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Research Article

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Antibacterial Activity of Cationic Proteins from Human Granulocytes

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ABSTRACT Human granulocytes contain several cationic proteins with a molecular weight of approximately 25,000, almost identical amino acid composition, and complete immunologic identity. These proteins possess a chymotrypsin-like protease activity at a neutral pH. The antibacterial activity of the cationic proteins has been studied. Bactericidal activities are found against both Gram-positive (*Streptococcus faecalis* and *Staphylococcus aureus*) and Gram-negative (*Escherichia coli* and *Pseudomonas aeruginosa*) organisms. Gram-positive bacteria are, however, the most sensitive. The pH-optimum is near neutrality, and the microbicidal activity shows an inverse relationship to the ionic strength, indicating an ionic interaction between the cationic proteins and the bacterial surface. The microbicidal effect of the cationic proteins is generally independent of the chymotrypsin-like activity of the same proteins since the activity against several bacterial species is heat stable while the chymotrypsin-like activity is heat labile. The surface properties of *S. aureus* that are determined by protein A do not seem to influence the susceptibility to cationic proteins. The properties of the Gram-negative envelope of *E. coli* that determine the susceptibility to the lytic action of serum do not influence the sensitivity to the action of cationic proteins. The present study shows that cationic proteins of human granulocytes represent one potential microbicidal mechanism that is independent of hydrogen peroxide and myeloperoxidase.

INTRODUCTION

Microbial killing in granulocytes follows ingestion of microorganisms into the cell. Killing takes place in the phagocytic vacuole after release of the microbicidal fac-

tors of the cytoplasmic granules. The increased production of hydrogen peroxide, linked to an increase in oxygen consumption and hexose monophosphate shunt activity, constitutes, together with myeloperoxidase, one microbicidal system (1). Information is, however, still scarce concerning the actual mechanism of bacterial killing in granulocytes (2). Prolonged intracellular survival of some microorganisms, e.g. *Staphylococcus aureus*, results under conditions of defective hydrogen peroxide production as in chronic granulomatous disease of childhood (CGD)¹ (3). Other organisms, e.g. Streptococci (4), Lactobacilli (5), and Pneumococci (6), are readily killed even in CGD granulocytes. Furthermore, individuals deficient in myeloperoxidase usually do not have excessive infections, and their granulocytes kill ingested bacteria, although at a reduced rate (7, 8). Therefore, alternative microbicidal mechanisms might exist that do not involve the myeloperoxidase-hydrogen peroxide system.

A bactericidal agent of rabbit granulocytes, phagocytin, was described by Hirsch (9) and shown by Zeya and Spitznagel (10) to consist of highly cationic proteins with different degrees of antimicrobial capacity against the bacterial species studied. Previous work from our laboratory has demonstrated cationic proteins to be present also in human granulocytes (11). One group of cationic proteins (components 1-4) shows molecular weights of 25,500-28,500, almost identical amino acid composition, and immunochemical identity. The other group of proteins (components 5-7) demonstrate molecular weights of 21,000-29,000 and immunological identity. Recently, Lehrer et al. (12) have identified a nonperoxidative fungicidal mechanism in human granulocytes. Fungicidal proteins with esterase activi-

¹ *Abbreviations used in this paper:* CGD, chronic granulomatous disease of childhood; HEPES, *N*-2-hydroxyethylpiperazine-*N'*-2-ethane sulfonic acid; KRP, Krebs-Ringer phosphate buffer.

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ties separated by micro-preparative electrophoresis seem to have properties identical with some of our purified cationic proteins. Similarly, Rindler and Braunsteiner (13) have also demonstrated the presence of cationic esterases in human granulocytes. It is the purpose of the present communication to report studies on the antimicrobial capacity of the cationic proteins obtained from human granulocytes.

METHODS

Microorganisms. *S. aureus* 502A, *Streptococcus faecalis* ATCC no. 8043, and clinical isolates from urine of *Escherichia coli*, *Pseudomonas aeruginosa*, and *S. faecalis* are used in the bactericidal assays. *S. aureus* 502A is maintained on CCY Medium (14) solidified with 1.5% Bacto agar, *S. faecalis* ATCC no. 8043 was cultured on Bacto micro-assay culture agar (Difco Laboratories, Detroit, Mich.), and the other organisms are maintained on blood agar. Stock cultures are passed every 2nd to 4th wk.

Cationic proteins. The cationic proteins are isolated from human leukocyte granule extracts as previously described (11). These proteins are numbered 1-7, and component 1 is the most cationic component with the highest electrophoretic mobility towards the cathode. In the present report only the group of the most cationic protein components 1-4 is studied. Components 2 and 3 are the major ones while components 1 and 4 usually are present only in small amounts. For the present purpose these components have not been completely purified. Instead two fractions are used; one fraction containing a mixture of components 1 and 2 (A) and the other containing a mixture of components 3 and 4 (B). The electrophoretic pattern on agarose gel of the fractions used is shown in Fig. 1.

Bactericidal assay. Bacteria are grown in Bacto antibiotic medium 3 (Difco Laboratories) at 37°C overnight, except for *S. faecalis* ATCC no. 8043 which is grown in Bacto folic assay medium fortified with 1 ng per ml pteroylglutamic acid. To obtain bacteria in the logarithmic growth phase, 0.1 ml of the overnight broth culture is transferred to 5 ml of fresh antibiotic medium 3 and incubated at 37°C. The OD of the broth cultures is followed at 628 nm, and the organisms are harvested during logarithmic growth, washed twice in 5 ml of calcium-free Krebs-Ringer phosphate buffer (KRP), pH 7.4, containing 0.1% gelatin, and resuspended in KRP to an OD of 0.23 at 628 nm, corresponding to $2.0\text{--}3.0 \times 10^8$ organisms per ml. Bacteria are then suspended in a modified Hanks' balanced salt solution to a concentration of about 2.5×10^8 organisms per ml, and a water solution of cationic protein is added. The final concentration of solutes in the incubation mixture is 67 mM NaCl, 3.5 mM NaHCO₃, 4.2 mM KCl, 1.0 mM CaCl₂, 0.32 mM MgSO₄, 0.39 mM MgCl₂, 0.27 mM Na₂HPO₄, 0.35 mM KH₂PO₄, 4.4 mM glucose, 16 mM N-2-hydroxyethylpiperazine-N'-2-ethane sulfonic acid (HEPES), pH 7.4, and gelatin 0.1%. The mixtures are incubated in 12 × 75-mm plastic tubes (Falcon Plastics, Division of BioQuest, Oxnard, Calif.) during rotation end-over-end at the rate of 20 rpm at 37°C. Controls are run without cationic protein. For determination of the number of surviving bacteria by the pour plate method, aliquots of 20-200 μl are removed at 0, 30, and 60 min and diluted in 8 ml KRP containing 0.1% gelatin. Samples are mixed with Bacto antibiotic medium 3 containing 1.5% Bacto agar, and the number of colony-forming units is determined after incubation overnight. The number of sur-

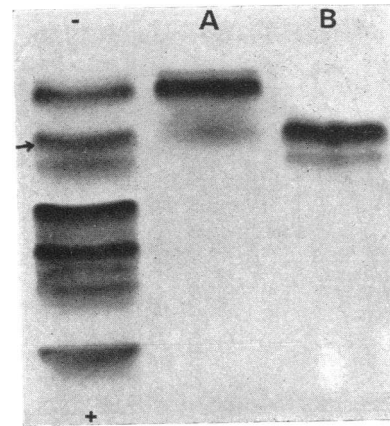


FIGURE 1 Agarose gel electrophoresis of the cationic protein fractions used in the present study (A and B). The cationic protein-enriched fraction obtained by gel chromatography (11) is shown for reference. The arrow indicates the electrophoretic mobility of lysozyme.

ving bacteria is also determined by OD readings at 628 nm. For this purpose, 4 ml of broth is added to the 60-min incubations, and the OD followed until a stationary growth phase is achieved. The time for attainment of an OD of 0.40 is determined. The time difference between cationic protein incubation and control incubation is divided by the mean generation time (MGT) obtained from the OD readings, and the number obtained is designed "MGT." The exponential growth rates are identical in control and cationic protein incubations. The number of surviving bacteria in the 60-min incubation mixture with cationic protein is calculated from the following formula: Surviving bacteria (cationic protein) = surviving bacteria (control) / $2^{(\text{MGT})}$. The number of surviving bacteria in the control incubation used in the formula is determined by colony count using the pour plate method.

RESULTS

The cationic protein fractions A and B both exhibit antibacterial activities against several species tested. Fig. 2 shows an experiment with *S. aureus* and *E. coli*. After incubation with cationic protein A for 60 min in the standard incubation medium, broth is added to the mixture, which is incubated at 37°C. Both colony counts and OD readings are used for determination of the bacteria concentration after different time periods.

The number of colony-forming units is significantly decreased after 60 min incubation with cationic proteins. The 60-min control incubations contain 1.2×10^8 (*S. aureus*) and 5.0×10^8 (*E. coli*) colony-forming units per ml whereas the figures for the cationic protein incubations are 9.2×10^4 (*S. aureus*) and 9.8×10^4 (*E. coli*). When calculated from OD readings according to the formula given in methods the number of surviving bacteria in the 60-min incubation with cationic protein is 5×10^4 (*S. aureus*) and 1.5×10^5 (*E. coli*) organisms per ml. The increase in OD, followed during the latter part of the

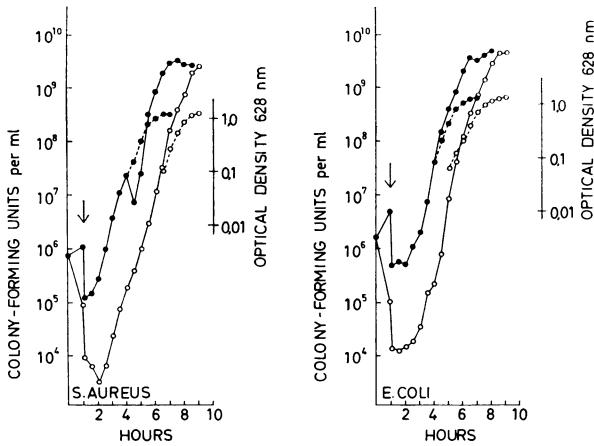


FIGURE 2 Antimicrobial effect of cationic protein A on *S. aureus* and *E. coli*. Bacteria are incubated with 54 $\mu\text{g/ml}$ (*S. aureus*) and 108 $\mu\text{g/ml}$ (*E. coli*) for 60 min. After that (indicated by arrow) 4 ml of broth is added both to control incubation (●—●) and to cationic protein incubation (○—○) and the mixtures are incubated at 37°C. The growth of the organisms is followed both by colony counting (—) and by OD readings (---).

exponential growth phase after addition of broth, is somewhat lower than the increase in colony-forming units, leading to an overestimation of the number of surviving bacteria when calculated from OD readings. On the other hand, the number of surviving bacteria after 60 min is underestimated if killing continues after the addition of broth as is the case for *S. aureus* (Fig. 2). Thus, identical results cannot be expected when the number of surviving bacteria is determined with colony counting and OD readings. In the experiments to be described below, aliquots are taken for colony counts after incubation of organisms with cationic proteins for 30 and 60 min. After addition of broth to the incubation mixtures the increase in OD is also followed and compared with the increase of OD of a control incubation with bacteria not subjected to the action of cationic protein as described in methods. In all instances the information obtained by colony counts and OD readings agreed fairly well. Below, only the results from the colony count method are documented.

In Fig. 3 the effect of pH on the antibacterial activity of cationic protein A is demonstrated. The pH of the standard incubation medium is varied from 6.0 to 7.8 with HEPES buffer. At pH 6-7 only slight killing is found. The activity is highly increased with a pH above 7. Identical results are obtained with *S. aureus* and *E. coli*. For the attainment of pH levels below 6.0 and above 7.8 2-(*N*-morpholino)ethane sulfonic acid buffer and Tris buffer, respectively, is used. At pH 5.0-6.0 the microbicidal activity is almost identical with that of pH 6.0-7.0. Increasing the pH above pH 7.8 gives a further

increase of the microbicidal activity with pH (results not shown).

The effect of the ionic strength on the antibacterial activity of cationic protein A is studied by the addition of increasing amounts of sodium chloride (Fig. 3). The microbicidal effect shows an inverse relationship to the ionic strength. At an ionic strength corresponding to 0.2 M sodium chloride or higher the microbicidal activity is abolished. In the standard incubation procedure for assay of antibacterial capacity an ionic strength corresponding to 0.09 M sodium chloride is chosen. This concentration is a little below the ionic strength of serum that corresponds to 0.12 M sodium chloride as measured by conductivity. It is also demonstrated that the antimicrobial activity is abolished when, after incubation for 30 min in the standard incubation medium, the concentration of sodium chloride is increased to 0.2 M (Fig. 4); the antibacterial activity is promptly inhibited by increasing the ionic strength.

The microbicidal activity of the cationic proteins against several microbial species are compared in Figs. 5 and 6. The effects of both cationic protein A (components 1 and 2) and B (components 3 and 4) are examined. It is found that the cationic proteins exhibit

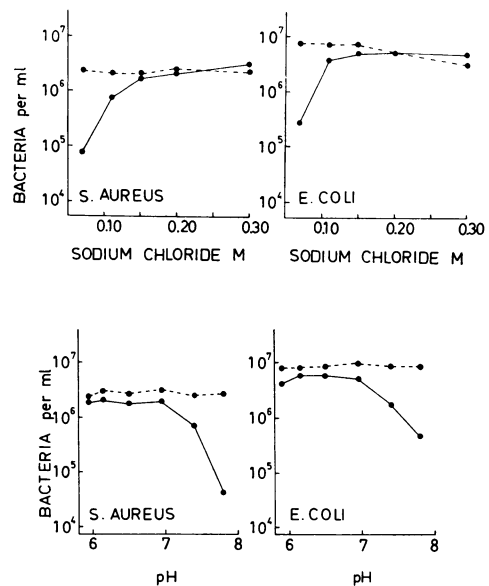


FIGURE 3 Effect of ionic strength and pH on the microbicidal activity of cationic protein A. For studies on ionic strength, bacteria are incubated with 80 $\mu\text{g/ml}$ (*S. aureus*) and 108 $\mu\text{g/ml}$ (*E. coli*) in the standard incubation medium where the NaCl concentration is varied (●—●). Controls are run without cationic protein (●---●). For studies of pH effect, bacteria are incubated with 50 $\mu\text{g/ml}$ (*S. aureus*) and 80 $\mu\text{g/ml}$ (*E. coli*) in the standard incubation medium where the pH is varied between 6 and 7.8 by the addition of HEPES buffer (●—●).

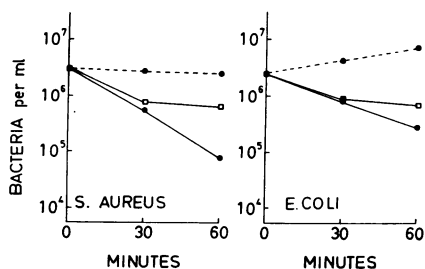


FIGURE 4 Effect of ionic strength on the microbicidal activity of cationic protein A. Bacteria are incubated with 80 $\mu\text{g/ml}$ (*S. aureus*) and 108 $\mu\text{g/ml}$ (*E. coli*) in the standard incubation medium (●—●). After 30 min of incubation NaCl is added to a final concentration of 0.2 M (□—□). Controls are also run without cationic protein (●---●).

an antibacterial effect against both Gram-positive (*S. faecalis* and *S. aureus*) and Gram-negative species (*E. coli* and *P. aeruginosa*). It is apparent that the Gram-positive bacteria studied are more sensitive to the action of the cationic proteins than are Gram-negative bacteria.

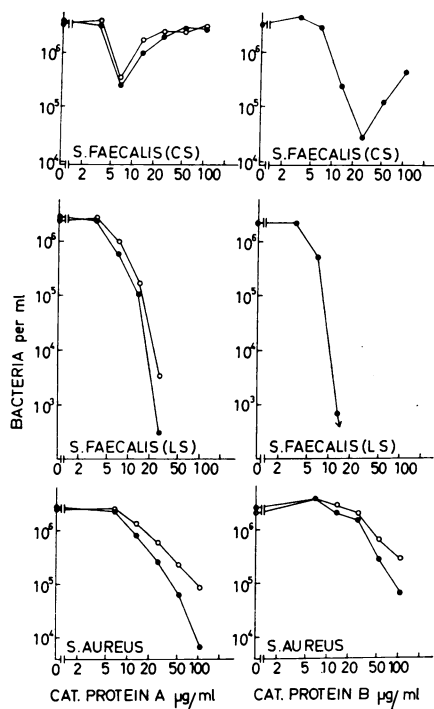


FIGURE 5 Antimicrobial effect of different concentrations of cationic protein A and B on *S. aureus*, *S. faecalis* ATCC 8043 (laboratory strain [LS]), and a clinical strain (CS) of *S. faecalis*. Surviving bacteria are determined by colony counting at 30 min (○—○) and 60 min (●—●) of incubation. The results shown to the left in the figure are from incubations with cationic protein A and results shown to the right are from incubations with cationic protein B.

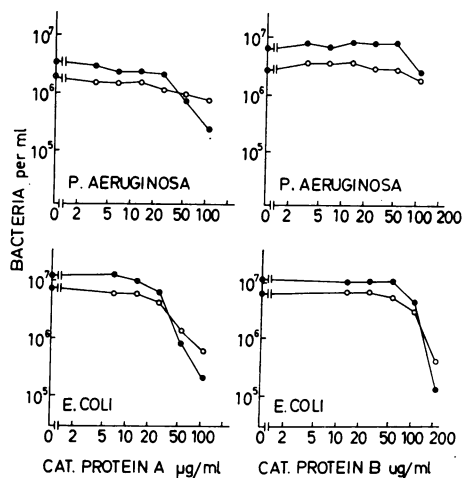


FIGURE 6 Antimicrobial effect of different concentrations of cationic protein A and B on *E. coli* and *P. aeruginosa*. Surviving bacteria are determined by colony counting at 30 min (○—○) and 60 min (●—●) of incubation. The results shown to the left in the figure are from incubations with cationic protein A and results shown to the right are from incubations with cationic protein B.

The two strains of *S. faecalis* react differently. At a low concentration of cationic protein (6.8 and 13.6 μg per ml) extensive reduction of the number of colony-forming units of both strains is achieved. By increasing the protein concentration the antibacterial effect against the laboratory strain is increased whereas the effect against the clinical strain is decreased; at a concentration of 54.4 μg per ml of cationic protein no antibacterial activity is found. The antibacterial effect against all bacteria tested except the clinical strain of *S. faecalis* is heat stable since heating of the cationic proteins at 90°C for 10 min does not abolish the effect. The effect against the clinical strain of *S. faecalis* is, however, heat labile.

The microbicidal effect of cationic proteins on three different strains of *S. aureus* (502 A, Cowan I, and Wood 46) is compared in Fig. 7. It is apparent that the strain Cowan I exhibits a higher sensitivity than the other two strains.

The antibacterial effects of cationic proteins against two strains of *E. coli* are compared in Fig. 8. One of the strains is sensitive to the bactericidal effect of serum since 5% serum causes an extensive reduction of colony-forming units while the other strain is resistant to the effect of serum. The antibacterial activity of cationic proteins is, however, similar for both strains.

Experiments were conducted to compare the microbicidal effect of cationic proteins with that of histones. Lysine-rich histones at a concentration of 200 $\mu\text{g/ml}$ cause a reduction in colony-forming units of *S. aureus*

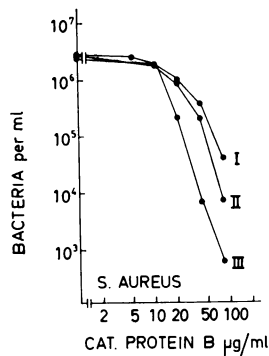


FIGURE 7 Antimicrobial effect of cationic protein B against three different strains of *S. aureus*. Surviving bacteria are determined by colony counting at 60 min (●—●). The strains are designated "I" (502 A), "II" (Wood 46), and "III" (Cowan I).

502 A from 2.3×10^6 to 1.6×10^5 while cationic protein A at a concentration of 107 $\mu\text{g/ml}$ gives a reduction from 2.3×10^6 to 6.5×10^8 U/ml. The microbicidal effect of arginine-rich histones is even lower than that of lysine-rich histones.

DISCUSSION

The present study has demonstrated a potential microbicidal system of the human granulocyte to consist of a group of cationic proteins previously isolated (11). So far the only significant difference found between the four protein components included is in electrophoretic mobility (11), indicating differences in ionic charge. Since the cationic proteins have a chymotrypsin-like esterase activity (15), it might be possible that the microbicidal effect is due to the enzyme properties of the proteins. It is found, however, that the antimicrobial

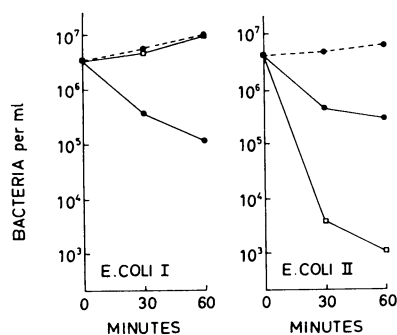


FIGURE 8 Antimicrobial effect of cationic protein A on two clinical isolates from urine of *E. coli* (designated "I" and "II"). Bacteria are incubated with 140 $\mu\text{g/ml}$ cationic protein A (●—●). A control is run without cationic protein (●---●). For comparison the effect of 5% human serum is also shown (□—□).

activity against several bacterial species is not inhibited by heating, which inhibits the esterase activity (15). Obviously the microbicidal effect demonstrated in these instances is independent of the esterolytic activity. Exception to the heat stability of the antimicrobial activity is shown, however, since the killing of one strain of *S. faecalis* is due to heat-labile properties of the cationic proteins. The dependence on the ionic strength with inhibition of the antimicrobial activity at a high salt concentration points to the possibility of an ionic interaction between the proteins and the bacterial surface relevant to the mechanism of killing. Thus, the sensitivity of the microorganisms may be determined by properties of the microbial envelopes. Generally, Gram-positive bacteria are also found more sensitive than Gram-negative organisms. The three strains of *S. aureus* employed show different susceptibility to the antibacterial action of the cationic proteins. Two of the strains, 502 A and Cowan I, possess staphylococcal protein A, whereas Wood 46 lacks this surface component. The surface properties that are determined by staphylococcal protein A probably have little influence on the susceptibility to cationic proteins as 502 A was more, and Cowan I less, sensitive than Wood 46. The properties of the Gram-negative envelope of *E. coli* that determine the susceptibility of the organism to the lytic action of human serum are obviously not of importance for the reaction with cationic protein since the two strains of *E. coli* compared show approximately the same sensitivity to cationic proteins but differ considerably in susceptibility to human serum.

The cationic proteins of rabbit granulocytes have been implicated in several biological activities. In addition to cytotoxic actions against bacteria (10), permeability-increasing capabilities have been shown (16). Zeya and Spitznagel (17) suggest that the antibacterial activity is mediated by damage to the cell membrane and/or an inhibition of aerobic respiration. It is well known that histones have microbicidal effects (18) as confirmed in the present study. Both histones and cationic proteins from rabbit granulocytes interfere with mitochondrial respiration (19). Cationic proteins of human granulocytes might have similar properties. Histones exhibit, however, a lower killing capacity than the granule proteins.

The microbicidal activity of the cationic proteins is independent of previously described systems responsible for bactericidal activity in human granulocytes, e.g., the hydrogen peroxide myeloperoxidase complex (1) or superoxide (20). The latter mechanisms are operative only when oxygen is available to the cell. As demonstrated by Mandell (21) anaerobic human granulocytes are able to kill several microorganisms like *Staphylococcus epidermidis*, *Enterococcus*, and *P. aeruginosa*

while organisms like *S. aureus* and *E. coli* are not killed normally by anaerobic phagocytes. Granulocytes from patients with CGD have an abnormal metabolic response to phagocytosis including a markedly reduced oxygen consumption and hydrogen peroxide generation (3). Nevertheless, such cells manifest an intact ability to kill some organisms, e.g., Streptococci (4), Lactobacilli (5), Pneumococci (6), and certain species of *Candida* (22). The explanation often given is that the bacteria mentioned do succumb after ingestion due to their own elaboration of hydrogen peroxide (5) substituting for defective production of this substance by the granulocyte. Recent work by Shoet et al. (23) demonstrated that CGD cells killed peroxide-positive wild pneumococci much more effectively than a peroxide-deficient mutant, suggesting that peroxidative mechanisms are required for intraphagocytic killing of these species. At the present time, it is not known whether the cationic proteins are involved in killing of microorganisms normally killed in CGD granulocytes. The second mechanism of candidacidal activity of human neutrophils described by Lehrer (12, 22) could be identical with the microbicidal system of the present work. The availability of this antimicrobial mechanism was suggested to provide an explanation for the ability of myeloperoxidase-deficient neutrophils and those from patients with CGD to kill certain species of *Candida* (22).

A pH optimum near neutrality of the antimicrobial activity raises questions as to the existing pH within the phagocytic vacuole of the granulocyte. Using indicator dyes, Metchnikoff (24) and Rous (25) conclude that the intracellular pH of various mammalian phagocytizing cells might be as low as 3.0. Recently, Jensen and Bainton (26), utilizing similar techniques, demonstrated temporal changes in the phagocytic vacuole of rat peritoneal exudate cells during phagocytosis; within 3 min pH dropped to 6.5, and within 7-15 min it dropped to 4.0. There is only one analogous study available on human granulocytes (27); a maximum pH depression in the phagocytic vacuole to 6.0-6.6 is reported. The small decrease thus found of intravacuolar pH in human cells as compared to other mammalian leukocytes indicates important species differences. It should be emphasized that several major granule components of human cells consist of proteases, e.g. elastase (28), collagenase (29), and the chymotrypsin-like activity of the cationic proteins (15), operating maximally at neutral pH. Such proteases are not found in rabbit or guinea pig granulocytes, which contain cathepsins with an acid pH-optimum. The operation of the cationic proteins of human granulocytes at a neutral pH would not be inconsistent with a proposed role for these substances in the interaction with ingested microorganisms. The anti-

bacterial activity might be exercised early during phagocytosis before a significant drop in pH has taken place.

Biochemical and morphological studies of granulocytes from several species have revealed two types of cytoplasmic granules, namely, peroxidase-positive (azurophil or primary granules) and peroxidase-negative (specific or secondary granules) (30-33). The cationic proteins of rabbit polymorphonuclear leukocytes are associated with the peroxidase-containing granules (32). Electrophoretic analysis carried out by Dewald et al. (34) on granule fractions obtained by isopycnic centrifugation showed that the most strongly cationic proteins of both human and rabbit granulocytes were confined to the azurophil granules. Recently we have shown the chymotrypsin-like cationic proteins of this study to be localized exclusively in the azurophil granules.⁹ Studies by Bainton (35) on rabbit granulocytes indicate that the two types of granules discharge their content in a sequential manner, specific granules fusing with the phagocytic vacuole before azurophil granules. The optimal conditions for the action of the cationic proteins in the phagocytic vacuole might vary with time after ingestion. Furthermore, recent studies by Leffell and Spitznagel (36) emphasize that relatively more of the specific granule proteins were released into the medium during phagocytosis, whereas more of the azurophil proteins were associated with the phagosomes.

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