

BIRDEM NEWS

(Birdem Med J 2015; 5(Supplement 1): 66-69)

Microbiology

Bacteria and Antibiogram Report
July to December 2014 (Vol 5, No. 2, 2015)

Table-I*Pattern of organisms isolated from different samples*

| Organisms | Blood | Urine | Respiratory Secretions | Pus | Wound | Total |
|----------------------------|-------|-------|------------------------|-----|-------|-------|
| Escherichia coli | 60 | 1365 | 46 | 214 | | 1685 |
| Klebsiella sp | 42 | 278 | 141 | 290 | | 754 |
| Acinetobacter sp | 35 | 17 | 293 | 93 | | 438 |
| Pseudomonas sp | 60 | 85 | 109 | 295 | | 549 |
| Staph aureus | 04 | 67 | 28 | 289 | | 191 |
| Gr D Non Enterococcus | 07 | 29 | - | 09 | | 38 |
| Salmonella sp (Typhoid gr) | - | - | - | - | | 29 |
| Proteus sp | 29 | 09 | 05 | 295 | | 249 |
| Enterobacter sp | - | 57 | 09 | 76 | | 145 |
| Candida sp | 03 | 231 | 128 | 10 | | 396 |
| Streptococcus sp | 27 | 41 | - | 14 | | 55 |
| Serratia marcescens | - | 01 | 01 | 04 | | 06 |
| Providencia sp | - | - | - | - | | - |
| Citobacter sp | 5 | 32 | 03 | 29 | | 64 |
| Coagulase negative Staph | - | 03 | - | - | | 03 |
| Morganella sp | - | - | - | - | | - |
| Haemo philus sp | - | 02 | - | - | | 02 |
| Streptococcus pneumoniae | - | - | - | - | | - |
| Flavobacterium sp | 01 | 02 | 03 | - | | 06 |
| Corynebacterium sp. | - | - | - | - | | - |
| Burkholderia pseudomallei | - | 01 | - | - | | 01 |

Table-II*Major organisms isolated from outdoor, indoor, ICU and ICHRI patients*

| Organisms | Outdoor N=2122 | Indoor N=2132 | ICUN=734 | ICHRIN=133 | SCABUN=14 | Total N=5230 |
|-----------------------|----------------|---------------|----------|------------|-----------|--------------|
| Ecoli | 761 | 831 | 57 | 30 | - | 1689 |
| Klebsiella | 305 | 335 | 104 | 17 | 01 | 762 |
| Acinetobacter | 63 | 66 | 286 | 26 | 02 | 443 |
| Pseudomonas | 254 | 179 | 106 | 10 | 02 | 551 |
| Staph aureus | 263 | 207 | 19 | 05 | - | 494 |
| Salmonella sp. | 22 | 04 | - | 01 | 01 | 28 |
| Gr D Enterococcus sp. | 65 | 114 | - | 01 | - | 170 |
| Gr D Non enterococcus | 13 | 21 | 04 | - | - | 38 |
| Candida sp. | 34 | 98 | 90 | 39 | 08 | 269 |
| Proteus sp. | 170 | 71 | 06 | 02 | - | 249 |
| Streptococcus sp. | 37 | 18 | 01 | - | - | 56 |
| Enterobacter sp | 91 | 49 | 03 | 02 | - | 145 |
| Citrobacter sp. | 32 | 31 | 01 | - | - | 603 |
| Serratia | 06 | - | - | - | - | 506 |

Table-III*ABST pattern of major gram negative organisms isolated*

| Antibiotics | % Resistant | | | | |
|-------------------|---------------------|-----------------------|--------------------------|------------------------|----------------------|
| | E. coli (N=1685) | Klebsiella (N=759) | Acinetobacter (N=440) | Pseudomonas (N=550) | Salmonella (N=29) |
| Imipenem | 5.1 | 19.4 | 90.1 | 58.3 | ND |
| Ceftriaxone | 69.7 | 35.4 | 96.1 | 78.8 | 0 |
| Ceftazidime | 69.7 | 66.1 | 94.8 | 63.8 | ND |
| Cefixime | 71.6 | 67.9 | 95.7 | 90.8 | 0 |
| Augmentin | 79.3 | 79.2 | 95.2 | 94.5 | ND |
| Piperacillin | - | - | 75.0 | 36.2 | ND |
| Tazo/piperacillin | 24.9 | 44.1 | 91.5 | 30.3 | ND |
| Amikacin | 9.9 | 29.6 | 89.5 | 49.0 | ND |
| Netilmicin | 8.7 | 28.5 | 78.0 | 54.0 | ND |
| Gentamicin | 24.1 | 37.4 | 90.0 | 58.4 | ND |
| Ciprofloxacin | 71.8 | 52.4 | 94.1 | 55.7 | 0 |
| Cotrimoxazole | 57.6 | 58.0 | 75.0 | 74.2 | 31.6 |
| Nitrofurantoin | 10.6 | 60.4 | 72.2 | 92.2 | ND |
| Colistin | 1.5 | 1.4 | 1.2 | 11.5 | ND |
| Nalidixic acid | ND | ND | ND | ND | 93.10 |
| Azithromycin | ND | ND | ND | ND | 27.59 |
| Chloramphenicol | 8.5 | 31 | 100 | 58.5 | 57.59 |
| Ampicillin | ND | ND | ND | ND | 37.93 |

Table-IV*ABST pattern of major Gram positive organisms isolated*

| Antibiotics | % Resistant | |
|----------------|-----------------------|-------------------------|
| | Staph aureus N=487 | Enterococcus (N=291) |
| Penicillin | ND | 27.0 |
| Amycillin | ND | 17.3 |
| Oxacillin | 28.8 | ND |
| Cephalexin | 30 | ND |
| Augmentin | 39.6 | ND |
| Amikacin | 3.1 | 61.1 |
| Netimicin | 0.6 | 41.9 |
| Gentamicin | 5.614.3 | 39.3 |
| Nitrofurantoin | 2.2 | 13.2 |
| Rifampicin | 3.9 | ND |
| Fusidic acid | 1.5 | ND |
| Erythromycin | 72.9 | 100 |
| Vancomycin | 0 | 0 |
| Cotrimoxazole | 21.0 | 97.6 |

ND=Not done

Table-V*ABST pattern of major Gram negative organisms of ICU*

| Antibiotics | % Resistant | | | |
|-------------------|-----------------|-----------------------|-------------------------|------------------------|
| | Ecoli (N=67) | Klebsiella (N=104) | Acinetobacter N=286) | Pseudomonas (N=106) |
| Imipenem | 22.2 | 52.4 | 100 | 81.6 |
| Ceftriaxone | 90.9 | 92.3 | 100 | 90.6 |
| Ceftazidime | 91.0 | 92.3 | 100 | 75.2 |
| Augmentin | 95.5 | 95.2 | 100 | 94.2 |
| Piperacillin | - | - | 100 | 71.4 |
| Tazo/piperacillin | 52.6 | 76.0 | 97.9 | 46.1 |
| Amikacin | 29.9 | 68.3 | 97.7 | 60.4 |
| Netilmicin | 34.3 | 69.2 | 92.7 | 77.1 |
| Gentamicin | 53.7 | 67.2 | 99.0 | 55.7 |
| Ciprofloxacin | 88.1 | 84.6 | 100 | 66.0 |
| Cotrimoxazole | 77.6 | 85.6 | 76.9 | 73.6 |
| Colistin | 0 | 1.0 | 0.37 | 14.9 |

Table-VI*Multidrug resistant organisms isolated from various samples*

| Organisms | Total isolated | Category of resistant organisms | N | % |
|------------------|-------------------|------------------------------------|------|-------|
| Staph aureus | 457 | MRSA | 141 | 30.85 |
| Salmonella sp. | 29 | NARST | 27 | 93.10 |
| Ecoli | 1685 | ESBL | 852 | 50.56 |
| Klebsiella sp. | 759 | ESBL | 1224 | 29.51 |
| Enterobacters sp | 145 | ESBL | 21 | 14.48 |
| Citrobacters sp | 64 | ESBL | 22 | 34.37 |
| Enterococcus sp. | 291 | VRE | 0 | 00 |

ABST=Antibiotic sensitivity test; NARST=Nalidixic acid resistant S. typhi & S paratyphi; ESBL=Extended spectrum beta lactamase; VRE=Vancomycin Resistant enterococcus; ND = Not done

