Non-invasive Detection of Co-infections in Hospitalized Patients with COVID-19 Using the Karius® Test, A Plasma-based Next-Generation Sequencing Test for Microbial Cell-free DNA

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Teno vrs

Adenovirus

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Background

Superinfections occur in up to 40% of the patients with COVID-19 but it is difficult to identify the pathogens through the usual invasive procedures due to patient instability and risk of exposure and pro-calcitonin is insensitive in this setting. 1-3 The Karius® Test may be used to rapidly and noninvasively identify fungal, viral and bacterial pathogens in one process, using a single plasma sample.4-6

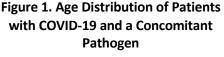
Methods

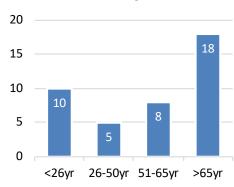
The Karius® Test (KT) is conducted in a CLIA certified/CAP-accredited laboratory in Redwood City, California. Microbial cell free DNA (mcfDNA) is extracted and next-generation sequencing (NGS) performed; microbial sequences are aligned to a curated database of > 1000 organisms. Organisms present above a statistical threshold are reported.4

Results

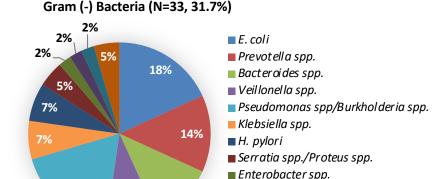
Fifty-three specimens from patients with COVID-19 were processed between 4/1/20 and 9/25/20. Twenty of 53 (38%) were female and 33 (62%) were male. 39 (74%) were adults; 14 (26%) were children or young adults (Figure 1). Twelve patients (22.6%) did not have a concomitant pathogen: (median age 41yr; min 2yr-max 68yr; p=0.057). The median age of those with a concomitant pathogen was 63vr (min 0.5 max 91vr); 37/41 (90.2%) had two or more co-pathogens. The number of concomitant pathogens was significantly higher among patients \geq 65yr (3.4, n=18) as compared to patients <65yr (2.2, n=23; p=0.009). The number of concomitant pathogens was similar between males (2.5, n=25) and females (3.1, n=18; p>0.05).

Results





24%



■ Hafnia spp.

Haemophilus spp.

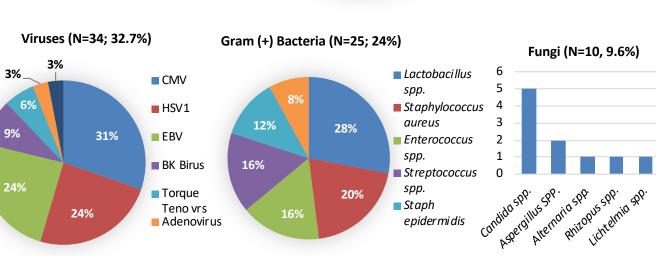
Fusobacteria spp.

11%

Staph

epider mi dis

9%



18%

Conclusions

Reactivation of herpesviruses, bacterial & fungal infections are common among hospitalized patients with COVID-19. The Karius® Test provides a rapid, non-invasive means to identify co-infecting pathogens using a single plasma sample.

1Chen Xi et al. Appl Microbiol Biotechnol 2020;104:7777 2Zhu X et al. Virus Res 2020; 285:198005. 3van Arkel et al. Am J Respir Crit Care Med. 2020; 202:132 4Blauwkamp et al. Nat Microbiol. 2019;4:663 Farnaes et al. Diagn Microbiol Infect Dis.;94:188-191 Rossoff et al. Open Forum Infect Dis. 2019; 6(8) The authors thank Varsha Baichwal for her contributions to the work