

Host and Bacterial Factors Involved in the Innate Ability of Mouse Macrophages To Eliminate Internalized Unopsonized *Escherichia coli*

TERRI S. HAMRICK, EDWARD A. HAVELL, JOHN R. HORTON, AND PAUL E. ORNDORFF*

Department of Microbiology, Pathology, and Parasitology, College of Veterinary Medicine,
North Carolina State University, Raleigh, North Carolina 27606

Received 25 May 1999/Returned for modification 16 July 1999/Accepted 12 October 1999

In an effort to better understand genetic and cellular factors that influence innate immunity, we examined host and bacterial factors involved in the nonopsonic phagocytosis and killing of *Escherichia coli* K-12 by mouse macrophages. Unelicited (resident) peritoneal macrophages from five different mouse strains, BALB/c, C57BL/6, CD-1, C3H/HeJ, and C3H/HeN, were employed. Additional macrophage populations were obtained from CD-1 mice (bone marrow-derived macrophages). Also, for BALB/c and C57BL/6 mice, peritoneal macrophages elicited with either thioglycolate or proteose peptone, bone marrow-derived macrophages, and macrophage-like cell lines derived from the two strains were employed. Two *E. coli* K-12 strains that differed specifically in their abilities to produce type 1 pili containing the adhesive protein FimH were examined. The parameters used to assess macrophage bacteriocidal activity were (i) the killing of internalized (gentamicin-protected) *E. coli* during the approximately 4-h assay and (ii) the initial rate at which internalized *E. coli* were eliminated. Data on these parameters allowed the following conclusions: (i) unelicited or proteose peptone-elicited peritoneal macrophages were significantly better at eliminating internalized bacteria than thioglycolate-elicited peritoneal macrophages, bone marrow-derived macrophages, or macrophage cell lines; (ii) the host genetic background had no significant effect upon the ability of unelicited peritoneal macrophages to kill *E. coli* (even though the mouse strains differ widely in their *in vivo* susceptibilities to bacterial infection); and (iii) the FimH phenotype had no significant effect upon *E. coli* survival once the bacterium was inside a macrophage. Additionally, there was no correlation between the bacteriocidal effectiveness of a macrophage population and the number of bacteria bound per macrophage. However, macrophage populations that were the least bacteriocidal tended to bind higher ratios of FimH⁺ to FimH⁻ *E. coli*. The effect of gamma interferon, fetal calf serum, and the recombination proficiency of *E. coli* were examined as factors predicted to influence intracellular bacterial killing. These had no effect upon the rate of *E. coli* elimination by unelicited peritoneal macrophages.

The mechanism by which bacteria are taken up and killed by macrophages and other cells of the reticuloendothelial system in the absence of normal or immune serum components has been under study for a number of years (39). This process has been referred to as nonopsonic phagocytosis. Nonopsonic processes differ in a number of respects from the opsonic mechanisms. One important difference is an increased rate of killing of opsonized internalized bacteria (3, 54, 59). Nonopsonic phagocytosis has been characterized as a primitive holdover from protozoal ingestion mechanisms (40). However, since some antibodies, particularly those directed against bacterial adhesins, actually prevent phagocytosis (62), it may be fortunate that this poorly understood mechanism is still in place.

One well-described bacterial ligand that mediates nonopsonic phagocytosis is the type 1 pilus (8). These pili are produced by many members of the *Enterobacteriaceae* (12) and promote bacterial adherence to the mucosal surfaces of a wide variety of hosts through a mannose-sensitive interaction with receptors on eucaryotic cells (12). This adherence is thought to allow the colonization of a number of host compartments (42) and pro-

mote interindividual spread (4). Type 1 pili also mediate adherence to phagocytic cells (39). The interactions between type 1 pili and neutrophils (58), mast cells (30–32), macrophages (3), and other leukocytes (45) have been some of the more carefully examined interactions between bacteria and phagocytes.

Early and more recent work on the nature of the interaction of type 1 pilated *Escherichia coli* cells with macrophages indicates that one minor component of the pili, the product of the *fimH* gene (FimH), is responsible for adherence (25). Whereas this adherence, in effect, tethers the pilated bacteria to macrophages and effectively increases the number of bacteria bound compared to the number of *fimH* mutants bound (25) and induces an oxidative burst (5, 15, 29, 41), reports on whether type 1 piliation actually results in an increased rate of killing compared to that for nonpilated or FimH⁻ cells, under defined *in vitro* conditions, are conflicting (5, 15, 16, 22, 25, 29). Reports agree that leukocyte-bound type 1 pilated cells are better protected against killing than opsonized FimH⁻ *E. coli* cells (3, 16, 55). This protection is due, at least in part, to a difference in the compartmentalization of opsonized *E. coli* versus that of FimH⁺ *E. coli* in bone marrow-derived macrophages (3). Direct comparisons of otherwise isogenic FimH⁺ and FimH⁻ bacteria (both unopsonized) suggest that there is a modest but statistically significant increase in the survivability of FimH⁺ over that of FimH⁻ *E. coli* in macrophages (25).

In order to better understand factors affecting innate host

* Corresponding author. Mailing address: Department of Microbiology, Pathology, and Parasitology, College of Veterinary Medicine, North Carolina State University, 4700 Hillsborough St., Raleigh, NC 27606. Phone: (919) 513-6207. Fax: (919) 513-6455. E-mail: Paul_Orndorff@ncsu.edu.

TABLE 1. Mouse and bacterial strains and cell lines used in this study

Organism or Cell line	Description and/or relevant phenotype	Source or reference ^a
Mice		
BALB/c	Inbred; <i>ity</i> ^S <i>lps</i> ^N ; sensitive to <i>Y. enterocolitica</i> , <i>L. monocytogenes</i> , and <i>S. enterica</i> serovar Typhimurium	11, 17, 48
C57BL/6	Inbred; <i>ity</i> ^S <i>lps</i> ^N ; resistant to <i>Y. enterocolitica</i> , <i>L. monocytogenes</i> , and <i>Legionella pneumophila</i>	11, 17, 65
CD-1	Outbred; <i>ity</i> ^R (presumed) <i>lps</i> ^N ; resistant to <i>S. enterica</i> serovar Typhimurium and <i>L. monocytogenes</i>	1, 2, 38
C3H/HeJ	Inbred; <i>ity</i> ^R <i>lps</i> ^D ; susceptible to <i>S. enterica</i> serovar Typhimurium and <i>Francisella tularensis</i>	21, 44, 61
C3H/HeN	Inbred; <i>ity</i> ^R <i>lps</i> ^N ; resistant to <i>S. enterica</i> serovar Typhimurium and <i>F. tularensis</i>	21, 48
Mouse cell lines		
J774	Macrophage-like cell line derived from BALB/c reticulum cell sarcoma	47a
IC-21	Macrophage cell line derived from simian virus 40 transformation of C57BL/6 peritoneal macrophages	34
Bacteria		
<i>E. coli</i>		
ORN115	<i>thr-1 leuB thi-1 Δ(argF-lac)U169 malA1 xyl-7 ara-13 mt1-2 gal-6 rpsL fhuA2 supE44 pilG Pil⁺ Hag⁺ (FimH⁺) Mal⁻</i>	56
ORN133	Same as ORN115 except <i>fimH::Kan Hag⁻ (FimH⁻) Mal⁻</i>	35
ORN175	Same as ORN115 except <i>Mal⁺</i>	25
ORN204	Same as ORN133 except <i>Mal⁺</i>	P1 transduction from ORN109 (19)
ORN205	Same as ORN115 except <i>recA13 Rec⁻</i>	P1 transduction from <i>E. coli</i> EC901 and EC931 (43)
ORN172	Same as ORN115 except <i>Δfim(BEACDEFGH) Pil⁻ Rec⁺</i>	63
ORN201	Same as ORN172 except <i>recA13 Rec⁻</i>	P1 transduction from <i>E. coli</i> EC901 and EC931 (48)
<i>L. monocytogenes</i>		
EGD 1/2a	Presumed wild type	37

^a References for mouse strains refer to their resistant properties. More-comprehensive listings of resistances and derivations of mouse strains can be found at the Jackson Laboratory, Bar Harbor, Maine.

susceptibility to bacterial diseases and also the role of type 1 pili in the nonopsonic phagocytic process, we examined resident peritoneal macrophages from five different mouse strains for their abilities to kill *E. coli* that were phenotypically either FimH⁺ or FimH⁻. Also, for some mouse strains, additional elicitation methods, anatomical sources, and derivation methods were examined to see if these factors affected macrophage function.

MATERIALS AND METHODS

Mouse strains and cell lines. Male BALB/c, C3H/HeN, C3H/HeJ, C57BL/6, and CD-1 mice 8 to 12 weeks of age were used in these experiments. The mice were purchased from either Charles River Laboratories (Wilmington, Mass.) or Taconic Farms (Germantown, N.Y.). Mice were maintained under pathogen-free husbandry conditions and were fed food and water ad libitum. Relevant genotypic and phenotypic differences in the mouse strains and macrophage-like cell lines are listed in Table 1.

Bacterial strains and growth conditions. Bacterial strains used in this study are shown in Table 1. All bacteria were grown in L broth (36) overnight with shaking to stationary phase. Prior to the bacteriocidal assay, bacteria were harvested by centrifugation (6,000 × g for 10 min), resuspended, and diluted to ca. 2 × 10⁸ cells/ml in phosphate-buffered saline (PBS). Media used to assess bacterial numbers both before and after macrophage killing assays consisted of maltose tetrarazolum agar (53) for *E. coli* (25) and L agar (36) for *Listeria monocytogenes*.

Isolation and cultivation of peritoneal macrophages. Peritoneal cells were isolated by intraperitoneal (i.p.) injection of 7 to 8 ml of RPMI 1640 medium (Gibco BRL, Grand Island, N.Y.) containing 5% fetal bovine serum (FBS), gentamicin (5 μg/ml), and heparin (5 U/ml) into mice that had been killed by cervical dislocation. Following i.p. injection, the mice were shaken to dislodge peritoneal cells and the lavage fluids were removed by syringe. The peritoneal cells were centrifuged (430 × g for 5 min), the resulting cell pellet was suspended in the aforementioned medium lacking heparin at a concentration of 5 × 10⁵ cells/ml, and 0.5 ml of the cell suspension was placed into each well of a 48-well

cell culture cluster plate (Costar, Cambridge, Mass.). Two hours later, the medium was removed and the plastic adherent cells were washed three times with 0.5 ml of Hanks balanced salt solution (HBSS) and then incubated in 0.5 ml of RPMI 1640 medium containing 5% FBS without antibiotics at 37°C in a humidified 5% CO₂ incubator. Eighteen to twenty-four hours later, the culture medium was aspirated and the cells were washed once with HBSS and then incubated with 0.25 ml of the above culture medium lacking antibiotics. Elicitation of proteose peptone- or thioglycolate-elicited inflammatory macrophages was done by injecting mice i.p. with 2 ml of either sterile 10% proteose peptone (Difco, Detroit, Mich.) or thioglycolate broth (Remel, Kansas City, Mo.). Three days later, the peritoneal cells were harvested as described above.

In some experiments, macrophages were treated with gamma interferon (IFN-γ) prior to use in the bacteriocidal assays. Immediately following the 2-h plastic adherence procedure, macrophages were incubated in medium containing 10 antiviral units of homogeneously pure recombinant mouse IFN-γ/ml (8.0 × 10⁶ U/mg of protein) for 18 to 24 h as previously reported (26, 47) and then used in the bacteriocidal assays. The recombinant mouse IFN-γ (lot 2271-54-F2) was the kind gift of Genentech, Inc. (South San Francisco, Calif.).

Cultivation of bone marrow-derived macrophages and macrophage cell lines. Bone marrow-derived macrophages were cultured as previously reported (20). IC-21 cells were grown in RPMI 1640–10% FBS. J774 cells were grown in Dulbecco minimal essential medium–Ham's F-12 mixture (1:1) with 10% FBS.

Macrophage bacteriocidal assay. Macrophages in 48-well tissue culture plates (~10⁵ to 2 × 10⁵ cells/well) containing 0.25 ml of RPMI 1640 medium were exposed to approximately 10⁶ bacteria added in 50 μl of PBS for 10 min at 37°C. After incubation, wells were washed four times (each wash was with 1 ml of PBS). After the final wash, 0.5 ml of prewarmed RPMI 1640 was added to each well. Gentamicin (2.5 μl of a 1-mg/ml stock) was added to selected wells to produce a final concentration of 5 μg of gentamicin/ml. The addition of 0.1 ml of 1.0% Triton X-100 lysed the macrophages and defined the end of the incubation. One to two minutes after the Triton X-100 additions, the contents of the wells were diluted and plated. The brief exposure of bacteria to gentamicin and Triton X-100 had no effect on bacterial viability. At each time point, the contents of (typically) four wells were plated: two wells contained macrophages, with each well having a mixture of FimH⁺ and FimH⁻ *E. coli* cells added at approximately

a 1:1 ratio. (Sometimes the two wells were duplicates; occasionally one well had some other treatment.) The two remaining wells did not have macrophages but were treated as if they did, both before and after the addition of bacteria. In one of the wells, gentamicin was added. In the last well gentamicin was omitted. In pilot experiments, we found that a small number of bacteria remained bound to the plastic of the wells without macrophages (1 to 10% of the macrophage-bound bacteria). The bacteria in each of these wells were used to assess the bacteriocidal kinetics of gentamicin when bacteria were "unprotected" and bacterial growth in the absence of gentamicin. Time points were spaced according to data from pilot experiments, which indicated that >90% of unprotected bacteria were eliminated after 20 to 25 min of exposure to gentamicin (establishing the basis for the first time point) and that the numbers of internalized bacteria were constant or tending upward after 4 h (establishing the basis for the end of the assay). The number of bacteria initially bound to macrophages was determined as described above but just prior to the addition of gentamicin.

Maltose tetrazolium agar allowed the convenient identification of FimH⁺ and FimH⁻ strains, which were genetically marked by their abilities to utilize maltose (25) (Table 1). The ratio of FimH⁺ to FimH⁻ *E. coli* cells obtained after macrophage exposure was normalized to the starting FimH⁺/FimH⁻ ratio to determine if there was any difference in the rate of macrophage killing based upon the FimH phenotype. FimH⁺ and FimH⁻ cells did not differ in their gentamicin sensitivities (tested in the control well lacking macrophages but with gentamicin) and did not differ noticeably in their growth rates (determined in wells lacking macrophages and gentamicin). Plating efficiencies on maltose tetrazolium agar, previously noted as slightly different depending upon the Mal phenotype (25), were not appreciably different under the present conditions.

Statistical analysis. Each experiment was typically performed with at least duplicate sets of wells. Standard deviations of the means of at least two separate experiments were calculated with the aid of the Microsoft Excel STDEV function. Standard error was calculated as the standard deviation divided by the square root of the number of experiments. All tabulated or illustrated values are the averages of at least two independent experiments. Regression analysis of the bacteriocidal curves was performed with the aid of the Microsoft Excel, version 5, trend line generator feature. Significant differences between means were determined by either Student's *t* test or analysis of variance. Both tests were provided by the Microsoft Excel, version 4, statistics package. The *F* statistic was also used to determine the probability of an accidental association of two variables in generating a trend line (Microsoft Excel, version 4). Statistically significant differences were defined as $P < 0.05$.

RESULTS

Analysis of macrophage killing curves. In order to compare the bacteriocidal activities of distinct macrophage populations from different anatomical sites of different mouse strains, a standard was needed. Pilot experiments revealed that the most consistent results came from conditions that produced a final ratio (after washing off unbound bacteria) of approximately 1 bound bacterium per 10 macrophages. This low multiplicity allowed direct comparisons between FimH⁺ and FimH⁻ bacteria in a single well under noncompetitive binding conditions and reduced potentially confounding effects associated with high multiplicities of infection (e.g., endotoxin effects). At 20 to 25 min after gentamicin addition, we began measuring the rate of killing. At least 90% of the bacteria still viable at this time had been internalized by macrophages (and were thus protected from gentamicin), but had not yet been killed. The remaining ca. 10% were external bacteria that had not been killed by gentamicin in the 25-min period. A considerable fraction ($84 \pm 41\%$ of the total originally bound; 10 experiments averaged) of the FimH⁺ and FimH⁻ *E. coli* cells initially bound (i.e., present immediately after washing) were still viable after 20 to 25 min of gentamicin exposure, even in the most bacteriocidal macrophage populations (unelicited peritoneal macrophages from BALB/c mice; see below). The fate of this "gentamicin-protected" population (set as 100%) over the next 4 h constituted our bacteriocidal curves. An example of one such curve, along with two parameters associated with the trend line through the points, is shown in Fig. 1. In this example, the points on killing curves represent values averaged from both FimH⁺ and FimH⁻ *E. coli*.

Parameters employed to assess differences in bacteriocidal activity between macrophage populations. Regression analysis of linear, exponential, and parabolic trend lines showed that

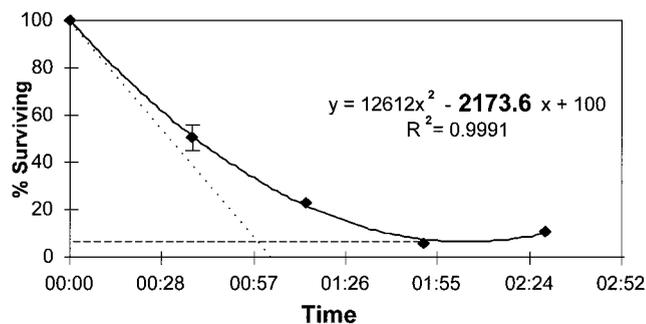


FIG. 1. Curve denoting the percentage of gentamicin-protected *E. coli* surviving over time. The trend line through the points (diamonds) describes a parabolic curve represented by the top equation. The instantaneous slope of the curve (dy/dx at $x = 0$) is depicted by the dotted line, and the numerical value is highlighted in the top equation. The y value at the vertex ($dy/dx = 0$) of the curve is illustrated by the dashed line. The R^2 statistic (coefficient of determination) denotes the degree to which the values of the experimentally derived points match those predicted by the equation shown. (R^2 values range from 0 to 1.0, with 1.0 being a perfect match.) Vertical bars represent standard errors from an experiment performed in triplicate.

the shape of most bacteriocidal curves most closely matched that of a parabola. This is reflected in the R^2 statistic (coefficient of determination), which gave the highest average value (a perfect correlation produces an R^2 value of 1.0) when 18 killing curves (employing unelicited peritoneal macrophages from different mouse strains and with values for FimH⁺ and FimH⁻ *E. coli* averaged) were compared (parabolic, $R^2 = 0.80 \pm 0.17$; linear (first three points), $R^2 = 0.78 \pm 0.24$; exponential, $R^2 = 0.74 \pm 0.16$). For the linear curve, only the first three time points were considered because the killing curves departed from linearity relatively rapidly. Data compiled from 15 different macrophage populations and 33 individual experiments revealed that the ordinate value of the vertex of the parabolic trend line accurately and precisely predicted the actual recorded minimum percentage of surviving *E. coli* cells (FimH⁺ and FimH⁻ cells averaged), further indicating the natural parabolic shape of the killing curves (Fig. 2A). Initial rate measurements (of both linear and parabolic curves) were less predictive of the ability of a macrophage population to eliminate internalized *E. coli* (reflected in the lower R^2 statistic; Fig. 2B and C). Nevertheless, rate measurements were linearly related to the minimal percent surviving. This linear relationship was confirmed by calculating the significance of the *F* statistic, which indicates the probability of erroneously concluding that a linear relationship exists (in all cases $P < 0.05$).

For the purpose of comparing killing curve parameters (Fig. 2), we excluded twelve experiments because a killing curve was not generated. That is, in these experiments, the ingested *E. coli* grew, generating a positively sloped linear trend line. While this exclusion of these curves was justified in drawing the conclusions about killing curve shape, this limited the use of values calculated from the parabolic curve (e.g., vertex) since not all experimental results could be analyzed this way. In view of this shortcoming, two curve parameters, percent eliminated and the initial linear rate of elimination, were used for all subsequent measurements. Values for both of these parameters could be derived from any curve, and the parameters could be applied to any macrophage population.

Comparison of the relative abilities of distinct macrophage populations from different mouse strains to eliminate internalized FimH⁺ and FimH⁻ *E. coli*. Peritoneal, bone marrow, and cell line macrophages derived from BALB/c and C57BL/6

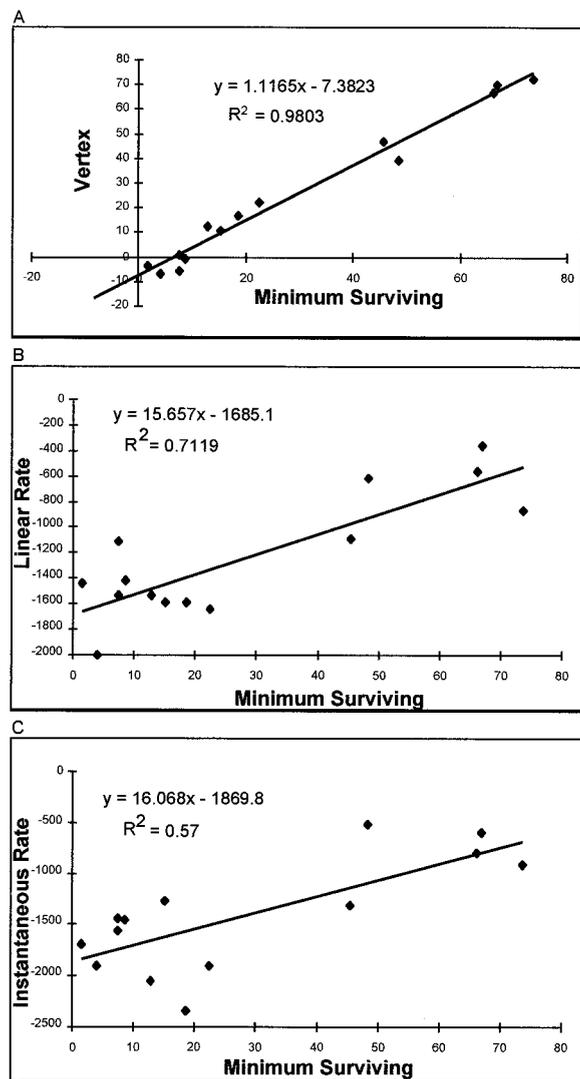


FIG. 2. Comparison of curve parameters for their abilities to predict the minimum percent of internalized bacteria surviving in a bacteriocidal assay. (A) Points represent the vertex values of curves describing the bacteriocidal assay versus the actual minimal percentage of bacteria surviving in the bacteriocidal assay. (B) Points represent the initial linear rate of elimination (in units of percentage of internalized bacteria eliminated per 24-h period) versus the actual minimal percentage of bacteria surviving in the bacteriocidal assay. (C) Points represent the instantaneous rate of bacterial elimination based upon a parabolic trend line best describing the points of the bacteriocidal assay (in arbitrary units of percentage of internalized bacteria eliminated per 24-h period) versus the actual minimal percentage of bacteria surviving in the bacteriocidal assay. Negative values on the ordinates in panels B and C reflect the negative slopes of the bacteriocidal assays. The top equation in each panel describes the curve derived from linear regression analysis. The R^2 value denotes the coefficient of determination as defined in the text.

mice were initially compared for their abilities to eliminate internalized FimH⁺ and FimH⁻ *E. coli*. When results indicated that elicitation did not enhance the rate or degree of killing of either FimH type of *E. coli*, elicitation methods were not employed for subsequent mouse strains examined (CD-1, C3H/HeJ, and C3H/HeN). For C3H/He strains, only resident peritoneal macrophages were tested. In all macrophage populations tested, there was no statistically significant difference in the degree or rate at which FimH⁺ and FimH⁻ *E. coli* cells

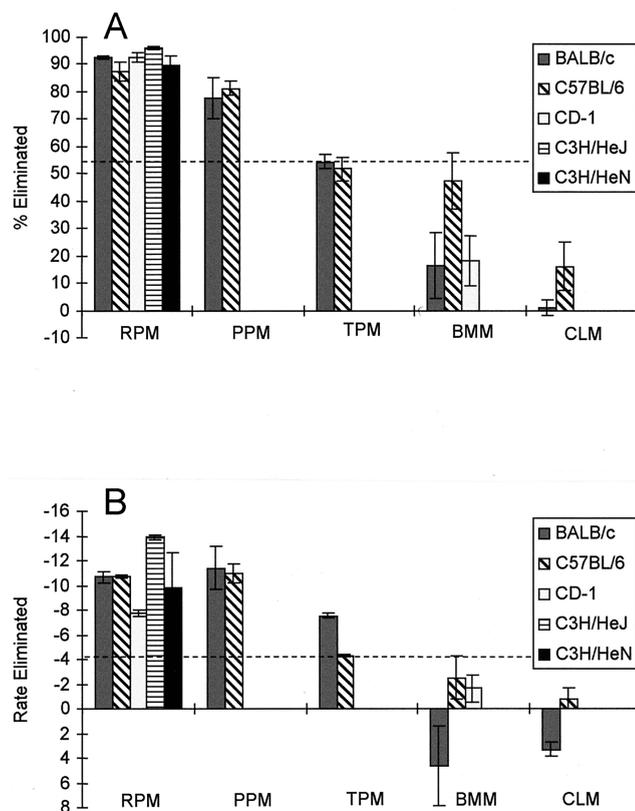


FIG. 3. Capacities of macrophage populations to eliminate internalized *E. coli*. (A) Percentages of internalized *E. coli* cells eliminated by macrophages from the mouse strains. (B) Initial rate of *E. coli* elimination in units of percent internalized bacteria eliminated per 10-min period. Negative values indicate a decrease in the internalized population; positive values denote an increase. RPM, resident (unelicited) peritoneal macrophages; PPM, proteose peptone-elicited peritoneal macrophages; TPM, thioglycolate-elicited peritoneal macrophages; BMM, bone marrow-derived macrophages; CLM, cell line macrophages. The cell line macrophages constitute the J774 line for BALB/c mice and IC-21 for C57BL/6 mice. Only unelicited peritoneal macrophages and bone marrow-derived macrophages were tested for CD-1 mice, and only unelicited peritoneal macrophages were tested for C3H/HeJ and C3H/HeN mice. Vertical lines represent the standard errors of the means. The dashed line in each panel divides two statistically distinct groupings: results extending above the line were (as a group) statistically distinct from those grouped below the line.

were eliminated. Consequently, average measurements of killing effectiveness include both FimH⁺ and FimH⁻ *E. coli*.

For BALB/c and C57BL/6 mice, unelicited and proteose peptone-elicited peritoneal macrophages were superior to thioglycolate-elicited macrophages, bone marrow-derived macrophages, and cell line macrophages in their ability to eliminate internalized *E. coli* (Fig. 3). This was especially apparent when the maximal number of ingested bacteria eliminated was used as a measure (Fig. 3A). Nevertheless, the rate measurement produced the same trend (Fig. 3B). One noticeable difference between the statistically significant groupings in Fig. 3A and B was that the BALB/c thioglycolate-elicited peritoneal macrophages showed a high initial rate of *E. coli* elimination that was not indicative of the relatively low total percentage of *E. coli* cells eliminated. Also, in some of the macrophage populations that killed poorly, *E. coli* actually grew (initially) before being reduced in number later in the assay (Fig. 3B). The outbred CD-1 strain exhibited the same general trend as the BALB/c and C57BL/6 mice, indicating the superiority of the resident peritoneal macrophages over bone

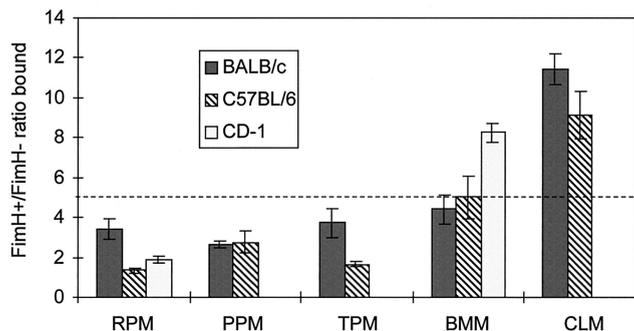


FIG. 4. Preferential binding of FimH⁺ *E. coli* over FimH⁻ *E. coli* by macrophages from different host strains, anatomical locations, and derivations. The ratio of FimH⁺ to FimH⁻ *E. coli* cells bound is shown on the vertical axis. The macrophage populations are as defined in the legend for Fig. 3. Only unelicited peritoneal macrophages and bone marrow-derived macrophages were tested for CD-1 mice. Vertical lines represent the standard errors of the means. The dashed line divides two statistically distinct groupings: results extending above the line are (as a group) statistically distinct from those grouped below the line.

marrow-derived macrophages (Fig. 3). The killing capacities of resident peritoneal macrophages from all five mouse strains tested were statistically indistinguishable regardless of the type of measurement employed (statistical analysis of data shown in Fig. 3).

***E. coli* binding characteristics of macrophage populations in comparison to their bacteriocidal abilities.** Whereas we could find no difference in the abilities of macrophage populations to eliminate FimH⁺ *E. coli* relative to FimH⁻ *E. coli*, they did differ in the binding of FimH⁺ and FimH⁻ cells (Fig. 4). The macrophage populations that bound the highest ratios of FimH⁺ to FimH⁻ bacteria (bone marrow and cell line macrophages) were the populations least able to kill *E. coli* once ingested. For BALB/c and C57BL/6 mice, this trend was statistically significant only in distinguishing the cell lines from the peritoneal and bone marrow-derived macrophages. For the outbred CD-1 mice, where just bone marrow and unelicited peritoneal macrophages were examined, there was a statistically significant difference between these two macrophage populations with respect to binding. There was no correlation between the absolute numbers of *E. coli* cells bound per macrophage and the bacteriocidal effectiveness. That is, the macrophage populations most effective at killing internalized *E. coli* did not bind significantly more of them. This was established in tests comparing unelicited macrophages (used as one statistical grouping) and the other macrophage populations (data not shown).

Examination of additional variables for macrophage bacteriocidal ability. The effect of serum in the assay, the recombination proficiency of the ingested bacteria (their ability to repair DNA), and the effect of IFN- γ treatment on macrophage bacteriocidal activity were examined. We tested the effect of FBS on macrophage bacteriocidal activity because *E. coli* antibodies, if elicited in utero, had the potential to serve as opsonizing antibodies. However, experiments done in the absence of serum had no significant effect upon the binding or the subsequent rate or degree of killing of FimH⁺ or FimH⁻ *E. coli*. This was true whether the serum was left out during the absorption period only or left out of the macrophage preparation protocol and killing assay entirely (data not shown).

Our results for *E. coli recA* mutants (we tested both fimbriate and nonfimbriate *recA* mutants) indicated that the rate and degree to which they were eliminated by BALB/c mouse unelicited macrophages did not differ significantly from those for

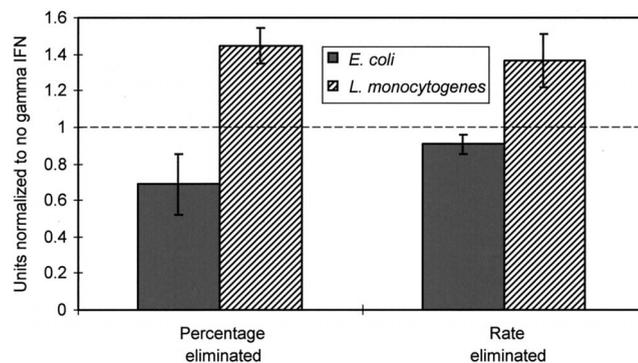


FIG. 5. Effect of the addition of 10 U of IFN- γ /ml to unelicited peritoneal macrophages prior to the bacteriocidal assay. IFN- γ was added as described in the text. The percentages of internalized *E. coli* and *L. monocytogenes* cells eliminated and the initial rates of elimination were recorded in three experiments for *L. monocytogenes* and four experiments for *E. coli*. In each experiment, the differences in degree and rate of elimination were normalized to the values obtained with no IFN- γ addition and then averaged (standard errors are denoted by the vertical lines). The dashed line denotes the normalized value (1) with no IFN- γ addition.

the parent (data not shown). Similarly, IFN- γ had no effect upon the rate or degree to which FimH⁺ and FimH⁻ *E. coli* cells were killed by unelicited macrophages from the BALB/c and C57BL/6 strains (Fig. 5). IFN- γ did have a significant effect upon the degree to which *L. monocytogenes* was killed by unelicited peritoneal macrophages (the lowest point in the killing curve); consequently the IFN- γ did appear to be having an effect on the macrophages. This effect of IFN- γ on macrophage killing of *L. monocytogenes* did not produce a difference in the initial elimination rate of these microorganisms.

DISCUSSION

In the experiments described herein, we compared the innate abilities of different macrophage populations from mice of different genetic backgrounds to kill internalized *E. coli*. Bacteriocidal activity was measured by two parameters (i) the initial elimination rate and (ii) the maximal percentage of internalized *E. coli* eliminated. FimH⁺ and FimH⁻ *E. coli* cells were used to assess the effects of different bacterial cell surface interactions upon the binding and elimination of the bacteria once internalized. Additional assays that examined physiological and genetic factors that might influence the process were employed.

Quantitative bacteriocidal measurements which involved measuring the percentage of internalized bacteria eliminated were developed. Employing this measurement effectively normalized the results so that data from macrophages with different binding and uptake kinetics could be compared. We chose to present two measurements, (i) the initial rate and (ii) the degree to which an internalized population was eliminated. Of these two, the measurement of maximal percentage eliminated was found to be the more reproducible. In particular, we often found that high initial elimination rates often did not result in a high percentage of internalized bacteria eliminated. A similar observation was made by van Dissel et al. (60) with opsonized *Salmonella enterica* serovar Typhimurium.

Using regression analysis, we found that the ordinate value of the vertex of a parabola describing the killing curves predicted the maximal percentage of internalized bacteria eliminated quite well. In contrast, van Dissel et al. (60) found that opsonized *S. enterica* serovar Typhimurium underwent an ex-

ponential decrease after being taken up by macrophages over the 90-min period of their assays. An exponential decrease would be expected if pure probability determined elimination rate (i.e., the macrophages acting as a simple bacteriocidal agent). We expect that the parabolic shape coincidentally best described a dynamic state in which most ingested bacteria were being killed by macrophages in an exponential fashion but in which other ingested bacteria were growing, protected from the gentamicin by macrophages incapable of killing them (or not killing all of them). As time progressed, the replicative power of the protected *E. coli* began to be witnessed as the upward slope of the parabola. Consequently, the parabolic shape may be simply indicative of a generally inefficient process. However, the killing curve shape is not a widely analyzed feature of bacteriocidal assays. Future and retrospective attention to the shapes of killing curves may reveal unappreciated mechanistic relationships.

Both the initial linear elimination rate and the degree to which internalized *E. coli* cells were eliminated supported the same order of macrophage bacteriocidal effectiveness, with resident and proteose peptone-elicited peritoneal macrophages being consistently the best, followed by, in declining order of effectiveness, thioglycolate-elicited peritoneal macrophages, bone marrow-derived macrophages, and macrophage-like cell lines. The killing curves of macrophage populations that eliminated approximately 60% (or less) of the ingested bacteria had significantly lower R^2 values (for linear or parabolic curves), and the curve parameters had higher standard deviations, than those of macrophage populations killing greater than 90% of the ingested bacteria. We found it difficult to draw any conclusions about the killing capacities of macrophage populations in this low-level killing category. The inability of thioglycolate-elicited macrophages to efficiently kill bacteria has been often reported (7, 18, 27, 57). Likewise, our observation that macrophage cell lines and bone marrow-derived macrophages were inefficient at killing internalized bacteria, compared to unelicited peritoneal macrophages, was consistent with those of others (9, 46–48). However, bone marrow-derived macrophages have been frequently used in bacterial phagocytosis assays (3, 46). When bone marrow-derived macrophages were examined with FimH⁺ *E. coli* (3), the degree to which FimH⁺ *E. coli* cells were eliminated was found to be similar to that reported here (FimH⁻ *E. coli* cells were not tested previously [3]).

Mouse genetic background had no effect on the in vitro bacteriocidal capacity of macrophages even though the strains of mice differ rather dramatically in their susceptibilities to gram-negative and gram-positive bacterial pathogens (33) (Table 1). We were somewhat surprised that there was no effect. However, susceptibility to bacterial infection, as it relates to macrophage bacteriocidal effectiveness, depends upon a number of factors, among them the type of bacteria under investigation (17, 51) and whether the bacteria are opsonized or not (1, 28, 60). Our results indicate that the innate susceptibility of mice to a variety of bacterial infections does not correlate with the ability of host macrophages to take up or kill unopsonized *E. coli*.

Whereas previous experiments have shown statistically significant differences between the survival of FimH⁺ *E. coli* and that of FimH⁻ *E. coli* in resident peritoneal macrophages (25), we witnessed no such differences here. The assay conditions, the criteria used to assess killing, and the magnitude of the differences shown in this earlier report leave open the possibility that there may be little difference between the killing of FimH⁺ and FimH⁻ *E. coli* once internalization has occurred. Consequently, our results support observations that piliation

has little effect on the outcome of nonopsonic phagocytosis as far as killing of ingested bacteria is concerned (5, 6, 16, 22). Our results leave open the possibility that FimH⁺ and FimH⁻ *E. coli* cells are directed to different vacuolar compartments once internalized (analogous to FimH⁺ and opsonized *E. coli*) (3) but indicate that this hypothetical difference in trafficking makes no effective difference in terms of killing, at least as measured by the methods we employed.

Whereas the killing of ingested bacteria was not influenced by the FimH⁺ or the FimH⁻ phenotype, macrophage populations were found to have distinctly different binding properties when the ratios of FimH⁺ to FimH⁻ *E. coli* bound were compared. We attribute these binding differences to different FimH receptor densities (3, 13) on the various macrophage populations relative to the densities of receptors that simply bind FimH⁻ *E. coli* via another mechanism or mechanisms (e.g., via lipopolysaccharide [64]). Interestingly, the macrophage populations least effective in killing ingested *E. coli* exhibited the highest relative binding of FimH⁺ *E. coli*. The biological significance of this is unknown. Since a number of different receptors have been proposed to have a role in binding FimH on phagocytic cells (3, 14, 24, 52), it may be that certain macrophage populations vary in the expression of receptor type as well as density.

Two factors that have been shown to affect the fate of bacteria internalized by macrophages are (i) recombination proficiency, shown to be a factor in the survival of (opsonized) *S. enterica* serovar Typhimurium (10) and (ii) the exogenous addition of IFN- γ , shown to affect the fate of unopsonized *L. monocytogenes* (47) and opsonized *S. enterica* serovar Typhimurium (23, 48), as well as the phagocytosis of unopsonized *E. coli* (49, 50; killing was not measured in these reports). Neither of these factors appears to affect the rate or degree to which unopsonized *E. coli* cells (FimH⁺ or FimH⁻) were killed as measured by our methods, although control experiments showed that IFN- γ did significantly improve the degree to which unopsonized *L. monocytogenes* was eliminated, as has been previously reported (47). Understanding the reason for the differential effect of IFN- γ may aid in understanding the molecular mechanisms by which certain bacteria are able to thwart macrophage killing. Under our assay conditions, the addition of heat-inactivated 5% FBS to unelicited peritoneal macrophages had no significant effect upon the binding or subsequent killing of FimH⁺ *E. coli* as has been previously reported (3). It may be that FBS effects are manifested only under certain conditions or with certain macrophage populations.

There are many complex biochemical events involved in the internalization and killing of bacteria by phagocytes. In vitro assays to quantify bacterial killing by macrophages can provide useful insights into host-pathogen interactions. However, there are numerous variables in such assays that make comparisons from different laboratories difficult. In the present study, we systematically examined several of the principal assay variables for their effects on the fate of phagocytized, unopsonized *E. coli*. We found that several factors previously reported to influence the microbicidal activity of macrophages, such as the presence or absence of serum, activation of the macrophage by IFN- γ , recombination proficiency of the bacteria, FimH phenotype, and the mouse strain background of the macrophage population had no effect. Of relevance for future studies was our finding that the anatomical source of macrophages and their derivation significantly influenced bacteriocidal activity.

ACKNOWLEDGMENTS

We thank Craig Altier for a critical reading of the manuscript and helpful suggestions.

This work was supported by grant AI 222223 from the Public Health Service and the State of North Carolina.

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Editor: J. T. Barbieri