

Microbiologic contamination present in mops and cloths used for cleaning procedures in firefighters' headquarters

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Outline

1. Background

2. Methods

3. Results

4. Take home messages

1. Background

Several occupational environments in Portugal have already been characterized regarding the *Aspergillus* genus prevalence, demonstrating its critical dissemination indoors (Viegas et al. 2019-2021).

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Aspergillus spp. prevalence in Primary Health Care Centres: Assessment by a novel multi-approach sampling protocol




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Air Quality, Atmosphere & Health
<https://doi.org/10.1007/s11869-019-00781-x>

Aspergillus spp. burden on filtering respiratory protective devices. Is there an occupational health concern?









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Article

Azole-Resistant *Aspergillus fumigatus* Harboring the TR₃₄/L98H Mutation: First Report in Portugal in Environmental Samples

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JOURNAL OF OCCUPATIONAL AND ENVIRONMENTAL HYGIENE
<https://doi.org/10.1080/15459624.2020.1834113>



SHORT REPORT



Aspergillus spp. presence on mechanical protection gloves from the waste sorting industry

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An assessment of Portuguese firefighters' ambulances identified hazardous levels of *Aspergillus* section *Fumigati* in ambulance air, which would be able to reach the alveoli (Viegas et al. 2021).



There were other relevant findings, such as:

- toxigenic fungi with clinical relevance found in ambulance air
- contamination of surfaces increased after cleaning at some sites
- mycotoxins detected in mops and electrostatic dust cloths

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Bioburden contamination and *Staphylococcus aureus* colonization associated with firefighter's ambulances

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This study aimed to characterize the microbial contamination (bacteria and fungi focusing on *Aspergillus* section *Fumigati*) through passive sampling methods in 11 firefighters' headquarters of Lisbon.

2. Methods

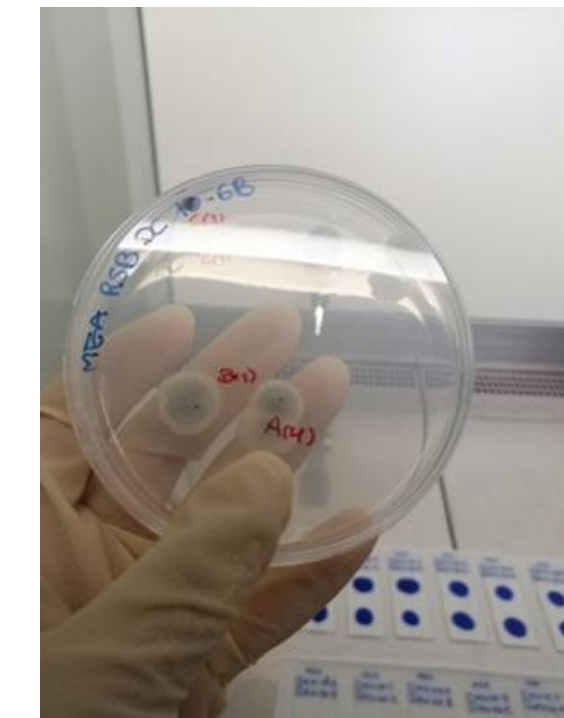
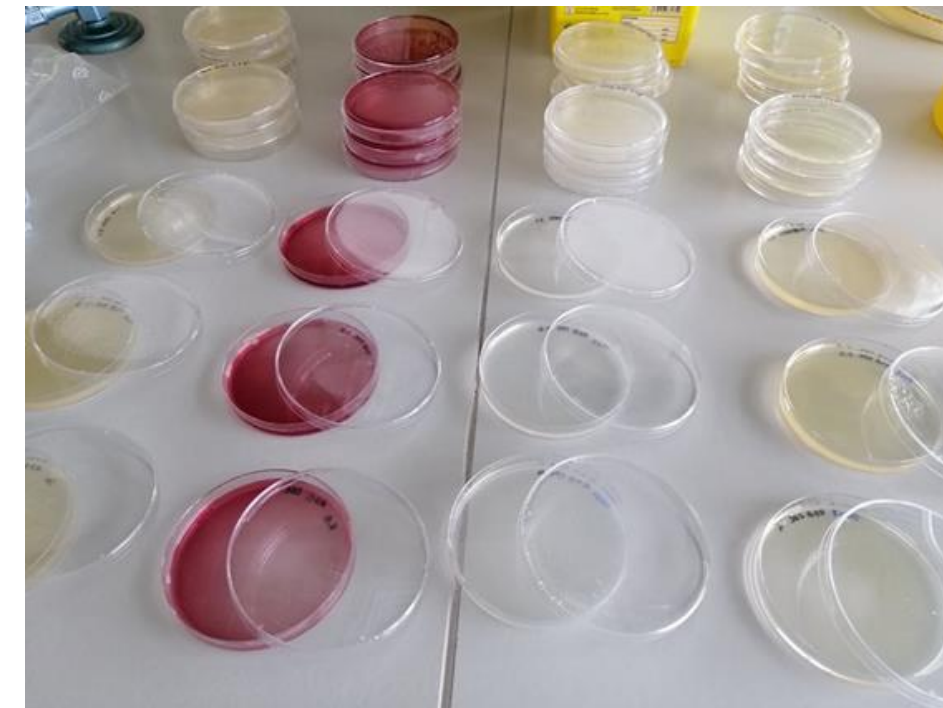


N = 14 mops



N = 25 cleaning cloths

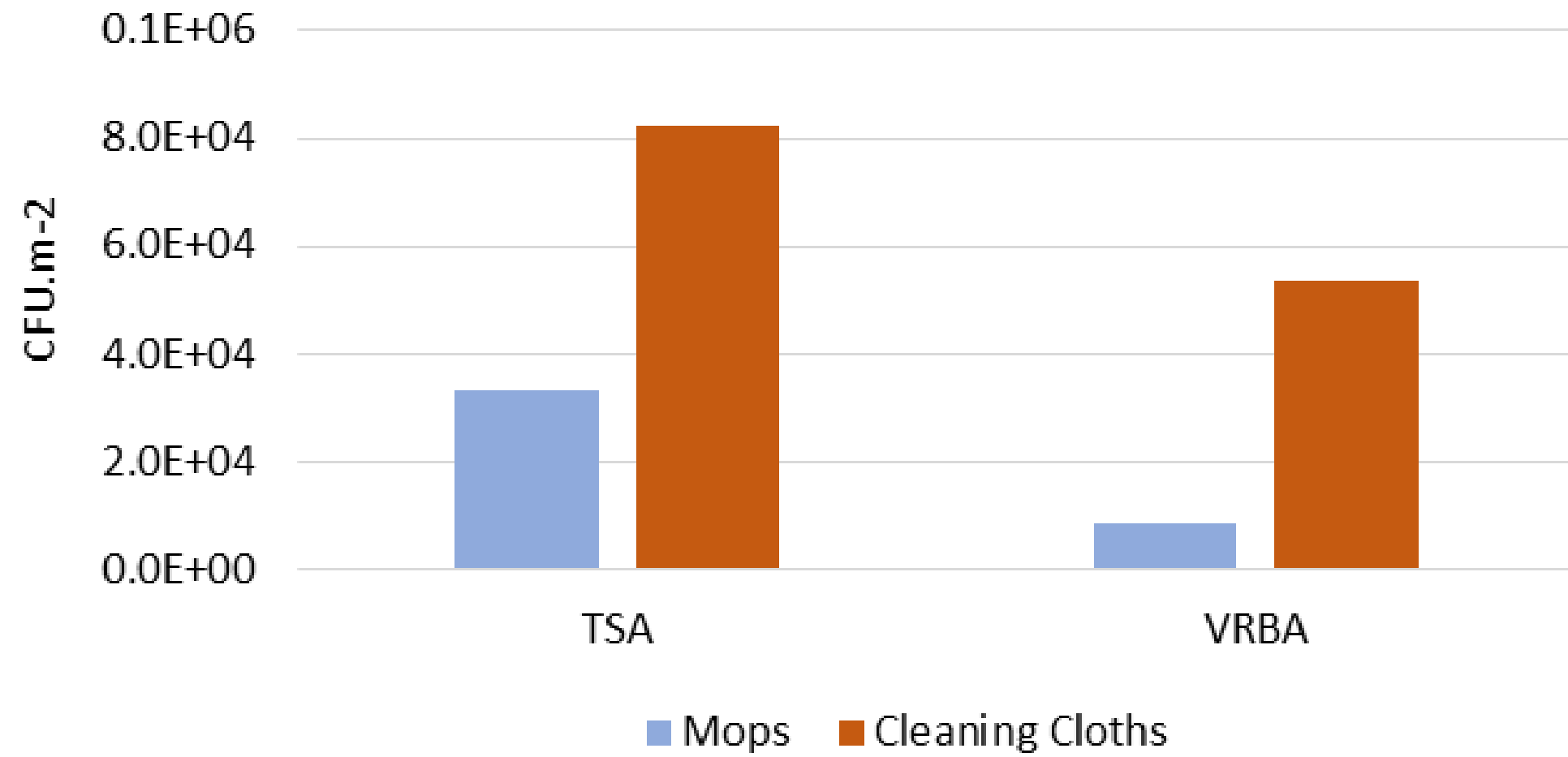
- Viable bioburden through culture-based methods and azole resistance profile were obtained.



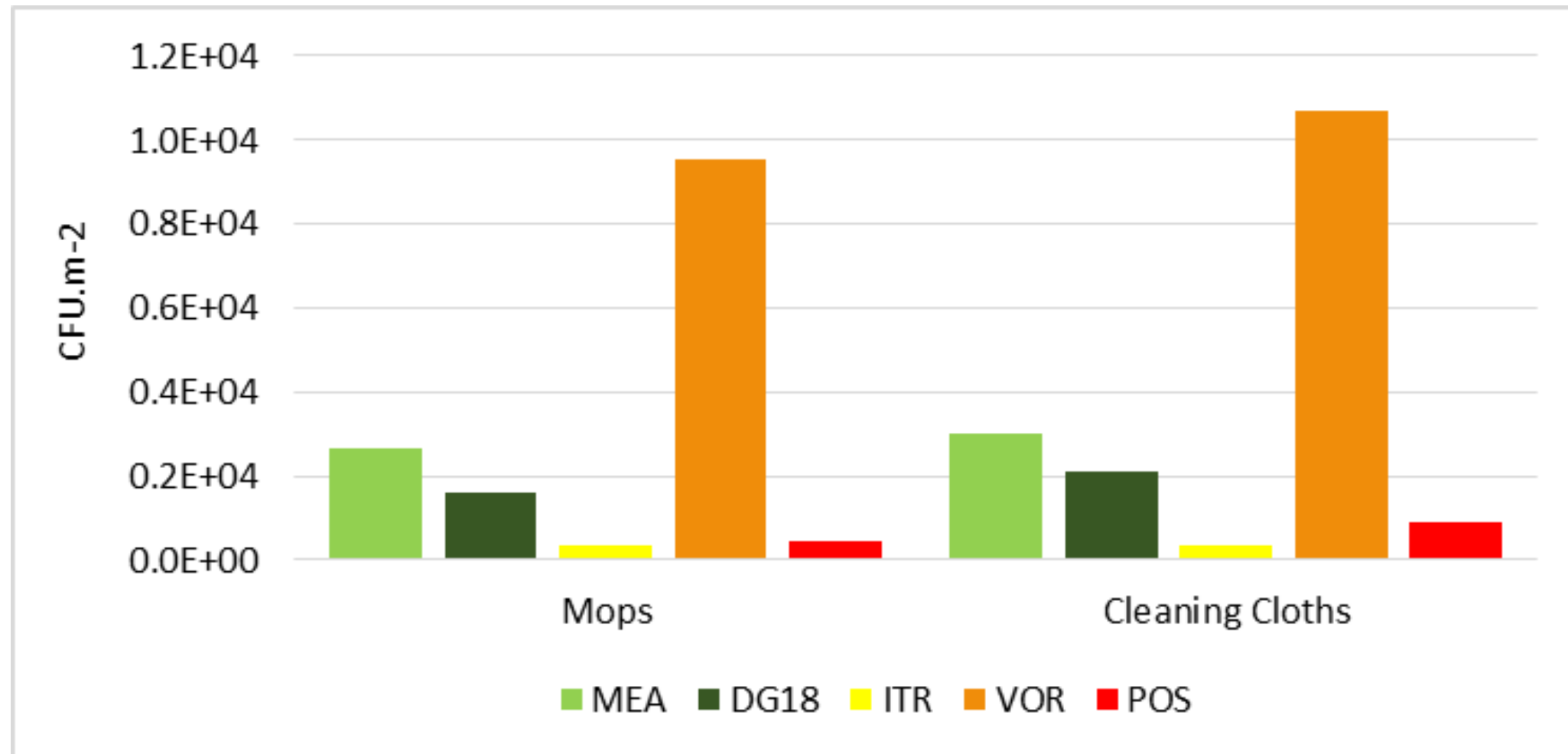
- Molecular tools (qPCR) were used for detection of *Aspergillus* sections, namely *Aspergillus* sections *Fumigati* and *Nidulantes*.

3. Results

Bacteria



Fungi



Mops:

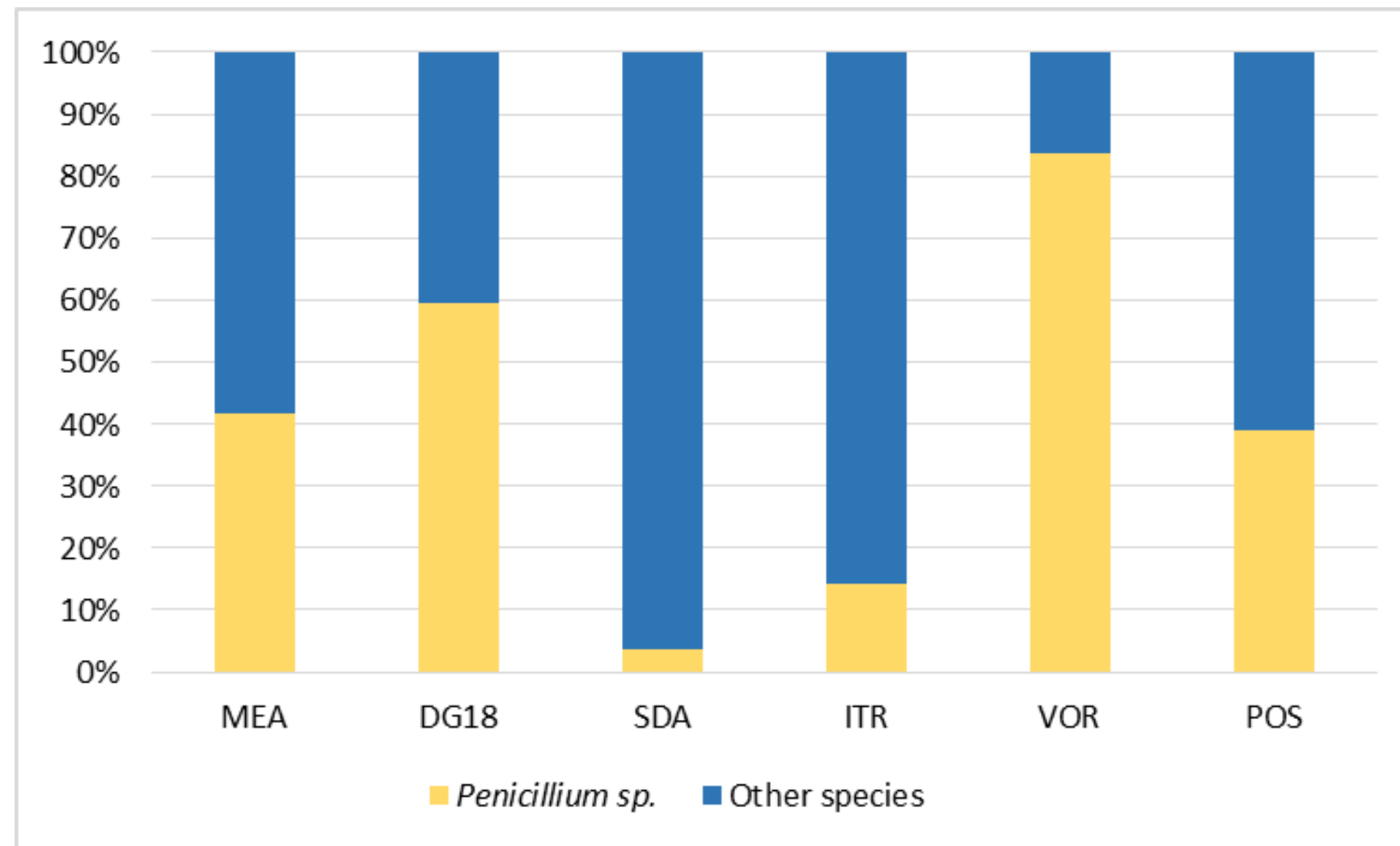
- bacteria ranged from 500 to 1.9x10⁵ CFU.m² on TSA and from 0 to 5.5x10⁴ CFU.m² on VRBA.
- fungi ranged from 1.0x10³ to 1.2x10⁴ CFU.m² on MEA and from 3.0x10³ to 1.2x10⁴ in DG18.

Cleaning cloths:

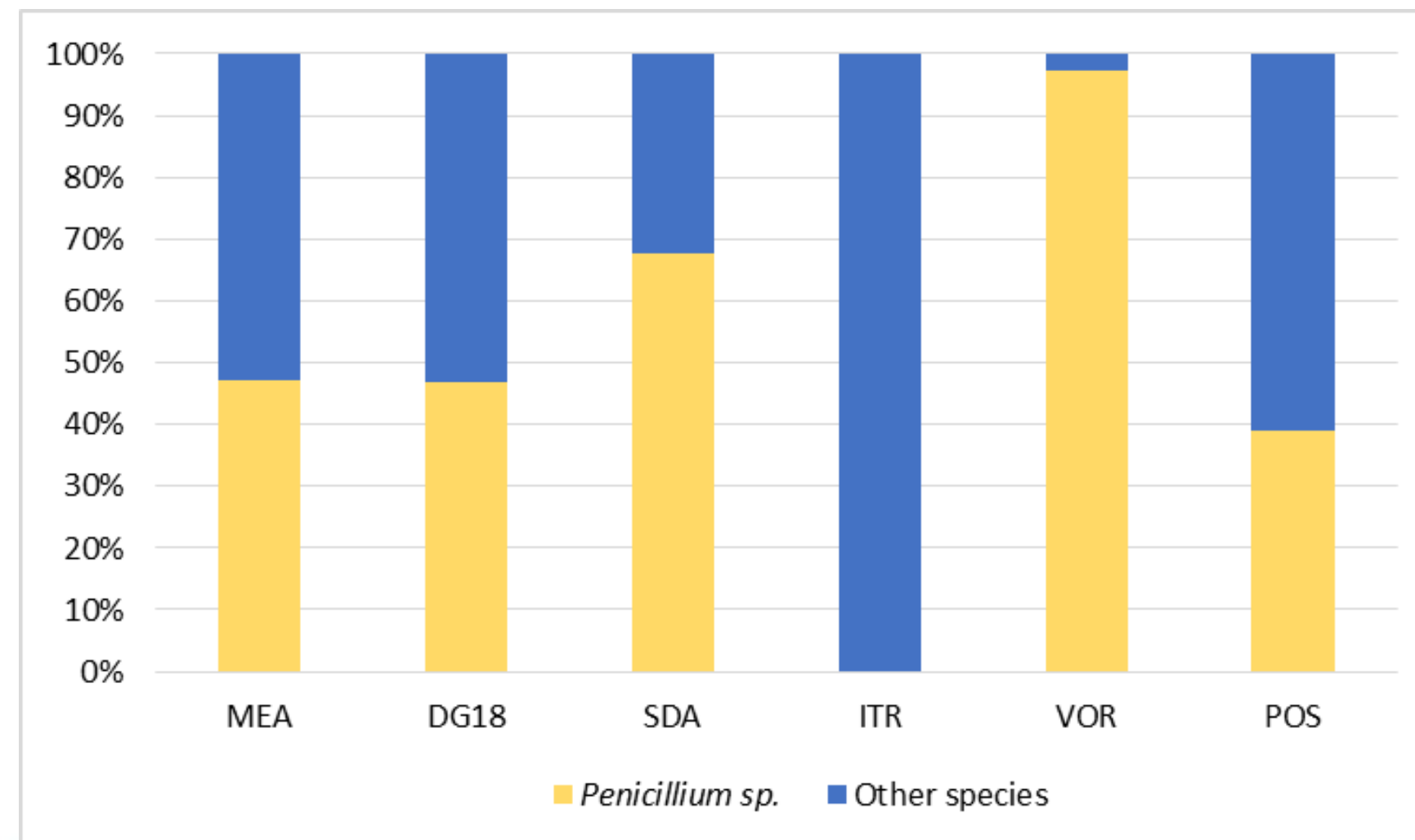
- bacteria ranged from 5.0x10² to 5.4x10⁵ CFU.m² on TSA and from 0 to 4.2x10⁵ CFU.m² on VRBA.
- fungi ranged from 3.0x10³ to 1.2 x10⁴ on MEA and from 1.0x10³ to 5.0x10³ on DG18.
- Mops presented higher bacterial contamination than cleaning cloths while the cleaning cloths presented higher fungal contamination than mops.
- Voriconazole was the supplemented media with higher fungal contamination.

3. Results

Mops



Cleaning cloths



- *Penicillium* sp. was the predominant genera in MEA, DG18 and in the azole resistance screening.
- *Aspergillus* section *Fumigati* was detected by qPCR:
 - Mops: 50% samples (n=14)
 - Cleaning Cloths: 60% samples (n=25).

4. Take home messages

The contamination found in mops and cleaning cloths suggest that :

- Cleaning procedures might be leading to cross-contamination through mops and cleaning cloths.
- The presence of fungi non-susceptible to azole drugs and detection of toxigenic *Aspergillus* section *Fumigati* might represent a health risk and must be further investigated.



Thank you for your attention

Acknowledgments

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