Table 1: Rape Resistance

| Microorganism | Isolates (n) | SXT | CIP | TGC | CTX | AMP | GM | ATM | CRO | VAN | Resistance Profile |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Klebsiella pneumoniae | 36 | 83% (44-97) | 33% (10-70) | 20% (4-62) | 0% (0-43) | 100% (57-100) | 67% (30-90) | 100% (21-100) | 0% (0-79) | 100% (21-100) | AMP-ATM-VAN (36/36) |
| Pasteurella Spp | 4 | 0% (0-0) | 0% (0-0) | 0% (0-0) | 0% (0-0) | 100% (21-100) | 100% (21-100) | 100% (21-100) | 0% (0-79) | 0% (0-0) | AMP-GM-ATM (4/4) |
| Erwinia Spp | 5 | 0% (0-0) | 0% (0-79) | 0% (0-79) | 0% (0-0) | 0% (0-0) | 0% (0-79) | 0% (0-79) | 0% (0-79) | 0% (0-0) |  |
| Pseudomonas cepacia | 16 | 100% (21-100) | 33% (6-79) | 0% (0-0) | 100% (34-100) | 0% (0-0) | 100% (44-100) | 100% (44-100) | 0% (0-56) | 100% (21-100) | SXT-CTX-GM-ATM-VAN (16/16) |
| Pseudomonas aeruginosa | 5 | 0% (0-0) | 0% (0-79) | 0% (0-0) | 100% (21-100) | 0% (0-0) | 0% (0-79) | 0% (0-79) | 0% (0-79) | 0% (0-0) | CTX (5/5) |
| Total Rape (n = 66) | 66 | 86% (49-97) | 27% (10-57) | 17% (3-56) | 38% (14-69) | 100% (61-100) | 67% (39-86) | 71% (36-92) | 0% (0-35) | 100% (34-100) | 70% MDR (7/10) |