

Clustering of neutrophil leucocytes in serum: possible role of C1q-containing immune complexes

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SUMMARY

Clustering activity for neutrophil granulocytes was generated in pooled normal human serum (NHS) by incubation of the serum with preformed IgG aggregates, but not in heat-treated NHS (56°C, 30 min), indicating that the function was complement-dependent. Judging from results of experiments with complement-deficient sera, and serum depleted of C1q, factor D and properdin, recruitment of the complement system beyond C1 was not required for induction of the activity. Zymosan treatment of NHS resulted in some neutrophil clustering activity, but recombinant C5a had a limited effect. C1q added to heat-treated NHS in conjunction with preformed IgG aggregates supported neutrophil clustering in a dose-dependent manner. The serum C1q inhibitor, a chondroitin 4-sulphate proteoglycan known to interact with the collagenous part of C1q, clearly reduced neutrophil clustering in heat-treated NHS supplemented with C1q and IgG aggregates. The C1q inhibitor also reduced the inherent neutrophil clustering activity of some sera from patients with systemic lupus erythematosus (SLE). Neutrophil clustering activity in SLE serum was earlier shown to be inversely related to the number of circulating neutrophils *in vivo*. Although the precise mechanisms remain unclear, we propose that C1q-containing immunoglobulin complexes mediate neutrophil clustering through C1q receptors, and that this might contribute to pathogenesis of immune complex diseases such as SLE.

Keywords neutrophil granulocytes neutrophil aggregation complement component C1q
immune complexes systemic lupus erythematosus

INTRODUCTION

Intravascular formation of neutrophil granulocyte aggregates is thought to be of pathogenetic significance in systemic lupus erythematosus (SLE), particularly with regard to development of central nervous system and pulmonary disease manifestations [1-3].

We have described a simple assay for *in vitro* assessment of neutrophil clustering or aggregation in serum [4]. Sera from patients with severe SLE gave increased clustering of neutrophils from healthy donors. The clustering activity varied with the clinical course of the disease and was inversely related to the number of circulating neutrophils in the patients. C1q-binding immune complexes and neutrophil clustering activity were correlated, and clustering activity could be generated in normal serum with preformed IgG aggregates, indicating that immune complexes in conjunction with complement could be responsible.

The purpose of the present study was to identify possible complement-related ligands for induction of neutrophil cluster-

ing in human serum with preformed IgG aggregates. C1 subcomponents together with C4 and C3 proteins are known to be fixed to complement-activating immunoglobulin complexes [5], and contribution of C5-derived peptides generated in the course of complement activation [6] was also considered. The results suggested that C1q-containing immunoglobulin complexes were critically involved, and that further recruitment of the complement system was not required in the reaction.

MATERIALS AND METHODS

Human sera

Sera were frozen in aliquots at -70°C within 4 h after blood sampling. Pooled normal human serum (NHS) from 20 healthy hospital staff members was used as a reference in all experiments. Sera from persons with homozygous deficiency of C1q, C4, C2, C5 or C7 were available in the laboratory. For some experiments, normal serum from one donor was depleted of C1q, factor D and properdin as described by Sjöholm *et al.* [7]. Where indicated, NHS was inactivated by heat treatment (56°C, 30 min). Zymosan treatment of serum was performed as described previously [8]. Sera with high granulocyte clustering

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activity from five patients with definite SLE [4] were also investigated. The patients fulfilled four or more of the ACR classification criteria for SLE [9]. Disease activity was estimated by SLEDAI scoring [10].

Complement components

C1q was purified according to Tenner *et al.* [11]. To remove possible aggregates, C1q preparations were consistently subjected to gel filtration on a TSK-4000 column (LKB, Bromma, Sweden) in an FPLC system (Pharmacia, Uppsala, Sweden). After gel filtration, C1q preparations were used within 2 days. Factor D and properdin were purified according to published methods [12,13]. Concentrations of the purified complement proteins were determined by electroimmunoassay, assuming a reference normal serum pool to contain C1q at 70 mg/l, properdin at 25 mg/l and factor D at 1 mg/l [6]. C1q inhibitor, a chondroitin-4 sulphate proteoglycan, was purified from 1.5 ml of outdated ACD plasma essentially as described by Silvestri *et al.* [14]. In brief, C1q inhibitor in complex with C1q was isolated from plasma by repeated euglobulin precipitation, and was then separated from C1q by chromatography on a 1.6 × 26 cm column of concanavalin A (Con A) Sepharose (Pharmacia) to which C1q was quantitatively bound [14]. The C1q inhibitor recovered in the effluent was dialysed overnight against 20 volumes of Tris 0.02 mol/l, NaCl 0.2 mol/l, pH 7.5, and was applied to a column of DEAE-Sepharose (Pharmacia) equilibrated in the same buffer. The inhibitor was eluted with a sharp NaCl gradient to 1 mol/l. C1q inhibitory activity was detected by haemolytic assay [15] and by single diffusion precipitation of purified C1q in agarose [14]. The sulphated glycosaminoglycan content of the C1q inhibitor preparation (0.92 mg/ml, total yield 2.8 mg) was kindly determined by Dr S. Björnsson (Lund, Sweden) according to a recently described procedure [16].

Other reagents

Commercial polyclonal IgG (Kabi, Stockholm, Sweden) at 10 g/l was aggregated by heat (63°C, 30 min). After treatment, the heat-aggregated IgG (HAIG) was centrifuged at 2000 g for 30 min, the supernatant being used for addition to serum. Recombinant human C5a was purchased from Sigma Chemical Co. (St Louis, MO).

Assay for measurement of neutrophil clustering

The assay for *in vitro* measurement of neutrophil clustering has been described in detail [4]. In short, neutrophils (0.01 ml, 10⁷ neutrophils/ml) from healthy donors were added to 0.04 ml of test serum, incubated at room temperature for 1 min, whereafter single and clustered cells were counted within a standardized area. In each of duplicate preparations 200 single cells were counted. Clustering activity was expressed as the number of clustered cells in per cent of the total number of counted cells.

To generate neutrophil clustering activity, the sera were incubated for 60 min at 37°C with HAIG at a final concentration of 1 g/l. In reconstitution experiments, purified complement proteins were first added to serum depleted of C1q, factor D and properdin, or to heat-treated NHS at physiological or specified concentrations. Sera from patients with SLE were used without pretreatment with HAIG. C1q inhibitor at specified concentrations was added to HAIG-treated sera or to SLE sera, followed by incubation at 37°C for 10 min before the addition of cells.

Table 1. Neutrophil clustering (%) in normal and complement-deficient sera with and without treatment with heat-aggregated IgG (HAIG)

| | Untreated serum | Serum treated with HAIG |
|-------------------------------|-----------------|-------------------------|
| Normal serum | 5 | 26 |
| C2-deficient serum | | |
| I | 7 | 36 |
| II | 10 | 35 |
| III | 12 | 54 |
| C4-deficient serum | ND | 51 |
| Hereditary angio-oedema serum | 12 | 26 |
| Properdin-deficient serum | ND | 32 |
| C3-deficient serum* | 10 | 32 |
| C5-deficient serum | 3 | 31 |
| C7-deficient serum | | |
| I | 13 | 42 |
| II | 5 | 38 |
| III | 9 | 28 |
| C8-deficient serum | 25 | 43 |

* Acquired C3 deficiency in a patient with C3 nephritic factor.

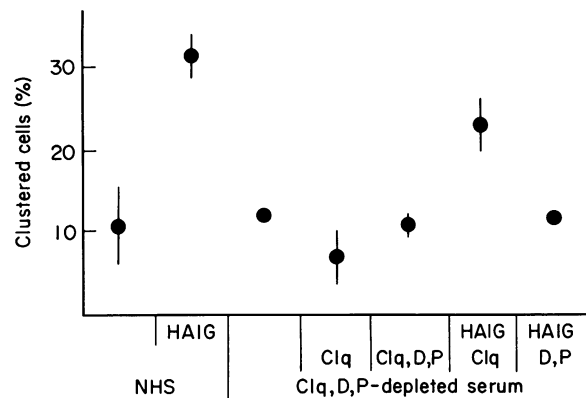


Fig. 1. Induction of neutrophil clustering with heat-aggregated IgG (HAIG) at 1 g/l in pooled normal human serum (NHS) and in normal serum depleted of C1q, factor D and properdin (C1q,D,P-depleted serum). Purified complement proteins were added to the depleted serum at physiological concentrations as indicated below the figure. Results of two separate experiments are shown. Neutrophil clustering activity is given as mean value and range.

RESULTS

Neutrophil clustering in complement-deficient and complement-depleted sera

In NHS the proportion of clustered cells varied between 5% and 25% of the total number of counted cells. In NHS treated with HAIG the number of clustered cells was consistently increased, and varied between 21% and 48%.

Addition of HAIG clearly promoted neutrophil clustering in sera from persons with homozygous deficiencies of C2, C5, C7 or C8 (Table 1). Similar findings were made in serum from a patient with hereditary angio-oedema and in serum from a

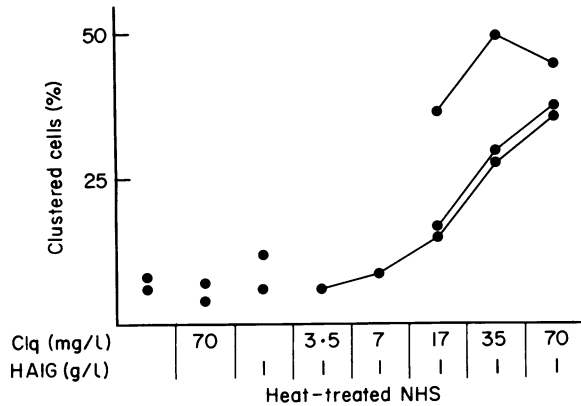


Fig. 2. C1q-dependent induction of neutrophil clustering activity in heat-treated (56°C, 30 min) pooled normal human serum (NHS). C1q and heat-aggregated IgG (HAIG) were added at the concentrations indicated below the figure. Results of three separate experiments are shown.

patient with acquired C3 deficiency. In addition, high neutrophil clustering activity was found in C4-deficient serum and properdin-deficient serum treated with HAIG.

Further experiments were carried out with normal serum artificially depleted of C1q, factor D and properdin (Fig. 1). The depleted serum did not support enhanced neutrophil clustering activity in the presence of HAIG. However, reconstitution with C1q, but not with factor D and properdin before addition of HAIG, produced a marked response with regard to neutrophil-clustering activity.

The results of experiments with complement-deficient and complement-depleted sera suggested that neutrophil clustering induced with HAIG required C1q or the C1 complex, and that further recruitment of the complement system was not necessary. In particular, the findings with C5-deficient serum argued against an important role of C5-derived peptides (C5a and C5a_{des arg}) in the assay. Zymosan-treated serum, known to be a

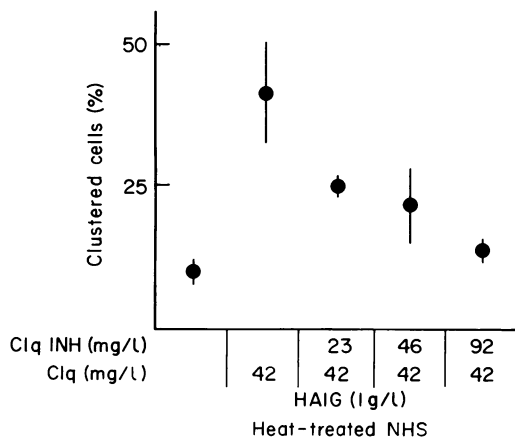


Fig. 3. Influence of C1q inhibitor (C1q INH) on neutrophil clustering in heat-treated (56°C, 30 min) pooled normal human serum (NHS) after incubation of the serum with C1q and heat-aggregated IgG (HAIG). Final concentrations of C1q INH, C1q and HAIG are given below the figure. Results of two separate experiments are shown. Neutrophil clustering activity is given as mean value and range.

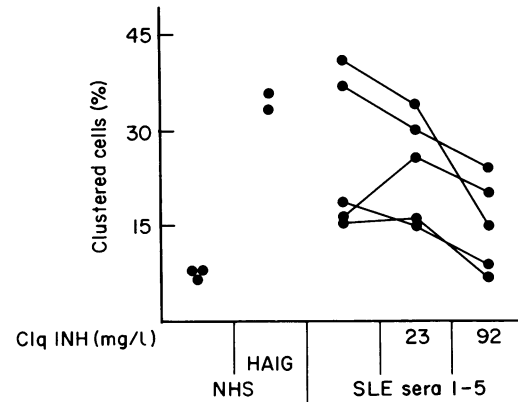


Fig. 4. Influence of C1q inhibitor (C1q INH) on neutrophil clustering in sera from five patients with active systemic lupus erythematosus (SLE) (SLE sera 1-5). C1q INH was added at the concentrations indicated below the figure. Results of included control experiments with pooled normal human serum (NHS) and heat-aggregated IgG (HAIG, 1 g/l) are shown to the left.

reliable source of C5-derived peptides [8], produced slightly less neutrophil clustering than normal serum activated with HAIG. Addition of recombinant C5a at a final concentration of 1-10 mg/l to normal serum resulted in very little neutrophil-clustering activity.

Neutrophil-clustering activity in heat-treated serum: effect of purified C1q

Background clustering in heat-treated NHS was similar to or less than that observed in NHS. Addition of HAIG to heat-treated NHS did not increase neutrophil clustering, nor did addition of C1q alone. When heat-treated NHS was reconstituted with C1q before addition of HAIG, C1q produced a dose-dependent increase of neutrophil-clustering activity in the serum (Fig. 2).

Inhibition experiments

In heat-treated NHS with C1q and aggregated IgG added, purified C1q inhibitor reduced neutrophil clustering in a dose-dependent manner (Fig. 3). Inhibitory doses were in the range 23-92 mg/l. The normal concentration of C1q inhibitor in serum is not known, but could be in the order of 20 mg/l assuming a 10% yield during the preparation procedure. This implies that the inhibitory doses were well beyond the normal concentration of C1q inhibitor.

Serum from five patients with active disease (SLEDAI scores 10-24) caused clustering from 17 to 43% of the total cell count. The morphology of cell clusters induced by SLE serum and by HAIG-activated serum was similar. In four of the five SLE sera, C1q inhibitor produced dose-dependent reduction of neutrophil clustering (Fig. 4).

DISCUSSION

Intravascular aggregation of neutrophil granulocytes is considered an important event in the inflammatory response, by promoting neutrophil contact for endothelial adhesion before transendothelial migration and accumulation of neutrophils at

extravascular sites [17]. Neutrophil expression of complement receptor type 3 (CR3), a member of the CD18 complex [18], appears to be critical for neutrophil aggregation and for adhesion to endothelium [19,20]. CR3-dependent neutrophil aggregation involves a cell surface counter-structure [21] that may be distinct from ICAM-1 and ICAM-2 [18]. Aggregation of neutrophils also requires cell activation as triggered by chemoattractants such as C5-derived peptides or other agents [19,21]. *In vivo*, intravascular chemoattractants and systemic complement activation induce neutropenia and neutrophil sequestration due to adherence of neutrophils to endothelium in the microvasculature [22,23].

Complement activation has been implied in neutrophil-mediated pathogenetic events associated with haemodialysis [24], the adult respiratory distress syndrome [25], and the 'post-perfusion syndrome' [26]. These studies have been focused on effects of C5-derived peptides. In the adult respiratory distress syndrome, neutrophil aggregating activity in plasma as ascribed to C5a was correlated with development of symptoms [25].

Sera from patients with SLE or Felty's syndrome have been reported to produce aggregation adherence to endothelium and superoxide generation of neutrophils *in vitro* [2]. Attempts to identify the serum factor(s) involved have been inconclusive [2-4].

In the present investigation, neutrophil aggregating activity in serum was assessed with a simple neutrophil clustering assay [4], whereas turbidimetry measurements have been employed in other studies [2,3,24,25]. Thus, findings might not be strictly comparable. Changes in light scattering are influenced by cell polarization and modification of subcellular structures [27]. Furthermore, cytochalasin B, an agent that enhances neutrophil aggregation responses [28] by interaction with actin [29], was used in the study of Abramson *et al.* [3]. In our assay, preformed IgG aggregates readily generated neutrophil clustering activity in normal serum, while no such effect was obtained in heat-treated serum. This argued against a role of cell activation mediated through Fc γ RII and Fc γ RIII [30], and clearly suggested a requirement for complement. It is questionable if the clustering effect of Zymosan-treated serum was due to C5-derived peptides, since recombinant C5a showed very limited activity.

The principal finding was that recruitment of the complement system beyond C1 was not necessary for efficient induction of neutrophil clustering. However, preformed IgG aggregates in combination with C1q produced marked activity, even in heat-treated serum. We conclude that the reaction was triggered by C1q-containing IgG complexes through interaction with neutrophil surface receptors for C1q [31].

C1q receptors recognize the collagenous part of C1q [32,33], and collagen structures of C-type lectins that resemble C1q [34]. The receptor binding site of C1q is masked by C1r and C1s in the C1q(C1r-C1s)₂ complex [32], but disassembly of the molecule by C1 inhibitor in conjunction with activation [35] exposes the ligand structure of fixed C1q. Like C1q receptors, the C1q inhibitor in serum [14] binds to the collagenous part of C1q [15]. C1q inhibitor virtually abolished neutrophil clustering elicited with C1q-containing complexes and the inherent neutrophil clustering activity of some SLE sera, indicating that the collagenous part of C1q was a significant ligand in the reaction.

To our knowledge, neutrophil clustering by a C1q-dependent mechanism has not been previously described. The pre-

vious finding of inverse correlation between neutrophil clustering activity and the number of circulating neutrophils in SLE [4] suggests that the activity is not a biologically inconsequential phenomenon. C1q can activate neutrophils as expressed by an oxidative response [36] with recruitment of partly unique signal transduction pathways [37]. It remains to be established if C1q-triggered neutrophil clustering resembles neutrophil aggregation induced by other stimuli in the requirement for continuous cell activation and CD18 leucocyte adhesion molecules [21], or if other mechanisms are involved.

The ligand requirements for induction of neutrophil clustering activity in serum strongly suggested involvement of C1q receptors, even if the precise mechanisms remain unclear. As proposed for platelets [38] and endothelial cells [39], C1q receptors might contribute to localization of C1q-containing immune complexes to neutrophil surfaces. Assuming that this leads to neutrophil clustering and enhanced margination of the cells, the findings relate to previous observations in disease conditions such as SLE [1-4]. Interestingly, C1 dissociation with possible exposure of immune complex-bound C1q is pronounced in active SLE [40,41]. We propose that C1q bound to immune complexes could be an important ligand for interaction with neutrophils in SLE, and perhaps also in other immune complex diseases.

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