Ascitic Antigen CA125: Study of Dissociation Forms and its Final - Stem Structure

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ABSTRACT

Aim: Study of dissociation forms of the ascites antigen CA125 and its stem structure – the final form of its dissociation.

Study Design: Research paper.

Place and Duration of Study: Department of biochemistry, N.I. Pirogov Russian National Research Medical University, laboratory of immunochemistry and laboratory immunology, N. N. Blokhin Russian Oncological Scientific Center.

Methods: Next materials were used in the work: ascitic fluid from patients with ovary cancer (14 patients), blood serum from patients with ovary cancer (28 patients), 40 donors (24 women and 16 men). Electrochemoluminiscent method, ELISA, electrophoresis and direct binding of labeled antibodies with antigen in polyacrylamide gel, reaction of precipitation, chromatography were used to study forms of dissociation of the antigen CA125.

Results: Dissociation forms of the antigen CA125 isolated from ascites fluid of patients

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with ovary cancer, demonstrated at native conditions of electrophoresis in polyacrylamide gel three main forms: gamma – globulin with molecular weight (MW) ~500 kDa, alpha-globulin with MW ~ 230 kDa and beta-globulin with MW ~110 kDa. All 3 forms bound I-MAB to CA125 and were isolated from the polyacrylamide gel during preparative electrophoresis. All these forms dissociated to a single protein band with MW 55 kDa that migrated in the albumin zone and did not bind I-MAB to CA125 under denaturing conditions of polyacrylamide gel-electrophoresis with 2% sodium dodecyl sulfate. This form also demonstrated a single protein band corresponding to MW 75 kDa in reducing conditions of electrophoresis. It has been established that the form with MW of 55 kDa represents a complex of three different polypeptides, which are identical to heavy chains IgG, albumin and unknown thermostable protein coupled to albumin. This thermostable protein was revealed in all immunoreactive forms of CA125 and all investigated commercially available samples of albumins in the reaction of precipitation with polyclonal antibodies. Antibodies to these proteins have also been revealed in commercial antisera to human serum proteins in a titer 1:40.

**Conclusion:** The final form of dissociation of immunoreactive forms of the antigen CA125 is represented by a complex of three different polypeptides with MW of 55 kDa that did not bind I-MAB to CA125.

**Keywords:** Forms of dissociative ascetic antigen CA125; final form dissociation of antigen CA125; thermostable protein coupled to albumin; bow-string effect; albumin spiral.

1. **INTRODUCTION**

Antigen CA125/MUC16 represents a slow reacting marker: is level is increased in the serum of patients with ovary cancer at III-IV stage of the disease and, predominantly, in patients with ascites [1]; in the more early phases of the evolution of tumorous process it uninformative [2]. Functions and devotion of this structure is still obscure, despite application of mass-spectrometry methods that are oriented on the increasing of specificity of a test for ovary cancer [3]. The nature of the immunoreactive epitope of the antigen CA125 is known and the existence of a final form of its dissociation is under discussion and remains to be elucidated [4]. Nevertheless, it has been shown that the final form of dissociation of the antigen CA 125 represents a complex of different polypeptides with MW of 55 kDa that includes polypeptides immunochemically identical to serum albumin (HSA), heavy chains of IgG (IGHC) and unknown thermostable protein coupled to albumin - TPC.A [5]. Difficulties in identification of this form are that it does not bind I-MAB to CA125, but reacts with polyclonal antibodies in the reaction of precipitation with A-HSA line identical to HSA, and with A-IGHC – line identical to IgG. With polyclonal antisera to donor serum proteins (AD) absorbed with HSA and IgG, this form reacts not only with TPC.A and donor sera, but also with all isoforms of CA125, which bind I-MAB to CA125. The antiserum lost immunoreactivity after its absorption by donor sera (5). It is necessary to note that A-HSA reveals up to 9 fragments and complexes of HSA with MW from 11 to 260 kDa in the serum and urine [6,7].

Use of specific units for measuring the serum level of CA125 did not allow estimating its real content in the sera of healthy donors. Only in 2001 the real concentration of the antigen in the sera of donors was determined as 5-10 mkg/ml with the use of monoclonal antibodies to CA125 [8]. It meant that CA125 is a minor serum protein. Concentration of the main immunoreactive form of ascitic CA125 in the donor sera determined with the use of polyclonal antibodies was equal to 500 mkg/ml (titer 1:128) that makes it possible to
consider this protein as typical serum protein [9]. A similar protein was revealed many times in tumors, ascites, sera and everywhere its serum origin was seen: glycoprotein «alpha-2H-ferroglycoprotein» [10], «macromolecular serum ferroglycoprotein» [12], «albumin-IgG complex» [13], «tumor-associated ascites IgG» [14], «IgG-like glycoferroprotein serum» [5] and «peroxidase active IgG-like glycoferroprotein ascites» [15].

Study of CA125 only with the use of labeled monoclonal antibodies (l-MAB) did not allow elucidating dissociative forms that did not bind l-MAB to CA125. The most noticeable work in this field is a very thorough investigation made by a group of M.T. de Los Frailes [16]. Immunoreactive CA125 was isolated from ascite of patients with ovary cancer by the use of immobilized MAB (OC125) to CA125 on Affigel 10. Under polyacrylamide gel electrophoresis in native conditions (n-PAGE), immunoreactive CA125 did not penetrate the gel and was at the start position in the separating PAG. Under mild denaturing conditions of SDS-PAGE with 0,1% SDS (these conditions are very close to natives ones) the immunoreactive form of CA125 demonstrated two proteins bands with MW of 205 and 55 kDa, and also a third form that did not penetrate the gel (MW ~500 kDa ). From this it follows that, from one side, the dominant form of CA125 partly dissociates to the final form of dissociation with MW of 55 kDa, and, from another side, re-associates, partly, with formation of a large molecular form that did not penetrate the gel and was at the starting position. In 2% SDS – PAGE authors observed only one band with MW of 55 kDa. In reducing conditions of SDS- ME-PAGE there also was on band, but with MW of 75 kDa. Only one band with MW of 205 kDa bound l-MAB to CA125 in Western blotting. The final form of dissociation with MW of 55 kDa did not bind antibodies. From this follows that the final form of dissociation may be only studied with the use of polyclonal antibodies. This unique structure has not attracted attention of researchers yet and is still opened for investigation. We consider the present work as continuation of works by M.T. de los Frailes with addition of polyclonal antibodies to study the final form of dissociation of the antigen CA125, which does not bind l-MAB to CA125.

Results of investigation of dissociative from of the antigen CA125 and its stem structure – the complex of polypeptides that do not bind l-MAB to CA125, are presented in this work.

2. MATERIALS AND METHODS

2.1 Isolation of Antigen CA125

Antigen CA125 was obtained from acid-soluble fraction of ascites and pooled donor serum from 50 women (SD50) [17]. Content of immunoreactive (IR) CA125 was determined by electrochemiluminiscent method at «Hoffmann-L Roche Elecsys 2010» (Germany) and by ELISA method with use of l-MAB OC 125 to antigen CA125. SD50, in which concentration of IR CA125 was equal to 34,6 U/ml или 5 mg/ml, ascites - 3140 U/ml or 448 mg/ml were used as control samples, and antigen CA125, isolated, according to Davis, from SD50 - 63,6 U/mg, serum of female patient B. (287 U/ml), GFP470 (48,3 U/mg), A230 (514 U/mg), CP55 (1,6 U/mg) and IgG (0,02 U/mg) as standard samples. Ascites without hemolysis were used in the work.

2.2 Isolation of GFP, CP55, the Complex of “TPC.A – Albumin” and IGHC

GFP was obtained from SD50 by gel-filtration of serum fraction insoluble under 30% saturation by ammonia sulfate with subsequent collection of fractions with MW ~ 500 kDa.
CP55 (zone of albumins) was isolated out of SD50 diluted 1:8 in tube variant of native PAGE. *Thermostable complex «ТПС.А- albumin»* was prepared by heating of albumin preparations – 2-10 mg/ml (100°C, 15 min). All isoforms of ascite GFP were isolated ex tempore, due to extremely fast variability of forms. A fraction with MW 55-60 kDa was isolated from IgG (2mg/ml) by gel-filtration under control of precipitating test-system for IGHC.

### 2.3 Renaturation of Albumin

5-10 mg/ml of BSA (Sigma, USA) dissolved in Tris-glycine buffer (0,05 mM tris, 4 mM of glycine, pH 8.3), were incubated in boiling water-bath for 15 min.; a pellet of denatured protein was separated by centrifugation, suspended in stock Tris-glycine buffer (50 mM tris, 0,4 M glycine, pH 8,3) and renaturated during 48 hours at + 37°C; ~ 0.5% of initial protein is renaturated.

### 2.4 Polyclonal Antiserum against Human Serum Proteins

Exclusively commercially available antiserums to human serum proteins were used in our work: (Russia; Binding Site Limited, UK; Sigma, USA), anti-IgG (Russia) and antiserum to gamma-, alpha131 and μ- chains of immunoglobulins G, A, M (Bio-Rad, USA). Rabbit polyclonal antiserum to antigen CA125, GFP and CP55 were prepared in our laboratory.

### 2.5 Polyacrylamide Gel Electrophoresis; Direct Binding of Labelled Antibodies to Antigen in PAG

Electrophoresis in slabs and tubes were performed under native, denaturing and reducing conditions. Slabs/tubes were incubated with commercial preparations of l-MAB (to antigen CA125 and AFP) and мечеными polyclonal antibodies to PSG for IFA (Vector Best, Russia) during 2 hour at +37°C after native PAGE, then washed three times by water for 15 min., incubated (+37°C, 15 min) with c 3,3,5,5, - tetramethylbenzidine (TMB) and washed by water. Another substrate - 3,3, diaminobenzidine tetrahydrochloride – is oxidized by all isoforms of ascite antigen CA125, with exclusion of the final dissociation form, without pre-incubation with l-MAB (Fig.1, b).

### 2.6 Determination of Peroxidase-Like Activity (PA) of Proteins in n-PAG

After PAGE slabs/tubes were incubated in DAB solution (Sigma, USA) with hydrogen peroxide (10 mg of DAB are dissolved ex tempore in 20 ml of 0,05 M tris-HC1 buffer, pH 7,5 and 0,3 ml of 3% H2O2 was added) during 15-30 min. at 37º C and were washed by water during 16-18 hours at + 4°C. HSA, alpha-fetoprotein (AFP) and placentaspecific-glycoprotein (PSG1), which do not oxidize DAB, were used as a control samples for PA specificity of CA125 isoforms.

### 2.7 Revealing of Proteins, Ferro - and Glycoproteins in PAG.

Proteins were stained by 0,25% Coomassie G-250 in ethanol : acetic acid : water (5:1:4) for 15 min and destained in water during 15-30 min. in boiling water-bath. To reveal ferroproteins plats/tubes of PAG were incubated during 16-20 hours in a mixture: 3% K4[Fe(CN)6]×3H2O : 10% HCl (1:1 ex tempore) and washed in 5% HCl [12]. Ferroproteins
are stained in a blue color. Fe$^{3+}$ in GFP and «antigen CA125» is bound weakly and is not practically determined after boiling. Glycoproteins were determined by Schiff-iodine acid [11].

2.8 A Reaction of Direct Precipitation of Murine I-MAB

Labeled monoclonal antibodies act as antigen in reaction of direct precipitation. Well-known property of cross-reaction of human and murine IgG is in the basis of the method [19]. To reveal murine I-Mabs any commercial antiserum to human serum proteins or anti-IgG were used on the line of precipitation with I-Mab was developed by a substrate [5].

2.9 A Reaction of Mixed Precipitation with m-MAB

Studies with use of this reaction have been performed and documented by the author of the method V.S. Poltoranina in G.I. Abelev's laboratory [19]. Also the method of direct precipitation of I-MAB in agar gel and the method of direct binding of I-MAB with antigen in PAG have been reproduced and the results have been confirmed by the reaction of mixed precipitation of I-MAB to CA125.

3. RESULTS AND DISCUSSION

High mobility of dissociation-association processes in GFP and its isoforms is mostly expressed in ascites of patients with ovarian cancer (Fig.1, a): from one side, GFP, A230 and B110 dissociate in partial to final form CP55 (Fig.1, a 2-4), from the other side, CP55 reassociates partly to GFP (Fig.1, a 5, 6) that reflects pulsating form of protein bodies existence – the property characteristic for many enzymes. PA results for ascites GFP isoforms till and after incubation with I-Mab to antigen CA125 were absolutely equal (Fig.1 b). the final form of GFP dissociation – CP55 does not demonstrate PA, but reassociated form of CP55 - GFP, does (Fig. 1, b 5).

After 7 months this form – CP55 represents up to 6 protein bands in 186 the tube variant of n-PAGE, and one of which is a sloping band (Fig.1, a b). Such an inclined protein band was observed by group of Frailes in the final form of dissociation of antigen CA125 with MW 55 kDa, but in plat variant of reducing PAGE [16].

Moreover, it has been established that the human, bovine and rat albumin samples isolated ex tempore from the serum demonstrated a protein with peroxidase-like activity that was readily stained for Fe$^{3+}$ at the start position in the separating gel under n-PAGE. IgG did not exhibit such properties ans was non identical to ascitic and serum GFP [5]). We observed the direct binding of I-MAB to CA125 with the antigen of female patient B. (287 E/ml of CA125) in case of the form B110 only, but in difference from ascitic CA125 forms, it did not exhibit peroxidase-like activity with DAB (Fig. 1, c). We did not observe peroxidase-like activity in samples of donor sera. 1-3 forms of the proteins exhibit direct binding of I-MAB to CA125 with antigen CA125 in different samples of ascites (Fig. 1, d11). No binding of antibodies to CA125 was revealed in the donor sera by this method. However, all isoforms of the ascites antigen CA125 are presented in the sera from healthy donors, but they did not bind I-MAB to the antigen CA125 and did not exhibit peroxidase (Table 1).

Fig. 1. Electrophoresis under non-denaturing conditions of tube PAG and direct binding of I-MAB with antigen in PAG

Table 1. Some properties of main forms of the antigen CA125

<table>
<thead>
<tr>
<th>Properties</th>
<th>G500 ascites</th>
<th>A230 ascites</th>
<th>B110 ascites</th>
<th>CP55 ascites</th>
<th>GFP serum</th>
<th>IgG serum</th>
<th>HSA serum</th>
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<tbody>
<tr>
<td>Binding of I-MAB to CA125 in ELISA</td>
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<tr>
<td>Direct binding of I-MAB to CA125 in PAG</td>
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<td>A dissociation form (kDa):</td>
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<td>A. in SDS-PAGE</td>
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<td>B. in SDS-ME-PAGE</td>
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<td>Peroxidase activity</td>
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<td>Staining for Fe&lt;sup&gt;3+&lt;/sup&gt;</td>
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<td>Staining for glycoproteins</td>
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</table>

GFP and serum CP55 are represented in the denaturing PAGE by a single protein band with MW ~ 55 kDa; also by a single band, but with MW ~ 75 kDa in reducing conditions of PAGE. Under these conditions, IgG represents 2 bands with MW ~55 and ~12 kDa, and the band with MW 75 kDa is absent here (Table 1). Molecular weight of GFP and TPC.A before and after reduction of S-S bonds by mercaptoethanol is the same and increases by ~ 20 kDa that HSA, BSA and rat albumin (Fig. 2, a 6-14) and also heated and renatured BSA (Fig. 2,
b) and HSA permanently demonstrate. We did not observe such phenomenon with IgG (Fig. 2, a2) and AFP.

![Electrophoresis of serum proteins in reducing conditions of denaturing 10% PAG in plats](image)

**Fig. 2. Electrophoresis of serum proteins in reducing conditions of denaturing 10% PAG in plats**

*a* – the “bowstring” effect: 1 – molecular weight markers (14.4 - 97 kDa); 2 – IgG; 3-5 – serum GFP; albumin). Lines 1, 5 contain 2% SDS and 5% ME; 2-4 - 2% SDS; 6-8 – serum CP55; 9-11 – BSA; 12-14 – rat albumin. 1, 2, 5, 8, 9, 14 lines contain 2% SDS and 5% - ME; 3, 4, 6, 7, 10 -13 lines contain 2% SDS; *b* – the spiral of albumin: 1 – thermostable BSA; 2-5 – renatured BSA: (in lines 2 and 4 – the spiral of albumin (magnified by 3,5 times).

The transitional form from low to high level of molecular weight of a structure is presented by sloping line under angle ~ 30° that links different height levels (Fig. 3, a). Expected number of horizontal bands is absent; because the bladder bowstring in projection hides the number of horizontal grades by united sloping line. The «bowstring» effect was reproduced with GFP, antigen CA125, CP55, TPC.A and all native, heated and renatured preparations of HSA and BSA and also with rat albumin, i.e. albumin in reducing conditions is “jumping up” (Fig. 2). Seemingly, this is a key stage in the mechanism of functional structures reconstruction on the basis of the stem complex «TPC.A–albumin», in which amount ratio is equal to ~1:30, and molar ~ 1:5 (seemingly, TPC.A has a molecular weight of 11.3 kDa).

Ability of this complex to form the «bowstring» in the tube variant of n-PAGE (Fig. 1, a, b), i.e. in the absence of exogenic thios, indicates, at the first glance, participation of mercaptalbumin, which accounts 2/3 molecules of albumin, in the process. However, the cysteine residues density in AFP and IGHC structures is higher than those in HSA, but they did form bowstring even in the presence of thios. It is also necessary to note that all albumin molecules (without exclusions) acquire a form with MW of 75 kDa in reducing conditions of
PAGE (Fig. 2, a). In renatured BSA the «bowstring» line rises on closed helix from 2 to 75 kDa (Fig. 2, b 2,4), and a ghost/traces of a second helix in the upper levels (Fig. 2, b 2,4), do not allow excluding its possible double nature.

Revealing of antigen TPC.A by reaction of precipitation in all serums and commercially available human preparations: HSA, IgG, and also in the antiserum to IGHC (Fig. 2, c), i.e. in places where HSA- depot for TPC.A evidences its dissemination. The effect of «bowstring», observed with BSA and rat albumin, indicates existence of functionally similar structure in animal proteins. Antibodies to TPC.A were revealed in all commercially available antisera to human serum proteins in a titer 1:8 (Sigma,USA) and in a titer 1 : 40 (Binding Site Limited, UK; Russia). It is necessary to note that the titer of antibodies to TPC.A in antisera from different Russian and UK firms was stably high and was equal, as a rule, to 1:40 in a minimum. High titer of antibodies has been also revealed in all antisera against serum fractions or native sera of healthy donors and patients. Hence, a priori, we could expect high titer of antibodies to TPC.A in all anti-HSA sera after absorption of antibodies against HSA.

We suggest that extracellular N-domain of CA125 (1-1638 a.a.) may represent a full primary structure the isoform A230 of serum, and the primary structure of TPC.A is accurately duplicated in its peptide motifs a. a. 421-524 and a. a. 641-742 [20]. The thermostable pair – «TPC.A-albumin», seemingly, is a stem structure if all CA 125-containing complexes up to supercomplexes with MW of 2700 kDa [18]. Number, weight and content of the complexes may depend on the number of bound toxic products of cell degradation (Fe ions, carbohydrates, heme groups etc.) that is clearly seen in ascites of ovary cancer patients. Seemingly, the ascites is a depot of toxins and a barrier for their penetration into the blood due to these complexes, which bind toxins and perform a sanitary role. For example, concentration of antigen CA125 in ascites (volume of 10 l) of the patient B. was equal to 9700 U/ml, and in the serum of the patient - 287 U/ml [2].

Hence, the final form of dissociation of all immunoreactive forms of the antigen CA125 is represented by a complex of polypeptides with MW of 55 kDa that includes IGCH, HSA and TPC.A, bud is not able to bind I-MAB to CA125. Ability of CP55 to re-associate and newly form peroxidase-active high molecular complexes with MW of 500 kDa (Fig.1, a5, b5) allows ranking CP55 among the stem structure of the antigen CA125. This unique structure is extremely interesting and it is impossible to name it as «dead»: it is always in movement. It is necessary to note the main difference of immunoreactive forms of ascitic CA125 from immunochemically similar serum forms – all immunoreactive forms of the ascites protein acquire peroxidase activity.

On the other side, identification of TPC.A highlights ambivalent nature of albumin, because it is difficult to explain heterogeneity, pulsating form of protein structure existence, reversible complexing and restoration of a denatured protein due to a helix principle only by «conformational adaptability» of a monovalent structure of albumin and this issue is of interest for further studies.

4. CONCLUSION

- Ascites antigen CA125 and IgG-like serum glycoprotein represent a complex immunochemically identical structures that includes IgG, HSA and TPC.A.
- All main isoforms of the antigen CA125 are presented in donor sera, but they do not bind I-MAB to CA125 and do not exhibit peroxidase activity.
• The final form of dissociation of all isoforms of the ascitic antigen CA125 and IgG-like donor sera glycoferroprotein is presented in denaturing SDS-ME-PAGE by a single protein band with MW of 55 kDa that represents a complex of three polypeptides: HSA, IGHC and TPC.A. This structure demonstrates a form with MW of 75 kDa in reducing SDS-ME-PAGE.

• Human, bovine and rat albumins demonstrate a bowstring effect in reducing SDS-ME-PAGE, and renatured BSA – bowstring twisted into a spiral.

• TPC.A is not identical to HSA and IGHC, but is tightly coupled to HSA. The pair «TPC.A-HSA», seemingly, is a stem structure for reconstruction of chelating supercomplexes that bind toxic products of cells degradation.

CONSENT

Not applicable.

ETHICAL APPROVAL

All authors hereby declare that “Principles of laboratory animal care” (NIH publication No. 85-23, revised 1985) were followed as well as specific laws of the Russian Federation where applicable. All experiments have been examined and approved by the appropriate ethics committee.

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COMPETING INTERESTS

Authors have declared that no competing interests exist.

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