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Corrigendum: Nosocomial and Community-Acquired Spontaneous Bacterial Peritonitis in patients with liver cirrhosis in China: Comparative Microbiology and Therapeutic Implications

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The original version of this Article contained errors. In the Abstract,

“The patients in the community-acquired SBP (n = 311) and the nosocomial SBP (n = 264) groups exhibited significant differences in clinical symptoms (P < 0.01)”.

now reads:

“The patients in the community-acquired SBP (n = 264) and the nosocomial SBP (n = 311) groups exhibited significant differences in clinical symptoms (P < 0.01)”.

In the Results section, under subheading ‘Basic patient information,’

“Patients were divided into either nosocomial (n = 264) or community-acquired groups (n = 311) according to the type of infection”.

now reads:

“Patients were divided into either nosocomial (n = 311) or community-acquired groups (n = 264) according to the type of infection”.

As a result, the values of the columns ‘Community-acquired (n, %)’ and ‘Nosocomial (n, %)’ in Table 2 were incorrect.

These errors have now been corrected in the PDF and HTML versions of the Article.

	Total (n, %)	Community-acquired (n, %)	Nosocomial (n, %)	P value
G – B	293	156	137	0.92
<i>Escherichia coli</i>	153 (26.6)	78 (50)	75 (54.7)	0.68
<i>Klebsiella pneumoniae</i>	75 (13.0)	35 (22.4)	40 (29.2)	0.66
<i>Acinetobacter baumannii</i>	7 (1.2)	5 (3.2)	2 (1.6)	0.15
<i>Pseudomonas aeruginosa</i>	4 (0.7)	2 (1.3)	2 (1.6)	0.82
<i>Enterobacter sp.</i>	17 (3.0)	12 (7.7)	5 (3.6)	0.03
<i>Serratia marcescens</i>	2 (0.3)	1 (0.6)	1 (0.7)	0.87
<i>Serratia fonticola</i>	2 (0.3)	2 (1.3)	0 (0.0)	0.11
<i>Stenotrophomonas maltophilia</i>	3 (0.5)	0 (0.0)	3 (2.2)	0.12
<i>Pseudomonas putida</i>	3 (0.5)	3 (1.9)	0 (0.0)	0.05
<i>Bacterium vulgare</i>	3 (0.5)	2 (1.3)	1 (0.7)	0.44
<i>Proteus mirabilis</i>	1 (0.2)	0 (0.0)	1 (0.7)	0.37
<i>Burkholderia cepacia</i>	3 (0.5)	2 (1.3)	1 (0.7)	0.44
<i>Acinetobacter lwoffii</i>	1 (0.2)	1(0.6)	0 (0.0)	0.26
<i>Salmonella</i>	4 (0.7)	3 (1.9)	1 (0.7)	0.22
<i>Shigella spp</i>	5 (0.9)	4 (2.6)	1 (0.7)	0.11
<i>Morganella morgani</i>	3 (0.5)	1 (0.6)	2 (1.5)	0.70
<i>Citrobacter</i>	3 (0.5)	1 (0.6)	2 (1.5)	0.70
<i>Haemophilus influenzae</i>	2 (0.3)	2 (1.3)	0 (0.0)	0.11
<i>Achromobacter xylosoxidans</i>	2 (0.3)	2 (1.3)	0 (0.0)	0.11
G+C	279	107	172	0.04
<i>Coagulase-negative staphylococcus</i>	131 (22.8)	56 (52.3)	75 (43.6)	0.68
<i>Staphylococcus aureus</i>	30 (5.2)	9 (8.4)	21 (12.2)	0.10
<i>Staphylococcus lugdunensis</i>	1 (0.2)	0 (0.0)	1 (0.6)	0.37
<i>Faecalis</i>	52 (9.0)	15 (14.0)	37 (21.5)	<0.01
<i>Faecium</i>	18 (1.7)	5 (4.7)	13 (7.6)	<0.05
<i>Other Enterococcus</i>	6 (1.0)	1 (0.9)	5 (2.9)	<0.05
<i>Streptococcus Pneumoniae</i>	1 (0.2)	1 (0.9)	0 (0.0)	0.26
<i>Streptococcus (other than S. pneumoniae)</i>	40 (7.0)	20 (18.7)	20 (11.6)	0.46
G+B	1	1	0	
<i>Ochrobactrum Anthropic</i>	1 (0.17)	1	0	0.26
<i>Anaerobic bacteria</i>	0 (0.0)	0	0	—
Fungus	2	0	2	
<i>Candida glabrata</i>	2 (0.3)	0	2	0.21

Table 2. Bacteria distributions in ascitic fluid.



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