

Table S2. Time to Organism Identification From Gram Stain (hr)

Organism Identified	Pre-implementation (n = 68)	Post-implementation (n = 69)	p value
Gram-positive organisms, median (IQR)			
Coagulase-negative staphylococcus	31.4 (23.7–36.1)	20.7 (6–25.7)	0.005*
<i>Staphylococcus aureus</i>	25.8 (22.2–31)	27.7 (7.8–33.1)	0.97
<i>Streptococcus</i> species	27.1 (22.7–30.2)	8.23 (4.6–17.1)	0.01*
<i>Kocuria (Micrococcus luteus) rhizophila</i>	26.3	2.3(2–5)	NA
Other Gram-positive organisms	21.9 (19.9–22.7)	22.6 (14–31.2)	NA
Gram-negative organisms, median (IQR)			
<i>Escherichia coli</i>	19.8 (17.9–29)	5.5 (3–7.4)	0.02*
<i>Enterobacter cloacae</i> complex	27.9 (21.7–28.8)	NA	NA
<i>Klebsiella</i> species	27.7 (25.4–29.7)	6.5 (5.8–7.1)	NA
Other Gram-negative organisms	23.5 (19.8–26.7)	10.1(5.5–24.4)	0.23

NA, not applicable

* < 0.05, considered significant.