Site code	Type code	Prevalent Infec.: # patients (% prev.)	Incident Infec.: # infec. (#patients, % incid.)	Causative Organisms
Bloodstream Infection	Laboratory-confirmed bloodstream infection	8/130 (6.2%)	29 (26/130, 20.0%)	Candida glabrata - [prev.]
				Coagulase-negative Staphylococcus species - [prev.]
				Other gram + species - Bacteroides uniformis - [prev.]
				Other gram + species - Clostridium tertium - [prev.]
				Other gram + species - Serratia marcescens - [prev.]
				Pseudomonas aeruginosa - [prev.]
				Vancomycin-susceptible Enterococcus faecalis - [prev.]
				Candida albicans - [inc.]
				Candida glabrata - [inc.]
				Coagulase-negative Staphylococcus species - [inc.]
				Enterobacter cloace - [inc.]
				Klebsiella pneumoniae - [inc.]
				Methicillin-resistant Staphylococcus aureus (MRSA) - [inc.]
				Methicillin-susceptible Staphylococcus aureus (MSSA) - [inc.]
				Other Klebsiella species [inc.]
				Other fungal species - Candida parapsilosis - [inc.]
				Other gram + species [inc.]
				Other gram + species - Enterococcus faecalis resistance not specified - [inc.]
				Other gram + species - Enterococcus species not specified - [inc.]
				Other gram + species - Enterococcus species, beta-lactamase negative - [inc.]
				Other gram + species - Staphylococcus aureus resistance unknown - [inc.]
				Other gram + species - Stenotrophomonas maltophilia - [inc.]
				Other gram - species - Citrobacter koseri - [inc.]
				Other gram - species - Enterobacter species not specified - [inc.]
				Vancomycin-susceptible Enterococcus faecalis - [inc.]
Central Nervous System Infection	Menitigitis or ventriculitis	0/130 (0.0%)	1 (1/130, 0.8%)	Vancomycin-resistant Enterococcus faecium - [inc.]
Cardiovascular System Infection	Mediastinitis	1/130 (0.8%)	0 (0/130, 0.0%)	(negative culture) - [prev.]
Cardiovascular System Infection	Arterial or venous infection	0/130 (0.0%)	1 (1/130, 0.8%)	Candida albicans - [inc.]

Site code	Type code	Prevalent Infec.:	Incident Infec.: # infec. (#patients, % incid.)	Causativa Organisms
- Site code	Type code	# patients (// prev.)	# IIIIec. (#pallerits, 76 IIIclu.)	Candida glabrata - [inc.]
Eye, Ear, Nose, Throat, or Mouth Infection	Oral Cavity (mouth, tongue, or gums)	0/130 (0.0%)	1 (1/130, 0.8%)	(no culture obtained) - [inc.]
Eye, Ear, Nose, Throat, or Mouth Infection	Sinusitis	1/130 (0.8%)	0 (0/130, 0.0%)	(no culture obtained) - [prev.]
Gastrointestinal System Infection	Gastroenteritis	1/130 (0.8%)	4 (4/130, 3.1%)	Clostridium difficile - [prev.]
				Clostridium difficile - [inc.]
Gastrointestinal System Infection	Gastrointestinal (GI) tract	1/130 (0.8%)	1 (1/130, 0.8%)	Other - Cytomegalovirus - [prev.]
				Clostridium difficile - [inc.]
Gastrointestinal System Infection	Intra-abdominal, not specified elsewhere	1/130 (0.8%)	0 (0/130, 0.0%)	Candida albicans - [prev.]
				Escherichia coli - [prev.]
				Other gram + species - Enterococcus species, strain 1 - [prev.]
				Other gram + species - Enterococcus species, strain 2 - [prev.]
Pneumonia	PNU1 - Clinically defined pneumonia	9/130 (6.9%)	11 (11/130, 8.5%)	(no culture obtained) - [prev.]
				Enterobacter aerogenes - [prev.]
				Methicillin-resistant Staphylococcus aureus (MRSA) - [prev.]
				Other fungal species - Yeast not specified - [prev.]
				Other gram + species - 1 + B hemolytic streptococcus - [prev.]
				Other gram + species - Haemophilus influenzae - [prev.]
				Other gram + species - Stenotrophomonas maltophilia - [prev.]
				Other gram + species - Streptococcus group C - [prev.]
				(negative culture) - [inc.]
				(no culture obtained) - [inc.]
				Escherichia coli - [inc.]
				Klebsiella pneumoniae - [inc.]
				Methicillin-resistant Staphylococcus aureus (MRSA) - [inc.]
				Other fungal species - Yeast not specified - [inc.]
				Other gram + species [inc.]
				Other gram + species - Citrobacter freundii - [inc.]
				Other gram + species - Lactose fermenter gram negative rod - [inc.]

		Prevalent Infec.:	Incident Infec.:	
Site code	Type code	# patients (% prev.)	# infec. (#patients, % incid.)	Causative Organisms
				Other gram + species - Micrococcus species - [inc.]
				Other gram + species - Non-lactose fermenter gram negative rod - [inc.]
				Other gram + species - Stenotrophomonas maltophilia - [inc.]
				Other gram - species - Enterobacter sakazakii - [inc.]
				Pseudomonas aeruginosa - [inc.]
Pneumonia	PNU2 - Pneumonia with specific lab findings	10/130 (7.7%)	8 (8/130, 6.2%)	Enterobacter aerogenes - [prev.]
				Enterobacter cloace - [prev.]
				Escherichia coli - [prev.]
				Klebsiella pneumoniae - [prev.]
				Methicillin-resistant Staphylococcus aureus (MRSA) - [prev.]
				Methicillin-susceptible Staphylococcus aureus (MSSA) - [prev.]
				Other Klebsiella species - Klebsiella not specified - [prev.]
				Other gram + species - Chryseobacterium indologenes - [prev.]
				Other gram + species - Stenotrophomonas maltophilia - [prev.]
				Pseudomonas aeruginosa - [prev.]
				Acinetobacter baumannii - [inc.]
				Escherichia coli - [inc.]
				Klebsiella pneumoniae - [inc.]
				Other gram + species - Serratia marcescens - [inc.]
				Other gram + species - Staphylococcus aureus resistance unknown - [inc.]
				Other gram + species - Stenotrophomonas maltophilia - [inc.]
				Other gram - species - Stenotrophomonas maltophilia - [inc.]
				Pseudomonas aeruginosa - [inc.]
Reproductive Tract Infection	Other infections of the male or female reproductive tract	1/130 (0.8%)	0 (0/130, 0.0%)	Vancomycin-resistant Enterococcus faecium - [prev.]
Surgical Site Infection	Deep incisional primary SSI	1/130 (0.8%)	0 (0/130, 0.0%)	Other gram + species - Proteus species - [prev.]
Surgical Site Infection	Gastrointestinal (GI) tract	1/130 (0.8%)	0 (0/130, 0.0%)	Candida albicans - [prev.]
				Candida glabrata - [prev.]
				Coagulase-negative Staphylococcus species - [prev.]
				Other gram + species - Enterococcus species, beta-lactamase negative - [prev.]

Site code	Type code	Prevalent Infec.: # patients (% prev.)	Incident Infec.: # infec. (#patients, % incid.)	Causative Organisms
Surgical Site Infection	Intra-abdominal, not specified elsewhere	5/130 (3.8%)	6 (5/130, 3.8%)	Candida albicans - [prev.]
				Candida glabrata - [prev.]
				Enterobacter aerogenes - [prev.]
				Enterobacter cloace - [prev.]
				Methicillin-resistant Staphylococcus aureus (MRSA) - [prev.]
				Other fungal species - Candida parapsilosis - [prev.]
				Other gram + species - Bacteroides fragilis - [prev.]
				Other gram + species - Lactobacillus species - [prev.]
				Vancomycin-resistant Enterococcus faecium - [prev.]
				Vancomycin-susceptible Enterococcus faecalis - [prev.]
				(no culture obtained) - [inc.]
				Candida albicans - [inc.]
				Candida glabrata - [inc.]
				Escherichia coli - [inc.]
				Klebsiella pneumoniae - [inc.]
				Methicillin-resistant Staphylococcus aureus (MRSA) - [inc.]
				Other gram + species - Enterococcus species not specified - [inc.]
				Other gram + species - Few streptococcus, viridans group - [inc.]
				Other gram + species - Rare gram positive cocci pairs - [inc.]
				Other gram - species - Few gram negative rods - [inc.]
				Other gram - species - Rare gram negative rod - lactose fermenter - [inc.]
				Pseudomonas aeruginosa - [inc.]
				Vancomycin-resistant Enterococcus faecium - [inc.]
Surgical Site Infection	Superficial incisional primary SSI	1/130 (0.8%)	1 (1/130, 0.8%)	Other gram + species - 2+ mixed gastro-intestinal flora - [prev.]
				Coagulase-negative Staphylococcus species - [inc.]
				Methicillin-susceptible Staphylococcus aureus (MSSA) - [inc.]
Surgical Site Infection	Skin	2/130 (1.5%)	0 (0/130, 0.0%)	(no culture obtained) - [prev.]
				Vancomycin-susceptible Enterococcus faecalis - [prev.]
Surgical Site Infection	Soft tissue	0/130 (0.0%)	1 (1/130, 0.8%)	Escherichia coli - [inc.]

Site code	Type code	Prevalent Infec.:	Incident Infec.: # infec. (#patients, % incid.)	Causative Organisms
Site code	туре соце	# patients (% prev.)	# Imec. (#palients, % Incid.)	Other gram + species - Citrobacter amalonaticus - [inc.]
				Other gram + species - Citrobacter freundii - [inc.]
				Vancomycin-resistant Enterococcus faecium - [inc.]
Surgical Site Infection	Unknown	0/130 (0.0%)	1 (1/130, 0.8%)	Enterobacter aerogenes - [inc.]
				Escherichia coli - [inc.]
				Vancomycin-resistant Enterococcus faecium - [inc.]
Skin and Soft Tissue Infection	Breast abscess or mastitis	0/130 (0.0%)	1 (1/130, 0.8%)	(negative culture) - [inc.]
Skin and Soft Tissue Infection	Soft tissue	1/130 (0.8%)	1 (1/130, 0.8%)	Coagulase-negative Staphylococcus species - [prev.]
				Candida albicans - [inc.]
Urinary Tract Infection	Asymptomatic bacteriuria	4/130 (3.1%)	14 (13/130, 10.0%)	Candida albicans - [prev.]
				Candida tropicalis - [prev.]
				Other fungal species - Yeast not specified - [prev.]
				Vancomycin-susceptible Enterococcus faecalis - [prev.]
				Candida albicans - [inc.]
				Candida glabrata - [inc.]
				Klebsiella pneumoniae - [inc.]
				Other Klebsiella species - Klebsiella not specified - [inc.]
				Other fungal species [inc.]
				Other gram + species [inc.]
				Pseudomonas aeruginosa - [inc.]
				Vancomycin-resistant Enterococcus faecium - [inc.]
				Vancomycin-susceptible Enterococcus faecalis - [inc.]
Urinary Tract Infection	Other infections of the urinary tract	0/130 (0.0%)	1 (1/130, 0.8%)	Other fungal species - Cadida not otherwise specified - [inc.]
Urinary Tract Infection	Symptomatic urinary tract infection	1/130 (0.8%)	17 (16/130, 12.3%)	Vancomycin-susceptible Enterococcus faecalis - [prev.]
				(negative culture) - [inc.]
				Candida albicans - [inc.]
				Candida glabrata - [inc.]
				Enterobacter cloace - [inc.]
				Escherichia coli - [inc.]
				Klebsiella pneumoniae - [inc.]
				Methicillin-resistant Staphylococcus aureus (MRSA) - [inc.]

011	Tomasada	Prevalent Infec.:	Incident Infec.:	Overethe Overether
Site code	Type code	# patients (% prev.) #	# infec. (#patients, % incid.)	Causative Organisms
				Other Klebsiella species - Klebsiella oxytoca - [inc.]
				Other fungal species - Yeast not specified - [inc.]
				Other gram + species [inc.]
				Other gram + species - Enterococcus faecalis resistance not specified - [inc.]
				Other gram + species - Enterococcus species not specified - [inc.]
				Other gram + species - Lactobacillus species - [inc.]
				Pseudomonas aeruginosa - [inc.]
				Vancomycin-resistant Enterococcus faecalis - [inc.]
				Vancomycin-resistant Enterococcus faecium - [inc.]
Overall total:		40/130 (30.8%)	99 (54/130, 41.5%)	