# Disparate Findings on the Role of Virulence Factors of Enterococcus faecalis in Mouse and Rat Models of Peritonitis

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Received 3 December 1997/Returned for modification 5 January 1998/Accepted 27 March 1998

The role of Enterococcus faecalis in polymicrobial peritonitis is still debated. Virulence factors expressed in some enterococcal strains might be involved in the pathogenicity of these organisms. To clarify their role, three of these virulence factors (cytolysin, gelatinase, and aggregation substance) were studied in six isogenic strains of E. faecalis expressing various combinations of these factors. Since the pathogenic effects of enterococci are only moderate, the expression of their virulence might vary from one animal species to another and from one type of infection to another. Therefore, we evaluated these effects in two animal models, i.e., a systemic infection in mice in which we assessed the virulence of the strains in 50% lethal dose studies and a model of compartmentalized infection in rats in which the microbiologic and inflammatory effects of the strains were evaluated in monomicrobial or polymicrobial infection. In mice, significant differences were observed in the cumulative survival curves depending on the virulence factors (P < 0.0001 [log rank test]). In rats, monomicrobial infection induced only mild changes. In polymicrobial peritonitis, the virulence factors mainly increased the inflammatory response while the changes observed in the microbiologic response were minimal. The combination of two virulence factors did not significantly increase the severity of infection either in the mice model or the polymicrobial rat model. These data argue for species and model dependence of the role of the virulence factors studied here and suggest that other important factors may be involved in the pathogenicity of enterococci.

Many issues remain unsolved regarding the pathogenic role of enterococci in the course of intraabdominal infection. From the previous reports, it appears that the enterococcal infections are rarely monomicrobial in nature, especially in surgical patients, suggesting the role of bacterial synergy (30, 31). Among the factors that might be involved, the virulence factors could be of interest but have been minimally studied (23, 25). Plasmids and conjugative transposons, which play a key role in the acquisition of drug resistance, may also carry these virulence traits (6), but their clinical relevance remains unclear.

Three virulence factors of Enterococcus faecalis might have the potential to increase the severity of intraabdominal sepsis. The plasmid-encoded cytolysin-bacteriocin (Cly) is the best studied of these factors. In vitro, this factor induces lysis of erythrocytes, polymorphonuclear neutrophils, and macrophages and could lead to a reduction in phagocytosis (29). Aggregation substance (Agg), which is closely linked to mating response to enterococcal sex pheromones (7), mediates adhesion of the bacteria to host cells, such as intestinal epithelial cells (34), renal tubular cells (27), and heart endothelial cells (17), and could be involved in the persistence of the organisms within the host tissues and fluids. Gelatinase (Gel) is a nonplasmid-encoded potential virulence factor. Although Gel strains of E. faecalis have been mainly studied in dental diseases (16), the properties given by this extracellular metalloendopeptidase (hydrolyzis of collagen, gelatin, and small peptides [18]) could increase bacterial dissemination. The expression of these virulence factors might vary from one animal species to another and from one type of infection to another, especially because the pathogenic role of E. faecalis seems to be moderate (36, 40, 41) and expressed only in polymicrobial infections (3, 10, 28, 30).

To address these issues, we evaluated the effects of six isogenic strains of *E. faecalis* containing various combinations of virulence factors in two animal models, i.e., a model of systemic infection in mice in which we assessed the virulence of enterococcal strains in 50% lethal dose studies and a model of compartmentalized infection in rats in which the microbiologic and inflammatory effects of the strains were evaluated in monomicrobial or polymicrobial infection in combination with *Escherichia coli* and *Bacteroides fragilis*, two species frequently associated in clinical samples.

## MATERIALS AND METHODS

Microorganisms. Six derivatives of E. faecalis OG1 were obtained from the University of Texas Medical School (Houston, Tex.) and the University of Michigan (Ann Arbor, Mich.). The four OG1x derivatives used in the present study are isogenic strains produced by mutagenesis with nitrosoguanidine from strain OG10 (14, 39). The Tn917 transposon was inserted into the PAD1 plasmid, leading to the following different phenotypes of OG1x: pAM714 with Cly and Agg (20, 21) and pAM9058 with Agg and pAM944 with Cly (5, 15). The two remaining strains (OG1RF and OG1SSP) are not isogenic with the previous four strains. OG1RF is a spontaneous mutant of OG1 with a Gel phenotype (14, 32). The OG1SSP strain is a derivative of OG1. The Tn915 transposon was inserted into the PCF10 plasmid, leading to the Gel and Agg phenotypes of the OG1SSP strain (12, 13, 35). All of these strains of E. faecalis had an intrinsic low-level resistance to aminoglycosides, and their characteristics are listed in Table 1. The resistance phenotype, which was encoded on the same plasmid as the virulence factor, was verified in each experiment. Gel and Cly expression was verified in vitro before inoculation according to the method described by Coque et al. (8). The strains of B. fragilis (AIP5-86) and E. coli (CB1496) used in the current study have previously been used in this model (30).

Animals. Female OF1 mice (Iffa Credo, L'Arbresles, France), weighing 20 to 25 g and housed 10 per cage, were used for 50% lethal dose  $(LD_{50})$  evaluation of the six strains of *E. faecalis*, which were injected intraperitoneally. Male Sprague-Dawley rats (Charles River, St-Aubin-les-Elbeufs, France), weighing 250 to 300 g and housed 5 per cage, were used for the peritontis model. All animals had access to chow and water ad libitum throughout the experiment. All of these experiments were performed according to current European regulations.

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TABLE 1. Microbiologic characteristics of the derivatives of *E. faecalis* inoculated

Strain	Plasmid content	Virulence factor	Antimicrobial resistance <sup>a</sup>	
OG1X	None	None	None	
OG1RF	None	Gel	RI + FA	
OG1X	pAM 9058 (Tn917)	Agg	EM	
OG1X	pAM 944 (Tn917)	Cly	EM	
OG1X	pAM 714 (Tn917)	Cly + Agg	EM	
OG1SSP	pCF10 (Tn915)	Gel + Agg	TC	

<sup>*a*</sup> All of these strains of *E. faecalis* had an intrinsic low-level chromosomal resistance to aminoglycosides. OG1RF had a constitutive chromosomal resistance to rifampin and to fucidic acid. Resistance to erythromycin and tetracycline was plasmid encoded. RI, rifampin; FA, fusidic acid; EM, erythromycin; and TC, tetracycline.

**Preparation of the microorganisms.** *B. fragilis* was grown and diluted anaerobically in prereduced thioglycolate broth. *E. coli* and the six strains of *E. faecalis* were grown in brain-heart infusion broth. The final inoculum was made when the bacteria were in the log phase of growth. The inocula were adjusted spectrophotometrically and bacteria were diluted to give the number of microorganisms required for bacterial challenge. Purity was assessed and counts were validated for each strain immediately before inoculation (mice model) or before mixing (rat model).

 $LD_{50}$  in mice. The virulence of each *E. faecalis* strain was evaluated by inoculation of increasing concentrations of microorganisms (10<sup>7</sup> to 10<sup>11</sup> CFU per ml) in mice. Nine groups of 10 animals were studied for each bacterium. The animals received intraperitoneally a 0.5-ml injection of the bacterial suspension. After inoculation, the animals were returned to their cages and daily mortality was recorded until day 7. Lethal dose curves were plotted, and LD<sub>50</sub>s were calculated according to the method described by Ike et al. (21).

**Intraabdominal infection in rats.** Semisolid agar medium was prepared by adding 2% (wt/vol) agar to the diluted broth cultures associated with barium sulfate (10% [wt/vol]). Aliquots (0.5 ml) of the final product were placed in double gelatin capsules for peritoneal implantation. Each *E. faecalis* strain was studied twice: once as a monomicrobial infection (10<sup>8</sup> CFU/ml, 12 animals in each group) and once as a polymicrobial infection (20 animals in each group) associated with *E. coli* (10<sup>8</sup> CFU/ml) and *B. fragilis* (10<sup>9</sup> CFU/ml).

**Implantation of inoculum.** The rats were anesthetized with an intramuscular injection of ketamine (30 mg/kg of body weight [Parke-Davis, Courbevoie, France]), and the gelatin capsule was inserted into the pelvic peritoneal cavity through a midline abdominal incision (40). The wound was closed with a musculoperitoneal layer and a skin layer by using interrupted nylon sutures.

Assessment of spontaneous outcome. After implantation of the inoculum, the animals were returned to separate cages; they were observed and weighed daily until sacrifice. No death was observed within 6 h of capsule implantation. In the six groups of animals receiving a monomicrobial inoculum, sacrifice was performed at 24 h after inoculation for four animals and at day 3 for four animals. In the groups receiving a polymicrobial inoculum, eight animals were sacrificed at 24 h and eight animals were sacrificed at day 3. In addition, an early sacrifice (6 h after inoculation) was planned in both models (monomicrobial and polymicrobial) for four animals in each group. The following pathogenicity criteria were studied: clinical criteria (body weight and mortality), microbiological criteria (positivity of blood cultures and bacterial counts in the peritoneal fluid), and the inflammatory response (peritoneal concentrations of phagocytes, tumor necrosis factor alpha [TNF-a], and interleukin-6 [IL-6] and serum concentrations of al-acid glycoprotein).

Sacrifice. Animals were killed with chloroform. Blood samples were obtained by aseptic percutaneous transthoracic cardiac puncture for qualitative blood culture on days 1 and 3 and for measurements of a1-acid glycoprotein concentrations in serum on day 3. After injection of 10 ml of cold phosphate-buffered saline (PBS) intraperitoneally, a midline laparotomy was performed and peritoneal fluid samples were recovered from all regions of the peritoneal cavity for bacterial and cell counts (days 1 and 3). Peritoneal fluid cytokine concentrations were specifically evaluated at 6 h after bacterial challenge. A dilution factor taking into account the fluid present in the peritoneal cavity prior to the injection of the 10 ml of PBS was applied in all calculations according to a technique previously described (26, 31). Blood cultures were inoculated immediately after collection (NR-7A; Becton-Dickinson, Le-Pont-de-Claix, France) and analyzed daily until day 5 for identification. Serial dilutions of the peritoneal fluid were made, and 0.1 ml of each dilution was spread on agar plates for colony counts. The limit of detection for each microbiological test was <1 log<sub>10</sub> CFU/ml. In cases of values below this threshold, the results were listed as  $\leq 1 \log_{10} \text{CFU/ml}$ . In the statistical analysis, these results were treated as 1 log<sub>10</sub> CFU/ml. Plates were incubated under appropriate conditions (aerobic and anaerobic) for 2 to 5 days. The selective medium used for detection of B. fragilis was Columbia agar base (Bio-mérieux, Charbonnières-les-Bains, France) with 5% sheep blood containing 75  $\mu$ g of kanamycin, 7.5  $\mu$ g of vancomycin, and 4  $\mu$ g of pefloxacin per ml. The selective media used for aerobic culture were Drigalski agar (Diagnostics Pasteur, Marnes-la Coquette, France) and bile-esculine-azide agar (Diagnostics Pasteur). Mueller-Hinton agar (Diagnostics Pasteur) was used for *E. faecalis* counts in the monomicrobial model.

**Peritoneal cell counts.** Total cell counts (polymorphonuclear neutrophils and macrophages) were made on an aliquot of the original peritoneal fluid by using a Malassez counting chamber.

**Cytokine assay.** Four 1- ml samples of the original peritoneal fluid recovered from all of the regions of the peritoneal cavity were centrifuged (300 × g for 15 min) and then divided into 200- $\mu$ l aliquots and stored at  $-80^{\circ}$ C until assaying. The samples were assayed in duplicate. TNF- $\alpha$  activity was measured with a Factor-Test-X rat TNF- $\alpha$  enzyme-linked immunosorbent assay kit (Genzyme Diagnostics, Cambridge, Mass.) according to a previously described technique (37). The sensitivity of the test, which was defined as the lowest concentration of standard which shows greater absorbance than the mean absorbance of the 0-pg/ml sample  $\pm$  2 standard deviations (SD), was 10 pg/ml. IL-6 activity was measured by means of a bioassay with the murine hybridoma cell line B9 according to a previously described technique (1). The limit of detection in peritoneal fluid was 0.2 ng/ml. The specificity of the response for IL-6 was assessed by using polyclonal rabbit antimurine IL-6 antibodies (Genzyme, Brussels, Belgium).

α**1**-acid glycoprotein assay. Blood samples were obtained by cardiac puncture and transferred to sterile glass tubes. After coagulation, the sera were collected and centrifuged at 500 × g for 5 min. The supernatant was divided into four aliquots and stored at  $-20^{\circ}$ C until the assay. Radial immunodiffusion was used for the assay which involved agar containing 3% (wt/vol) immune serum and 3% (wt/vol) polyethylene glycol 6000 (Fluka, Malakov, France). The values of α1acid glycoprotein were determined with a monoclonal rabbit anti-rat α1-acid glycoprotein antibody (4). A value of less than 200 mg/liter was considered normal. The limit of detection was 25 mg/liter.

Statistical analysis. Results are expressed as means  $\pm$  SD. The LD<sub>50</sub>s in mice were compared by a Kaplan-Meier analysis by using a log rank test. Continuous parameters were compared by an analysis of variance analysis, followed (in the case of significance) by limited comparisons between the control group without virulence factor (*E. faecalis* OG1X) and the other groups by using Fisher's least significant procedures. A chi square test was used for quantitative data. A *P* value of  $\leq 0.05$  was considered significant.

## RESULTS

LD<sub>50</sub>s in mice. *E. faecalis* was classified into three groups according to the LD<sub>50</sub> (Table 2). The first group, which was characterized by the lowest LD<sub>50</sub>, included *E. faecalis* OG1X(pAM 714) and OG1X(pAM 944). An intermediate LD<sub>50</sub> was evidenced for *E. faecalis* OG1RF. Finally, the highest LD<sub>50</sub>s were observed with *E. faecalis* OG1X, OG1X(pAM 9058), and OG1SSP(pCF10). With a Kaplan-Meier model, a statistically significant difference was observed within the groups of animals in the delay before death (Fig. 1). Mice inoculated with *E. faecalis* OG1X(pAM 714), OG1X(pAM 944), and OG1RF strains had a significantly higher and earlier mortality than those receiving *E. faecalis* OG1X, OG1X(pAM 9058), and OG1SSP(pCF10), in which a low and delayed mortality was observed.

**Peritonitis model in rats. (i) Effect of inoculum on survival.** No mortality was observed in the rat peritonitis model.

(ii) Effect of *E. faecalis* strains on body weight. In the polymicrobial model, maximal loss of weight occurred in most groups

TABLE 2.  $LD_{50}s$  in mice according to the derivative of *E. faecalis* inoculated

Mouse group <sup>a</sup>	E. faecalis strain	Virulence factor	LD <sub>50</sub> <sup>b</sup>
1	OG1X	None	10.5
2	OG1RF	Gel	9.9
3	OG1X(pAM 9058)	Agg	10.5
4	OG1X(pAM 944)	Cly	9.4
5	OG1X(pAM 714)	Cly + Agg	9.6
6	OG1SSP(pCF10)	Gel + Agg	10.6

<sup>a</sup> Each group was composed of 90 mice.

<sup>b</sup> Expressed in log<sub>10</sub> CFU per milliliter.



FIG. 1. Cumulative survival curves of mice (expressed as percentages of survivors) after inoculation of  $\log_{10}$  CFU of each *E. faecalis* strain per ml \*, P < 0.0001 by a Kaplan-Meier analysis with a log rank test.

on day 1, except for the animals receiving the OG1X(pAM 9058) and OG1RF strains, in which maximal weight loss was reported on day 2 (Fig. 2). Progressive recovery was observed afterward. The greatest weight loss was observed in animals receiving the OG1X strain, while the least weight loss was seen with the OG1SSP(pCF10) strain. In the monomicrobial model, there was no difference in the variations of weight within the groups (data not shown). When comparing polymicrobial and monomicrobial models, a significant difference was observed in weight variations in the 3 days of the study (P < 0.01 in every case).

(iii) Effect of *E. faecalis* strains on blood cultures. All animals had positive blood cultures at day 1. A significant decrease in the frequency of positive blood cultures was noted between days 1 and 3 (Table 3). However, the different strains of *E. faecalis* did not modify the frequency of bacteremia of the other organisms. The frequency of bacteremia due to *E. faecalis* was more marked in the polymicrobial model than in the monomicrobial one; 75 of 90 rats (80%) had *E. faecalis*-positive blood cultures in the polymicrobial model versus 26 of 48 rats (55%) in the monomicrobial model, the different strains of *E. faecalis* had similar frequencies of positive culture (data not shown).

(iv) Effect of *E. faecalis* strains on peritoneal cultures. The bacterial titers within the peritoneal cavity between days 1 and 3 are displayed in Table 4. In the monomicrobial model, no difference was observed on day 1 or 3 or between days 1 and 3 in the different groups of animals. The peritoneal concentrations of *E. faecalis* were significantly higher on day 1 in the polymicrobial model than in the monomicrobial model in animals receiving *E. faecalis* OG1X, OG1X(pAM 9058), OG1X (pAM 714), and OG1SSP(pCF10) (data not shown). On day 3, only the animals inoculated with *E. faecalis* OG1X(pAM 9058) had significantly higher *E. faecalis* peritoneal concentrations in the polymicrobial model than in the monomicrobial model.

(v) Effect of *E. faecalis* strains on peritoneal cell counts. In the polymicrobial model, an increased peritoneal cell count was observed between days 1 and 3 for all groups (Fig. 3), while an increase was noted only for the OG1SSP(pCF10) strain in the monomicrobial model. At day 1, the peritoneal cell counts were similar in each group in the polymicrobial and monomicrobial models, except for the group of animals receiving the OG1SSP(pCF10) strain, in which an increased cell count was noted in the polymicrobial model compared to the monomicrobial inoculum (P < 0.05). On day 3 in the polymicrobial model, the peritoneal cell count was significantly increased in



FIG. 2. Changes in body weight expressed as percentages of baseline (means  $\pm$  SD) in rats receiving one of the strains of *E. faecalis* combined with *E. coli* and *B. fragilis.* \*, *P* < 0.01 compared with changes in the animals receiving the OG1X strain; †, *P* < 0.01 compared with changes in the animals receiving the OG1X strain.

animals receiving the OG1X(pAM 944) strain compared to the other groups (P < 0.01). Moreover, increased peritoneal cell counts were noted at day 3 in the polymicrobial model compared to the monomicrobial model in the groups receiving the OG1X(pAM 9058), OG1X(pAM 944), and OG1SSP(pCF10) strains (P < 0.05, P < 0.01, and P < 0.05, respectively).

(vi) Effect of *E. faecalis* strains on cytokine levels. The animals inoculated with *E. faecalis* OG1X(pAM 714) and OG1RF had higher concentrations of TNF- $\alpha$  in the peritoneal fluid than those receiving *E. faecalis* OG1X (P < 0.05, P < 0.05) and OG1X(pAM 944) (P < 0.05, P < 0.05) (Fig. 4). The animals receiving *E. faecalis* OG1RF had higher peritoneal concentrations of IL-6 than those inoculated with *E. faecalis* OG1X (P < 0.01), OG1X(pAM 944) (P < 0.001), OG1X(pAM 9058) (P < 0.05), and OG1SSP(pCF10) (P < 0.05). Moreover, animals given *E. faecalis* OG1X(pAM 714) had higher concentrations of IL-6 in the peritoneal fluid than those receiving *E. faecalis* OG1X(pAM 944) (P < 0.05).

(vii) Effect of *E. faecalis* strains on  $\alpha$ 1-acid glycoprotein concentrations. The concentrations of  $\alpha$ 1-acid glycoprotein in plasma were significantly higher in animals receiving a polymicrobial inoculum than in those receiving *E. faecalis* alone (*P* < 0.01), except for animals inoculated with *E. faecalis* OG1X(pAM 9058) and OG1X(pAM 714) (Fig. 5). In the

TABLE 3. Numbers of infected animals with positive blood cultures at sacrifice in the polymicrobial model according to the bacterium inoculated

Bacterium	No. of infected animals positive (% of total)		
	Day 1	Day 3	
E. coli	45 (98)	35 (73) <sup>c</sup>	
B. fragilis	48 (100)	$39(81)^c$	
E. faecalis <sup>a</sup>	57 (81)	$44(62)^{b}$	

<sup>a</sup> Including all of the strains of E. faecalis.

 $^{b}P < 0.05$  compared to day 1.

 $^{c}P < 0.01$  compared to day 1.

Group <sup>b</sup>	Titer					
	Day 1 study <sup>a</sup>			Day 3 study <sup>a</sup>		
	E. coli	B. fragilis	E. faecalis	E. coli	B. fragilis	E. faecalis
OG1X	$6.5 \pm 1.6$	5.8 ± 1.3	$5.6 \pm 0.5$	$4.1 \pm 1.7^{c}$	$2.8 \pm 1.8^{c}$	$4.7 \pm 0.9^{c}$
OG1RF	$4.1 \pm 0.7$	$5.2 \pm 0.5$	$4.5 \pm 0.9$	$3.8 \pm 0.9$	$4.8 \pm 1.4$	$4 \pm 0.5$
OG1X(pAM 9058)	$4.9 \pm 1.3$	$5.5 \pm 0.6$	$4.7 \pm 0.7$	$3.6 \pm 0.8^{\circ}$	$4.5 \pm 1.3$	$4 \pm 0.5^{c}$
OG1X(pAM 944)	$4.6 \pm 1.1$	$5.8 \pm 0.6$	$4.5 \pm 1.1$	$3.5 \pm 0.9^{c}$	$4.8 \pm 1.2^{d}$	$4 \pm 0.6$
OG1X(pAM 714)	$6.4 \pm 1.8$	$6.2 \pm 1.3$	$5.2 \pm 0.4$	$3.5 \pm 1.9^{d}$	$4.6 \pm 1.7$	$3.6 \pm 1.2^{d}$
OG1SSP(pCF10)	$4.9 \pm 0.8$	$5.5 \pm 0.8$	$4.5 \pm 0.8$	$3.7 \pm 0.6^d$	$4.7 \pm 1.5$	$3.9\pm0.5$

TABLE 4. Bacterial titers within the peritoneal cavity at sacrifice in animals receiving a polymicrobial inoculum

<sup>*a*</sup> Bacterial titers are expressed in  $\log_{10}$  CFU per milliliter  $\pm$  SD.

<sup>b</sup> Each group was composed of eight rats on day 1 and eight rats on day 3.

 $^{c}P < 0.05$  versus day 1 study for the same bacterium.

 $^{d}P < 0.01$  versus day 1 study for the same bacterium.

polymicrobial model, only the animals given *E. faecalis* OG1SSP(pCF10) had higher  $\alpha$ 1-acid glycoprotein concentrations than the other groups (P < 0.05). There was no difference within groups in the concentrations of  $\alpha$ 1-acid glycoprotein in plasma in the monomicrobial model.

## DISCUSSION

The literature on the pathogenic role of E. faecalis strains is somewhat unclear. Their potential pathogenic effects are quite subtle and are not evidenced in nondiscriminative models of infection. In addition, the results of an experimental model of enterococcal infection seem to differ from one model to another, as illustrated by our results.

The direct injection of microorganisms within the peritoneum represents a model of systemic sepsis induced within a very short period of time (9, 33). This model allows a good approach to identify quickly the most pathogenic strains. However, this experimental design may not resemble the development of infection in patients, which commonly occurs over days. Therefore, the information provided by the peritoneal implantation of a septic capsule is of interest. This model, which represents a compartmentalized sepsis with prolonged infectious and inflammatory responses, allows study over a prolonged period of time of the relationships between the host and the offending organisms and among the various organisms inoculated. Taking these features into consideration, we chose these two models in order to study different aspects of the role of virulence factors of enterococci.

Several experimental investigations, mainly performed with rats, have demonstrated the synergistic role of enterococci (2, 11, 28, 30). In these studies, the enterococcal strains did not express any known virulence factor. Nevertheless, increased bacteremia (10, 30) or increased mortality (2, 10) was reported when enterococcus was part of the inoculum. Our current results confirm these observations of bacterial synergy. Moreover, we demonstrated that the pathogenicity of *E. faecalis* in rats is minimal when inoculated alone, an issue frequently suggested in clinical studies but rarely assessed. All of the parameters that we tested were in agreement in demonstrating



FIG. 3. Peritoneal cell counts expressed in cells per cubic millimeter (means  $\pm$  SD) measured at days 1 and 3, according to the strain of *E. faecalis* inoculated in rats receiving a polymicrobial (P) inoculum (one of the six strains of *E. faecalis* plus *B. coli* plus *B. fragilis*) or a monomicrobial (M) inoculum (one of the six strains of *E. faecalis*). \*, P < 0.05 changes observed between days 1 and 3;  $\dagger$ , P < 0.01 between days 1 and 3.



FIG. 4. Concentrations of TNF- $\alpha$  (A) and IL-6 (B) within the peritoneal fluid expressed in nanograms per milliliter (means ± SD) 6 h after inoculation of a polymicrobial inoculum (one of the six strains of *E. faecalis* plus *E. coli* plus *B. fragilis*). \*, *P* < 0.05 compared to *E. faecalis* OG1X and OG1X(pAM 944) strains; †, *P* < 0.05 compared to *E. faecalis* OG1X and OG1X(pAM 944) strains; 0.05 compared to *E. faecalis* OG1X, OG1X(pAM 944), Strains; \$, *P* < 0.05 compared to *E. faecalis* OG1X, pAM 944), OG1SP(pCF10) strains; \$, *P* < 0.05 compared to *E. faecalis* OG1X, OG1X(pAM 945), and OG1SP(pCF10) strains; \$, *P* < 0.05 compared to *E. faecalis* OG1X and OG1X(pAM 945), OG1X(pAM 944).



FIG. 5. Concentrations of  $\alpha$ 1-acid glycoprotein in serum expressed in milligrams per liter (means ± SD) measured at day 3 after inoculation of one of the six isogenic strains of *E. faecalis* according to whether the strain of *E. faecalis* was inoculated alone (monomicrobial) or combined in a polymicrobial inoculum. \*, P < 0.05 compared to *E. coli* and *B. fragilis* (polymicrobial); †, P < 0.01 compared to polymicrobial inoculum.

that polymicrobial infection and monomicrobial infection significantly differed. On the other hand, there was divergence between the parameters according to the strain tested in the rat polymicrobial model.

Our observations with the mouse model poorly predicted the effects of enterococci in rats. To the best of our knowledge, the virulence of a microorganism is rarely assessed in two different models in the same study, raising difficulties in the interpretation of the results of previous studies. The rat model has been largely used to evaluate the mechanisms of bacterial synergy (2, 11, 28, 30), and the information obtained seems to be clinically relevant (40, 41). Since we have previously demonstrated that enterococcal pathogenicity is expressed in a dosedependent fashion (30, 31), the results reported with the rat model could be related to an insufficient concentration of enterococci in the inoculum. In addition, virulence factors might not be expressed in rats or might not be important in the model. Only few studies assessing the virulence of enterococci are available, and these are always performed with monomicrobial infections in mouse and rabbit models (5, 21, 24). It is not possible from the data available to determine whether virulence factors are expressed in intraabdominal infections or at what level and whether this expression is similar in poly- and monomicrobial infections.

In the mouse model, the highest mortality was observed for animals given the strain with the cytolysin factor. This is in agreement with the results obtained by Ike et al. (21), who reported that inoculation of a Cly-producing strain of *E. faecalis* caused an increased mortality and a 90% decrease in LD<sub>50</sub> compared to a control strain. Moreover, in previous studies performed with models of endophthalmitis (24, 38) or endocarditis, disease severity was markedly increased after inoculation of an enterococcal strain producing Cly (5). In clinical practice, the Cly phenotype, although difficult to detect, seems to be common, as reported by Ike et al. (22), who tested 97 clinical isolates of *E. faecalis* and showed that 60% were hemolytic. The data reported by Huycke et al. also suggested the pathogenicity of Cly in cases of enterococcal bacteremia in which a fivefold increased risk of death was observed in paIn our models of the mouse and rat, Agg did not seem to exert a significant virulence. In the mouse model, the mortality level induced by strains mating Agg and no Cly was low and delayed. The effects of adherence and persistent infection, which were linked to the properties of Agg (17, 27, 34), were not observed in our rat model, suggesting that Agg has only a minor role to play in the severity of peritoneal infection. Agg is the most difficult factor to assess in vivo, since there is no clinical marker of its activity. Only a molecular approach could demonstrate its activation, as previously performed by other authors (8).

Gelatinase is probably the least-studied virulence factor. In the mouse model, this factor induced an increased mortality compared to the control strain OG1X, as well as a 68% decrease of the LD<sub>50</sub>. In the rat model, the Gel strain seemed to be the least pathogenic, with only moderate weight loss and a mild peritoneal cellular reaction. The pathogenicity of Gel strains of *E. faecalis* in intraabdominal infections is purely hypothetical and has never been demonstrated. However, since a relationship between Gel and gentamicin resistance has been reported in one study (8), therapeutic difficulties could be expected in the presence of this factor.

The effect obtained with the OG1X strain (control strain) in the polymicrobial model is unclear. Significant and prolonged weight loss combined with an increased frequency of positive blood cultures was noted in animals receiving this strain. Reversion of this mutant to a wild expression of gelatinase might be involved, although this strain acts differently from the OG1RF strain. A lower level of interaction of the other OG1 derivatives with *E. coli* and *B. fragilis* might also be involved. In this setting, the gelatinase may damage the virulence factors of these organisms. Even though the cytolysin demonstrates bacteriocin activities that might be deleterious to other bacteria, such properties have not been reported with gelatinase. On the other hand, OG1X was generated with nitrosoguanidine, a technique which could induce a mutation in other genes.

The combination of two virulence factors (Cly and Agg or Agg and Gel) gave results different from those reported with a single factor. A combination of Cly and Agg generated the lowest  $LD_{50}$  in the mouse model. These results suggest a possible interaction between Cly and Agg, the mechanism of which remains to be elucidated. Similar synergy has been previously reported with the combination of Cly and Agg, which caused an increased level of mortality in a rabbit model of endocarditis (5). In contrast, the combination of Agg and Gel did not seem to result in increased pathogenicity. In the rat model, the effects of the combinations were minimal, but the inflammatory response was increased, as assessed by the concentrations of  $\alpha$ 1-acid glycoprotein in plasma.

In conclusion, our study confirms the previously reported mechanisms of bacterial synergy between enterococci and other organisms. However, the marked discrepancy in results obtained from mice and rats renders any extrapolation to clinical practice difficult. The use of one virulence factor only minimally influenced the course of the disease. In contrast, the combination of two factors (Cly and Agg) seemed to be responsible for the more severe peritoneal infection which resulted in mice. In view of our results, other components shared by these strains may have a greater influence on the pathogenicity of enterococci. A good candidate could be the bacterial wall and, more specifically, some of its constituents, such as peptidoglycan or lipoteichoic acid, as has previously been suggested with experimental peritonitis (31). Vol. 66, 1998

We thank B. Murray, Texas University, Houston, and D. B. Clewell, Michigan University, Ann Arbor, for providing the strains of *E. faecalis*; and C. Poüs, Faculté de Pharmacie de Chatenay-Malabry, for performing the  $\alpha$ 1-acid glycoprotein assay.

This work was supported by a grant from the French Foundation for Medical Research.

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