Conclusions: Findings suggest that patients with certain easily assessed characteristics, such as mechanical ventilation, tracheostomy, or stool incontinence or who require bathing assistance, may be associated with CPO positivity during screening. Further data collection and analysis of such risk factors may provide insight for the development of more targeted admission and contact screening strategies.

Funding: None

Disclosures: None

Presentation Type: Poster Presentation

Evaluation of Sampling Methods for Detection of Pathogens from Steel Surfaces; Contact Plates, Foam Swabs, and Flocked Swabs

Rolieria West, Centers for Disease Control and Prevention; Laura Rose, Centers for Disease Control and Prevention; Judith Noble-Wang, Centers for Disease Control and Prevention

Background: Contaminated healthcare surfaces can serve as reservoirs for the transmission of pathogens. Sensitive sampling methods are needed to investigate sources of pathogens for implementing effective disinfection strategies and thereby preventing environmental transmission. Conventional approaches employ swabs to sample environmental surfaces. Contact plates represent an alternative approach for sampling healthcare surfaces that does not require lab processing, though little is known about their performance. A contact plate is an agar plate that is overfilled with selective or nonselective media. It can be gently applied to the surface, then simply incubated at a temperature optimal for target organism(s), thus saving time and resources. Methods: In this study, contact plates containing trypticase soy agar with 5% sheep blood (TSAI1), foam swabs, and flocked swabs were evaluated for their ability to recover 4 pathogens that persist on healthcare surfaces. Stainless-steel coupons (4 in²) were inoculated with the following pathogens (10⁵ CFU): Acinetobacter baumannii (AB, strain type 12), carbapenemase-producing KPC+ Klebsiella pneumoniae (KP; ATCC BAA-1705); methicillin-resistant Staphylococcus aureus (MRSA; ATCC 43300); and vancomycin-resistant Enterococcus faecalis (VRE; Van A + 256). The plates were allowed to dry 1 hour. Sampling with CPs was performed in 2 ways; (1) a single contact plate was used to sample 1 stainless-steel surface and (2) a composite was collected by 3 sequential contact-plate samples of the same stainless-steel surface. The contact plates were then incubated at 35±1°C. Foam and flocked swabs were premoistened with phosphate-buffered saline + 0.02% polysorbate 80 (PBST) and were used to sample the stainless-steel coupons. Swabs were held for 1 hour and processed by sonication and vortexing in 5 mL of PBST, then the eluent was cultured and CFU counted. Mean percentage recoveries (%R) relative to the inoculum were calculated and compared. Results: When the %R for all 4 pathogens were pooled, the composite contact-plate sampling method yielded the highest, (P < .05) (66.0%; SD, 0.22), followed by the single contact plate method (39.7%; SD, 0.12), foam swab (32.9%; SD, 0.18), and flocked swab (20.3%; SD, 0.20). The composite contact plate method yielded the highest %R for VRE (102.1%; SD, 0.17), and the lowest %R was observed when using flocked swabs to recover KP (6.3%; SD, 0.05). Conclusions: The contact-plate composite method may provide investigators with minimal environmental microbiology capacity an alternative method for environmental sampling and detection of organisms from surface areas (≤4 in 2) with low bioburden.

Funding: None

Disclosures: None

Presentation Type: Poster Presentation

Evaluation of Surgical Site Infections After Change in Surgical Prophylaxis in VAD Patients

Emily Min, University of Maryland School of Medicine; Timileyin Adediran, University of Maryland School of Medicine; Kerri Thom, University of Maryland School of Medicine; Emily Heil, University of Maryland School of Pharmacy

Background: In October 2013, the University of Maryland Medical Center established a formal antibiotic prophylaxis protocol for patients undergoing ventricular assist device (VAD) placement, replacing a previous system of various broad-spectrum antibiotic combinations typically for prolonged durations based on surgeon preference. This new protocol consisted of a standardized regimen of 72 hours of vancomycin and ceftriaxone after the procedure. The objective of this project was to evaluate the rate of surgical site infection (SSI) related to VAD placement to ensure that implementing the new protocol did not cause an increase in SSI rates. Methods: The study was a retrospective cohort study of patients who had undergone VAD placement before the protocol change (January 1, 2011, to October 1, 2013) and after the change (October 1, 2013, to November 15, 2015). The primary outcomes were the difference in SSI rate before and after the protocol change using CDC NHSN definitions. Pertinent data points of interest included reason for VAD placement, duration/type of antibiotics used, delayed sternal closure, SSI, characterization of infection (bloodstream, driveline, or pocket), organism identified on culture and mortality at 30 days and 1 year. SSI rates were assessed using the Fischer exact test, and descriptive statistics were used for other outcome variables. Results: In total, 75 patients were included before the protocol and 46 after the protocol change. Overall, 27% and 17% of patients were on therapeutic antibiotics prior to the VAD placement, respectively (P = .233). Also, 8 (6.6%) patients in the preintervention group had an SSI compared to 1 patient (0.8%) in the postintervention group (P = .15). Adherence to the protocol was suboptimal, with 27% of patients in the postintervention group receiving non–protocol-adherent antibiotics and 65% of patients receiving antibiotics >96 hours postoperatively. When evaluating the patients collectively, SSI rates were the same when antibiotics were discontinued <72 hours postoperatively versus when antibiotics were continued beyond 72 hours postoperatively or were not given at all postoperatively (3.1% vs 10.7% vs 0%; P = .243). SSI rates were also no different among patients who received ceftazolin monotherapy (0%), vancomycin and ceftriaxone (2.7%), vancomycin and piperacillin tazobactam (2%), and other antibiotic combinations (7.7%) for surgical prophylaxis (P = .1). Conclusions: No change in SSI rates was noted after a protocol change narrowing the spectrum and duration of antibiotic prophylaxis was implemented. Evaluation of optimal surgical prophylaxis in this patient population is difficult due to low event rates and frequent therapeutic indications for antibiotics outside the standard prophylaxis. Despite these challenges, this study supports the safety of studying SSI prophylaxis reduction
in the VAD population. Further studies are reasonable and warranted.

**Funding:** None

**Disclosures:** None

Doi:10.1017/ice.2020.779

---

**Presentation Type:** Poster Presentation

**Evaluation of the bioMérieux EPISEQ-CS Software for wgMLST-Based Bacterial Strain Typing**

Amorce Lima, Tampa General Hospital; Steven Bruzek, Tampa General Hospital; Amanda Lasher, Tampa General Hospital; Grant Vestal, Tampa General Hospital; Suzane Silbert, Tampa General Hospital

**Background:** Whole-genome sequencing (WGS) is becoming the method of choice for outbreak analysis of microbial pathogens. However, the main challenge with WGS for microbial strain typing is the conversion of raw sequencing data to actionable results for epidemiology and surveillance analysis. We evaluated the bioMérieux EPISEQ-CS, a cloud-based WGS data analysis software for outbreak detection to compare the results for 4 groups of different species previously characterized by strain typing and commonly isolated in hospital-acquired infections. **Methods:** In total, 30 methicillin-resistant *Staphylococcus aureus* (MRSA), 15 *Clostridiodes difficile* (CDIFF), 17 *Pseudomonas aeruginosa* (PSA), and 10 *Acinetobacter baumannii* (ACB) isolates were included in this study. All isolates had been previously characterized by rep-PCR using the DiversiLab system (bioMérieux, France) and saved at ~70°C. Before testing, samples were thawed and plated, and DNA extraction was performed on the QIAcube (Qiagen, Hilden, Germany) using the DNEasy Ultra Clean Microbial kit extraction protocol. DNA libraries were prepared using the Nextera DNA Flex Kit and sequenced on the Illumina iSeq100 platform according to manufacturer’s recommendations (Illumina, San Diego, CA). Generated sequences were uploaded into EPISEQ-CS, and wgMLST-based analysis was performed. We compared clusters generated by the DiversiLab system and EPISEQ-CS. **Results:** DiversiLab identified 9 MRSA clusters among 30 isolates. EPISEQ-CS reclassified 14 of 30 isolates into 5 MRSA clusters and the remaining 16 isolates were unrelated. DiversiLab identified 2 CDIFF clusters among 15 isolates. EPISEQ-CS reclassified 3 isolates into 1 CDIFF cluster and determined the remaining 12 to be unrelated. DiversiLab identified 5 PSA clusters among 17 isolates, whereas EPISEQ-CS reclassified all 17 isolates as unrelated. DiversiLab identified 2 ACB clusters among 10 isolates, whereas EPISEQ-CS reclassified 2 ACB isolates into 1 cluster and determined 8 to be unrelated. Analysis using Simpson’s diversity index (D) suggested that the EPISEQ-CS showed increased diversity when compared to DiversiLab clustering across all bacterial species analyzed in this study. **Conclusions:** EPISEQ-CS enabled a comprehensive wgMLST analysis, including quality control and comparative epidemiological analysis, thereby providing a more reliable method for bacterial strain typing. As WGS becomes more affordable and applicable to routine epidemiological surveillance, EPISEQ-CS provides an informative tool in the monitoring of hospital-acquired infections.

**Funding:** None

**Disclosures:** None

Doi:10.1017/ice.2020.780

---

**Presentation Type:** Poster Presentation

**Evolution and Associated Factors of Hand Hygiene Compliance in the Surgical Areas of a Tertiary-Care Hospital**

Natividad Algado-Sellés, Alicante Institute for Health and Biomedical Research (ISABIAL). Epidemiology Unit, Preventive Medicine Department, General University Hospital of Alicante, Alicante, Spain; Paula Gras-Valentí, Alicante Institute for Health and Biomedical Research (ISABIAL). Epidemiology Unit, Preventive Medicine Department, General University Hospital of Alicante, Alicante, Spain; Juan Gabriel Mora-Muriel, Alicante Institute for Health and Biomedical Research (ISABIAL), Epidemiology Unit, Preventive Medicine Department, General University Hospital of Alicante, Alicante, Spain; Pablo Chico-Sánchez, Alicante Institute for Health and Biomedical Research (ISABIAL), Epidemiology Unit, Preventive Medicine Department, General University Hospital of Alicante, Alicante, Spain; Edith Leutscher-Vasen, Alicante Institute for Health and Biomedical Research (ISABIAL), Epidemiology Unit, Preventive Medicine Department, General University Hospital of Alicante, Alicante, Spain; Marina Fuster-Pérez, General University Hospital of Alicante; Sandra Canovas-Javeaga, Alicante Institute for Health and Biomedical Research (ISABIAL), Epidemiology Unit, Preventive Medicine Department, General University Hospital of Alicante, Alicante, Spain; Cesar O. Villanueva-Ruiz, Alicante Institute for Health and Biomedical Research (ISABIAL), Epidemiology Unit, Preventive Medicine Department, General University Hospital of Alicante, Alicante, Spain; Natali J. Jimenez-Sepulveda, Alicante Institute for Health and Biomedical Research (ISABIAL), Epidemiology Unit, Preventive Medicine Department, General University Hospital of Alicante, Alicante, Spain; Ana S. Lameiras-Azevedo, Alicante Institute for Health and Biomedical Research (ISABIAL), Epidemiology Unit, Preventive Medicine Department, General University Hospital of Alicante, Alicante, Spain; Maria Hernandez-Maldonado, Alicante Institute for Health and Biomedical Research (ISABIAL), Epidemiology Unit, Preventive Medicine Department, General University Hospital of Alicante, Alicante, Spain; Jose Sanchez-Paya, Epidemiology Unit, Preventive Medicine Department, Alicante University General Hospital, Alicante, Spain

**Background:** Hand hygiene (HH) is the most important measure for preventing healthcare-associated infections. The objective is to gain insight into the evolution of the degree of compliance with recommendations (DCR) on HH and its associated factors in the surgical areas of a tertiary-care hospital. **Methods:** This observational, cross-sectional study, was repeated over time, with direct observation of the DCR on HH during the daily activity of healthcare workers in surgical areas: general surgery, urology, vascular surgery, traumatology, neurosurgery, thoracic surgery, heart surgery, pediatric surgery, otorhinolaryngology, gynecology and obstetrics, ophthalmology. Over 14 years (from 2005 to 2018), 15,946 HH opportunities were registered, together with different additional variables (age, sex, professional position, surgical area . . .). The χ² test was used to study the association and the crude, and adjusted odds ratios were used to quantify its magnitude. **Results:** The DCR on HH in surgical areas was 49.7% (95% CI, 48.9%–50.5%), and in the group of nonsurgical areas it was