

years, our contamination rates were well below the CLSI recommendation; however, a significant reduction in blood culture contamination was observed after the use of a Red Top discard tube (0.8% vs. 1.9%) (Figures 1–3).

Fig. 1

| | 2017 (ChloraPrep + Red Top discard) | | | 2016 (ChloraPrep) | | |
|-----------|-------------------------------------|--------------|----------------|-------------------|--------------|----------------|
| | Volume | Contaminated | % Contaminated | Volume | Contaminated | % Contaminated |
| June | 1356 | 10 | 0.74 | 1059 | 25 | 2.36 |
| July | 1404 | 9 | 0.64 | 1227 | 24 | 1.96 |
| August | 1228 | 10 | 0.81 | 1240 | 17 | 1.37 |
| September | 1304 | 12 | 0.92 | 1643 | 34 | 2.07 |
| October | 1291 | 13 | 1.01 | 1433 | 31 | 2.16 |
| November | 1192 | 9 | 0.76 | 1424 | 25 | 1.76 |
| December | 1358 | 10 | 0.74 | 1232 | 22 | 1.79 |
| Total | 9233 | 73 | 0.8 | 9258 | 178 | 1.9 |

Fig. 2

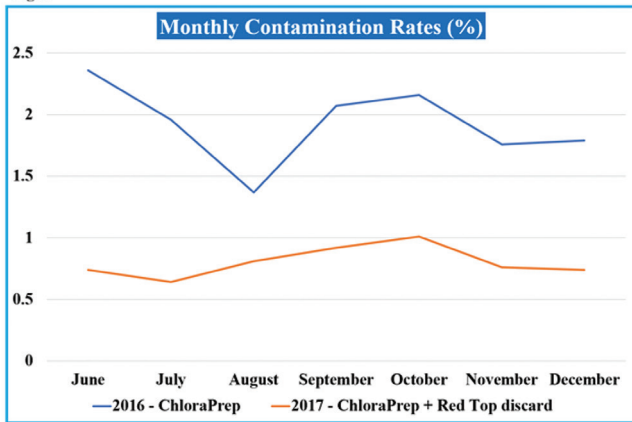
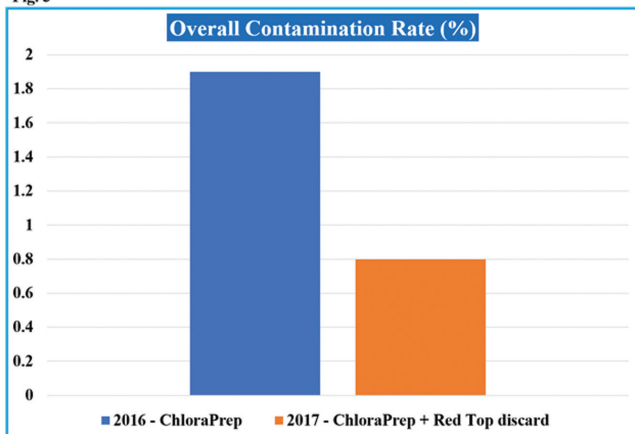


Fig. 3



Conclusion. The cost of a standard blood draw with Red Top tubes is minimal (few cents) while a single collection using an initial specimen diversion device (ISDD) can range from \$15 to \$18. During the course of this study, the use of a standard Red Top discard cost approximately \$456 (2017); if an ISDD was used instead, this would have generated \$136,995 in healthcare cost. At our institution, we were able to keep our contamination rates below 1% after the implementation of a standard Red Top discard tube. This suggests that the use of a Red Top discard prior to blood culture collection is an effective means for reducing and maintaining a low blood contamination rate.

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158. The Effect of Insurance Coverage on Appropriate Selection of Hospital Discharge Antibiotics for *Staphylococcus aureus* Bacteremia

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Background. Inappropriate or inadequate antimicrobial therapy of *Staphylococcus aureus* bacteremia (SAB) is associated with worsened outcomes. The impact of insurance coverage on appropriate selection of antibiotics is poorly understood. In patients diagnosed with SAB, we assessed the impact of insurance coverage on appropriate selection and duration of antibiotics at discharge.

Methods. We analyzed 273 patients who were diagnosed with SAB during their hospitalization at Nebraska Medicine and were discharged on antibiotics in 2015 and 2016. Antimicrobial therapy was deemed inappropriate if (i) total treatment duration was less than 14 days, (ii) oral delivery route was used, (iii) vancomycin was used to treat methicillin sensitive *S. aureus* in non-β-lactam allergic patients, or (iv) any penicillin or cephalosporin was used to treat methicillin-resistant *S. aureus*. Insurance was categorized broadly into (i) no insurance, (ii) Medicaid, (iii) Medicare, and (iv) Commercial. We collected data on a suite of additional variables that included: type of infectious disease (ID) inpatient consult and location of discharge. Logistic regression was used to determine the odds of being prescribed inappropriate therapy in univariate and multivariate analyses and likelihood ratio tests (LRT) were used to evaluate the strength of evidence.

Results. In unadjusted models, not having insurance was associated with inappropriate antimicrobial therapy (Reference Group: Any insurance; OR No insurance 4.71; LRT $P = 0.027$). Two additional risk factors for inappropriate therapy were identified in unadjusted models: discharge location (Reference Group: Nursing assistance; OR Home without assistance 3.37; 95% CI 1.34–8.46; LRT $P = 0.008$) and lack of an ID consult (Reference Group: Academic team; OR Not consulted 26.8, 95% CI 7.66–93.8; LRT $P < 0.001$).

Conclusion. We found strong evidence that not having insurance, being discharged to home without assistance, and not having an inpatient ID consult are risk factors for being prescribed inappropriate antimicrobial therapy for SAB upon hospital discharge; however, the sparsity of outcomes prevents us from drawing causal inferences. This study adds to the extensive body of evidence that has shown that uninsured patients tend to have suboptimal therapeutic choices.

Disclosures. All authors: No reported disclosures.

159. Genomic Epidemiology of MRSA at Intake to a Large Inner-City Jail: Evidence for Community Transmission Networks?

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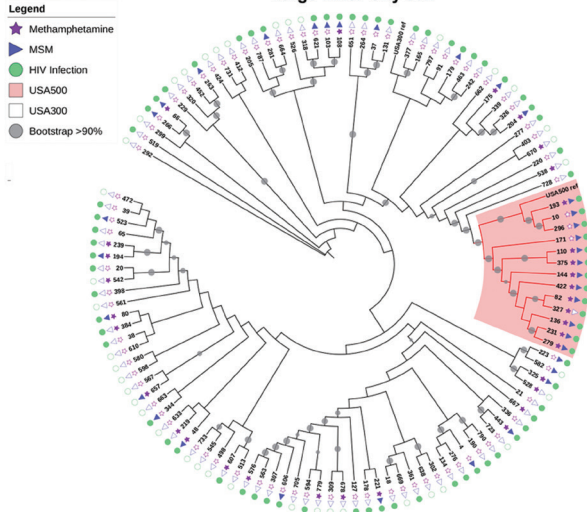
Background. USA300 is endemic in the community, with congregate settings potentially facilitating spread. The impact of community MRSA transmission networks on importation of MRSA into urban jails is unknown. We examined MRSA colonization isolates entering the jail and determined whether there are community transmission networks for MRSA that precede incarceration.

Methods. HIV-infected and HIV-negative males incarcerated at the Cook County Jail were enrolled within 72 hours of intake. Surveillance cultures (nares, throat, and groin) were collected to determine prevalence of MRSA colonization. A survey was administered to identify predictors of colonization. Whole-genome sequencing (WGS) and phylogenetic analysis were integrated with epidemiologic data to identify community transmission networks.

Results. A total of 800 males were enrolled (83% AA and 9% Hispanic); 58% were HIV-infected. The prevalence of MRSA colonization at intake was 19%. In multivariate analysis, methamphetamine use (METH), unstable housing, and prior jail incarceration were significant predictors of MRSA. Among HIV patients, injection drug use and HIV care at outpatient Clinic A that emphasize comprehensive care to the LGBTQ community were significant predictors of MRSA. Of the 31 (45%) patients with care at Clinic A, 14 had MRSA colonization. We sequenced 145 isolates from unique individuals, with 102 and 13 closely related to USA300 and USA500 reference genomes, respectively. USA300 strains from intake were diverse (median pairwise SNV distance = 109), with several small clusters noted. WGS revealed the high prevalence of MRSA in Clinic A was not due to clonal spread but rather an intermingling of distinct community transmission networks (strains were highly diverse; median pairwise SNV distance = 410). We did identify a 13-member community transmission network underlying spread of USA500 (figure). Members of this network were more likely to be HIV-infected ($P < 0.004$), MSM ($P < 0.001$), and METH ($P < 0.001$).

Conclusion. A high proportion of individuals enter jail already colonized with MRSA and colonization risk factors provide clues to community reservoirs for MRSA. WGS extended epidemiologic analysis and revealed community transmission networks that could be a potential focus for an intervention.

Genomic Epidemiology of MRSA Colonization Isolates at Entrance to a Large Inner-City Jail



Disclosures. All authors: No reported disclosures.

160. Reduction in the Spread of Hospital-Associated Infections Among Pediatric Oncology Patients in an Animal-Assisted Intervention Program from a Canine Decolonization Procedure

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Background. Animal-assisted interventions (AAI), the use of animals as a complementary therapy in holistic patient care, has shown many positive outcomes. However, therapy animals can serve as mechanical vectors of hospital-associated infections (HAI), e.g., methicillin-resistant *Staphylococcus aureus* (MRSA). This pilot study assessed for transmission of HAIs among therapy animals, patients, and the hospital environment. We tested the effectiveness of a novel decolonization protocol for therapy dogs to reduce the risk of transmission of HAIs and enhance AAI program sustainability. Our hypothesis was that HAI transmission occurs from positive child to child, with the dog as an intermediary fomite.

Methods. Before and after child-animal interaction, we sampled patients, dogs, and the environment, and collected vital statistics and survey data from patients. MRSA was detected in samples by culture and molecular testing. Therapy dog handlers performed normal pre-visit practices for 2 control visits, then switched to a decolonization protocol (chlorhexidine-based shampoo prior to the visit, and chlorhexidine wipes on the fur during the visit) for 2 intervention visits.

Results. We evaluated 45 children and 4 therapy dogs over 13 visits. Children had decreased blood pressure and heart rate, and reported improved mental health scores post visit. MRSA conversion was identified from 10.2% of the children and 38.5% of the dogs, while 93% of the environmental samples were MRSA positive both pre and post. Patients that interacted closely with the dog had 8.01 times higher odds (95% CI 1.1–15.2) of MRSA conversion compared with patients who barely interacted with the dog. When stratified by intervention group, the MRSA conversion odds ratio of close interaction was 0.93 (95% CI 0.1–10.8) when the dog was decolonized versus 9.72 (0.9–99) when not decolonized.

Conclusion. This study showed the potential for AAI visits to improve physiological and mental health of pediatric outpatients. A risk of HAI exposure to patients from interaction with the dog was found, but this effect was nullified by the decolonization procedure. Future research is needed to increase the safety of this valuable alternative therapy.

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161. Prevalence and Risk Factors for *Candida auris* Colonization Among Patients in a Long-term Acute Care Hospital—New Jersey, 2017

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Background. *Candida auris* can be transmitted in healthcare settings, and patients can become asymptotically colonized, increasing risk for invasive infection and transmission. We investigated an ongoing *C. auris* outbreak at a 30-bed long-term acute care hospital to identify colonization for *C. auris* prevalence and risk factors.

Methods. During February–June 2017, we conducted point prevalence surveys every 2 weeks among admitted patients. We abstracted clinical information from medical records and collected axillary and groin swabs. Swabs were tested for *C. auris*. Data were analyzed to identify risk factors for colonization with *C. auris* by evaluating differences between colonized and noncolonized patients.

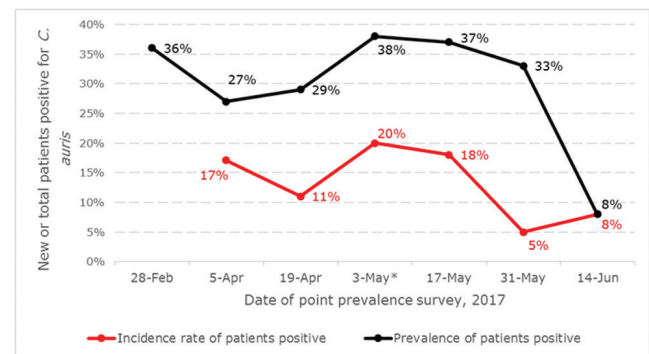
Results. All 101 hospitalized patients were surveyed, and 33 (33%) were colonized with *C. auris*. Prevalence of colonization ranged from 8% to 38%; incidence ranged from 5% to 20% (figure). Among colonized patients with available data, 19/27 (70%) had a tracheostomy, 20/31 (65%) had gastrostomy tubes, 24/33 (73%) ventilator use, and 12/27 (44%) had hemodialysis. Also, 31/33 (94%) had antibiotics and 13/33 (34%) antifungals during hospitalization. BMI for colonized patients (mean = 30.3, standard deviation (SD) = 10) was higher than for noncolonized patients (mean = 26.5, SD = 7.9); $t = -2.1$; $P = 0.04$. Odds of colonization were higher among black patients (33%) vs. White patients (16%) (odds ratio [OR] 3.5; 95% confidence interval [CI] 1.3–9.8), and those colonized with other multidrug-resistant organism (MDRO) (72%) vs. noncolonized (44%) (OR 3.2; CI 1.3–8.0). Odds of death were higher among colonized patients (OR 4.6; CI 1.6–13.6).

Conclusion. Patients in long-term acute care facilities and having high prevalences of MDROs might be at risk for *C. auris*. Such patients with these risk factors could be targeted for enhanced surveillance to facilitate early detection of *C. auris*. Infection control measures to reduce MDROs' spread, including hand hygiene, contact precautions, and judicious use of antimicrobials, could prevent further *C. auris* transmission.

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Figure 1: Incidence and prevalence of *Candida auris* colonization by point prevalence survey (PPS), at a long-term acute care hospital, New Jersey, February 28 and June 14, 2017 ($N = 101$).



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162. Association Between Antibiotic Use and Multidrug-Resistant Organism Detection in Advanced Cancer Patients on Palliative Chemotherapy

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Background. Data suggest end-of-life antibiotics predispose to multidrug-resistant organism (MDRO) acquisition in intensive care units (ICUs). Less is known regarding antibiotics and MDRO acquisition in other palliative care populations.

Methods. We conducted a nested case-control study of advanced cancer patients aged ≥ 65 years started on palliative chemotherapy from January 2016 to September 2017 at Yale New Haven Hospital. We identified patients with (cases) and without (controls) new MDRO detected from clinical or surveillance cultures from the first hospitalization after starting palliative chemotherapy. All patients had no history of MDRO, and 3 controls were randomly selected per case. Antibiotic use was defined as