

The K1 Capsule Is the Critical Determinant in the Development of *Escherichia coli* Meningitis in the Rat

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Abstract

Although *Escherichia coli* strains possessing the K1 capsule are predominant among isolates from neonatal *E. coli* meningitis and most of these K1 isolates are associated with a limited number of 0 lipopolysaccharide (LPS) types, the basis of this association of K1 and certain 0 antigens with neonatal *E. coli* meningitis is not clear. The present study examined in experimental *E. coli* bacteremia and meningitis in newborn and adult rats whether or not the K1 capsule and/or 0-LPS antigen are critical determinants in the development of meningitis. Rats received subcutaneously a K1 *E. coli* strain (018+K1+) or mutants lacking either the K1 capsule (018+K1-) or 0 side-chain (018-K1+). 12–24 h later, blood and cerebrospinal fluid (CSF) specimens were obtained for quantitative cultures. The isolation of *E. coli* from CSF was observed in both newborn and adult rats infected with K1+ strains regardless of LPS phenotype (018+ or 18-) who also developed a high degree of bacteremia (e.g., > 10⁴ CFU/ml of blood). In contrast, none of the newborn and adult rats infected with 018+K1- and developing bacteremia of > 10⁴ were found to have positive CSF cultures. These findings indicate that the presence of the K1 capsule and a high degree of bacteremia are key determinants in the development of *E. coli* meningitis, suggesting that there may be specific binding sites present in the brain which have an affinity for the K1 capsule and thus may be responsible for the entry of K1-encapsulated *E. coli* into the meninges. (*J. Clin. Invest.* 1992. 90:897–905.) **Key words:** *Escherichia coli* • hematogenous meningitis • lipopolysaccharide • K1 capsular polysaccharide • pathogenesis

Introduction

The mortality and morbidity associated with neonatal gram-negative bacillary meningitis have remained significant despite advances in antimicrobial chemotherapy and supportive care (1, 2). Both clinical and experimental data indicate limited efficacy with antimicrobial chemotherapy alone (3, 4). A contributing factor may be the incomplete development of host defense mechanisms in neonates (5, 6). Thus, adjunct therapy

in the form of active or passive immunotherapy is a logical next step. The feasibility of this approach, however, is limited by inadequate understanding of host-microbial factors associated with the increased susceptibility of neonates to gram-negative bacilli.

Escherichia coli is the most common gram-negative organism that causes meningitis during the neonatal period (1, 2). Given the plethora of *E. coli* serotypes (e.g., 103 capsular or K, 170 0 lipopolysaccharides or LPS, and 56 flagellar or H antigens) (7), it is striking that *E. coli* strains possessing the K1 capsular polysaccharide are predominant (~80%) among isolates from neonatal *E. coli* meningitis (8–10). It is also of interest that only a few somatic antigens (e.g., 018, 07, 01, 016) have been associated with most of the K1 strains isolated from the cerebrospinal (CSF)¹ (11). The basis of this association of K1 and certain 0 antigens with neonatal *E. coli* meningitis is not clear.

We have previously shown that experimental *E. coli* bacteremia and meningitis in newborn rats have important similarities to human newborn *E. coli* infection. Hematogenous infection of meninges can be reliably established without the need for administration of adjuvant or direct inoculation of bacteria into CSF (4, 12–17). With development of techniques for atraumatic collection of blood and CSF specimens, we have used this model to determine whether or not the K1 capsule and/or 0-LPS antigen are critical determinants in the development of *E. coli* meningitis.

Methods

Bacterial strains. A serum-resistant *E. coli* strain possessing the K1 capsular polysaccharide, C5 (serotype 018:K1:H7) was originally obtained from Dr. R. Bortolussi, Dalhousie University, Halifax, Nova Scotia, Canada. This strain was isolated from the CSF of a newborn infant with meningitis (12). The presence of the K1 capsule was confirmed by lytic sensitivity to the K1-specific bacteriophages (18–20) as well as by the antiserum agar technique with equine antiserum to group B meningococcus (11). *Salmonella typhimurium* strain X3604 cya::Tn 10 was obtained from Dr. R. Curtiss III, Department of Biology, Washington University, St. Louis, MO.

K1-negative derivatives. Unencapsulated mutants of strain C5 were obtained by selection for resistance to the K1-specific bacteriophages (phages A, B, C, D, E) as previously described (18–20). Each clone was then reexamined for the loss of capsule production using the K1-specific bacteriophages as well as the antiserum agar technique. Previous studies in our laboratories have shown no detectable differences in LPS and outer membrane protein complex between parents and K1- clones when examined by SDS-PAGE (20–22). Parent K1+ strains and their K1- mutants also possess identical hemolysin, biochemical

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1. **Abbreviations used in this paper:** CSF, cerebrospinal fluid; Tc^R, tetracycline resistance (gene).

reactions, and binding patterns to homologous LPS monoclonal antibody (by Western blot using whole organisms or LPS). In addition, parent K1+ strain and K1- mutants were found to have identical genotypes when examined by the multilocus enzyme electrophoresis (kindly performed by Dr. Richard Wilson at *E. coli* Reference Center, Pennsylvania State University, State College, PA). K1- mutants were stable and reversion to their K1+ status has not been observed either after > 10 years of storage at -70°C or after > 10 passages in vitro and in vivo rats.

Rough mutants. Two different rough mutants of strain C5 were generated. The first type of rough mutant was obtained by treatment with *N*-methyl-*N*-nitro-*N*-nitrosoguanidine as described previously (19). Briefly, strain C5 grown overnight in 10 ml of fresh broth was centrifuged and resuspended in 0.5 ml of fresh broth. 0.1 ml of *N*-methyl-*N*-nitro-*N*-nitrosoguanidine (4.0 µg/ml) was then added. After 20 min of incubation at 37°C, 4.5 ml of additional broth was added and incubation was continued for another 5 hs, at which time the bacterial suspension was diluted and plated on agar. Most isolates that survived this treatment were found to retain the K1+ phenotype. Potential mutants that produced rough colony appearance were selected and confirmed as producing rough LPS by SDS-PAGE and sensitivity to rough-specific bacteriophages (19, 20, 22). These rough mutants were also found to have different outer membrane protein complex compared to the parent strain C5 (22).

The second type of rough mutants was prepared by mutagenesis with transposon Tn 10 (23, 24). *E. coli* auxotrophic donor C600, carrying the temperature-sensitive F₁₈ 114 lac plasmid into which the tetracycline (Tc)-resistant transposon Tn 10 had been inserted, was mated with strain C5 at 32°C. Selection for transconjugants that inherited F₁₈ 114 lac::Tn 10 was achieved by plating mating mixtures on minimal salts glucose agar supplemented with Tc (20 µg/ml). After incubation at 32°C for 48 h, several Tc^R strain C5 transconjugants were isolated. One of these strain C5 derivatives harboring the F₁₈ 114 lac::Tn 10 plasmid was chosen for transposition mutagenesis. To select for transposition of Tn 10, this transconjugant was serially cultured at a restrictive temperature (42°C) and plated on Trypticase soy Agar (BBL Microbiology Systems, Cockeysville, MD) containing Tc (20 µg/ml). After incubation at 37°C overnight, Tc^R derivatives were examined for rough colonial morphology via a binocular microscope. Tentative rough mutants were picked and scored for their LPS phenotype via spot tests for sensitivity to rough specific bacteriophages. Unlike smooth strain C5 which is resistant to these phages, rough mutants were sensitive to phages BR2, Ffm, and BR60 (18–20).

Two rough mutants (strain XYL derived from nitrosoguanidine mutagenesis and strain 2513 derived from transposon mutagenesis) chosen for animal studies were found to differ from the parent strain only at one locus when examined by the multilocus enzyme electrophoresis, which in each case was a null allele, reflecting a lack of enzyme activity at the loci (suggestive of isogenicity according to Dr. Richard Wilson, Pennsylvania State University).

Southern blot analysis of Tn 10 derived mutants. EcoRI- and HindIII-digested *E. coli* DNA was separated by electrophoresis through a 1.0% agarose gel in TBE buffer (0.05 M Tris base/0.05 M boric acid/0.001 M EDTA, pH 8.3). DNA was transferred to neutral nylon membrane by capillary action with 10× SSC (1.5 M NaCl/0.15 M Na citrate), and then fixed to the nylon membrane by exposure to 12,000 J/cm² UV. The filter was prehybridized with hybridization buffer (6× SSC, 5× Denhardt's solution, 10% dextran sulfate, 1% SDS, and 100 µg denatured Salmon sperm DNA) at 65°C for 1 h. The filter was probed with random primer labeled tetracycline resistance (Tc^R) gene DNA at 65°C, with 2× SSC, 0.5% SDS, 1× SSC, 1% SDS and then 0.1× SSC, 1% SDS. X-ray film was exposed overnight at -70°C with the nylon membrane.

The Tc^R gene DNA was isolated by polymerase chain reaction from *S. typhimurium* cya::Tn 10. Five microliters of a log-phase bacterial suspension was diluted into 45 µl of distilled H₂O, boiled for 4 min, and then rapidly cooled on ice. To 0.5 µl of target DNA Tc^R gene primers (100 pM 5'-GCTCTAATGCGTGT-3', 5'-GATCCAAGAGAAC-

CAAC-3'), 200 µM dATP, 200 µM dGTP, 200 µM dTTP, and 100 µM dCTP, and 1 U of AmpliTaq polymerase (Perkin-Elmer Cetus Corp., Emeryville, CA) were mixed in reaction buffer (25 mM Tris HCl, pH 9.3/50 mM KCl/2 mM MgCl₂/1 mM 2-mercaptoethanol). The Tc^R gene primers flank bp 1478–3098 of Tn 10 (23). This-1620 bp amplified contains a single EcoRI site 2,204 bp, and a single HindIII site 3,072 bp. The reaction mix was heated to 94°C in a DNA thermal cycler (Perkin-Elmer Cetus Corp.) for 5 min, then five cycles of reannealing at 40°C for 3 min, DNA synthesis at 72°C for 1 min, and denaturation at 94°C for 1 min. This was followed by 25 cycles of reannealing at 60°C for 2 min, DNA synthesis at 72°C for 1 min and denaturation at 94°C for 1 min. After DNA amplification, a single 2,600-bp product was detected by agarose gel electrophoresis.

Animal model for *E. coli* infection. *E. coli* bacteremia and meningitis were induced in 5-d-old rats by a previously described method (4). Briefly, outbred, pathogen-free, pregnant Sprague-Dawley rats with timed conception were purchased from Charles River Breeding Laboratories (Wilmington, MA); the rats delivered in our vivarium 5–7 d after arrival. Each adult rat and her pups (average litter size of 10 with range of 8–16) were housed in a solid-polypropylene opaque cage under a Small Animal Isolator (model 1894, Forma Scientific, Inc., Marietta, OH).

At 5 d of age, all members of each litter were randomly divided into three groups to receive subcutaneously 4–10× LD₅₀ of each *E. coli* strain. LD₅₀'s in 5-d-old-rats were 30 CFU for strain C5, 2.8 × 10⁶ CFU for the unencapsulated mutant and 5.0 × 10⁶ to 1.0 × 10⁷ CFU for the rough mutants (strains XYL and 2513). 24 h after inoculation, blood and CSF specimens were obtained as described previously (4, 12–17) for quantitative cultures. In blood and CSF specimens obtained from animals infected with strain 2513, agar containing Tc (20 µg/ml) was used for cultures. Immediately after obtaining blood and CSF specimens, the brains of selected animals were fixed in 10% formaldehyde and examined for the evidence of leptomeningitis. Bacteria were identified using Brown and Brenn stain.

We also examined whether or not the K1 capsule and 0-LPS were the key factors in the development of *E. coli* meningitis in adult rats (weighing ~ 300 g). Briefly, pathogen-free Sprague-Dawley adult rats received intravenously 4 × LD₅₀ of each *E. coli* strain. LD₅₀'s in adult rats were 5.2 × 10⁸ CFU for strain C5, 3.7 × 10⁹ CFU for the unencapsulated mutant and 5.1 × 10⁹ to 1.0 × 10¹⁰ CFU for the rough mutants (strains XYL and 2513). 12–24 h after inoculation, blood and CSF specimens were obtained for quantitative cultures.

To confirm that all isolates derived from animals maintained their status of unencapsulation and rough LPS, one colony each of blood and CSF isolates from each animal were retested for the K1 capsule with the K1-specific phages and the antiserum agar technique and for rough LPS with the rough specific bacteriophages.

Results

Studies in infant rats. Table I summarizes the prevalence of meningitis (defined as positive CSF cultures) in 5-d-old rats with varying degrees of bacteremia caused by strain C5 or its derivatives lacking the K1 capsule (018+K1-) or 0-LPS (018-K1+). Bacterial counts in blood 24 h after subcutaneous inoculation varied widely, ranging from 1 × 10³ CFU/ml to 2 × 10⁹ CFU/ml of blood. At this time, some animals were also found to have positive CSF cultures and bacterial counts in CSF varied from < 5 × 10³ (lower limit of detection) to 2 × 10⁹ CFU/ml of CSF. *E. coli* isolates from the blood and CSF of animals infected with 018+K1- or 018-K1+ were tested for their K1- or 018- status. All isolates maintained their status of unencapsulation or rough LPS, respectively, and reversion to the status of their parent strain (018+K1+) did not occur in vivo.

Table 1. Development of Meningitis (Defined as Positive CSF Culture) in Newborn Rats with Varying Degrees of Bacteremia

Bacteremia CFU/ml of blood	Number of animals with positive CSF culture/number of animals with positive blood culture			
	Strain C5* (018+K1+)	C5 mutant* (018+K1-)	Strain XYL* (018-K1+)	Strain 2513* (018-K1+::Tn 10)
< 10 ⁴	0/9	0/8	0/11	0/5
10 ⁴ to < 10 ⁵	2/20 (10%)	0/22	0/8	1/7 (14%)
10 ⁵ to < 10 ⁶	10/25 (40%)	0/19	5/16 (31%)	3/15 (20%)
10 ⁶ to < 10 ⁷	4/10 (40%)	0/10	7/15 (46%)	5/18 (28%)
> 10 ⁷	19/23 (83%)	0/9	5/6 (83%)	8/13 (62%)

* See Methods for description of strains.

The isolation of *E. coli* from CSF was observed in animals infected with K1+ strains and also developing a high degree of bacteremia (e.g., > 10⁴ CFU/ml of blood). Overall 35/78 (45%) of animals infected with strain C5 (018+K1+), 17/45 (38%) of animals infected with strain XYL (018-K1+) and 17/53 (32%) of animals infected with strain 2513 (018-K1+::Tn 10) and also developing bacteremia of > 10⁴ CFU/ml of blood were found to have positive CSF cultures. For the unencapsulated mutant (018+K1-), 60 animals developed a similar level of bacteremia (i.e., > 10⁴ CFU/ml of blood); however, none had positive CSF cultures (*P* < 0.001 compared to 018+K1+ or 018-K1+ strain).

We next examined the brains of selective animals with positive CSF cultures. Fig. 1 was derived from a 5-d-old rat infected with strain C5 (018+K1+), which was found to have bacterial counts of 4 × 10⁷ CFU/ml of CSF, and Fig. 2 from an animal infected with strain 2513 (018-K1+::Tn 10) and possessing CSF bacterial counts of 5 × 10⁷ CFU/ml. As shown, the histopathological examinations of the brains of such rats with positive CSF cultures revealed the presence of gram-negative rods in the subarachnoid space predominantly around the perivascular areas (Figs. 1 and 2), but not in the chroid plexus (Fig. 3). Of interest, a concomitant response of inflammatory cells (e.g., PMN, macrophages) was lacking at this stage of *E. coli* meningitis (Figs. 1 and 2).

Studies in adult rats. Fig. 4 illustrates the development meningitis (defined as positive CSF cultures) in adult animals with varying degrees of bacteremia caused by different *E. coli* strains. The isolation of *E. coli* from CSF was again observed in animals developing a high degree of bacteremia caused by K1+ strains regardless of LPS phenotype; 3/5 (60%) of adult rats infected with strain C5 (018+K1+) and 5/10 (50%) of adult animals infected with strain XYL (018-K1+) and also developing bacteremia > 10⁴ CFU/ml of blood were found to have positive CSF cultures. In contrast, none of the 10 animals infected with 018+K1- and developing bacteremia > 10⁴ CFU/ml of blood were found to have positive CSF cultures (*P* < 0.05 compared to 018+K1+ or 018-K1+ strain). As shown in studies with newborn rats, *E. coli* isolates obtained from adult animals infected with 018+K1- or 018-K1+ mutants were also tested and no reversion to the status of their parent strain was found.

Copy number of Tn 10. One critical question for strain 2513 was that there might be multiple Tn 10 insertions affecting other virulence factors and thereby allowing strain 2513 to

enter the CNS in the presence of a high degree of bacteremia. The presence of Tn10 was detected in *E. coli* strain 2513 (018-K1+::Tn10) by PCR amplification of the Tn 10 Tc^R gene. A 1.6-kb DNA fragment was amplified from the Tn 10 mutagenized *E. coli* strain 2513 (018-K1+::Tn 10) and from *S. typhimurium* strain ×3604 (cya::Tn 10) but not from *E. coli* parent strain C5 (018+K1+). Because Tc^R gene primers amplify an internal 1.6-kb fragment within Tn 10, one cannot determine if *E. coli* strain 2513 (018-K1+::Tn10) contained one or more copies of Tn 10. To address this question, genomic DNA digested with restriction endonucleases EcoR1, EcoR1/Pst1, Bam H1, Hind III, and Bgl 2 (lanes 2-6; Fig. 5) was probed with Tc^R DNA amplified from *S. typhimurium* strain ×3604 (cya::Tn10). The Tc^R probe detected the predicted internal 2.6-kb Bgl 2 DNA fragment (lane 6; Fig. 5) and 4.9-kb Hind III DNA fragment (lane 5; Fig. 5). Tn 10 contained a single Bam H1 site located to the left of Tc resistance gene and a single EcoR1 site spanned by the Tc^R probe (Fig. 5). Only two EcoR1 (lane 2, 15.1 and 5.6 kb; Fig. 5) and one Bam H1 (lane 4, 8.9 kb; Fig. 5) fragments hybridized to the Tc^R probe. A Pst 1 site was mapped to the 15.1-kb EcoR1 fragment (lane 3, 7.8 kb; Fig. 5) by digesting *E. coli* 2513 (018-K1+::Tn 10) with both EcoR1 and Pst 1. These results showed that *E. coli* strain 2513 (018-K1+::Tn 10) contained a single Tn 10 insertion. *E. coli* parent strain C5 (018+K1+) did not hybridize with the Tc^R probe.

Discussion

Inadequate knowledge of the pathogenesis and pathophysiology has contributed to the significant mortality and morbidity associated with neonatal bacterial meningitis. For example, it is not clear why *E. coli* strains possessing the K1 capsule are predominant among isolates from neonatal *E. coli* meningitis and also why most of these K1 isolates are clonally associated with a limited number of O types.

We and others have shown the importance of both K and O surface determinants in conferring serum resistance (21, 22, 25), which is the essence of bacteremia. In contrast, for the development of meningitis, a sequelae of *E. coli* bacteremia, the K1 capsule, but not the LPS phenotype, appears to be the critical determinant. This is true not only with K1+ *E. coli*, where strains with rough LPS can induce meningitis (11), but also with group B meningococcus, where the LPS is typically of a rough phenotype (26). Thus, the O serogroups that are associated with the K1 encapsulated *E. coli* strains causing meningitis in neonates (e.g., 018, 07, 01, 016) may be relevant to the pathogenesis of bacteremic stage of disease, but not to the meningitis. K1 strains possessing certain O types may resist clearance from the bloodstream, thus allowing replication to a level that is required for meningeal invasion. Several studies in humans and experimental animals suggest a relationship between the magnitude of bacteremia and the development of meningitis (27-31). Dietzman et al. (30) reported a higher incidence of *E. coli* meningitis in neonates who had bacterial counts in blood > 10³ CFU/ml (6/11 or 60%) compared to those bacterial counts in blood < 10³ CFU/ml (1/19 or 5%). As shown in the present study, a high degree of bacteremia was a primary determinant for meningeal invasion by K1 *E. coli* strains regardless of status of LPS phenotype (smooth or rough). These findings suggest that one of the reasons for the close association

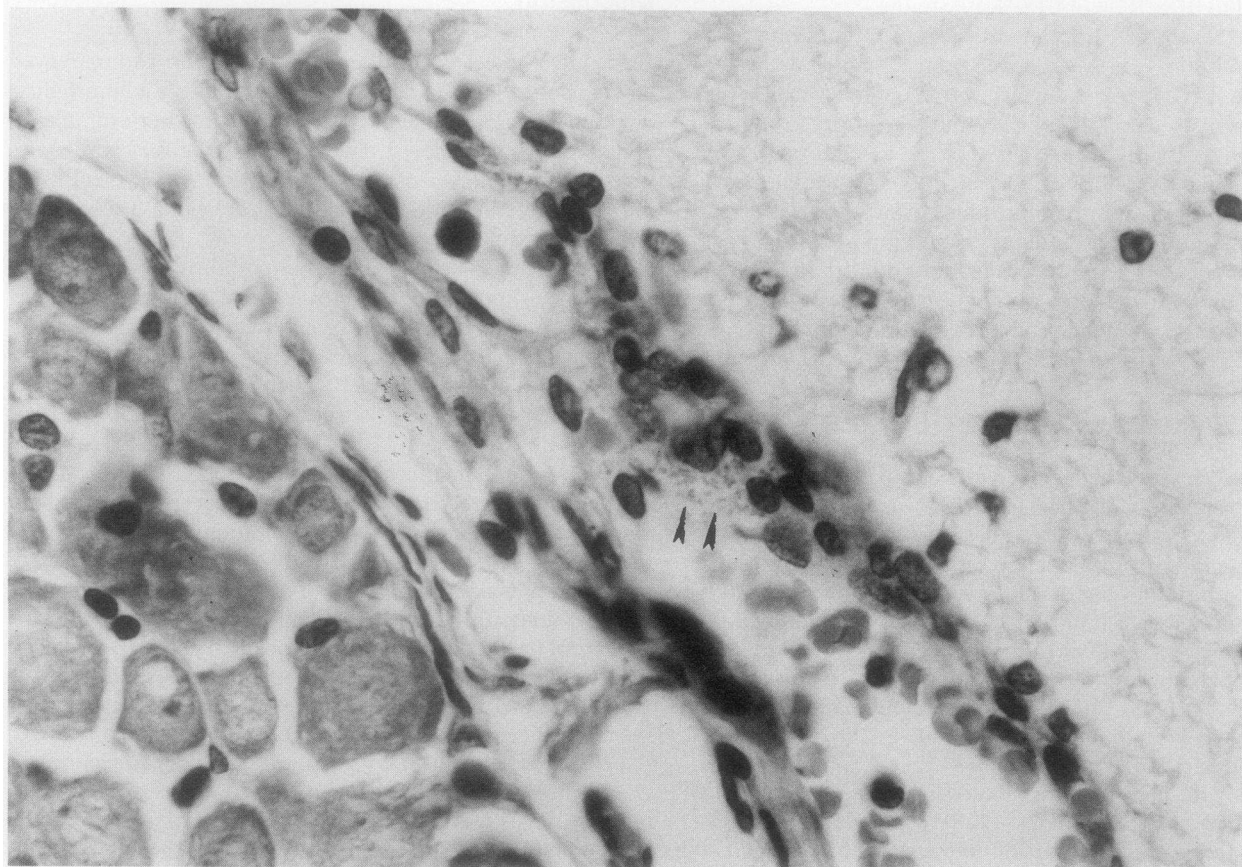


Figure 1. (Top) Coronal section of the base of the brain at frontal level with original magnification of 30 from a 5-d-old rat infected with E. coli strain C5 (O18+K1+) and was found to have meningitis (defined as positive CSF culture). (Bottom) 630 magnification of the circled portion of upper panel shows the presence of bacteria (arrows) without any evidence of inflammatory cells in the subarachnoid space.

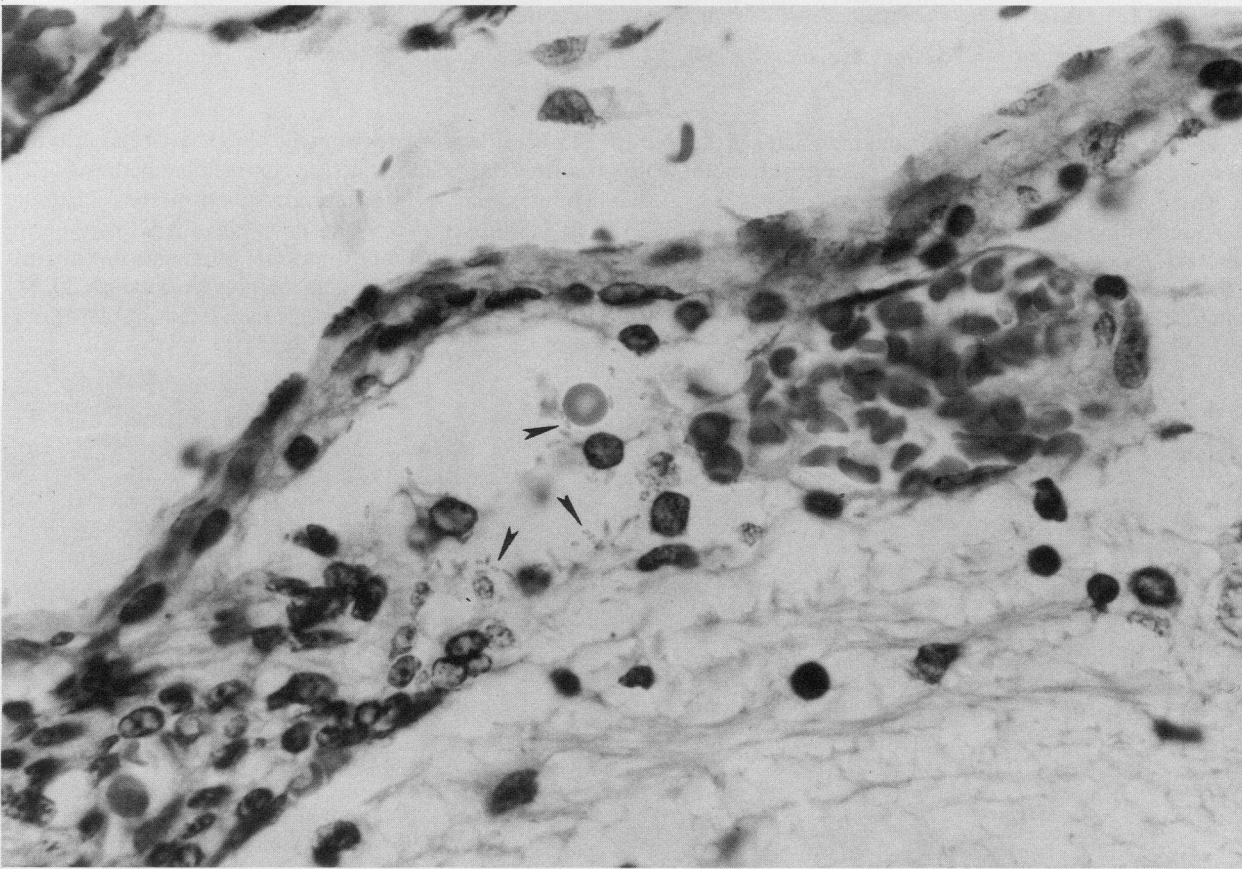
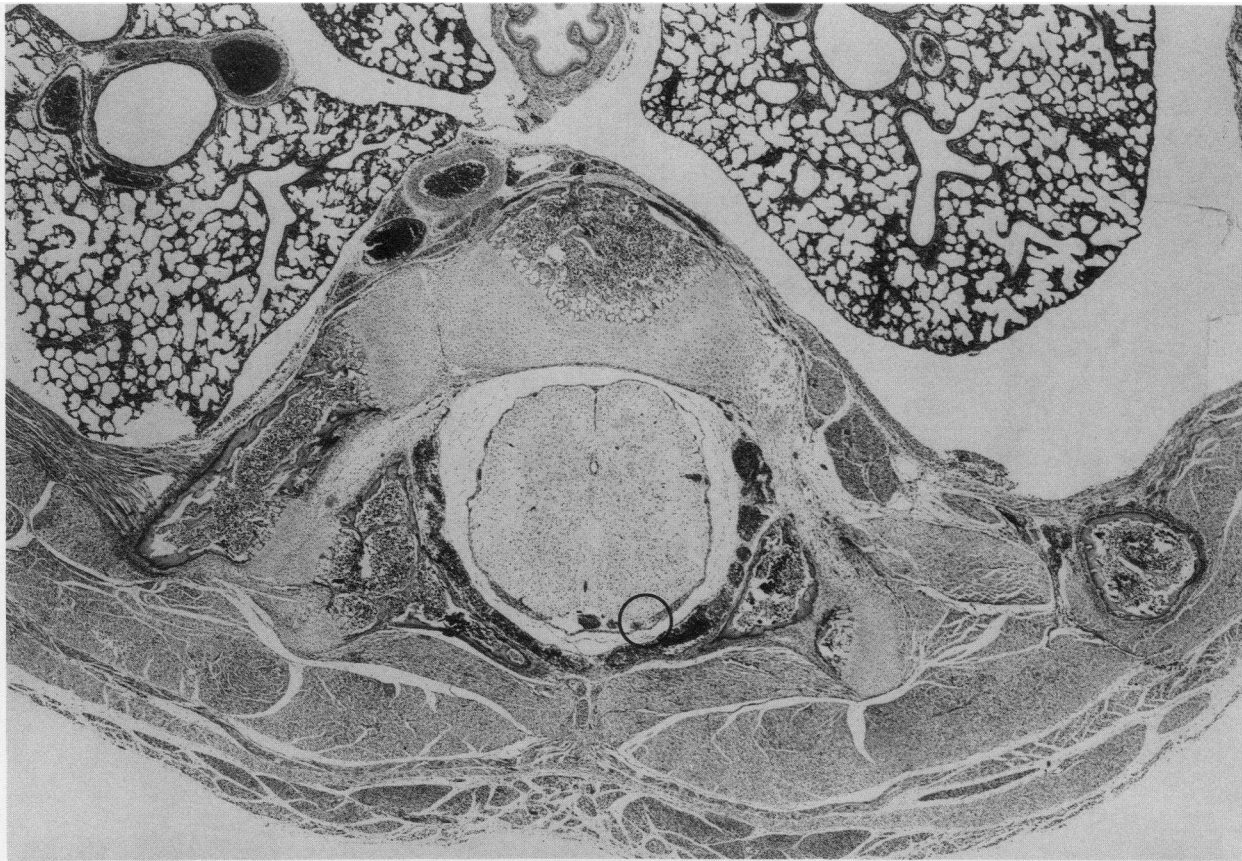


Figure 2. (Top) Coronal section of spinal cord at thoracic level with original magnification of 16 from a 5-d-old rat infected with a rough mutant of strain C5 (018-K1+) and also was found to have meningitis (defined as positive CSF culture). (Bottom) 630 magnification of the circled portion of upper panel shows the presence of bacteria (arrows) without inflammatory cells in the subarachnoid space.

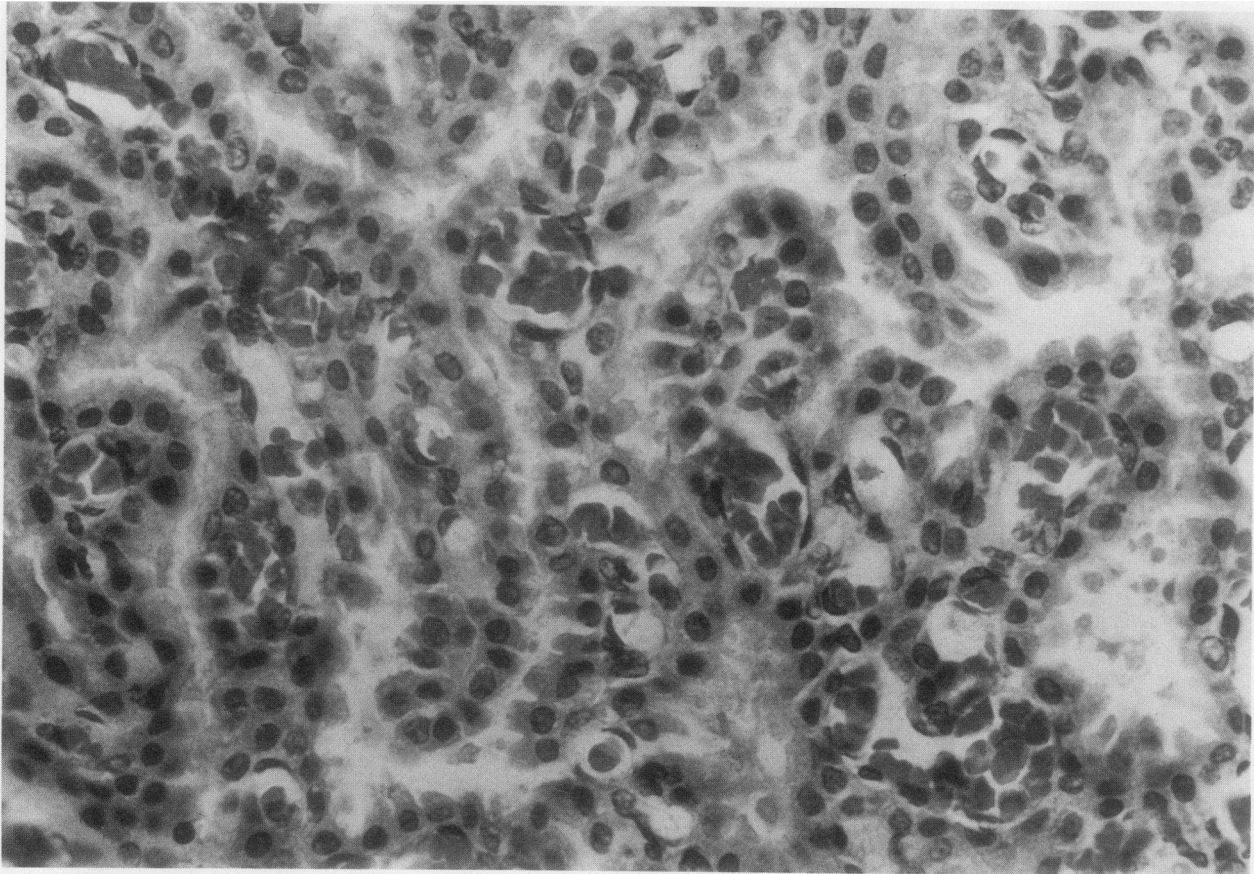


Figure 3. Choroid plexus of lateral ventricle from the same rat as in fig. 1 (430 magnification) shows no bacteria.

of certain K1 *E. coli* serotypes with neonatal meningitis is the ability to escape from host defenses and to achieve a threshold level of bacteremia necessary for meningeal invasion.

Previous studies to correlate the association of certain microbial factors of *E. coli* with the ability to replicate in vivo are not clear. For example, K1 capsular polysaccharides have been shown to increase the serum resistance of rough *E. coli* strains

(18–22). K1 *E. coli* isolates from the blood have been shown to be opsonized only via the classical complement pathway (32), but in humans there is little natural antibody to the K1 capsule (33). In addition, *E. coli* strains possessing the K1 capsule are considerably more resistant to normal human serum in vitro (20, 34) and as shown here and in our previous studies (20, 34) were more virulent in newborn rats than unencapsulated mu-

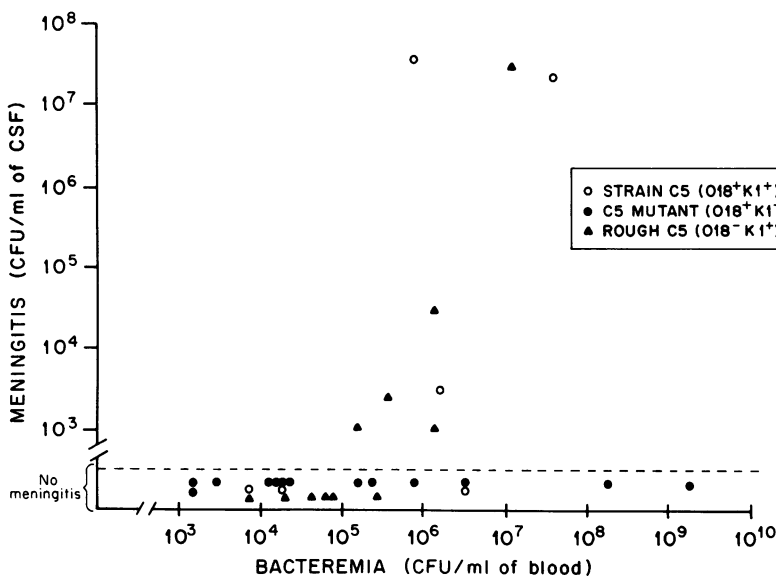


Figure 4. Relationship between the magnitude of bacteremia and the development of meningitis (defined as positive CSF culture) among adult rats developing varying degrees of bacteremia caused by *E. coli* strain C5 (O18+K1+), its unencapsulated (O18+K1-) or rough mutants (O18-K1+).

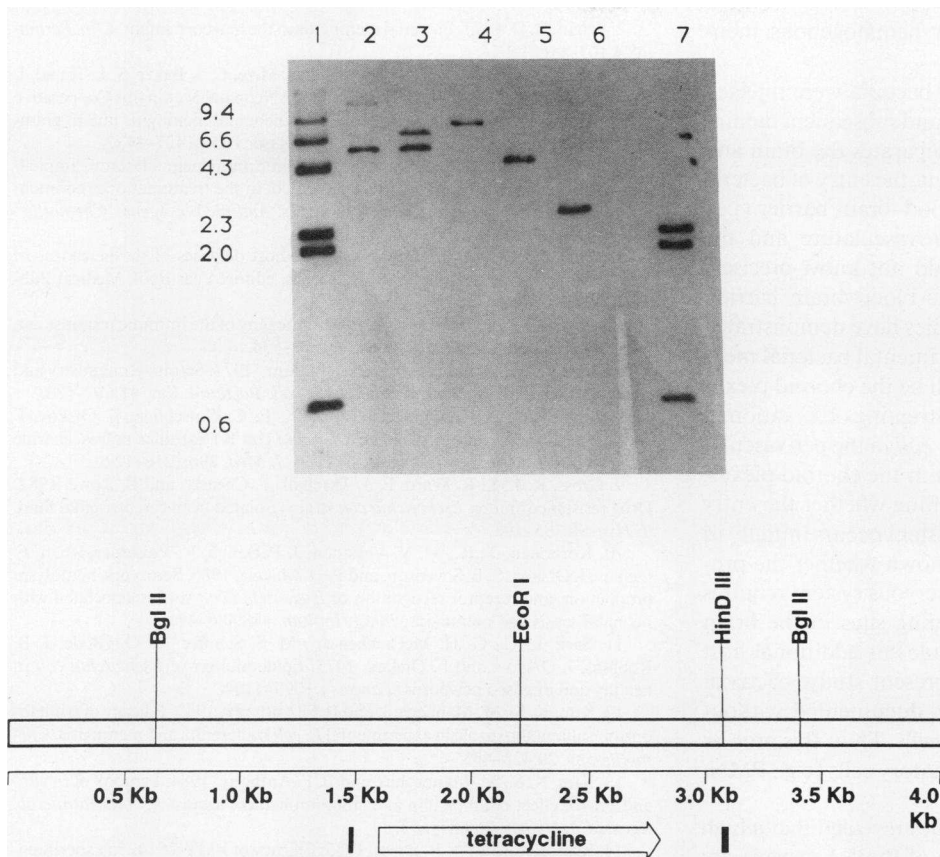


Figure 5. Hybridization of genomic DNA isolated from *E. coli* strain 2513 (018-K1+::Tn 10) with Tc^R probe. Genomic DNA was digested with restriction endonucleases EcoRI, EcoRI/PstI, BamHI, Hind III, and Bgl II, separated by agarose gel electrophoresis, and blotted to nylon membranes. Fluorescein-11 dUTP (Amersham International, Bucks, UK) labeled Tc^R probe DNA hybridized to the nylon membrane was detected with phosphatase-labeled anti-fluorescein antibody and luminol substrate (Amersham International). Lanes 1 and 7, Hind III digested Lambda molecular weight standards. Lane 2, EcoRI; lane 3, EcoRI/PstI; lane 4, BamHI; lane 5, Hind III, and lane 6, Bgl II. Restriction endonuclease map of tetracycline resistance gene of Tn 10 base on DNA sequence (23). The black boxes represent the PCR primers used to amplify the Tc^R probe DNA.

tants. This suggests that for the demonstration of serum resistance and in vivo virulence the K1 capsule is required. In contrast, other investigators suggest that other microbial surface factors may also affect virulence. Smooth LPS antigen (rather than K antigen) has been shown to be a virulence factor for serum resistance of *E. coli* (35, 36). The concept that some smooth LPS types can function as virulence determinants, however, does not apply to all instances of meningitis and therefore poses a paradox. For instance, ~20% of K1 *E. coli* isolates from neonatal meningitis spontaneously agglutinate in saline, suggesting rough LPS phenotype (11). In the present study, we obtained two different rough mutants. Although these rough mutants were considerably less virulent compared to the parent smooth LPS strain (i.e., ~10³-fold greater LD₅₀'s compared to the parent strain), they were able to induce meningitis in the presence of a high degree of bacteremia provided the K1 capsule was also present (Table I). These findings suggest that a smooth LPS phenotype is not required for meningeal invasion by K1-encapsulated *E. coli* strains. Thus, one of the reasons for the close association of certain O serotypes or rough LPS of K1 *E. coli* with neonatal meningitis may be that these LPS phenotypes of *E. coli* have evolved with the K1 capsule for protection from serum bacteriolysis. This concept is based on the previous demonstration that the K1 capsule is associated with a relatively limited number of O serotypes, particularly those LPS phenotypes that are unable to protect the bacteria from killing by normal human sera in the absence of capsule (21, 22, 37).

One intriguing observation was that the induction of a high degree of bacteremia alone was not sufficient for the development of meningitis by *E. coli*. This was demonstrated by our

inability to induce meningitis in newborn as well as in adult animals infected with the unencapsulated (K1-) mutant despite the production of a high degree of bacteremia (up to 10⁸ to 10⁹ CFU/ml of blood). These findings indicate that the presence of the K1 capsule is a critical element that is required in addition to the achievement of a high degree of bacteremia for the development of *E. coli* meningitis. The K1 capsule, which is a homopolymer of sialic acid, is immunochemically identical to group B meningococcal capsular polysaccharide (38). These bacteria, which are of diverse evolutionary background and have little genetic homology, share a common biochemical capsule and both lead to an infection of the same anatomic site, the meninges. This suggests some specific interaction of the capsule with host brain tissue, which may be responsible for the entry of K1-encapsulated *E. coli* into the meninges. The two rough mutants used in these studies, strains XYL and 2513, were derived by different methods (nitrosoguanidine and transposon mutagenesis). In that a great many genetic loci are involved in the synthesis of LPS (39), it is likely that these two different methods of mutagenesis altered different sites of LPS synthesis and consequently, strains XYL and 2513 were probably two different mutants. While these two mutants may differ in some characteristics not examined by us, nevertheless, both mutants did exhibit a rough LPS phenotype but retained the K1 capsule, were identical by multilocus enzyme electrophoresis and behaved similarly in animal studies. In addition, Southern blot analysis of strain 2513 revealed a single Tn10 insertion, suggesting that strain 2513 stems from a single lesion in the genesis of LPS without affecting other virulence factors. The data obtained with strain 2513 (Table I) further support the idea that the K1 capsule, not the LPS pheno-

type, is an important determinant for hematogenous meningitis.

In our experimental animal model, bacteria were injected subcutaneously, resulting in bacteremia and subsequent meningitis. Because the blood-brain barrier separates the brain and CSF from the intravascular compartment, the entry of bacteria should have occurred at the sites of blood-brain barrier (i.e., the choroid plexus, the cerebral microvasculature and the arachnoid membrane). We presently do not know precisely how K1 encapsulated *E. coli* cross the blood-brain barrier. Inasmuch as previous morphologic studies have demonstrated an intact arachnoid membrane in experimental bacterial meningitis (40, 41), the most likely sites will be the choroid plexus and/or the cerebral vasculature. One intriguing observation in this respect was the demonstration of *E. coli* in the perivascular area of the subarachnoid space, but not in the choroid plexus. Additional studies are needed to determine whether the entry of K1 *E. coli* into the central nervous system occurs initially in the cerebral vasculature. It is also unknown whether the process of bacterial entry into the central nervous system requires specific structure(s) (e.g., specific binding sites in the brain which have an affinity for the K1 capsule) or additional host factor(s). However, as shown in the present study, bacterial entry into the subarachnoid space was documented without concomitant presence of inflammatory cells. Thus, this process does not appear to require host inflammatory cells (e.g., PMN, macrophages).

In summary, our experimental studies revealed that a high degree of bacteremia and the presence of the K1 capsule are important determinants in the development of *E. coli* meningitis. Based on these findings, it appears that the prevention of bacterial multiplication in the blood that is required for bacterial entry into the central nervous system would be one potential approach to the prevention of *E. coli* meningitis. This approach may be feasible because administration of functionally active antibodies alone or in combination with antibiotics have been shown to decrease the level of bacteremia (42-45). In addition, the number of O antigens common in neonatal *E. coli* meningitis are limited (11) and techniques are available for preparing nontoxic O polysaccharide conjugate vaccines (46) as well as hyperimmune globulin containing opsonic antibodies against several O antigens. By using hyperimmune globulin, a large quantity of functional anti-O LPS antibody can be administered in a comparatively small volume, thus possibly circumventing at least some of the adverse reactions noted upon giving large volumes of nonhyperimmune globulin preparations (47). Further investigations on the role of immunotherapy (e.g., hyperimmune globulin) for controlling neonatal *E. coli* sepsis and meningitis are therefore warranted.

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