor complex, Clr-Cls, can be obtained from ACD-human plasma by DEAE-Sephadex A-50 chromatography and gel filtration on Sephadex G-200 (unpublished data). So, it seems likely that the Cl macromolecule

is partly dissociated by the chelating action of anti-coagulant citrate into Clq and the Clr-Cls complex and then activated during polyethylene glycol fractionation. Activation of Cl by polyethylene glycol was not unexpected matter, since polyethylene glycol was known as a non-immunological effector of complement system in human serum [18].

In relation to the present observation, Valet and Cooper have recently observed the affinity of Clr for Cls to be so high as to form a stable Ca^{2+} -dependent complex [19].

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