

**GLND - Summary of Nosocomial Infections
By Cultured Organism**

Class	Organism	A	A	B	B
		Prevalent Infec.: # patients (% prev.)	Incident Infec.: # infec. (#patients, % incid.)	Prevalent Infec.: # patients (% prev.)	Incident Infec.: # infec. (#patients, % incid.)
Gram+ Bacteria	Methicillin-susceptible Staphylococcus aureus (MSSA)	2/64 (3.1%)	0 (0/64, 0.0%)	1/66 (1.5%)	3 (2/66, 3.0%)
	Methicillin-resistant Staphylococcus aureus (MRSA)	2/64 (3.1%)	2 (2/64, 3.1%)	2/66 (3.0%)	3 (3/66, 4.5%)
	Coagulase-negative Staphylococcus species	2/64 (3.1%)	6 (5/64, 7.8%)	2/66 (3.0%)	4 (4/66, 6.1%)
	Vancomycin-susceptible Enterococcus faecalis	3/64 (4.7%)	2 (2/64, 3.1%)	2/66 (3.0%)	0 (0/66, 0.0%)
	Vancomycin-resistant Enterococcus faecalis	0/64 (0.0%)	0 (0/64, 0.0%)	0/66 (0.0%)	1 (1/66, 1.5%)
	Vancomycin-susceptible Enterococcus faecium	0/64 (0.0%)	0 (0/64, 0.0%)	0/66 (0.0%)	0 (0/66, 0.0%)
	Vancomycin-resistant Enterococcus faecium	1/64 (1.6%)	6 (4/64, 6.3%)	1/66 (1.5%)	4 (3/66, 4.5%)
	Streptococcus pneumoniae	0/64 (0.0%)	0 (0/64, 0.0%)	0/66 (0.0%)	0 (0/66, 0.0%)
	Clostridium difficile	1/64 (1.6%)	4 (4/64, 6.3%)	0/66 (0.0%)	1 (1/66, 1.5%)
	Other gram + species	6/64 (9.4%)	16 (8/64, 12.5%)	7/66 (10.6%)	10 (6/66, 9.1%)
	-- (not specified)	0/64 (0.0%)	1 (1/64, 1.6%)	0/66 (0.0%)	3 (2/66, 3.0%)
	-- Chryseobacterium indologenes	1/64 (1.6%)	0 (0/64, 0.0%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- Citrobacter amalonaticus	0/64 (0.0%)	1 (1/64, 1.6%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- 1 + B hemolytic streptococcus	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	1/66 (1.5%)	0 (0/66, 0.0%)
	-- Citrobacter freundii	0/64 (0.0%)	2 (1/64, 1.6%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- 2+ mixed gastro-intestinal flora	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	1/66 (1.5%)	0 (0/66, 0.0%)
	-- Bacteroides fragilis	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	1/66 (1.5%)	0 (0/66, 0.0%)
	-- Clostridium tertium	1/64 (1.6%)	0 (0/64, 0.0%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- Bacteroides uniformis	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	1/66 (1.5%)	0 (0/66, 0.0%)
	-- Enterococcus faecalis resistance no	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	0/66 (0.0%)	2 (1/66, 1.5%)
	-- Enterococcus species, beta-lactamas	1/64 (1.6%)	1 (1/64, 1.6%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- Enterococcus species not specified	0/64 (0.0%)	2 (2/64, 3.1%)	0/66 (0.0%)	1 (1/66, 1.5%)
	-- Enterococcus species, strain 1	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	1/66 (1.5%)	0 (0/66, 0.0%)
	-- Few streptococcus, viridans group	0/64 (0.0%)	1 (1/64, 1.6%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- Enterococcus species, strain 2	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	1/66 (1.5%)	0 (0/66, 0.0%)
	-- Haemophilus influenzae	1/64 (1.6%)	0 (0/64, 0.0%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- Lactobacillus species	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	1/66 (1.5%)	1 (1/66, 1.5%)
	-- Non-lactose fermenter gram negative	0/64 (0.0%)	1 (1/64, 1.6%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)

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	-- Lactose fermenter gram negative rod	0/64 (0.0%)	1 (1/64, 1.6%)	0/66 (0.0%)	1 (1/66, 1.5%)
	-- Proteus species	1/64 (1.6%)	0 (0/64, 0.0%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- Micrococcus species	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	0/66 (0.0%)	1 (1/66, 1.5%)
	-- Rare gram positive cocci pairs	0/64 (0.0%)	1 (1/64, 1.6%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- Serratia marcescens	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	1/66 (1.5%)	1 (1/66, 1.5%)
	-- Staphylococcus aureus resistance un	0/64 (0.0%)	2 (1/64, 1.6%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- Stenotrophomonas maltophilia	2/64 (3.1%)	3 (3/64, 4.7%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- Streptococcus group C	1/64 (1.6%)	0 (0/64, 0.0%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	== Any Gram+ Bacteria ==	10/64 (15.6%)	36 (17/64, 26.6%)	13/66 (19.7%)	26 (14/66, 21.2%)
Gram- Bacteria	Pseudomonas aeruginosa	0/64 (0.0%)	6 (6/64, 9.4%)	1/66 (1.5%)	2 (2/66, 3.0%)
	Escherichia coli	1/64 (1.6%)	7 (6/64, 9.4%)	1/66 (1.5%)	0 (0/66, 0.0%)
	Acinetobacter baumannii	0/64 (0.0%)	0 (0/64, 0.0%)	0/66 (0.0%)	1 (1/66, 1.5%)
	Enterobacter cloace	2/64 (3.1%)	1 (1/64, 1.6%)	1/66 (1.5%)	2 (2/66, 3.0%)
	Enterobacter aerogenes	1/64 (1.6%)	1 (1/64, 1.6%)	3/66 (4.5%)	0 (0/66, 0.0%)
	Klebsiella pneumoniae	0/64 (0.0%)	7 (4/64, 6.3%)	2/66 (3.0%)	3 (3/66, 4.5%)
	Other Klebsiella species	1/64 (1.6%)	1 (1/64, 1.6%)	0/66 (0.0%)	3 (2/66, 3.0%)
	-- (not specified)	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	0/66 (0.0%)	1 (1/66, 1.5%)
	-- Klebsiella not specified	1/64 (1.6%)	1 (1/64, 1.6%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- Klebsiella oxytoca	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	0/66 (0.0%)	2 (2/66, 3.0%)
	Other gram - species	0/64 (0.0%)	4 (3/64, 4.7%)	0/66 (0.0%)	2 (2/66, 3.0%)
	-- Citrobacter koseri	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	0/66 (0.0%)	1 (1/66, 1.5%)
	-- Enterobacter sakazakii	0 (0/ 64, 0.0%)	0 (0/ 64, 0.0%)	0/66 (0.0%)	1 (1/66, 1.5%)
	-- Enterobacter species not specified	0/64 (0.0%)	1 (1/64, 1.6%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- Few gram negative rods	0/64 (0.0%)	1 (1/64, 1.6%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- Rare gram negative rod - lactose fe	0/64 (0.0%)	1 (1/64, 1.6%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	-- Stenotrophomonas maltophilia	0/64 (0.0%)	1 (1/64, 1.6%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	== Any Gram- Bacteria ==	5/64 (7.8%)	27 (14/64, 21.9%)	7/66 (10.6%)	13 (10/66, 15.2%)
Fungal Species	Candida albicans	2/64 (3.1%)	11 (11/64, 17.2%)	3/66 (4.5%)	3 (3/66, 4.5%)
	Candida glabrata	1/64 (1.6%)	5 (5/64, 7.8%)	1/66 (1.5%)	4 (2/66, 3.0%)
	Candida tropicalis	1/64 (1.6%)	0 (0/64, 0.0%)	0/66 (0.0%)	0 (0/66, 0.0%)
	Other fungal species	1/64 (1.6%)	5 (4/64, 6.3%)	3/66 (4.5%)	1 (1/66, 1.5%)
	-- (not specified)	0/64 (0.0%)	1 (1/64, 1.6%)	0/66 (0.0%)	1 (1/66, 1.5%)
	-- Candida parapsilosis	0/64 (0.0%)	1 (1/64, 1.6%)	1/66 (1.5%)	0 (0/66, 0.0%)

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	-- Yeast not specified	1/64 (1.6%)	2 (1/64, 1.6%)	2/66 (3.0%)	0 (0/66, 0.0%)
	-- Cadida not otherwise specified	0/64 (0.0%)	1 (1/64, 1.6%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
	== Any Fungal Species ==	3/64 (4.7%)	21 (16/64, 25.0%)	7/66 (10.6%)	8 (5/66, 7.6%)
Other	Cytomegalovirus	1/64 (1.6%)	0 (0/64, 0.0%)	0 (0/ 66, 0.0%)	0 (0/ 66, 0.0%)
Any Infection		17/64 (26.6%)	84 (31/64, 48.4%)	21/66 (31.8%)	47 (21/66, 31.8%)