

2020. Source investigation, contact tracing, and serial testing revealed 3 additional patients, and 8 HCWs. One HCW also had a community exposure. Patient median age was 60 years (range, 48–68) and all were male. In total, 11 samples (4 patients and 7 HCWs) were sequenced. Using WGS, cluster 1 was confirmed to be an outbreak: WGS showed 0–5 mutations in between samples. Cluster 2 was also an outbreak: WGS showed less diversity (0–3 mutations) and ruled out the HCW with a community exposure (20 mutations of difference). **Conclusion:** Whole-genome sequencing confirmed the outbreaks identified using classic epidemiologic methods. Serial testing allowed for early outbreak detection. Early outbreak detection and implementation of control measures may decrease outbreak size and genetic diversity.

Funding: No

Disclosures: None

Antimicrobial Stewardship & Healthcare Epidemiology 2021;1(Suppl. S1):s10–s11

doi:10.1017/ash.2021.19

Presentation Type:

Poster Presentation - Top Poster Award

Subject Category: COVID-19

COVID-19 Vaccination of HCWs in the First Phase of a Large-Scale Mass Vaccination Program within a Healthcare System

Kimberly Korwek; E. Jackie Blanchard; Julia Moody; Katherine Lange; Ryan Sledge; Maria Benedetti and Kenneth Sands

Background: The approval of the first SARS-COV-2 vaccines for COVID-19 were accompanied by unprecedented efforts to provide vaccination to healthcare workers and first responders. More information about vaccine uptake in this group is needed to better refine and target educational messaging.

Methods: HCA Healthcare used federal guidance and internal experience to create a systemwide mass vaccination strategy. A closed point-of-dispensing (POD) model was developed and implemented. The previously developed enterprise-wide emergency operations strategy was adapted and implemented, which allowed for rapid development of communications and operational processes. A tiering strategy based on recommendations from the National Academies was used in conjunction with human resources data to determine vaccine eligibility for the first phase of vaccination. A comprehensive data and reporting strategy was built to connect human resources and vaccine consent data for tracking vaccination rates across the system. **Results:** Vaccination of employed and affiliated colleagues began December 15, 2020, and was made available based on state-level release of tiers. Within the first 6 weeks, in total, 203,544 individuals were eligible for vaccine based on these criteria. Of these, 181,282 (89.1%) consented to and received vaccine, 19,788 (9.7%) declined, and 2,474 (1.2%) indicated that they had already been vaccinated. Of those eligible, the highest acceptance of vaccine was among the job codes of specialists and professionals (n = 7,914 total, 100% consent), providers (n = 23,335, 99.6%), and physicians (n = 3,218, 98.4%). Vaccine was most likely to be declined among job codes of clerical and other administrative (n = 12,889 total, 80.1% consent), clinical specialists and professionals (n = 22,853, 81.0%), and aides, orderlies and technicians (n = 17,803, 82.6%). Registered nurses made up the largest eligible population (n = 56,793), and 89.5% of those eligible consented to receive vaccination. Average age among those who consented was slightly older (48.3 years) than those that declined (44.7 years), as was length of employment tenure (6.9 vs 5.0 years).

Conclusion: A large-scale, closed POD, mass vaccination program was able to vaccinate nearly 200,000 healthcare workers for SARS-CoV-2 in 6 weeks. This program was implemented in acute-care sites across 20 different US states, and it was able to meet the various state-level requirements for management of processes, product, and required reporting. The development of a standardized strategy and custom, centralized monitoring and reporting facilitated insight into the characteristics of early vaccine adopters versus those who decline vaccination. These data can aid in the refining and targeting of educational materials and messaging about the SARS-CoV-2 vaccine.

Funding: No

Disclosures: None

Antimicrobial Stewardship & Healthcare Epidemiology 2021;1(Suppl. S1):s11

doi:10.1017/ash.2021.20

© The Author(s), 2021. Published by Cambridge University Press on behalf of The Society for Healthcare Epidemiology of America. This is an Open Access article, distributed under the terms of the Creative Commons Attribution licence (<http://creativecommons.org/licenses/by/4.0/>), which permits unrestricted re-use, distribution, and reproduction in any medium, provided the original work is properly cited.

Presentation Type:

Poster Presentation - Top Poster Award

Subject Category: Disinfection/Sterilization

High-Level Disinfection in Ambulatory Care: Overcoming the Barriers of a Decentralized System through Auditing and Education

Sonja Rivera Saenz

Background: High-level disinfection (HLD) of semicritical instruments in a multispecialty ambulatory care network has the potential for increased risk due to the decentralized instrument reprocessing and lack of a sterile processing department. Attention to HLD practices is an important part of device-borne outbreak prevention. **Method:** An HLD database was developed to identify specific departments and locations where HLD occurred across a 30-medical practice ambulatory care network in eastern Massachusetts, which included otolaryngology, urology, endoscopy, and obstetrics/gynecology departments. Based on qualitative feedback from managers and reprocessing staff, this database centralized information that included the supply inventory including manufacturer and model information, HLD methodology, standard work, and listing of competency evaluations. The infection control team then led audits to directly observe compliance with instrument reprocessing and a monthly-driven HLD calendar was developed to enforce annual competencies. **Result:** The results of the audits demonstrated variability across departments with gaps in pre-cleaning, transportation of used instruments, the dilution of enzymatic cleaner, and maintenance of quality control logs. Given the uniqueness of shape and size of various ambulatory locations, proper storage and separation between clean and dirty spaces were common pitfalls. Auditing also revealed different levels of staff understanding of standard work and variable inventory management. Centralized education sessions held jointly by the infection control team and various manufacturers for the reprocessing staff helped to create and reinforce best practices. **Conclusion:** Decentralized HLD that occurs across multiple ambulatory care sites led to gaps in instrument reprocessing and unique challenges due to variable geography of sites, physical space constraints, and an independent approach to procuring medical supplies. Through the auditing and feedback of all areas that perform HLD, an effective and sustainable strategy was created to ensure practice improvement. Streamlining standard work, seeking direct input from frontline staff, and collective educational events were critical to our success in the ambulatory setting.

Funding: No

Disclosures: None

Antimicrobial Stewardship & Healthcare Epidemiology 2021;1(Suppl. S1):s11

doi:10.1017/ash.2021.21

Presentation Type:

Poster Presentation - Top Poster Award

Subject Category: Long-Term Care

Coronavirus in Wisconsin Nursing Homes: A Longitudinal Analysis of the First 10 Months of the Pandemic

Cameron Gmehlin; Frida Rivera; Jorge Ramos-Castaneda; Liliana Pezzin; Edmund Duthie and L. Silvia Munoz-Price

Background: The COVID-19 pandemic has disproportionately affected nursing home residents, and emerging evidence suggests quality, location, resident demographics, and staffing levels may be related to COVID-19 incidence within facilities. We describe the distribution of COVID-19 cases in Wisconsin nursing homes from January 2020 to October 2020, the effect of rural versus urban locations on COVID-19 incidence, and the temporal changes in COVID-19 incidence. **Methods:** We constructed a database using the Center for Medicaid and Medicare Services' (CMS) publicly available data. Variables obtained per facility included location, number of beds, ownership type, average census, 5-star ratings (overall, quality, health, staffing, and nurse staffing categories), number of COVID-19 cases, resident Medicaid/Medicare share, area deprivation index, and social vulnerability index. Nursing homes were divided into tertiles based on total COVID-19 cases for descriptive analysis (zero cases, 1–7 cases, >7 cases). Demographic and clinical variables were reported as frequencies, mean