

Glucose-Nonfermenting Gram-Negative Bacilli Associated with Clinical Veterinary Specimens

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Glucose-nonfermenting gram-negative bacilli (NFB) have been recognized recently as opportunistic pathogens of humans. With few exceptions, strains of NFB have not been considered important enough to be identified when isolated from animals. In this study, all NFB isolated during a 1-year period in a clinical veterinary microbiology laboratory were identified to determine their prevalence. Of the 347 strains of NFB obtained, the most common species were *Pseudomonas aeruginosa*, *Acinetobacter calcoaceticus*, *Bordetella bronchiseptica*, and *Pseudomonas pseudoalcaligenes*. Of all clinical veterinary specimens submitted for cultures, 10% contained nonfermenters.

Glucose-nonfermenting gram-negative bacilli (NFB) have had much attention recently in human medicine (5, 10, 15, 16). Most species of NFB were once thought to be contaminants when cultured from human specimens, but many have been shown to be opportunistic pathogens of humans (4, 8, 13).

In veterinary medicine, however, glucose-nonfermenting species are not considered to be animal pathogens. Most veterinary microbiology laboratories do not routinely identify NFB other than *Pseudomonas aeruginosa*, *Bordetella bronchiseptica*, and *Moraxella bovis*. For this reason, little is known about NFB associated with clinical specimens from animals or their role in animal disease.

This study was undertaken to determine the identity and the prevalence of glucose-nonfermenting species in clinical veterinary specimens submitted to our laboratory during a 1-year period. It was hoped that this would be a first step in determining the role of these organisms in infections of animals.

MATERIALS AND METHODS

Clinical isolates of NFB. All strains of NFB isolated in the Clinical Microbiology Laboratory of Texas A & M University Teaching Hospital between 1 January 1980 and 1 January 1981 were collected for identification. Information concerning animal species, specimen type, clinical history, and other bacteria isolated was recorded for each isolate of NFB.

Identification of isolates of NFB. The isolates of NFB were identified with a series of 30 conventional identification tests which were prepared, performed, and

interpreted according to recognized methods (9-11, 16, 17). Unless specifically noted, all identification tests were incubated aerobically at 35°C and interpreted at 24, 48, and 96 h. The following identification tests were performed on each unknown isolate: OF dextrose, OF lactose, OF maltose, OF xylose, OF fructose, OF mannitol, OF galactose, OF rhamnose, OF mannose, cytochrome oxidase, hemolysis, nitrate reduction (to nitrite or nitrogen gas or both), lysine and ornithine decarboxylase, arginine dihydrolase, esculin hydrolysis, citrate utilization, DNase, hydrogen sulfide production, indole production, *o*-nitrophenyl-beta-D-galactopyranosidase, growth on MacConkey agar, growth on salmonella-shigella agar, growth on cetrimide agar, growth at 42°C, pigment production (25°C), motility (25°C), and flagellar morphology. The identification of each isolate was made by comparing the results of these tests with published reactions for all species of NFB (5-7, 10, 15, 18).

Quality control for identification system. The reliability of the identification system was tested to ensure that these tests could accurately identify isolates of NFB. Twenty-seven known strains of NFB (22 species) were obtained from G. L. Gilardi (Hospital for Joint Diseases, New York). These known isolates were tested with the previously described set of identification tests. The identification of each known isolate obtained with our system was compared with that made by Gilardi. In each case, the identifications were the same.

RESULTS

Glucose nonfermenters were isolated from 347 (10%) of the 3,474 specimens submitted to the clinical microbiology laboratory during the 1-year study period. Six genera of NFB were found in the veterinary specimens (Table 1). *Pseudomonas* was the most prevalent species of glucose nonfermenters, accounting for 76.9% of all strains recovered. *Acinetobacter* (14.5%) was next in frequency, followed by *Bordetella*

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TABLE 1. Prevalence of 347 strains of NFB in 3,474 clinical veterinary specimens

Species (no.)	% All isolates	No. of isolates from:				
		Equine	Canine	Bovine	Feline	Other ^a
<i>Pseudomonas aeruginosa</i> (233)	67.4	111	104	4	8	7
<i>Acinetobacter calcoaceticus</i> (50)	14.5	27	17	2	2	2
<i>Bordetella bronchiseptica</i> (17)	4.9	8	9	0	0	0
<i>Pseudomonas pseudoalcaligenes</i> (8)	2.3	7	0	0	0	1
<i>Pseudomonas stutzeri</i> (5)	1.4	1	2	2	0	0
<i>Pseudomonas maltophilia</i> (4)	1.2	3	0	1	0	0
<i>Pseudomonas alcaligenes</i> (4)	1.2	3	1	0	0	0
<i>Pseudomonas putida</i> (4)	1.2	3	0	1	0	0
<i>Moraxella phenylpyruvica</i> (4)	1.2	1	3	0	0	0
CDC M-5 (3)	0.9	0	3	0	0	0
<i>Alcaligenes faecalis</i> (2)	0.6	1	0	1	0	0
<i>Pseudomonas fluorescens</i> (2)	0.6	1	0	1	0	0
<i>Pseudomonas diminuta</i> (2)	0.6	2	0	0	0	0
<i>Pseudomonas testosteroni</i> (2)	0.6	0	0	2	0	0
<i>Pseudomonas putrefaciens</i> (1)	0.3	0	0	0	0	1
CDC VE-2 (1)	0.3	1	0	0	0	0
CDC IIF (1)	0.3	0	0	0	0	1
<i>Alcaligenes odorans</i> (1)	0.3	0	1	0	0	0
<i>Moraxella nonliquifaciens</i> (1)	0.3	0	1	0	0	0
<i>Pseudomonas</i> sp. (1)	0.3	1	0	0	0	0

^a Includes specimens from birds, reptiles, pigs, goats, and rabbits.

(4.9%), *Moraxella* (2.4%), *Alcaligenes* (0.9%), and *Flavobacterium* (0.3%).

Table 1 also shows the prevalence of the species of NFB cultured from animal specimens. *P. aeruginosa* (67.4%) was the most commonly isolated glucose-nonfermenting species. *Acinetobacter calcoaceticus* (14.5%) was next in frequency, followed by *B. bronchiseptica* (4.9%) and *Pseudomonas pseudoalcaligenes* (2.3%). Twenty different species of NFB were cultured from the animal specimens.

NFB were found in specimens representing nine animal species (Table 1). Isolates of NFB were cultured most commonly from equine specimens (49.4%). Canine samples yielded 40.4% of all of the isolates. Bovine (3.5%), feline (2.9%), and other animal species (3.5%) provided the remaining isolations of NFB.

DISCUSSION

Strains of NFB were found in 10% of the clinical veterinary specimens. This isolation rate indicates that these bacteria may be more prevalent in animal specimens than is generally believed. There are no other reports on the prevalence of NFB species in clinical veterinary materials, but the prevalence of glucose nonfermenters in human specimens has been reported to be 12, 15, and 16% (14, 16).

In this study, 67.4% of NFB cultured from animals were *P. aeruginosa*, which is comparable to the 66% reported from human specimens by Pickett (14). The second most prevalent species found in the clinical veterinary speci-

mens was *Acinetobacter calcoaceticus* (14.5%), which has been reported to be the second most prevalent species of NFB cultured from humans (14).

The third most frequent species of NFB isolated from clinical veterinary materials was *B. bronchiseptica* (4.9%). This organism is a recognized pathogen of dogs and swine (1, 2). The majority of these isolates were from canine specimens (52.9%), but it is unusual that 47.1% of these strains were isolated from equine specimens (3, 12). *B. bronchiseptica* has rarely been reported from human specimens, accounting for less than 1% of all NFB strains (14).

The data presented on the distribution of glucose-nonfermenting species by animal species were influenced by the rate of specimen submission from the various animal species. In our laboratory, equine and canine specimens accounted for most of the submissions. These distribution data are of interest in that they may reveal general trends about the occurrence of a particular NFB species in animal specimens.

In this investigation, no etiological role was assigned to any of the isolates of NFB. The intent was to determine the prevalence of NFB species associated with veterinary specimens, since there are few reports on the species of NFB isolated from animals. Many of the species that we isolated have not been reported from animals. We have shown that NFB species do occur at a significant rate in animal specimens. Hopefully, this will lead to studies on the possible role of NFB species as animal pathogens.

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