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reported in Riyadh region (50.36%), and minority was in Jazan region (0.12%). The generation of infection was determined from three main sources; Primary infection was 24.16% (camel contact and/or consumption of their raw milk), Secondary infection was 38.34% (hospital acquired 92.16% and community acquired 13.48%) and un-identified source 37.5% of all cases.

MERS infection caused 368 death among 829 cases (case fatality rate CFR = 44.4%) for all the period. The mortality rate according to the source of infection was; un-identified source (44%), secondary infection (32.6%, including 97.5% hospital acquired), and primary infection (23.4%).

**Conclusion:** Incident and mortality rates were found significantly decreased from 2015 to 2017. Age group and source of infection demonstrated high risk population. Identification of MERS epidemiological characteristics support the control plan and preventive measures for improving public health impact in Saudi Arabia.

https://doi.org/10.1016/j.jiph.2018.10.098

**Descriptive Epidemiology, Clinical Characteristics and Outcomes for Middle East Respiratory Syndrome Coronavirus (MERS-CoV)Infected Patients in Al Ain – Abu Dhabi Emirate**

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**Introduction:** MERS-CoV was first identified in 2012 and highest incidence was in Saudi Arabia, followed by Republic of Korea and UAE. The majority of reported cases are males. Patients with comorbidities have higher mortality rate. The source of MERS-CoV infection is linked to camel exposure, nosocomial and human-to-human transmission. The clinical course of MERS-CoV infection can vary from asymptomatic disease to septic shock with multi-organ failure (MOF). We studied the clinical characteristics and outcomes of MERS-CoV infected patients in Al Ain – UAE.

**Method:** A retrospective multicenter chart review study for MERS-CoV confirmed cases by RT-PCR at Tawam and Al Ain Hospitals over 6 years period (2012 – 7/2018). Epidemiological, demographic, clinical and laboratory data were collected and analyzed using descriptive analysis.

**Results:** A total of 58 individuals were identified, 41 males (70.6%) with median age of 43.5 years. The majority of patients were asymptomatic carriers or having mild symptoms (34 [58%]) and were previously healthy. The comorbid conditions were hypertension (18 [31%]), diabetes mellitus (11 [18.9%]), coronary artery disease (6 [10.3%]), and chronic kidney disease (9 [15.5%]). The risk factors for MERS-CoV infection were camel contact (58.6%), travel (8.6% [Oman, 2; Saudi Arabia, 3]) and human-to-human contact with MERS-CoV infected patient or nosocomial transmissions (40 [68.9%]). Common symptoms at presentation were fever 46.5%, myalgia 41.3%, respiratory symptoms 41.3%, and gastrointestinal abnormality to evaluate the difference. Most of these risk factors had statistical significant of the difference except neonate and the presence of VP- shunt. The following organisms were yielded; from gram positive cocci (70% of all organisms) were coagulase –ve Staphylococcus (15% of BCs), Staphylococcus epidermidis (15% of BCs) and both Streptococcus pneumoniae and viridans (3.6% of BCs for both), from gram –ve rods (14% of BCs) were Klebsiella pneumoniae (5% of BCs) and Acinetobacter spp. (3.5% of BCs) and from non-bacteria organism was Candida spp. (6% of BCs).

https://doi.org/10.1016/j.jiph.2018.10.100

**Latent Tuberculosis Screening Using T-Spot-TB Test Among People Living With HIV: A Retrospective Study Of 190 Patients At A Tertiary Care Hospital In Dubai, UAE**

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**DHA**

**Background:** It is estimated that one third of 33.3 million people living with HIV worldwide are infected with TB. Overall mortality is twofold higher for HIV/TB co-infected individuals compared to those with isolated HIV infection. Consequently, diagnosis of latent TB infection (LTBI) and provision of chemotherapy to those testing positive is strongly recommended. This article reviewed the epidemiology of LTBI among adult people living with HIV following up in a tertiary care center in Dubai, United Arab Emirates (UAE).

**Methods:** A retrospective study included all HIV-infected patients who were screened for LTBI using T-Spot test at Rashid hospital, Dubai from January 2016 until December 2016, through medical records. Patient with active TB either during study period or in past were excluded.

**Results:** 209 patients with HIV were evaluated and 190 were included in analysis. 52 (27%) were female and 138 (73%) were male. 140/190 (74%) of patients were UAE national, 24/190 (12%) were African and 15/190 were from other Arab countries. T-Spot was reactive in 33/190 (17%) of cases, 155/190 (82%) were nonreactive and only 2/190 (1%) patients had an indeterminate result. There was no statistically significant association between low CD4+ T-cell count, viral load and T-Spot reactivity. Over 70% of patients in both arms have suppressed HIV viral activity with viral load below 50 copies/ml. A CD4 cell count less 200 cells/mm3 was observed in less than 30% of cases in each group, whereas over 30% of patients in each group had a CD4 cell count of more than 500 cells/mm3. Twenty one (68%) patients with reactive T-Spot were offered preventive therapy.

https://doi.org/10.1016/j.jiph.2018.10.099

**The yield of blood cultures performed at Tawam Hospital for all children from all department**

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All blood cultures (BCs) were taken from January 2013 up to June 2013 for children from birth up to 15 years old at Tawam hospital. A retrospective cross-sectional descriptive analytic study was performed to them. One hundred seventy one (6%) were yielded from 3015 BCs. These organisms were classified into true bacteremia (62%) and false positive cultures (38%). The true bacteremia was compared with negative cultures by gender, age groups, fever, and the presenting condition of the child, oncology patient, immunodeficiency patient, and any child having indwelling catheter or other devises and any cardiovascular abnormality to evaluate the difference. Most of these risk factors had statistical significant of the difference except neonate and the presence of VP- shunt. The following organisms were yielded; from gram positive cocci (70% of all organisms) were coagulase –ve Staphylococcus (15% of BCs), Staphylococcus epidermidis (15% of BCs) and both Streptococcus pneumoniae and viridans (3.6% of BCs for both), from gram –ve rods (14% of BCs) were Klebsiella pneumoniae (5% of BCs) and Acinetobacter spp. (3.5% of BCs) and from non-bacteria organism was Candida spp. (6% of BCs).