Antibacterial effect of copper sulfate against multi-drug resistant nosocomial pathogens isolated from clinical samples

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ABSTRACT
Background and Objective: With the emergence of antibiotic resistance and the hospital acquired infection, the interest for antimicrobial agents has recently increased again in public health. Copper is recommended as a supplementary method of increasing biological safety in the hospital environment. The objective of this study was to determine the antibacterial activity of copper sulfate salts on strains of bacterial pathogens isolated from different clinical pictures in different health establishment in Algeria.

Methods: A total of 25 different bacterial isolates (16 Enterobacteriaceae, 5 Staphylococci, and 4 Pseudomonas) were tested for susceptibility to copper sulfate using minimum inhibitory concentration (MIC-Cu) and minimum bactericidal concentrations (MBC-Cu) determinations. All isolates were also tested for susceptibility to six antibiotics.

Results: Antibiotic susceptibility studies revealed that 100% of isolates were resistant to one or more antibiotics. Fifty two percent of isolates were very susceptible to copper sulfate, with MICs ranging from 100 to 200 µg/ml. MBC-Cu = 1600 µg/ml showed the best bactericidal effect against the great majority of studied bacteria (52%). A good bactericidal activities of copper sulfate were recorded against Proteus vulgaris and Staphylococcus aureus (MBC/MIC=1). The Gram-negative bacteria isolates which were copper resistant also showed a high resistance to chloramphenicol (r=0.78) and Trimethoprim (r=0.61). Furthermore, the strains that were no-susceptible to three different antimicrobial classes (Escherichia coli, Staphylococcus saprophyticus) were not resistant to copper sulfate.

Conclusion: Copper sulfate salts has significant antibacterial activity against multi-drug resistant nosocomial pathogens.

KEYWORDS: Antibacterial activity, Copper sulfate, Nosocomial pathogens, Minimum bactericidal concentration, Minimum inhibitory concentration, Multi-drug resistance.

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INTRODUCTION

An alarming increase in antibiotic resistance among hospital pathogens has revived interest in alternative methods of reducing bioburden in healthcare facilities, focusing on the environment within hospitals.1 One alternative to be used as effective teat disinfection may be a copper. Living organisms requires copper at low concentrations as cofactors for metalloproteins and enzymes; however at high concentrations, Cu (II) induces an inhibition of growth in bacteria, and has a toxic effect...
on most microorganisms. The use of antimicrobial copper was accepted for the first time in 2008 by the United States Environmental Protection Agency. It has been shown that copper surfaces, or surfaces coated with this metal, have a 90 to 95% lower bacterial load, reducing the transmission of nosocomial infections. Moreover, recent findings that physiological Cu may be harnessed as a direct antimicrobial in innate immune cells.

Considering these antecedents, the aim of this study was to describe the antibiotic resistance and the susceptibility to copper sulfate salts of frequent nosocomial pathogens (Enterobacteriaceae, Pseudomonas and Staphylococci), isolated from various clinical samples from Algerian patients.

METHODS

Bacterial isolates included twenty five different strains isolated from patients from government hospital and the direction of the health of Guelma, Algeria. The strains were 7 isolates of Escherichia coli, two Citrobacter freundii, two Klebsiella oxytoca, one Citrobacter diversus, one Yersinia enterocolitica, one Edwardsiella tarda, one Proteus vulgaris, one Salmonella typhimurium, two Pseudomonas fluorescens, two Pseudomonas aeruginosa, two Staphylococcus aureus, one Staphylococcus xylosus, one Staphylococcus epidermidis and one Staphylococcus saprophyticus. The strains were isolated from blood cultures, wounds, faces and endotracheal sites of patients.

Antibiotic susceptibility was determined by the standard disk-diffusion method on Muller-Hinton (MH) agar plates (Lab. Pvt. Mumbai, India). For Enterobacteriaceae and Pseudomonas, susceptibility tests with six antibiotics namely amoxicillin (20µg), cefotaxime (5µg), imipenem (10µg), gentamicin (30µg), chloramphenicol (30µg) and trimethoprime (5µg). Staphylococci strains were tested susceptibility to penicillin G (1U), gentamicin (10µg), vancomycin (30µg), erythromycin (15µg), chloramphenicol (30µg) and trimethoprime (5µg).

The determination of the MIC-Cu was done by the broth dilution method. Copper (II) sulfate pentahydrate (CuSO₄·5H₂O) (Merck Millipore, Germany) were used to prepare 50 g/1 stock solution. This stock solution was filter-sterilized and used for preparation of the final concentrations. By two-fold dilutions, concentration of Cu²⁺ was ranged from 12.5 to 1600 μg/ml. After standardization of the inoculums to 0.5 McFarland, 1ml of the diluted inoculums were added to 1ml of each metal concentration except for the sterility control. Subsequently, the tubes were incubated in an oven at 37°C for 24 hour. All the experiments were performed in duplicate. MIC-Cu determination was done visually. Microbial growth was considered as positive in the tubes that showed any increase in turbidity or growth at the bottom. The isolates were considered resistant if the MIC-Cu values exceeded that of the Escherichia coli K12 ATCC 10798 (for Enterobacteriaceae isolates) Pseudomonas aeruginosa ATCC 27853 (for Pseudomonas isolates) and Staphylococcus aureus ATCC 25932 (for Staphylococci isolates) strains which were used as the control.

To determine the minimum bactericidal concentration (MBC-Cu), a 10 µl from those tubes, which did not show any visible growth in MIC-Cu assay, was cultured on Mueller-Hinton agar (MHA; Oxoid, UK) and incubated at 37°C for 18 to 24 h. The lowest concentration of copper producing no growth was considered to be the minimum bactericidal concentration (MBC-Cu).

Statistical analysis was done using SPSS 25.0, one-way ANOVA test was conducted to identify the significant differences between the isolated bacteria. The correlations between the copper resistance and antibiotic resistance of different resistant strains are evaluated by the Pearson correlation coefficient (r) (p≤0.05).

RESULTS

The obtained result revealed high level of multi-drug resistance among the isolates (Table-I). All the Enterobacteriaceae isolates were completely resistant to amoxicillin and imipenem. It was observed that 13 (86.67%) Enterobacteriaceae were resistant to trimethoprim while nine (60%) and four (26.67%) were resistant to chloramphenicol and cefotaxime respectively. Pseudomonas isolates were completely (100%) resistant to amoxicillin, imipenem and trimethoprim. No Enterobacteriaceae and Pseudomonas strains were found to be resistant to gentamicin. All Staphylococci isolates, were resistant to penicillin, however, 4 (80%) and 2 (40%) were resistant to cefotaxime, vancomycin and erythromycin, respectively. The overall patterns of antimicrobial resistance showed that the major profiles included AMC/IPM/C/TMP which occurred in 28% (7/25) of bacterial isolates.

The forty percent of isolates (10/25) were inhibited by 200 µg/ml of copper sulfate. Also, the concentration 400 µg/ml of copper sulfate could
inhibit 40% of isolates (10/25); however, there were two isolates (8%) that required 800 μg/ml of copper (E12: *Klebsiella oxytoca*, S4: *Staphylococcus aureus*). No growth was observed for any bacteria at 1600 μg/ml of copper (Fig.1A). Comparison between *Enterobacteriaceae*, *Pseudomonas* and *Staphylococci* isolates to the same concentration of copper showed that the p-values were not significantly different (p>0.05).

The results for *Enterobacteriaceae* isolates showed that 200 μg/ml and 400 μg/ml of copper inhibited 43.75% (7/16) and 37.5% (6/16) respectively (Fig.1B). *Staphylococci* isolates showed slightly high MIC-Cu values than those observed for *Pseudomonas* isolates, 400 μg/ml of copper inhibited 60% (3/5) and 25% (1/4) of isolates of *Staphylococci* and *Pseudomonas*, respectively. Comparing MICs-Cu of studied bacteria and that of the standard strains, resistance for copper was found among 43.75% of the studied *Enterobacteriaceae* (Fig.2A), one isolate of *Pseudomonas aeruginosa* (P4) (Fig.2B) and 80% of *Staphylococci* strains (Fig.2C).

The MBC-Cu equal to 1600 μg/ml of copper showed the best bactericidal effect against studied bacteria (52%: 13/25) (Table-II). However, a concentration of 800 μg/ml of copper was sufficient to kill 36% (9/25) of isolated strains. MBC-Cu against *Proteus vulgaris* (E15) and *Staphylococcus aureus* (S5) were found to be 400 μg/ml of copper.

Table-I: Resistance patterns of the bacteria isolates.

| Resistance patterns | No of antibiotics | No of isolate | Resistant bacteria |
|--------------------|------------------|---------------|-------------------|
| P                  | 1                | 1             | *Staphylococcus aureus* |
| P/TMP              | 2                | 2             | *Staphylococcus saprophyticus, Staphylococcus aureus* |
| AMC/IPM            | 2                | 1             | Escherichia coli |
| AMC/IPM/TMP        | 3                | 4             | *Escherichia coli, Yersinia enterocolitica, Klebsiella oxytoca, Salmonella typhimurium.* |
| AMC/IPM/TMP        | 3                | 2             | *Pseudomonas fluorescens, Pseudomonas aeruginosa* |
| AMC/IPM/C/TMP      | 4                | 7             | Escherichia coli, Citrobacter freundii, Citrobacter diversus, Klebsiella oxytoca. |
| AMC/CTX/IPM/TMP    | 4                | 2             | *Escherichia coli* |
| AMC/IPM/C/TMP      | 4                | 1             | *Pseudomonas fluorescens* |
| AMC/CTX/IPM/TMP    | 4                | 1             | *Pseudomonas aeruginosa* |
| P/VA/E/TMP         | 4                | 1             | *Staphylococcus epidermidis* |
| AMC/CTX/IPM/C/TMP  | 5                | 2             | *Edwardsiella tarda, Proteus vulgaris.* |
| P/GM/VA/E/C/TMP    | 6                | 1             | *Staphylococcus xylosus* |

P: Penicillin, TMP: Trimethoprim, AMC: Amoxicillin, IPM: Imipinem, C: Chloramphenicol, VA: Vancomycin, E: Erythromycin, CTX: Cefotaxime, GM: Gentamicin.

Fig.1: Minimum inhibitory concentration of copper (MIC-Cu) value distribution for the most prevalent isolates from Algerian patients.a, b, c, d Different letters indicate no significant differences (p≤0.05) between *Enterobacteriaceae*, *Pseudomonas* and *Staphylococci* isolates under the same copper concentration according to one-way ANOVA test.
μg/ml. As shown in Table-III, the copper sulfate salts demonstrated bactericidal or bacteriostatic effects.

**DISCUSSION**

Multi-drug resistant bacteria are serious threat in clinical health settings and very challenging to treat infectious disease. From current study it was found that 84% of clinical isolates have multi-drug resistance pattern. The antibiotic resistance patterns gave interesting information indicating a selective multi-drug resistant on the clinical isolates from Algerian patients, reflects indiscriminate use of antibiotics in human medicine.

All Gram-negatives bacteria in our research found to be resistant to imipinem which is an important antibiotic for the treatment of infections caused by Gram-negatives bacteria. From recent study, it was found that carbapenem have lost their activity against *Enterobacteriaceae* and *Pseudomonas* because of two main mechanisms, carbapenemase activity and loss of porin function. Resistance of studied isolates to the chloramphenicol is unacceptably high, particularly for *Pseudomonas* isolates (100%). Chloramphenicol is an antibiotic that is rarely used due to its well-described toxicity profile. It is conceivable that high resistance rates might have been driven by overuse of this antibiotic in the study health establishment. Many studies in *Pseudomonas* confirmed the role of the efflux pump in tolerance to chloramphenicol. This study demonstrated the resistance of *Staphylococcus xylosus* and *Staphylococcus aureus*. 

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**Table-II: Minimum bactericidal concentrations of copper sulfate against an important nosocomial pathogens.**

| Bacterial species                  | MBC-Cu (µg/ml) *mean±standard error |
|-----------------------------------|-------------------------------------|
| **Gram (-)**                      |                                     |
| *Escherichia coli* K12            | 400 ±00                              |
| ATCC 10798                        |                                     |
| *Escherichia coli*                | 1286±560*                           |
| *Citrobacter freundii*            | 1200±566*                           |
| *Citrobacter diversus*            | 1600±00*                            |
| *Klebsiella oxytoca*              | 1600±566*                           |
| *Yersinia enterocolitica*         | 1600±00*                            |
| *Edwardsiella tarda*              | 800±00*                             |
| *Proteus vulgaris*                | 400±00*                             |
| *Salmonella Typhimurium*          | 1600±00*                            |
| *Pseudomonas aeruginosa* ATCC 27853 | 1600±00*                        |
| *Pseudomonas fluorescens*         | 800±00*                             |
| *Pseudomonas aeruginosa*          | 1600±00*                            |
| **Gram (+)**                      |                                     |
| *Staphylococcus aureus* ATCC 25932 | 1600±00*                        |
| *Staphylococcus xylosus*          | 800±00*                             |
| *Staphylococcus epidermidis*      | 800±00*                             |
| *Staphylococcus saprophyticus*    | 1600±00*                            |
| *Staphylococcus aureus*           | 600±283*                            |

Values followed by different letters in a column are not significantly different (p>0.05) according to ANOVA tests. * Significant difference between Gram-positives and Gram-negatives bacteria after one-way ANOVA at a significant level of p ≤ 0.05.
Table-III: Report MBC/MIC.

| Strains                  | Copper MBC/MIC | Antibacterial activity |
|--------------------------|----------------|------------------------|
| *Escherichia coli* K12 ATCC 10798 | 2±0.0          | bactericidal           |
| *Escherichia coli*       | 5.6±5.1        | bacteriostatic         |
| *Citrobacter freundii*   | 10±8.4         | bacteriostatic         |
| *Klebsiella oxytoca*     | 4.5±4.9        | bacteriostatic         |
| *Citrobacter diversus*   | 4±0.0          | bacteriostatic         |
| *Yersinia enterocolitica*| 8±0.0          | bacteriostatic         |
| *Edwardiella tarda*      | 4±0.0          | bacteriostatic         |
| *Proteus vulgaris*       | 1±0.0          | bactericidal           |
| *Salmonella Typhimurium* | 8±0.0          | bacteriostatic         |
| *Pseudomonas aeruginosa* ATCC 27853 | 8±0.0 | bacteriostatic |
| *Pseudomonas fluorescens*| 6±2.8          | bacteriostatic         |
| *Pseudomonas aeruginosa* | 6±2.8          | bacteriostatic         |
| *Staphylococcus aureus* ATCC 25932 | 8±0.0 | bacteriostatic |
| *Staphylococcus xylosus* | 2±0.0          | bactericidal           |
| *Staphylococcus epidermidis* | 2±0.0    | bactericidal           |
| *Staphylococcus saprophyticus* | 8±0.0 | bacteriostatic |
| *Staphylococcus aureus* | 1±0.0          | bactericidal           |
| *Salmonella Typhimurium* | 8±0.0          | bacteriostatic         |
| *Pseudomonas aeruginosa* ATCC 27853 | 8±0.0 | bacteriostatic |
| *Pseudomonas fluorescens*| 6±2.8          | bacteriostatic         |
| *Pseudomonas aeruginosa* | 6±2.8          | bacteriostatic         |

Table-IV: Matrix of correlation between copper resistance and antibiotic resistance (r > 0.5 appear in bold type).

| Resistance to | Cu | AMC | IPM | C | TMP | Resistance to | Cu | P | TMP |
|---------------|----|-----|-----|---|-----|---------------|----|---|-----|
| Cu            | 1  |     |     |   |     | Cu            | 1  |   |     |
| AMC           | 0.142 | 1  |     |   |     | P             | 0.0123 | 1  |
| IPM           | 0.142 | **0.812** | 1  |   |     | TMP           | 0.0123 | **0.9121** | 1 |
| C             | **0.781** | 0.012 | 0.012 | 1 |     |               |     |   |     |
| TMP           | **0.618** | 0.0245 | 0.0245 | **0.6056** | 1 |     |

Cu: Copper, AMC: Ampicillin, IPM: Imipinem, C: Chloramphenicol, P: Penicillin, TMP: Trimethoprime, *: Enterobacteriaceae and Pseudomonas strains, **: Staphylococci strains.
negative bacteria (One-way ANOVA, p<0.05). The reason for the difference in sensitivity between Gram-positive and Gram-negative bacteria might be ascribed to the differences in morphological constitutions between these microorganisms. The best bactericidal activity of copper sulfate was observed against Proteus vulgaris, Staphylococcus aureus, Staphylococcus xylosus and Staphylococcus epidermidis. The other strains tested recorded bacteriostatic activity. These results could be explained by the differences between strains tested.

All other compounds of copper (copper oxide, copper acetate and copper nitrate) demonstrated bacteriostatic or bactericidal properties (results not shown), but the greatest antimicrobial effectiveness for all bacteria tested was observed with copper sulfate. From the clinical point of view, the use of copper sulfate (salts or impregnated in textile or liquids) with confirmed bacteriocidal properties against multi-drug resistant nosocomial pathogens as an antibacterial agent should be very important, because there is a problem of infections and epidemics caused by these bacteria in Algerian hospitals.

One concern in use of copper sulfate as a bactericidal agent often voiced is the potential for development of resistance as has happened with antibiotics. This study could clearly demonstrate that 80% Staphylococci, 43.75% Enterobacteriaceae and one Pseudomonas aeruginosa strains are considered resistant to copper. This study supports previous studies suggesting that most Gram negative bacteria and S. aureus also encode a multi-copper oxidase, which is required for the periplasmic oxidation of Cu(I) to Cu(II).

In terms of the potential for cross-resistance between copper and clinical antibiotics, the present study showed that there is good correlation among the proportion of copper-resistant Gram-negative bacteria and chloramphenicol resistance and trimethoprim resistance, with r=0.78 and r=0.61, respectively. The correlation of copper-resistant Gram-positive bacteria with penicillin resistance and trimethoprim resistance showed lower values (r=0.0123) (Table-IV). Correlation between copper resistance and resistance to chloramphenicol, trimethoprim and beta-lactams was related to four main strategies:
1. Reduction of membrane permeability
2. Rapid efflux of the metal and antibiotic
3. Alteration of cellular target
4. Drug and metal sequestration.

Caille et al. suggested that the presence of copper could increase resistance to imipenem in P. aeruginosa because of possible coregulation.

More recently, resistance gene to copper linked to resistance genes to beta-lactam, chloramphenicol, tetracycline, fluoroquinolones and vancomycin have been found.

CONCLUSION

The copper sulfate salts has significant antibacterial activity against multi-drug resistant nosocomial pathogens. The study confirmed the effective activity (bacteriocidal or bacteriostatic) of the copper sulfate salts. This study has suggested that there was probably some correlation between the phenotype of antibiotic resistance and copper resistance.

Recommendations: Further studies are needed to assess the microbial response to copper sulfate salts in vivo. However, it is necessary to comprehensively assess the role of copper-resistance as a selective force in maintaining and propagating the antibiotic-resistance.

Declaration of Interest: None.

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Author’s Contribution:

LB: Conceived & design, statistical analysis and editing of manuscript.
LB, SA: Data collection and manuscript writing.
MB, LB: Conceptualization, methodology and writing revisions.
RO: Review and final approval of manuscript.