

2023-022 Investigation of a COVID-19 Cluster in the Basement of a Local Hospital in Singapore from 30 August to 16 September 2021

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Background	Results
 Primary case of the cluster developed symptoms on 30 August 2021. (Figure 1) 	 58% of those affected were outsourced maintenance staff
 Symptom onset in 72% of secondary cases occurred from 1 – 3 September. 	 Substantial time spent in common areas such as workshop, rest area and male toilet likely caused propagation of infection Household transmission was also likely as some staff lived together across 6 households in rented apartments
 Subsequent sporadic cases were identified till 16 September. (Figure 1) Of the 57 staff in this cluster. 6 did not work in the 	

- basement
 - Visited the basement or had social interactions with staff who worked there.
- All staff were fully vaccinated at the time of investigation.

Figure 1. Epidemic Curve showing onset of illness in CGH Staff with epidemiological and phylogenetic linkage (n=57)



- Transmission to other areas in the basement
 - > Visits by maintenance staff
 - > Poor ventilation in basement
 - Lack of safe distancing
 - > Unmasked interactions in crowded common areas
- 32 phylogenetically sequenced cases belonged to the same parent node, supporting findings from the investigation. (Figure 2)

Figure 2. Part of the phylogenetic tree showcasing selected cases from the cluster that were sequenced

- with B1 staff
- Yellow cases work in B1
- Blue cases work in other parts of CGH but had visits to B1 or social interactions
- 32 out of 57 staff cases were sequenced.

C+ on 2nd Sept

■ BME ■ Maintainence ■ Lab ■ FM ■ ES ■ DFS ■ ALPS ■ Renal Dialysis Unit ■ Inpatient Ops

Methods

- Site visit conducted to identify risk factors of transmission
- CCTV footage from the corridors and staff clock in/clock out areas was reviewed.
- Interviews conducted with selected staff to understand
 - > Staff movements
 - > Type & nature of interactions
 - \succ Staff activities surrounding the time of the outbreak.
- Further details were obtained with assistance provided by in-house staff.

- All of the above cases belonged to the same parent node (Node 2691)
- Sequences that fall under the branch of the main node, with SNP differences, indicate an ancestraldescendant relationship

Conclusion

- Air and droplet, fomite and multiple mode **exposures** accounted for the transmission of infection within this cluster
- Poor ventilation facilitated rapid spreading of infection

Recommendations

- Air handling unit (AHU) filters and exhaust mechanisms upgraded to improve ventilation
- Increased disinfection of common areas daily
- Non-touch clock in/clock out points introduced

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Split team arrangements to access high traffic points

Improved education

Enforcement of infection prevention and control

measures

Staff to be up-to-date on COVID-19 vaccination