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1 Table 2: Microbiological characteristics at study entry for the MITT population by treatment

2

Pathogen	Iclaprim q12h (N =8)	Iclaprim q8h (N =7)	Vancomycin (N = 6)
Positive Respiratory Culture	8 (100.0%)	7 (100.0%)	6 (100.0%)
Gram-positive			
Staphylococcus aureus	7	5	3
MRSA	1	3	2
MSSA	6	2	1
Streptococcus agalactiae	0	0	1
Streptococcus beta hemolytic	0	1	0
Streptococcus group C	0	0	1
Streptococcus group F	1	0	0
Streptococcus pneumoniae	1	2	2
Gram-negative			
Acinetobacter baumannii	2	2	1
Citrobacter freundii	0	0	1
Enterobacter cloacae	1	0	0
Escherichia coli	1	1	0
Klebsiella pneumoniae	3	1	2
Moraxella catarrhalis	1	1	0
Morganella morganii	1	0	0
Pseudomonas aeruginosa	1	1	0

Positive Blood Culture	3 (37.5%)	0 (0%)	2 (33.03%)	3
Staphylococcus aureus	1	0	1	4
MRSA	0	0	1	5
MSSA	1	0	0	6
Staphylococcus species, (coagulase negative)	2	0	2	7
Streptococcus viridians	0	0	1	8
Enterobacter cloacae	1	0	0	9
Enterococcus faecalis	1	0	0	

- 10 Note: Number and percentages do not add up to total or 100% because some patients had more
- 11 than one pathogen identified from either respiratory or blood cultures.