

Linezolid-resistant *Staphylococcus epidermidis* at a tertiary hospital, Copenhagen, Denmark

Investigating an outbreak of Healthcare-Associated Infections (HAI) between 2017-2021

REGION

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HEALTHCARE-ASSOCIATED BLOODSTREAM INFECTIONS¹

- *S. epidermidis* accounts for more than 75% of CoNS found in clinical specimens
- It is essential to distinguish between common contaminations and significant infections
- *S. epidermidis* possesses different determinants to facilitate survival, biofilm formation, adhesion to tissue and prosthetic surfaces and even serve as a reservoir for the transfer of virulence-related genetic information to *S. aureus*.
- Patients particularly at risk of significant CoNS infections include those with prosthetic devices and immunocompromised hosts.
- CoNS is the most common cause of nosocomial bloodstream infections, approximately one-third of those found in ICUs, with 15-25% having significant bloodstream infections.
- Health care-associated infections affect 7.1 % of all hospitalised patients in tertiary hospitals in the European Union.
- In the USA, 55% of ICU patients have a CVC.
- Studies point towards CoNS as responsible for at least 12% of all cases; among these pathogens, antimicrobial infections are a severe problem.
- CVC-associated bloodstream infections have a crude excess mortality rate in adult patients in ICU of 23.6% for those with central line-associated bloodstream infections.
- A German study estimated an additional 12.5 days of hospital stay.
- Estimated costs of HAI are approximately 7 billion euros annually.

INTRODUCTION

Linezolid-resistant *Staphylococcus epidermidis* (LRSE) is increasingly reported throughout the western world and is considered an important pathogen for immunocompromised patients and patients with prosthetic devices.

METHOD

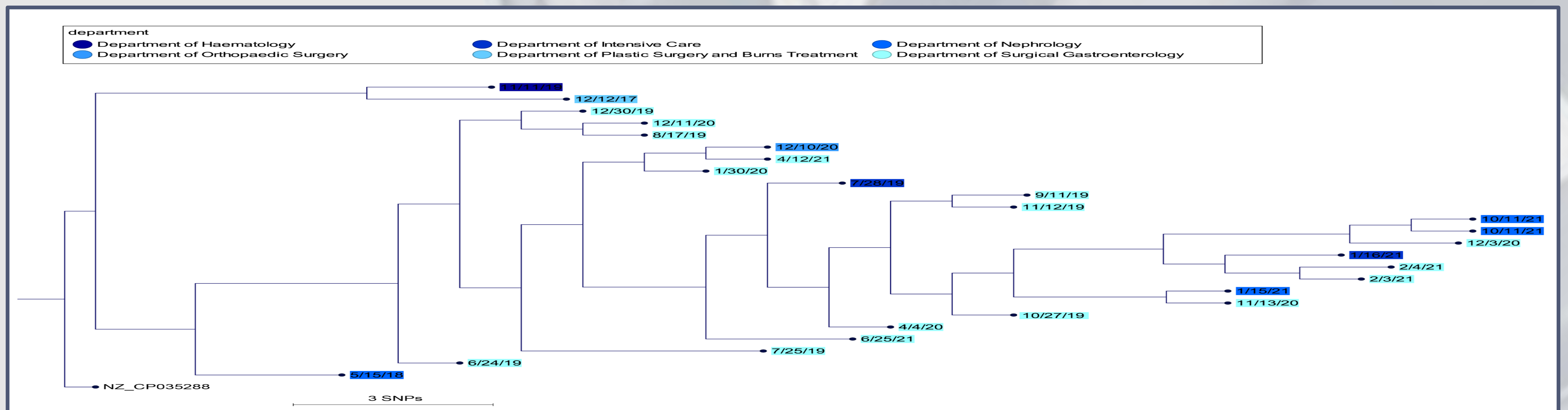
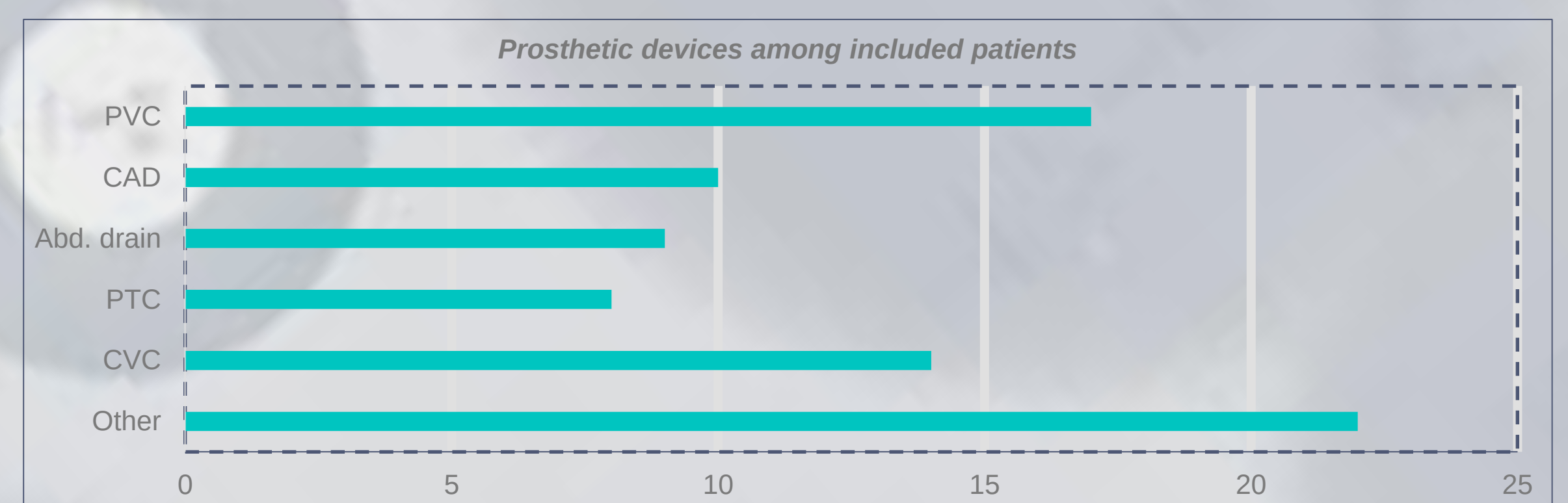
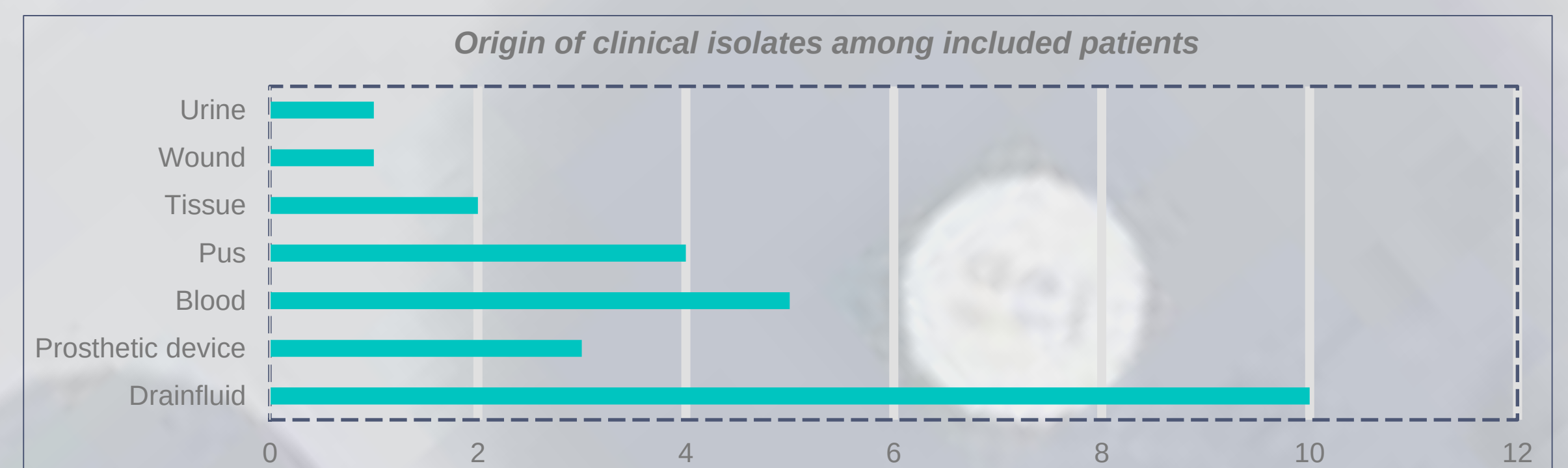
Between 2017 and 2021, 26 clinical isolates were identified by routine cultivation and susceptibility testing, followed by whole-genome sequencing (WGS) on an Illumina platform. Bacdist performed SNP-analysis and phylogenetic tree. Environmental swabs and linezolid (LZD) consumption were analysed during the outbreak.

RESULTS & DISCUSSION

All isolates of LRSE were confirmed to belong to the clone ST22 with 1-60 SNP variations. Most patients had multiple prosthetic devices, which are thought to play an important portal of entry for LRSE, though LRSE was not found on any environmental swabs. During the outbreak, 0.15% of hospitalised patients at University Hospital of Copenhagen, Rigshospitalet were given LZD, and the Department of Surgical Gastroenterology constituted 21.4% of the overall consumption of Rigshospitalet.

CONCLUSION

The molecular epidemiology of the outbreak showed an HAI outbreak with the same clone of LRSE. LZD consumption and lack of infection control are thought to play an important role.



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