

Microorganisms Identified in Central Line-Associated Blood Stream Infections in Intensive Care Units in Illinois Hospitals

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A variety of organisms are found to contribute to central line-associated bloodstream infections (CLABSIs). Below is a chart showing the organisms identified in such infections in intensive care units (ICUs) in Illinois in 2011. Note that some infections have more than one organism present. Out of 510 microorganisms identified in 447 central line-associated bloodstream infections, the most common were *Enterococcus* spp., coagulase-negative *Staphylococcus* and *Candida* spp. MRSA accounted for 4.5% of these infections.

Pathogen	Number of Isolates	Percent of Infections
Coagulase-negative <i>Staphylococcus</i> (CNS)	97	19.0
Methicillin-resistant <i>Staphylococcus aureus</i> (MRSA)	23	4.5
Methicillin-susceptible <i>Staphylococcus aureus</i> (MSSA)	32	6.3
<i>Enterococcus</i> species	97	19.0
Vancomycin-resistant <i>Enterococcus</i> (VRE)	45	
Vancomycin-susceptible <i>Enterococcus</i>	44	
Other <i>Enterococcus</i> (Vancomycin susceptibility unknown or intermediate)	8	
<i>Candida</i> species	82	16.1
<i>C. albicans</i>	34	
Other <i>Candida</i> species	48	
<i>Enterobacter</i> species	30	5.9
<i>Klebsiella</i> species	31	6.1
<i>Pseudomonas</i> species	20	3.9
<i>Acinetobacter</i> species	21	4.1
<i>Escherichia coli</i>	22	4.3
Other gram-negative rods	33	6.5
Other pathogens	22	4.3
Total	510	100

Microorganisms Identified in CLABSIs in IL ICUs, 2011

