

ERRATA

CLINICS 2007;62(1):47-54

Page 52, 1º Column, Line 22

Replace: Quintanilha et al ^{20,21} for Quintanilha et al ²¹⁻²²

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Table 4 - Mean Concentration and Prevalence (%) of Microorganisms in each Site of the LDT

| Microorganisms | ÍLEUM | | CECUM | | COLON | | | | | | RECTUM | | | |
|--|-------|------|-------|------|-----------------|----------------|------------------|-----------------|------------------|-----------------|---------------|--------------|-----|------|
| | MC | % | MC | % | Ascending MC | Ascending % | Transverse MC | Transverse % | Descending MC | Descending % | Sigmoid MC | Sigmoid % | MC | % |
| <i>Bacillus</i> sp | 1 | 9.5 | 2 | 4.2 | 5 | 4.2 | 3 | 4.2 | | | 1 | 4.3 | 3 | 5.0 |
| <i>Bacteroides</i> sp (pig.) | 5 | 4.8 | | | | | | | | | 5 | 4.3 | 1 | 5.0 |
| <i>Bacteroides</i> sp (npg) | 3 | 47.6 | 5 | 29.2 | 4 | 37.5 | 5 | 12.5 | 5 | 12.5 | 5 | 47.8 | 5 | 65.0 |
| <i>Bacteroides</i> sp | 2 | 4.8 | | | | | 2 | 4.2 | | | | | 3 | 5.0 |
| <i>Bifidobacterium</i> sp | 2 | 9.5 | | | 3 | 4.2 | 4 | 4.2 | 4 | 8.3 | 4 | 8.7 | 3 | 5.0 |
| <i>Candida</i> sp | 3.5 | 9.5 | 2 | 20.8 | 3 | 20.8 | 2 | 29.2 | 4 | 16.7 | 3 | 34.8 | 3 | 35.0 |
| <i>Clostridium ramosum</i> | 1 | 4.8 | | | | | | | 3 | 4.2 | | | | |
| <i>Clostridium</i> sp (gel -) | 4 | 66.7 | 4 | 45.8 | 5 | 50.0 | 4.5 | 58.3 | 5 | 54.2 | 5 | 69.6 | 7 | 60.0 |
| <i>Clostridium</i> sp (gel+) | | | 2 | 4.2 | 4 | 4.2 | | | | | | | 3 | 5.0 |
| <i>Clostridium</i> sp | 1 | 4.8 | 4 | 45.8 | 7 | 4.2 | | | | | | | 5 | 5.0 |
| <i>Corynebacterium</i> sp | 3 | 57.1 | 3 | 54.2 | 3.5 | 50.0 | 4 | 41.7 | 4 | 45.8 | 5 | 60.9 | 5 | 65.0 |
| <i>E.coli</i> | 5 | 47.6 | 5 | 37.5 | 5 | 45.8 | 4.5 | 50.0 | 5 | 37.5 | 5 | 69.6 | 7 | 80.0 |
| <i>Enterobacter cloacae</i> | | | 4 | 4.2 | 1 | 4.2 | | | | | | | | |
| <i>Enterobacter</i> sp | 4 | 28.6 | 4 | 20.8 | 7 | 37.5 | 4 | 29.2 | 5 | 45.8 | 5 | 52.2 | 7 | 35.0 |
| <i>Enterococcus faecalis</i> | 4.5 | 9.5 | 3 | 4.2 | 4 | 4.2 | 3 | 4.2 | 1 | 4.2 | 4 | 8.7 | | |
| <i>Enterococcus faecium</i> | | | | | 5 | 4.2 | | | | | | | | |
| <i>Enterococcus</i> sp | 4 | 38.1 | 2.5 | 33.3 | 3.5 | 33.3 | 3 | 58.3 | 5 | 58.3 | 5 | 34.8 | 5 | 60.0 |
| <i>Eubacterium lentum</i> | 4 | 4.8 | | | | | | | | | | | | |
| <i>Eubacterium</i> sp | 3.5 | 9.5 | | | | | | | | | | | | |
| <i>Fusobacterium fusiformes</i> | 3 | 4.8 | | | | | | | | | | | | |
| <i>Fusobacterium</i> sp | 3.5 | 19.0 | 2.5 | 25.0 | 3 | 20.8 | 4 | 16.7 | 4 | 8.3 | 3 | 21.7 | 4 | 55.0 |
| <i>Klebsiella pneumoniae</i> | | | | | | | | | 5 | 4.2 | | | | |
| <i>Klebsiella</i> sp | 5 | 76.2 | 4 | 54.2 | 6 | 62.5 | 5 | 54.2 | 5 | 41.7 | 7 | 69.6 | 7 | 65.0 |
| <i>Lactobacillus acidophilus</i> | 4 | 4.8 | 3 | 4.2 | | | | | | | | | | |
| <i>Lactobacillus</i> sp | 4 | 33.3 | 2 | 29.2 | 2.5 | 33.3 | 3 | 37.5 | 3 | 25.0 | 2 | 43.5 | 4 | 70.0 |
| <i>Leptotrichia</i> sp | | | | | 1 | 4.2 | | | | | 1 | 4.3 | | |
| <i>Peptococcus anaerobius</i> | 2 | 4.8 | 4 | 4.2 | 2 | 4.2 | | | | | 1 | 4.3 | | |
| <i>Peptococcus assachalyticus</i> | 2 | 4.8 | | | | | | | | | | | | |
| <i>Peptococcus</i> sp | 2 | 28.6 | 2.5 | 33.3 | 3 | 25.0 | 3 | 33.3 | 3 | 25.0 | 3 | 56.5 | 3 | 35.0 |
| <i>Peptostreptococcus</i> sp | 3 | 9.5 | | | 2 | 8.3 | | | 1 | 4.2 | 5 | 13.0 | | |
| <i>Propionibacterium</i> sp | 3 | 52.4 | 3 | 20.8 | 4 | 33.3 | 3 | 50.0 | 3 | 29.2 | 5 | 26.1 | 5 | 30.0 |
| <i>Proteus</i> sp | 3.5 | 19.0 | 4 | 12.5 | 5 | 33.3 | 5 | 37.5 | 5 | 25.0 | 7 | 34.8 | 7 | 30.0 |
| <i>Pseudomonas</i> sp | 3 | 9.5 | | | | | 1 | 4.2 | | | | | | |
| <i>Rodothorula</i> sp | 4 | 14.3 | 2 | 20.8 | 1 | 4.2 | 1 | 12.5 | 3 | 16.7 | 1 | 4.3 | 3.5 | 20.0 |
| <i>Selenomonas</i> sp | | | | | | | | | | | | | 2 | 5.0 |
| <i>Staphylococcus</i> sp | 5 | 4.8 | 2 | 4.2 | | | | | | | | | | |
| <i>Staphylococcus</i> sp coag - | 2 | 33.3 | 2 | 20.8 | 1.5 | 8.3 | 2 | 20.8 | 2 | 8.3 | 3 | 21.7 | 3 | 45.0 |
| <i>Staphylococcus</i> sp (coag+) | | | | | | | | | | | | | 3 | 5.0 |
| <i>Streptococcus/</i> gama hem | | | | | | | | | 2 | 8.3 | | | | |
| <i>Streptococcus</i> sp/alfa hemolítico grupo Viridans | 4 | 14.3 | 4 | 16.7 | 2 | 8.3 | 3.5 | 8.3 | | | 2 | 13.0 | 3 | 5.0 |
| <i>Veillonella</i> sp (Gel -) | | | | | | | 3 | 8.3 | | | | | | |
| <i>Veillonella</i> sp | 4 | 90.5 | 2.5 | 83.3 | 4 | 70.8 | 4 | 62.5 | 5 | 75.0 | 5 | 95.7 | 5 | 90.0 |

MC = Mean concentration (UFC / Log₁₀), % = prevalence

Table 5 - Association between microorganisms/sites of the lower digestive tract

| Microbiota | Lower Digestive Tract site |
|---|----------------------------|
| <i>Klebsiella</i> sp, <i>Clostridium</i> sp (gel-), <i>Veillonella</i> sp | All sites |
| <i>Enterobacter</i> sp | Sigmoid* |
| <i>Cândida</i> sp | Sigmoid and Rectum* |
| <i>Enterococcus</i> sp e <i>Lactobacillus</i> sp | Rectum* |

sp = species, *p < 0,05 or < 5%