Synthesis, Characterization, and Biological Evaluation of Chromium(III) Complexes of Alanine and Valine

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Abstract

Two metal-amino acid complexes, Cr(III)-alanine and Cr(III)-valine, were synthesized and characterized by IR spectroscopy, powder X-ray diffraction (XRD) analysis, magnetic susceptibility, and molar conductivity measurements. Molar conductivity measurements indicated that the composition of the metal complexes corresponds to a metal-amino acid ligand ratio of 1:3. The IR spectra indicated that the amino acids act as bidentate ligands with coordination involving the carboxyl oxygen and the nitrogen of the amino group. Magnetic susceptibility measurements revealed a six-coordinate local symmetry around the Cr(III) ions which depicted that the complexes were paramagnetic with magnetic moment values ranging from 5.10 to 6.00 BM. Powder XRD studies confirmed that the amino acid complexes were crystalline with monoclinic crystal structure. The in vitro biological activity of the investigated chromium(III) complexes with alanine and valine was tested against Bacillus subtilis, Staphylococcus aureus, Salmonella typhi, Pseudomonas aeruginosa, and Escherichia coli. All the microorganisms were standardized using 0.5 McFarland standard. The antimicrobial studies showed that the ligands were biologically active with an inhibition zone range of 10-17 mm and their metal complexes showed significantly enhanced antimicrobial sensitivity with an inhibition zone range of 12-21 mm. The standard drug showed slightly better activity with an inhibition zone range of 24-38 mm.

Keywords: Synthesis, Characterization, XRD, Biological activity

1. Introduction

In general, bacteria have the genetic ability to cause and acquire resistance to drugs, which are used as therapeutic agents. Although a large number of new antibiotics have been produced in the last three decades, the resistance of microorganisms to these drugs has increased (1). The microbial resistance presents a problem and the use of antimicrobial drugs in the future is still uncertain. Therefore, the measures to reduce or solve this problem include controlling the use of antibiotics, developing the research to better understand the genetic mechanisms of resistance, and continuing researching on how to develop new drugs, either synthetic or natural. The ultimate goal is to offer appropriate and efficient antimicrobial drugs to the patient. For centuries, people have used copper, iron, chromium, cobalt, and other transition metal ions complexes to inhibit the growth of harmful microbes (1). The complexes of transition metals with amino acids have been widely studied for their anti-microbial properties. They have been evaluated against several microorganisms with promising results. In addition to their ability to combat infection or neoplastic disease, these new agents should exhibit selective toxicity, chemical stability, and optimum rates of bio-transformation and elimination (2).

However, amino acids are absorbed well from intestinal lumen by specific active transport mechanisms. They display significant biological activity (3,4) and easily form stable complexes with most transition metal ions (5). The structure of organic compounds has been observed to undergo a significant modification in the addition of metals. In particular, amino acid and their metal complexes have attracted a great deal of attention as anti-cancer, anti-tubercular, anti-convulsant, insecticidal, antibacterial, anti-fungal, anti-biotic, and anti-inflammatory agents (6). The structure of alanine and valine are represented in Figs. 1 and 2, respectively. It is in the light of this challenge that the development of new drugs that are resistant to microorganisms cannot be overemphasized.
2. Materials and Methods

2.1. Synthesis and Experimentation

The reagents and solvents were of analytical grade which included Cr(NO₃)₃·9H₂O, ethanol, methanol, Dimethyl formide, Dimethyl sulphoxide, and Amino acids (Valine and Alanine). Clinical bacteria isolates (Bacillus subtilis, Staphylococcus aureus, Salmonella typhi, Pseudomonas aeruginosa, and Escherichia coli) were obtained from Ahmadu Bello University Teaching Hospital, Shika, Zaria, Nigeria. Ciprofloxacin (May and Barker) was used as the positive control.

Additionally, griffin water bath (E-550 model), capillary tube, thermometer (0-360°C), Fourier-transform infrared spectrophotometer (Cary 630 model, Agilent technologies), UV-visible spectrophotometer (Cary 300 model), melting point apparatus (Gallen Kamp), digital analytical balance (Sartorius ED 224S), magnetic susceptibility balance (Sherwood Scientific, Cambridge, UK) and conductivity meter (Jenway 4020) were used in the study. The X-ray powder diffraction analysis was carried out using Xpert-Pro X-ray diffractometer with Cu Kα radiation (λ = 1.54056 Å).

Moreover, funnel, measuring cylinder, water bath, round bottom flask, digital weighing balance, oven, furnace, melting point apparatus, and magnetic stirring hot plate are the equipment and apparatus used for the research.

2.2. Preparation of Ligand Solution

The ligands were prepared by stirring amino acid (BDH, 1.2 g, 2.0 mmol of valine, 1.0 mmol of alanine) with warm distilled water (20 mL, 20 g) in 30% NaOH (BDH, 0.33 mL) until the amino acid dissolved completely (7).

2.3. Preparation of Precursor Solution

The precursor solution was prepared by stirring the mixture of the metal salt (BDH, Cr(NO₃)₃·9H₂O salt) with distilled water (25 mL, 25 g) (7).

2.4. Synthesis of Complexes

The complexes were prepared by stirring the mixtures of precursor solution (20 mL) with amino acid solution (25 mL) at 60°C for 2 hours. A precipitate was formed, washed with ethanol, recrystallized in methanol and kept in a desiccator (7) as represented in formulas (1) and (2).

\[
\text{CrCl}_3 \cdot 9\text{H}_2\text{O} + 3\text{C}_3\text{H}_7\text{NO} \rightarrow \left[\text{Cr}(\text{C}_3\text{H}_7\text{NO})_3\right]_3 + 9\text{H}_2\text{O} + 3\text{HNO}_3
\]

Cr(III)-Valine Complexation Reaction   \hspace{1cm} (1)

\[
\text{CrCl}_3 \cdot 9\text{H}_2\text{O} + 3\text{C}_3\text{H}_7\text{NO} \rightarrow \left[\text{Cr}(\text{C}_3\text{H}_7\text{NO})_3\right]_3 + 9\text{H}_2\text{O} + 3\text{HNO}_3
\]

Cr(III)-Alanine Complexation Reaction   \hspace{1cm} (2)

3. Results and Discussion

3.1. Physical Properties of the Synthesized Complexes and the Ligands

The percentage yield of the synthesized complexes ranged between 50 and 70%, the melting points were found to be within the range of 258–298°C for the ligands and 230–290°C for the complexes. The complexes were colored and this is due to the fact that most of the transition metal complexes are colored and the color is observed due to d-d transition in the visible region. Analytical data and the properties of the compounds are shown in Table 1.

3.2. FTIR Results

The characteristic frequencies of the expected functional groups are depicted in Table 2. The assignment of peaks was done based on standard references previously published by several authors. The IR spectra of the ligands (alanine and valine) showed strong absorption bands at 1320 and 1330 cm⁻¹, respectively, which was due to carbon-nitrogen vibration (υ C-N), while in the metal complexes chromium(III)-alanine and chromium(III)-valine, the υ C-N absorption bands were obtained at 1398 and 1167 cm⁻¹, respectively. The new weak intensity bands in the region 541–571 cm⁻¹ and 359–489 cm⁻¹ in the spectra of the complexes were assigned to υ(M-O) and υ(M-N) stretching vibrations, respectively (8-10).

The absorption band of the amine group (N-H) ranged between 3448 and 3383 cm⁻¹, suggesting the possibility of the coordination of ligand through the nitrogen atom of the amine group. The N-H stretching vibration at 3119 cm⁻¹ in the ligand was shifted to a higher frequency in the complex, which is an indication that the coordination of the metal ion with the ligand took place via the nitrogen atom (11,12). The absorption frequency band at 1624 cm⁻¹ was ascribed to the C=O stretching vibration in the spectrum of the ligand, and it was shifted to 1578 cm⁻¹.
and 1584 cm⁻¹ in the complexes, which is an indication of the involvement of this group in metal-ligand bond formation (9). The major peak frequencies are depicted in Table 2.

However, for chromium-alanine complex, the N-H stretching vibration appears at 3416 cm⁻¹, and the lower frequency at 3052 cm⁻¹ in the free ligand. This indicates the formation of M-N bond via the amine group. The absorption frequency band at 1608 cm⁻¹ was attributed to C=O stretching in the spectrum of the free ligands and was shifted to a higher frequency at 1615 cm⁻¹ in the chromium-alanine complex. This evidence suggests the formation of M-O bond via the carboxylate group in the ligand. Moreover, the involvement of oxygen and nitrogen in the acid and amino groups in the ligand was further supported by the appearance of band around 432-435 cm⁻¹ for the M-OH stretching and around 535-700 cm⁻¹ for M-N stretching (4,8).

### 3.3. UV-Visible Results

The assignments have been done based on standard references and earlier studies published by several authors (13-15). The absorption bands in all the complexes located between 200 and 400 nm were attributed to the organic moiety and the absorption peaks above 400 nm were attributed to the formation of metal-ligand bond. In all the complexes, the n→π* characteristic band assigned to C=O bond appeared at 250–370 nm, while it appeared at 270–290 nm in the ligands spectrum, which also supported the involvement of carboxylate ion in the complex formation. The bands attributed to π→π* transitions in the complexes appeared at 270–390 nm, whereas in the free ligands, they were found at 270–280 nm. The presence of absorption band within 269–278 nm in the complexes was due to n→π* transition that was found at 269–278 nm in the free ligands. The presence of π→π*, n→π*, and n→σ* band in all the complexes indicated the presence of the functional groups of the parent ligands in the complex (C=O and NH₃).

A large shifting of the absorption band in all the complexes caused the appearance of a new band for d-d electronic transition, which indicates the possibility of forming metal-ligand coordination bond in the complexes. Transition complexes are generally colored and their colors are observed due to the absorption of light in the visible region. Therefore, the bands appearing at above 400 nm in all the complexes were clearly observed due to d-d electronic transitions that occurred in the complexes (15) as seen in Table 3.

### 3.4. Magnetic Susceptibility

The complexes of Cr(III) are high spin paramagnetic compounds as suggested by their magnetic moment values (16–20). The values of the magnetic moments of these complexes were in the range of 4.3–5.2 as represented in Table 4, which are comparable with the values reported for octahedral chromium(III) complexes (17-19). The value of the magnetic moments of these types of complexes was favorable for the formation of an octahedral complex. Therefore, the proposed geometry for all these complexes was octahedral.

### 3.5. Molar Conductivity

The molar conductivity values of Cr-alanine and Cr-valine were 25 and 30 cm²·Ω⁻¹·mol⁻¹, respectively, as shown in Table 5. This value showed that these complexes are not electrolytic in nature (20).

### 3.6. X-Ray Diffraction Results

The diffractogram of Cr(III) complex of valine had ten reflections with maxima at θ = 14.10° corresponding to a d value of 2.384 Å. However, the diffractogram of Cr(III) complex of alanine had only three reflections with maxima at θ = 10.20° corresponding to a d value of 2.60 Å as depicted in Figs. 3 and 4, respectively. However, a summary of the crystal information is presented in Table 6 and the spectra are represented in Figs. 3 and 4, respectively.

### 3.7. Metal Content Analysis

The percentage composition of the metal ions in the complexes ranged from 12.70 to 21.35 %, as presented in Table 7.

### 3.8. Water Content Analysis

The percentages of the water content of Cr-alanine and Cr-valine were 4.2% and 5.20%, respectively, as shown in Table 8.

### 3.9. Biological Activity

The results of the inhibitory activity (sensitivity test) showing the inhibition zones (mm) were obtained within
Characterization of complexes of Alanine and Valine

Table 3. Electronic Transition (nm) of the Ligands and Complexes

| Complex/Ligands | π→π* | n→π* | n→π* | d-d |
|----------------|-------|-------|-------|-----|
| Alanine        | 250   | 269   | 270   |     |
| Valine         | 260   | 278   | 280   |     |
| CrIII-alanine  | 333   | 347   | 298   | 566 |
| CrIII-valine   | 345   | 389   | 368   | 546 |

Table 4. Result of Magnetic Measurement of the Complexes

| Complex        | X x 10^4 (cgs) | X_α x 10^4 (cgs) | μ_α-calculated/observed (BM) |
|----------------|----------------|------------------|-----------------------------|
| CrIII-AL       | 1.60           | 6.45             | 3.87(3.95)                  |
| CrIII-VL       | 1.86           | 5.90             | 3.87(3.78)                  |

Table 5. Molar Conductivities Values of the Synthesized Complexes

| Compounds/Ligands | Molar Conductivity (cm² Ω⁻¹ mol⁻¹) |
|-------------------|-------------------------------------|
| C₆H₅NO₂ (alanine) |                                   |
| C₆H₅NO₂ (valine) |                                   |
| [Cr(C₆H₅NO₂)₂]²⁺ | 25                                 |
| [Cr(C₆H₅NO₂)₂]²⁺ | 30                                 |

Table 6. Summary of Crystal Data of the Complexes

|                      | Cr-AL                  | Cr-VL                  |
|----------------------|------------------------|------------------------|
| a, b, c (Å)          | 6.37, 10.30, 15.25     | 5.12, 8.89, 12.0       |
| α, β, γ              | 90, 90, 78             | 90, 90, 120            |
| Size (Å)             | 132                    | 156                    |
| System               | Monoclinic             | Monoclinic             |
| Geometry             | Octahedral             | Octahedral             |

Table 7. Metal Content Analysis of the Complexes

| Complexes | Weight of Complex (g) | Weight of Oxide (g) | Gravimetric Factor | % of metal Calculated | % of Metal Observed |
|-----------|-----------------------|---------------------|--------------------|-----------------------|---------------------|
| CrIII-AL  | 0.10                  | 0.0222              | 0.8725             | 18.60                 | 19.20               |
| CrIII-VL  | 0.10                  | 0.0245              | 0.8725             | 20.65                 | 21.35               |

Table 8. Water Content Analysis of the Complexes

| Complexes | Initial Weight (g) | Final Weight (g) | Loss in Weight (g) | % Of Water Calculated | % Of Water Observed |
|-----------|--------------------|------------------|--------------------|-----------------------|---------------------|
| CrIII-AL  | 0.20               | 0.1958           | 0.0042             | 5.40                  | 4.20                |
| CrIII-VL  | 0.20               | 0.1948           | 0.0052             | 7.10                  | 5.20                |

Note: AL=Alanine, VL=Valine

4. Conclusion

The complexes of Cr(III) with alanine and valine have been synthesized in basic aqueous medium and characterized. The molar conductivity measurements showed a ratio of 1:3. The IR spectra showed that the amino acids acted as bidentate ligands with coordination involving the carboxyl oxygen and the nitrogen of the amino group. Magnetic susceptibility measurements suggested a six-coordinate local symmetry around the pathogen and were more active than the parent ligand, suggesting the enhanced lipophilicity of the complexes on coordination as reported in previously published works (21-23). However, the organism appeared to be more susceptible to Cr-alanine with greater activity shown by the complex than the ligand as reported in the literature (23,24), and the standard drug (ciprofloxacin) showed better activity against all the Gram-negative bacteria. Therefore, this outcome suggests the engagement of these complexes in the development of new antibacterial agents for therapeutic applications. All the complexes showed potency against *Staphylococcus aureus*. This outcome is fascinating since *S. aureus* causes food poisoning and is highly resistant to most antibiotics.

Generally, the Gram-positive bacteria proved to be more susceptible to the amino acid complexes than the Gram-negative bacteria. The weak antibacterial activity against gram-negative bacteria may be ascribed to the presence of an outer membrane which poses hydrophilic polysaccharide chains as a barrier to the amino acid complexes (23,24). Therefore, the comparative studies of the ciprofloxacin and metal complexes indicated that the complexes showed antimicrobial activity against the studied microbial strains as reported by Kabbani et al (24).

Additionally, from the above analyses, the proposed structures for these complexes are shown in Figs. 5 and 6.

A comparative evaluation of the antibacterial activity of the complexes and ciprofloxacin was carried out against five test organisms. The result indicated that both complexes and ciprofloxacin were active against all the test organisms. All other complexes showed potency against the pathogen and were more active than the parent ligand, suggesting the enhanced lipophilicity of the complexes on coordination as reported in previously published works (21-23). However, the organism appeared to be more susceptible to Cr-alanine with greater activity shown by the complex than the ligand as reported in the literature (23,24), and the standard drug (ciprofloxacin) showed better activity against all the Gram-negative bacteria. Therefore, this outcome suggests the engagement of these complexes in the development of new antibacterial agents for therapeutic applications. All the complexes showed potency against *Staphylococcus aureus*. This outcome is fascinating since *S. aureus* causes food poisoning and is highly resistant to most antibiotics.

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Additionally, from the above analyses, the proposed structures for these complexes are shown in Figs. 5 and 6.
**Table 9.** The Diameter of Zone of Inhibition (mm) at Varying Concentration (mg/mL) of the Complexes and Ciprofloxacin

| Test organism/ concentration (mg/mL) | Cr(III)-VL Complex | Cr(III)-AL Complex | Ciprofloxacin |
|-------------------------------------|--------------------|--------------------|---------------|
| S. aureus                           | 12.5               | 100                | 12.5          |
| B. Subtilis                         | 15                 | 17                 | 10            |
| E. coli                             | 16                 | 13                 | 12            |
| S. typhi                            | 13                 | 15                 | 14            |
| P. aeruginosa                       | 13                 | 15                 | 12            |

**Table 10.** Minimum Inhibitory Concentration (MIC) of the Complexes (mg/L)

| Test Organism | Cr(III)-VL | Cr(III)-AL |
|---------------|------------|------------|
| S. aureus     | 12.5       | 25         |
| B. Subtilis   | 12.5       | 25         |
| E. coli       | 25         | 25         |
| S. typhi      | 25         | 50         |
| P. aeruginosa | 50         | 50         |

Note: AL, alanine; VL, valine.

**Table 11.** Minimum Bactericidal Concentration (MBC) of the Complexes (mg/L)

| Test organism | Cr(III)-VL | Cr(III)-AL |
|---------------|------------|------------|
| S. aureus     | 25         | 25         |
| B. Subtilis   | 25         | 50         |
| E. coli       | 50         | 50         |
| S. typhi      | 50         | 100        |
| P. aeruginosa | 100        | 100        |

Note: AL, alanine; VL, valine.
revealed that all the complexes were paramagnetic. Metal content and water content analyses showed that the complexes contained no water of crystallization. Powder XRD studies confirmed that the amino acid complexes were crystalline in nature and that they largely crystallized in monoclinic fashion. Moreover, it suggested that the bonding mode in the complexes was similar. In general, the complexes can be represented by the formula [M(II)\(\mathbf{L}_2\)] (where M=Cr(III), L=Valine anion, alanine anion). The complexes were stable in the air and soluble in DMF and DMSO. The low molar conductance values (10\(^3\) M) of solutions in DMSO indicated that all the complexes behaved as non-electrolytes. The antimicrobial studies suggested that the amino acid ligands were biologically active and their metal complexes showed significantly enhanced antimicrobial sensitivity against the studied microbial strains in comparison to the free ligands.

**Conflict of Interests**
There is no conflict of interest.

**Author's Contribution**
The authors were all supportive and help in the research.

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