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PS07.05 (137)
Microflora in patients with community-acquired pneumonia and hospital environment in Blagoveshchensk during pandemic
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Purpose: An increase of community-acquired pneumonia (CAP) incidence has been detected during COVID-19 pandemic. Aim: to conduct analysis of bacterial microflora isolated from patients suffering from CAP and hospital environment in hospitals.

Methods & Materials: Bacteriological survey of sputum obtained from 210 hospitalized patients with CAP was conducted from December 2020 – March 2021 in the Blagoveshchensk city. A simultaneous sanitary-bacteriological control of hospital environment was performed (210 environment samples and 24 air samples). The research was conducted during 12 weeks (6 cycles, one cycle lasted two weeks). Bacteriological assessment was performed via classical method. Identification of isolated pathogens and evaluation of drug-resistant strains were performed with bacteriological analyzer “Multiscan”.

Results: Examination of sputum that was obtained from 151 patients with CAP revealed 43 isolates and 8 bacterial species in 32 patients (21.2%). Candida spp. fungi (51.16±7.82%) and Gram-positive microflora without drug-resistant strains (39.53±7.46%) were prevailing in the structure of detected isolates Gram-negative enterobacteria were presented by one species of Klebsiella pneumoniae with multiple drug-resistance – ESBL (4.65±3.21%) and non-fermenter Gram-negative bacteria – one species of Pseudomonas aeruginosa (2.33±2.30%). Evaluation of 210 environmental samples showed bacterial microflora in 49 samples (23.3% cases) that was comparable to frequency detection of bacterial pathogens from patients (21.2%). Microflora identified from environment was presented by 12 pathogenic species including Gram-positive microflora (S. aureus, S. epidermidis, S. hominis, E. faecalis, E. Faecium) – 55.0%, Gram-negative enterobacteria (K. pneumoniae, E. coli, E. Agglomerans, E. cloacae, C. freundii) – 38.8%, non-fermenter Gram-negative bacteria (A. baumannii, A. Iwoffii) – 6.1%. Should be noted that during first three cycles bacterial contamination was detected in 19 out of 90 samples (21.1%) including 3 cases with multiple drug-resistance (3.3%). During following three cycles, bacterial contamination was revealed in 25.0% of samples (30 out of 120 samples) including 13 samples that had multiple drug-resistant bacteria (10.8%). Evaluation of 24 air samples showed no evidence of microbial contamination.

Conclusion: Dynamic surveillance revealed wide circulation of bacterial pathogens in the hospital as well as build up of epidemiologically significant pathogen strains in hospital environment during observation period. These findings indicate considerable risk of nosocomial infections formation among patients.

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PS07.06 (441)
Post-vaccination COVID-19 among Healthcare workers in a Tertiary Care Hospital, Malaysia
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Purpose: To describe the sociodemographic, comorbidities, indication of RT-PCR testing, number of doses of vaccine received, COVID-19 category among the post-vaccination healthcare workers.

Methods & Materials: A retrospective secondary data analysis was conducted at University of Malaya Medical Centre (UMMC), a tertiary referral and teaching hospital with more than 6000 HCWs located in Kuala Lumpur, Malaysia. Immunisation Programme at UMMC started on 1st March 2021, prioritising medical and non-medical frontline staff. All HCWs were included in the study if they have received either one or completed both doses of BNT162b2 (Pfizer-BionTech) vaccine and was subsequently inflicted with COVID-19 between 1st March until 31st May 2021. HCW were excluded whose gestation was not between 14-33 weeks at the time of the first dose and individuals who had a history of severe anaphylaxis reaction.

Results: A total of 39 HCWs tested positive for SARS CoV-2 through RT-PCR. 6(15.4%) of them tested positive after their first dose while 33 (84.6%) staff were confirmed to be infected after receiving the second dose. For the job category, 21(53.8%) nurses, 7(18%) physicians, 6 attendants (15.4%) and 5(12.8%) supporting staff were infected. 21 were indicated for testing because of close contact with positive SARS CoV-2 patients, whether they were symptomatic or not during the time of testing. The balance of the HCW were tested because they presented with symptoms ranging from mild flu and sore throat to cough and fever. 11 of the HCW (28.2%) have underlying comorbidity declared prior to vaccination and the rest did not declare any previous underlying medical illness. 33.3%(n=13) of the HCW tested positive were in category 1 and 66.7%(n=26) in category 2. Both categories did not require any hospital admission and they were closed monitoring throughout the maximum of 10 days for isolation at home. They were only admitted if unable to isolate themselves at home.

Conclusion: Well-organized surveillance system is essential for early detection of infected HCW. COVID-19 vaccination among HCW did not only prevent them from getting seriously ill, but also reduced the need for ICU admission and eventually reduced the burden of the disease in the country.

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PS07.07 (171)
Analysis of the influence of age and location of contacts on the frequency of transmission of SARS-CoV2 in society
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**Purpose:** Analysis of the frequency of transmission SARS-CoV2 in different age groups and locations allows to identify risks of transmission, influencing on the spread of COVID-19 in the population and to strengthen the control on pandemic

**Methods & Materials:** Epidemiological analysis of contacts with primary cases of COVID-19 and secondary cases of infection for 6 months of 2020 in the different age groups was performed on the data of epidemiological surveillance system in 190,856 COVID-19 patients and 146,996 their contacts

**Results:** Patients with mild form of disease had the main proportion of contact - 50.6 %, moderate form - 45.8%. The proportion of contacts in patients with severe form was only 3.6%.

Group aged 41-64 years had the highest number of cases (44.4%) and contacts (44.5%), aged 18-40 years - 30.2% of cases and 34.2% of contacts; 0-6 years - 2.7% of cases and 1.9% of contacts, 7-17 - 4.0% of the cases and 3.1% of the contact. The number of contact persons was directly related to the number of secondary cases of the disease in all age groups: at the age of 0-6 years - 3.5 %, 7-17 years - 5.8 %, 18-40- 33.9%, 41-64 - 42.1%, 65 and older -14.7%.

Patients of all age groups mainly contacted with healthy people in domestic focuses: the age group 0-17 years accounted for 65.9% of contacts, 18-40 years - 63.9%, 41-64 years - 64.2% over 65 years - 65.18%, respectively.

The frequency of contacts at work was the highest at the age of 18-40 years - 9.0% and 41-64 years - 9.0%, but were significantly less than contacts in everyday (family) life - 63.8% and 64.2%, respectively.

**Conclusion:** The most active transmission of infection is carried out by patients aged 18-64 years, carrying the disease in mild and moderate-severe form. Secondary transmission of COVID-19 most of all is active in household foci in all age groups. In the age groups of 18-64 years, the transmission of infection in the workplace is also important. These results of analysis can be used to optimize prevention measures against COVID-19.

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**PS07.08 (310)**

**Determination of SARS-CoV-2 Contamination in a Neonatal Intensive Care Unit (NICU) Environment Using Droplet Digital PCR (ddPCR)**

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**Purpose:** Neonatal infections with SARS-CoV-2 are thought to be less contagious than in older children and adults. The transmission of SARS-CoV-2 from neonates and their environment has not been well studied. Droplet Digital PCR (ddPCR) is an emerging and sensitive technology that can aid infection control investigations. We sought to document surface contamination within the immediate environment of a preterm neonate with congenital COVID-19 using ddPCR.

**Methods & Materials:** On day 5 of life, a total of 23 environmental samples were collected in Eswabs (Amies media) based on proximity to the neonate, from the inside (7) and outside (16) of the neonate’s incubator for ddPCR analysis. Samples were extracted, using an in-house method and each extract was run for reverse-transcription ddPCR measurement using the Bio-Rad SARS-CoV-2 ddPCR Kit. The 96-well RT-ddPCR ready plate was loaded into the QX200 Droplet Reader (Bio-Rad, Pleasanton, CA). The fluorescence intensity of each droplet was measured, and droplets were determined to be positive or negative for gene targets (N1, N2).

**Results:** All samples collected from outside of the incubator were negative. These included: a stethoscope hanging outside of the incubator, nearby keyboard/mouse, wireless phone receiver, barcode scanner, blood culture bottles, pens/pencils, light switches, weigh scale, countertop/shelf, cart with drawers and incubator port release clips. Samples collected from inside the incubator were positive for SARS-CoV-2. These results reported in copies per microlitre (cp/µl) extract included: the swaddle cloth (0.4 N2), sheets behind the neonate’s head (11.4 N1, 16.9 N2), cardiorespiratory and saturation monitor leads and cables near the neonate’s head (2.8 N1, 4.5 N2), near the neonate’s feet (2.1 N1, 3.7 N2), and nametags hanging on a panel (1.0 N1, 1.2 N2). The highest levels were noted from the neonate’s drool (25.2 N1, 35.2 N2).

**Conclusion:** The presence of SARS-CoV-2 was confirmed by ddPCR in environmental samples inside the incubator confirming the ability of the neonate to spread the virus in close quarters. No virus was identified outside of the incubator which suggests appropriate hand hygiene and disinfection of environmental surfaces. ddPCR appears to be a useful tool for investigating the potential role of fomites in COVID-19 transmission

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**PS07.09 (666)**

Clinical and epidemiological assessment of the epidemic process of covid-19 in hostels depending on the type of their planning arrangement

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**Purpose:** During the COVID-19 pandemic, people living in dormitories became one of the most vulnerable groups of the population involved in the process of focal morbidity.

**Aim:** To analyze the clinical and epidemiological features of the manifestation of the epidemic process of COVID-19 in hostels, depending on the type of their planning arrangement.

**Methods & Materials:** A prospective analytical study of 350 foci of COVID-19 formed in hostels in Moscow in the period from 04.12.2020 to 06.23.2020 with 3,228 cases in total.

**Results:** The epidemic process in the corridor-type dormitories (CTB) had an earlier development, the first foci of COVID-19 began to form as early as 04/12/2020, which was 7 days ahead of the formation of foci in block-type dormitories (BCD). The average growth rate in the corridor-type dormitories was 8.4%, which was 5.3 times higher than the corresponding indicator in block-type dormitories. The prevalence rate was 1.5 times higher in the corridor-type dormitories. From 04/12/2020 to 06/23/2020, the number of COVID-19 infected in the block-type dormitories was significantly less than in the corridor-type dormitories: there were 4 cases of infection per 100 guests of the block-type dormitories, while in the corridor-type dormitories there were 10 cases of COVID-19 per...