

Result overview table--continued from previous page					
Sample Name	Sample ID	Organism (best match)	Score Value	Organism (second-best match)	Score Value
<u>D4</u> (+++)(A)	670 (standard)	Staphylococcus epidermidis	<u>2.22</u>	Staphylococcus epidermidis	<u>2.22</u>
<u>D5</u> (+++)(A)	36 (standard)	<u>Escherichia coli</u>	<u>2.43</u>	<u>Escherichia coli</u>	<u>2.24</u>
<u>D6</u> (+++)(A)	36 OKO MEM (standard)	Enterococcus faecium	<u>2.49</u>	Enterococcus faecium	<u>2.40</u>
<u>D7</u> (+++)(A)	37 (standard)	<u>Escherichia coli</u>	<u>2.45</u>	<u>Escherichia coli</u>	<u>2.24</u>
<u>D8</u> (+++)(A)	37 OKO MEM (standard)	Enterococcus faecium	<u>2.40</u>	Enterococcus faecium	<u>2.32</u>
<u>D9</u> (+++)(A)	128 1 (standard)	<u>Acinetobacter baumannii</u>	<u>2.32</u>	<u>Acinetobacter baumannii</u>	<u>2.32</u>
<u>D10</u> (+++)(A)	2 (standard)	Staphylococcus epidermidis	<u>2.20</u>	Staphylococcus epidermidis	<u>2.12</u>
<u>D11</u> (+++)(A)	MC (standard)	<u>Acinetobacter baumannii</u>	<u>2.34</u>	<u>Acinetobacter baumannii</u>	<u>2.29</u>
<u>D12</u> (+++)(A)	145 MC OKR (standard)	Providencia stuartii	<u>2.24</u>	Providencia stuartii	<u>2.22</u>
<u>E1</u> (+++)(A)	MC RAZL (standard)	Providencia stuartii	<u>2.54</u>	Providencia stuartii	<u>2.50</u>
<u>E2</u> (+++)(A)	CNA BIJ (standard)	Providencia stuartii	<u>2.41</u>	Providencia stuartii	<u>2.37</u>
<u>E3</u> (+++)(A)	CNA SIV (standard)	Providencia stuartii	<u>2.50</u>	Providencia stuartii	<u>2.46</u>
<u>E4</u> (+++)(A)	147 MC (standard)	<u>Acinetobacter baumannii</u>	<u>2.36</u>	<u>Acinetobacter baumannii</u>	<u>2.17</u>
<u>E5</u> (+++)(A)	CNA KR (standard)	Staphylococcus aureus	<u>2.56</u>	Staphylococcus aureus	<u>2.50</u>
<u>E6</u> (+++)(A)	CNA SIT (standard)	Corynebacterium striatum	<u>2.40</u>	Corynebacterium striatum	<u>2.36</u>
<u>E7</u> (+++)(A)	672 (standard)	<u>Raoultella ornithinolytica</u>	<u>2.22</u>	<u>Raoultella ornithinolytica</u>	<u>2.01</u>
Result overview table--continued on next page					



Result overview table--continued from previous page

Sample Name	Sample ID	Organism (best match)	Score Value	Organism (second-best match)	Score Value
<u>F12</u> (+++)(C)	3-1 (standard)	<u>Klebsiella oxytoca</u>	<u>2.21</u>	<u>Klebsiella oxytoca</u>	<u>2.12</u>
<u>G1</u> (+++)(A)	4917-1-1 (standard)	Micrococcus luteus	<u>2.03</u>	Micrococcus luteus	<u>1.94</u>
<u>G2</u> (-)(C)	X-1 (standard)	no peaks found	<u>0.00</u>	no peaks found	<u>0.00</u>
<u>G3</u> (+++)(A)	4664-1-1 (standard)	Staphylococcus haemolyticus	<u>2.18</u>	Staphylococcus borealis	<u>1.97</u>
<u>G4</u> (+++)(C)	2-1 (standard)	Staphylococcus epidermidis	<u>2.35</u>	Staphylococcus epidermidis	<u>2.24</u>
<u>G5</u> (+++)(C)	3-1 (standard)	Staphylococcus epidermidis	<u>2.22</u>	Staphylococcus epidermidis	<u>2.00</u>
<u>G6</u> (+++)(A)	KP 51-1 (standard)	<u>Klebsiella pneumoniae</u>	<u>2.24</u>	<u>Klebsiella pneumoniae</u>	<u>2.23</u>
<u>G7</u> (+++)(A)	KP 52-1 (standard)	<u>Klebsiella pneumoniae</u>	<u>2.12</u>	<u>Klebsiella pneumoniae</u>	<u>2.02</u>
<u>G8</u> (+++)(A)	4763-1-1 (standard)	<u>Citrobacter freundii</u>	<u>2.30</u>	<u>Citrobacter freundii</u>	<u>2.27</u>
<u>G9</u> (+++)(C)	2-1 (standard)	<u>Klebsiella pneumoniae</u>	<u>2.38</u>	<u>Klebsiella pneumoniae</u>	<u>2.37</u>
<u>G10</u> (+++)(C)	3-1 (standard)	Enterococcus faecalis	<u>2.39</u>	Enterococcus faecalis	<u>2.35</u>
<u>G11</u> (+++)(C)	4-1 (standard)	Enterococcus faecium	<u>2.52</u>	Enterococcus faecium	<u>2.48</u>