

**Nosocomial Infections - Cultured Organisms**  
**With adjudications applied (n = 130)**

<i>Class</i>	<i>Organism</i>	<i>Prevalent Infec.: # patients (% prev.)</i>	<i>Incident Infec.: # infec. (#patients, % incid.)</i>
<i>Gram+ Bacteria</i>	Methicillin-susceptible Staphylococcus aureus (MSSA)	3/130 (2.3%)	3 (2/130, 1.5%)
	Methicillin-resistant Staphylococcus aureus (MRSA)	4/130 (3.1%)	5 (5/130, 3.8%)
	Coagulase-negative Staphylococcus species	4/130 (3.1%)	10 (9/130, 6.9%)
	Vancomycin-susceptible Enterococcus faecalis	5/130 (3.8%)	2 (2/130, 1.5%)
	Vancomycin-resistant Enterococcus faecalis	0/130 (0.0%)	1 (1/130, 0.8%)
	Vancomycin-susceptible Enterococcus faecium	0/130 (0.0%)	0 (0/130, 0.0%)
	Vancomycin-resistant Enterococcus faecium	2/130 (1.5%)	10 (7/130, 5.4%)
	Streptococcus pneumoniae	0/130 (0.0%)	0 (0/130, 0.0%)
	Clostridium difficile	1/130 (0.8%)	5 (5/130, 3.8%)
	Other gram + species	13/130 (10.0%)	26 (14/130, 10.8%)
	-- (not specified)	0/130 (0.0%)	4 (3/130, 2.3%)
	-- 1 + B hemolytic streptococcus	1/130 (0.8%)	0 (0/130, 0.0%)
	-- 2+ mixed gastro-intestinal flora	1/130 (0.8%)	0 (0/130, 0.0%)
	-- Bacteroides fragilis	1/130 (0.8%)	0 (0/130, 0.0%)
	-- Bacteroides uniformis	1/130 (0.8%)	0 (0/130, 0.0%)
	-- Chryseobacterium indologenes	1/130 (0.8%)	0 (0/130, 0.0%)
	-- Citrobacter amalonaticus	0/130 (0.0%)	1 (1/130, 0.8%)
	-- Citrobacter freundii	0/130 (0.0%)	2 (1/130, 0.8%)
	-- Clostridium tertium	1/130 (0.8%)	0 (0/130, 0.0%)
	-- Enterococcus faecalis resistance not specified	0/130 (0.0%)	2 (1/130, 0.8%)
	-- Enterococcus species not specified	0/130 (0.0%)	3 (3/130, 2.3%)
	-- Enterococcus species, beta-lactamase negative	1/130 (0.8%)	1 (1/130, 0.8%)
	-- Enterococcus species, strain 1	1/130 (0.8%)	0 (0/130, 0.0%)
	-- Enterococcus species, strain 2	1/130 (0.8%)	0 (0/130, 0.0%)
	-- Few streptococcus, viridans group	0/130 (0.0%)	1 (1/130, 0.8%)
	-- Haemophilus influenzae	1/130 (0.8%)	0 (0/130, 0.0%)
	-- Lactobacillus species	1/130 (0.8%)	1 (1/130, 0.8%)
	-- Lactose fermenter gram negative rod	0/130 (0.0%)	2 (2/130, 1.5%)
	-- Micrococcus species	0/130 (0.0%)	1 (1/130, 0.8%)
	-- Non-lactose fermenter gram negative rod	0/130 (0.0%)	1 (1/130, 0.8%)
	-- Proteus species	1/130 (0.8%)	0 (0/130, 0.0%)
	-- Rare gram positive cocci pairs	0/130 (0.0%)	1 (1/130, 0.8%)
	-- Serratia marcescens	1/130 (0.8%)	1 (1/130, 0.8%)
	-- Staphylococcus aureus resistance unknown	0/130 (0.0%)	2 (1/130, 0.8%)
	-- Stenotrophomonas maltophilia	2/130 (1.5%)	3 (3/130, 2.3%)
	-- Streptococcus group C	1/130 (0.8%)	0 (0/130, 0.0%)
	== Any Gram+ Bacteria ==	23/130 (17.7%)	62 (31/130, 23.8%)
<i>Gram- Bacteria</i>	Pseudomonas aeruginosa	1/130 (0.8%)	8 (8/130, 6.2%)
	Escherichia coli	2/130 (1.5%)	7 (6/130, 4.6%)
	Acinetobacter baumannii	0/130 (0.0%)	1 (1/130, 0.8%)
	Enterobacter cloacae	3/130 (2.3%)	3 (3/130, 2.3%)
	Enterobacter aerogenes	4/130 (3.1%)	1 (1/130, 0.8%)

**Nosocomial Infections - Cultured Organisms**  
**With adjudications applied (n = 130)**

<i>Class</i>	<i>Organism</i>	<i>Prevalent Infec.: # patients (% prev.)</i>	<i>Incident Infec.: # infec. (#patients, % incid.)</i>
	Klebsiella pneumoniae	2/130 (1.5%)	10 (7/130, 5.4%)
	Other Klebsiella species	1/130 (0.8%)	4 (3/130, 2.3%)
	-- (not specified)	0/130 (0.0%)	1 (1/130, 0.8%)
	-- Klebsiella not specified	1/130 (0.8%)	1 (1/130, 0.8%)
	-- Klebsiella oxytoca	0/130 (0.0%)	2 (2/130, 1.5%)
	Other gram - species	0/130 (0.0%)	6 (5/130, 3.8%)
	-- Citrobacter koseri	0/130 (0.0%)	1 (1/130, 0.8%)
	-- Enterobacter sakazakii	0/130 (0.0%)	1 (1/130, 0.8%)
	-- Enterobacter species not specified	0/130 (0.0%)	1 (1/130, 0.8%)
	-- Few gram negative rods	0/130 (0.0%)	1 (1/130, 0.8%)
	-- Rare gram negative rod - lactose fermenter	0/130 (0.0%)	1 (1/130, 0.8%)
	-- Stenotrophomonas maltophilia	0/130 (0.0%)	1 (1/130, 0.8%)
	== Any Gram- Bacteria ==	12/130 (9.2%)	40 (24/130, 18.5%)
<i>Fungal Species</i>	Candida albicans	5/130 (3.8%)	14 (14/130, 10.8%)
	Candida glabrata	2/130 (1.5%)	9 (7/130, 5.4%)
	Candida tropicalis	1/130 (0.8%)	0 (0/130, 0.0%)
	Other fungal species	4/130 (3.1%)	6 (5/130, 3.8%)
	-- (not specified)	0/130 (0.0%)	2 (2/130, 1.5%)
	-- Cadida not otherwise specified	0/130 (0.0%)	1 (1/130, 0.8%)
	-- Candida parapsilosis	1/130 (0.8%)	1 (1/130, 0.8%)
	-- Yeast not specified	3/130 (2.3%)	2 (1/130, 0.8%)
	== Any Fungal Species ==	10/130 (7.7%)	29 (21/130, 16.2%)
<i>Other</i>	Cytomegalovirus	1/130 (0.8%)	0 (0/130, 0.0%)
	== Any Infection ==	38/130 (29.2%)	131 (52/130, 40.0%)