



Synthesis, Biological and In Silico Studies of a Tripodal Schiff Base Derived from 2,4,6-Triamino-1,3,5-triazine and Its Trinuclear Dy(III), Er(III), and Gd(III) Salen Capped Complexes

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Abstract: A tripodal Schiff base ligand, 2,4,6-Tris(4-carboxybenzimino)-1,3,5-triazine (MT) and its trinuclear Dy(III), Er(III), and Gd(III) complexes were synthesized. These were characterized using UV-visible, IR, ¹H, and ¹³C NMR spectroscopies, elemental analysis, and molar conductivity measurements. The spectral studies indicate that the ligand is hexadentate and coordinates to the Ln(III) ions through the oxygen atoms of the carboxylic group. The trinuclear complexes were characterized as being bridged by carboxylate anions to the Dy(III), Er(III), and Gd(III) salen centers and displaying a coordination number of six. Biological studies revealed that MT is more active against the test micro-organisms relative to the trinuclear complexes. Acute toxicity studies indicate that MT is safe and has a wide range of effective doses (ED₅₀). In vivo antimalarial studies indicate that MT could serve as an effective antimalarial agent since it has parasitemia inhibition of 84.02% at 50 mg/kg and 65.81% at 25 mg/kg, close to the value (87.22%) of the standard drug—Artesunate. Molecular docking simulation studies on the compounds against SARS-CoV-2 (6Y84) and *E. coli* DNA gyrase (5MMN) revealed effective binding interactions through multiple bonding modes. The binding energy calculated for Er(III)MT-6Y84 and Er(III)MT-5MMN complexes showed active molecules with the ability to inhibit SARS-CoV-2 and *E. coli* DNA gyrase.

Keywords: tripodal schiff base; trinuclear Ln(III) complexes; antimicrobial activity; antimalarial activity; docking

1. Introduction

The emergence of resistant strains of the malaria parasite has necessitated the continued search for other effective, safe, and cheap antimalarial agents. The resistance of the malaria parasite to chloroquine and other anti-malarials has driven scientists into an intensive search for more effective agents against the scourge. Currently, there are reports of parasite resistance to the newly developed Artemisinin in some regions [1]. Additionally,



Citation: Oruma, U.S.; Ukoha, P.O.; Uzoewulu, C.P.; Ndefo, J.C.; Ugwuoke, S.C.; Ukwueze, N.N.; Eze, T.E.; Ekowo, L.C.; Eze, F.U.; Chinaegbomkpa, U.V.; et al. Synthesis, Biological and In Silico Studies of a Tripodal Schiff Base Derived from 2,4,6-Triamino-1,3,5-triazine and Its Trinuclear Dy(III), Er(III), and Gd(III) Salen Capped Complexes. *Molecules* 2021, 26, 4379. https://doi.org/10.3390/ molecules26144379

Academic Editors: Antonella Dalla Cort and Jan Janczak

Received: 19 May 2021 Accepted: 14 July 2021 Published: 20 July 2021

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the WHO warns that COVID-19 disruption to anti-malaria programs could cause deaths in Sub-Saharan Africa from malaria to double [2]. Thus, there is an urgent need for increased efforts in antimalarial drug discovery especially in Africa [3].

1,3,5-Triazine(*s*-triazine) analogs demonstrate a wide range of biological applications such as antimalarial, antimicrobial, antiviral, anticancer, antituberculosis, anti-HIV, antileishmanial, anti-inflammatory agents, insecticidal, and herbicidal [4–10]. Moreover, they are mostly used in the pharmaceutical, plastic, textile, and rubber industries [11–13]. They have also been used as dyestuffs, optical bleaches, explosives, and surface-active agents [14–17]. Additionally, cycloguanil, a derivative of 1,3,5-trazine, has been noted to be a cyclic metabolite of the antimalarial drug proguanil [18].

The *s*-triazine core has been applied in the development of less toxic bioactive compounds [19]. Consequently, melamine (2,4,6-triamino-*s*-triazine) based dendrimers are widely used in cancer and in many biological treatments [20]. Additionally, in the field of neuroleptics, *s*-triazines showed the most potent antimethamephetamine action with less toxicity than chlorpromazine [21]. Gupta and co-workers have reported thioetherlinked *s*-triazine to 1,2,4-triazine derivative bearing piperidine substituent as potent against *Leishmania donovani* with less toxicity compared with standard pentamidine and sodium stibogluconate, respectively [22]. Comparison of *s*-triazine based drugs with other nontriazine based drugs has shown that *s*-triazine based drugs are more potent [19].

The literature has shown that *s*-triazine has been combined with other heterocycles with the aim of developing more potent drug-like molecules [23–27]. However, there is no report of s-triazine scaffolds containing lanthanides. In recent years, lanthanides and lanthanide compounds have attracted a great deal of interest because they have applications in medicinal inorganic chemistry and in material science [28]. In medicine, lanthanide complexes are exploited as contrast agents for magnetic resonance imaging (MRI) and as radiotherapeutic drugs [28,29]. Due to the catalytic, magnetic, and luminescent properties of lanthanide complexes with organic ligands, they have found application in electroluminescent devices and diodes, lasers, cathode ray tubes, sensors, dosimeters, biological fluoro-immunoassays, imaging agents, organic light emitting diodes (OLEDS), display application, decoration purposes, and telecommunication [30,31]. Lanthanide complexes with desired functions have been designed and synthesized using different kinds of organic ligands [32,33]. Moreover, lanthanides have an interesting, but not well understood, biological role in living organisms as trace elements [34]. Our group recently reported neodymium(III) and thallium(III) complexes of N-benzothiazol-2-yl)-4-chlorobenzenesulphonamide as potential antibacterial agents [35]. Recently, Taha et al. (2011) [36] reported that Nd, Dy, Sm, Pr, Gd, Tb, and Er complexes with bis-(salicylaldehyde)-1,3-propylenediimine Schiff base ligand possessed high antibacterial activity against Shigella dysenteriae, Pseudomonas aeruginosa, and Proteus vulgaris (Gram-negative bacteria).

In view of the noted biological properties of *s*-triazines and lanthanide(III) complexes, we synthesized and characterized 2,4,6-Tris(4-carboxybenzimino)-1,3,5-triazine (MT) and its trinuclear Dy(III), Er(III), and Gd(III) salen capped complexes. Their in vitro antimicrobial, in vivo antimalarial activities, and acute toxicity were also investigated. Additionally, in silico studies were carried to have a view of the interaction of the synthesized compounds with SARS-CoV-2 (6Y84) and *E.coli* DNA gyrase (5MMN).

2. Results and Discussion

2,4,6-Tris(4-carboxybenzimino)-1,3,5-triazine (MT) was synthesized by the reaction of 2,4,6-triamino-1,3,5-triazine and 4-carboxybenzaldehyde (Scheme 1). MT was identified using FTIR, ^IH, and ¹³C-NMR elemental analysis. It is soluble in acetone, chloroform, ethylacetate, DMF, and DMSO.



Scheme 1. Synthesis of MT.

 $[Dy/Er/Gd(salen)]_2O$ were used as Ligand complexes (LC) because they were able to coordinate to MT. The reaction of the LCs with MT gave rise to the tripodal trinuclear complexes, $[{Dy/Er/Gd(salen)}_3(MT)].3H_2O$ (Scheme 2). These tripodal trinuclear complexes are the first examples of *s*-triazine based trinuclear complexes bridged to the Dysprosium(III), Erbium(III), and Gadolnium(III) centers by COO-. All complexes are stable at room temperature and soluble in DMSO and DMF but insoluble in water. The analytical data of MT and its trinuclear complexes are in good agreement with the proposed molecular formula as shown in Table 1. Molar conductivity measurements in methanol at room temperature show that the compounds are non-electrolytes [37].



Scheme 2. Synthesis of the tripodal trinuclear complexes; M = Dy, Er, Nd.

Table 1. Elemental and	physical data of 2,4,	6-Tris(4-carboxybe	enzimino)-1,3,5-triazin	e(MT) and its I	Ln(III) comple	exes
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					Elemental Analysis % Calc. and Found					
Compound	Color	$\Lambda_{\rm m}$ (Ω^{-1} cm ² mol ⁻¹)	Yield g (%)	M.p. (°C)	(С	H	I]	N
		(12 chi hioi)			Calc.	Found	Calc.	Foun	dCalc.	Found
$C_{27}H_{18}O_6N_6$ (MT)	White	-	(2.25) 86.21	346 ^a	62.07	61.95	3.45	3.40	16.09	15.90
C ₇₅ H ₅₇ N ₁₂ O ₁₂ Dy ₃ .3H ₂ O (Dy(III)MT)	Yellow	4.60	(0.27) 58.70	325 ^a	48.43	48.30	3.39	3.50	9.04	8.80
C ₇₅ H ₅₇ N ₁₂ O ₁₂ Er ₃ .3H ₂ O (Er(III)MT)	Yellow	6.70	(0.28) 59.57	324 ^a	48.08	48.10	3.37	3.50	8.97	8.60
C ₇₅ H ₅₇ N ₁₂ O ₁₂ Gd ₃ .3H ₂ O (Gd(III)MT)	Yellow	18.40	(0.30) 65.22	352 ^a	48.86	48.90	3.42	3.60	9.12	8.90

^a = decomposition temperature.

2.1. Electronic Spectra

The UV/Vis absorption spectra of MT and its complexes (10^{-4} moldm⁻³) were carried out in methanol at room temperature, see Table 2 and Figure S1. The spectra are shown in supporting documents. The absorption band of MT showed two bands at 233 and 291 nm assigned to π - π * transitions of the benzene rings. In the Ln complexes, these bands are red shifted, supporting the coordination of MT to the Ln ions (Figures S2–S4).

Table 2. Electronic absorption data of 2,4,6-tris(4-carboxybenzimino)-1,3,5-triazine (MT) and its Ln(III) complexes.

<u> </u>	λ	max		Band Assignment	
Compound -	nm	cm^{-1}	$\epsilon \times 10^{3} \text{ (mol}^{-1} \text{dm}^{3} \text{cm}^{-1}\text{)}$		
МТ	233	42,918	2.88	π-π*	
IM I	291	34,364	1.82	π – π *	
Dy(III)MT	348	28,736	10.9	n-π*	
Er(III)MT	347	28,818	9.72	n-π*	
Gd(III)MT	350	28,571	9.82	n-π*	

2.2. Infrared Spectra

The relevant stretching frequencies of MT and its Ln(III) complexes are shown in Table 3, Figures S5–S8. The absorption band due to the carboxylic acid C=O was observed at 1674 cm⁻¹ in MT [38,39]. This band shifted to higher frequencies of about 22–28 cm⁻¹ in the complexes, suggesting coordination of the ligand complexes via the carboxylic acid C=O of MT. This was further supported by the vibrations of the COO⁻ group observed at 1410–1412 cm⁻¹ in the complexes and at 1391 cm⁻¹ in the ligand [40]. The IR spectrum of MT showed two medium bands for C=N(a) and C=N(b) at 1501 and 1573 cm⁻¹, respectively. However, in the complexes, three bands were observed: C=N(a) bands at 1592–1596 cm⁻¹, C=N(b) bands at 1628–1636 cm⁻¹, and C=N(c) bands at 1536–1545 cm⁻¹ respectively. The C=N(a) and C=N(b) stretching vibration in the complexes shifted to higher wavenumbers in comparison to the same transition in the ligand indicating delocalization of the double bond of the tripodal Schiff base in coordinating with the ligand complexes. While the C=N(c) band, which was absent in the tripodal Schiff base ligand but present in the complexes, confirms that the ligand complexes were actually capped to the tripod. Similar observation has been made in literature [38–40]. Bands in the range of 579–571 cm⁻¹ in the tripodal-trinuclear complexes were assigned to ν (Ln–O) [36,41] while bands in the range of 489–437 cm⁻¹ were assigned to v (Ln–N) [42].

Table 3. IR band assignments (cm	-1) for 2,4,6-tris(4-carbox	ybenzimino)-1,3,5-triazine (MT)	and its Ln(III) metal complexe
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Compound	νC-H ar	νC=0	vC=N	νC-C	νCOO-	vC-N	vLn-O	vLn-N
MT	-	1674(s)	1501(m) a 1573(m) b	1422(m)	1391(m)	1168(m)	-	-
Dy(III)MT	3129(sh)	1698(m)	1594(m) a 1631(s) b 1542(m) c	1475(m) 1448(m)	1410(s)	1151(m) 1125(m)	579(m)	489(m) 437(m)
Er(III)MT	3158(w)	1696(m)	1596(s) a 1632(s) b 1543(s) c	1477(m)	1410(s)	1151(m) 1125(w)	575(m)	439(m)
Gd(III)MT	3125(m)	1702(m)	1592(m) a 1636(m) b 1540(s) c	1483(m)	1412(s)	1153(m)	571(m)	437(m)

Where a = from triazine ring; b = from azomethine linkage; c = from salen.

2.3. ¹H and ¹³C-NMR Spectra

The ¹H and ¹³C-NMR spectra of MT and its Ln(III) complexes are presented in Tables 4 and 5 respectively. The ¹H NMR spectrum of MT revealed a singlet peak at 10.17 ppm due to carboxylic proton (Figure S9). This peak disappeared in the complexes. The signal due to azomethine protons was observed between 9.77 and 8.29 ppm in the compounds. The signals in the range of 6.25–7.99 ppm in the compounds were assigned to aromatic protons. The signal due to ethylene protons appeared only in Dy(III)MT at 5.82 ppm. The signal due to DMSO was observed at 2.46 ppm in Dy(III)MT (Figure S10). The spectra for Er(III)MT and Gd(III)MT showed only solvent peak while that of Dy(III)MT showed only three peaks, probably due to the extent of paramagnetism of the Ln (III) ions [42,43].

Table 4. ¹H-NMR data of 2,4,6-tris(4-carboxybenzimino)-1,3,5-triazine (MT) and its Ln(III) Complexes (ppm).

Compound	OH _{Carboxylic}	CH=N	Haromatic	CH ₂ =CH ₂	DMSO
MT	10.17(1H,s)	9.77(1H,s) 8.29(1H,s)	7.99– 6.25(4H,m)	-	2.50
Dy(III)MT	-	-	7.65(4H,m)	5.82(4H,s)	2.46
Er(III)MT	-	-	-	-	-
Gd(III)MT	-	-	-	-	-

Table 5. ¹³C-NMR data of 2,4,6-tris(4-carboxybenzimino)-1,3,5-triazine (MT) and its Ln(III) Complexes (ppm).

Compound	Carboxylic Carbon	Azomethine Carbon	Carbons on Triazine Ring	Aromatic Carbons	DMSO Peak
MT	193.47	167.47, 165.75	139.07, 136.98	130.32, 129.92	39.89
Dy(III)MT	-	-	-	-	39.58
Er(III)MT	-	-	-	-	-
Gd(III)MT	-	-	-	-	-

The ¹³ C-NMR of MT gave signal at 193.47 ppm attributed to carboxylic carbon (Figure S11) [44]. The signal due to azomethine carbon was observed at 165.75 and 167.47 ppm in MT [45]. The signal at 139.07 and 136.98 ppm in MT has been assigned to carbons on the triazine ring. Carbons on the benzene ring are present at 130.32 and 129.92 ppm in MT. The spectra of the complexes revealed only the peak due to DMSO because of their paramagnetic nature (Figure S12) [42,43].

2.4. In Vitro Antimicrobial Activity

The results of the in vitro antimicrobial screening carried out on the compounds are given in Table 6. Ciprofloxacin, Tetracycline, Gentamicin, and Fluconazole were used as positive control while sterile DMSO served as negative control. These drugs have been chosen because they have the same mechanism of action, which is by inhibiting nucleic acid synthesis [46]. The structures of these drugs are shown in Supplementary Materials (Figure S13). Ciprofloxacin ($C_{17}H_{18}FN_3O_3$) belongs to fluoroquinolnes and inhibits bacteria growth by preventing Deoxyribonucleic acid (DNA) synthesis before mitosis. Tetracycline ($C_{22}H_{24}N_2O_8$) inhibits the multiplication of bacteria by binding to a subunit of the ribosomes, thereby inhibiting protein and nucleic acid synthesis and consequent death of the bacterium [47,48]. Gentamycin ($C_{21}H_{43}N_5O_7$) belongs to the class of aminoglycosides and acts by preventing protein synthesis, thereby inhibiting the synthesis of nucleic acids (DNA replication or RNA synthesis) and causes death of the bacterium. Fluconazole is an antifungal drug ($C_{13}H_{12}F_2N_6O$) and belongs to synthetic triazoles. Fluconazole inhibits fungal cytochrome P–450, an enzyme responsible for fungal sterol synthesis, thereby causing fungal cell walls to weaken [48].

			50 µg/mL			
Compound	B. c (ATCC 14579)	S. a (ATCC 6538P)	<i>P. a</i> (ATCC 9027)	E. c (ATCC 6749)	С. а	A. n
MT	5	15	12	6	31	21
Dy(III)MT	5	1	1	3	2	15
Er(III)MT	2	2	5	5	3	14
Gd(III)MT	3	9	2	8	2	12
25 μg/mL						
MT	-	10	7	2	19	15
Dy(III)MT	-	-		-	-	2
Er(III)MT	-	-	-	-	-	-
Gd(III)MT	-	3	-	4	-	-
			12.5 μg/mL			
MT	-	5	3	3	6	10
Dy(III)MT	-	-	-	-	-	-
Er(III)MT	-	-	-	-	-	-
Gd(III)MT	-	-	-	-	-	-

Table 6. Inhibition zone diameter (IZD in mm) of the compounds against typed strains (ATCC CULTURES) micro-organisms.

Key: B. c = Bacillus cereus, S. a = Staphylococcus aureus, P. a = Pseudomonas aeruginosa, E. c = Escherichia coli, C. a = Candida albicans. A. n = Aspergillus niger, (-) = no zone of inhibition observed.

The results obtained (Table 6) show that the compounds inhibited the growth of *Bacillus cereus, Staphylococcus aureus, Pseudomonas aeruginosa, Escherichia coli, Candida albicans,* and *Aspergillus niger* with inhibition zone diameter (IZD) in the range of 2–5, 1–15, 1–12, 2–8, 2–31, and 2–21 mm respectively. This reflects that the compounds exhibit higher activity against fungi (*Candida albicans* and *Aspergillus niger*) relative to the bacteria strains used. Among the test bacteria, the compounds were most active against *Staphylococcus aureus* followed by *Pseudomonas aeruginosa*. It was observed from the results (Table 6) that the activity of MT is higher than that of the trinuclear complexes. Hence, it could be inferred that the activity of the trinuclear complexes was not enhanced after anion coordination.

The inhibition zone diameter (IZD in mm) of the controls is displayed in Table S1. From Table S1, the inhibition zone diameters (IZD) of the controls are higher than that of the compounds.

The minimum inhibitory concentration (MIC) of the compounds and controls against *Bacillus cereus, Staphylococcus aureus, Pseudomonas aeruginosa, Escherichia coli, Candida albicans,* and *Aspergillus niger* are displayed in Table 7. From Table 7, the MIC of the compounds is found to be in the range of 50 for *Bacillus cereus,* 5.57–50 for *Staphylococcus aureus,* 6.27–50 for *Pseudomonas aeruginosa,* 7.3–50 for *Escherichia coli,* 2.6–50 for *Candida albicans,* and 2.3–50 for *Aspergillus niger.*

			MIC (µg/mL)			
Compound	<i>B. c</i> (ATCC 14579)	S. a (ATCC 6538P)	P. a (ATCC 9027)	<i>E. c</i> (ATCC 6749)	С. а	A. n
MT	50	5.57	6.27	7.3	2.60	2.30
Dy(III)MT	50	50	50	50	50	25
Er(III)MT	50	50	50	50	50	50
Gd(III)MT	50	25	50	25	50	50
		C	ontrols (µg/m	L)		
Т	1.90	1.80	0.63	2.15	2.10	0.58
F	6.25	6.25	6.25	2.80	0.64	0.74
СР	1.50	0.70	0.92	0.65	2.00	6.25
G	1.40	2.70	0.71	2.60	2.50	0.64

Table 7. Minimum inhibitory concentration (MIC) of the compounds and controls against test bacteria and fungi.

Legend: T = Tetracycline, F = Fluconazole, CP = Ciprofloxacin, G = Gentamycin; B. c = Bacillus cereus, S. a = Staphylococcus aureus, P. a = Pseudomonas aeruginosa, E. c = Escherichia coli, C. a = Candida albicans, A. n = Aspergillus niger.

From Table 7, the MIC of the controls is found to be in the range of $1.40-6.25 \ \mu\text{g/mL}$ for *Bacillus cereus*, 0.70–6.25 $\ \mu\text{g/mL}$ for *Staphylococcus aureus*, 0.63–6.25 $\ \mu\text{g/mL}$ for *Pseudomonas aeruginosa*, 0.65–2.80 $\ \mu\text{g/mL}$ for *Escherichia coli*, 0.64–2.50 $\ \mu\text{g/mL}$ for *Candida albicans*, and 0.58–6.25 $\ \mu\text{g/mL}$ for *Aspergillus niger*.

The MIC of MT against *Staphylococcus aureus* was 5.57 μ g/mL while that of Gentamycin was 2.70 μ g/mL. Among all the test compounds, MT was found to be the most active against *Candida albicans* and *Aspergillus niger* (MIC = 2.60 and 2.30 μ g/mL, respectively). However, the standard antifungal drug Fluconazole was more active against *Candida albicans* and *Aspergillus niger* (MIC = 0.64 and 0.74 μ g/mL, respectively) relative to MT.

2.5. Acute Toxicity

The acute toxicity test was recorded for MT alone as shown in Table S2. MT was selected because it showed higher sensitivity against the test micro-organisms. For mice administered with MT, no animal died within 24 h after administration (Table S2). This implies that MT is safe and has a wide range of effective doses (ED_{50}).

2.6. In Vivo Antimalarial Studies

The suppressive antiplasmodic effect of methanolic solutions of MT samples on albino mice are displayed in Table S3 while Table 8 gives the percentage parasitemia inhibition.

Table 8. Percentage of parasitemia inhibition.

Drug/Dose (mg/kg)	% Parasitemia	%Inhibition (PC)
MT 25	10.6667 ± 0.88192	65.81
MT 50	5.0000 ± 1.52753	84.02
Artesunate 5	4.00 ± 0.58	87.22
Dist. Water 5 mL/kg	31.33 ± 3.38	00.00

Artesunate is an antimalarial drug with a molecular formular of $C_{19}H_{28}O_8$ and was used as standard because of the presence of –COOH group in both Artesunate and MT. The structure is shown in Figure S14. Its mechanism of action is by inhibiting cytochrome

oxidase of the malaria parasite surface membrane—food vacuole membrane and mitochondrial membrane—thereby blocking the supply of nutrients from the host cell cytoplasm [49]. It was observed from Table S3 that the effect of the samples on weight (Wt), PCV, and Hb of the infected mice treated did not show an orderly pattern of dose dependent effect. However, the effect was significant compared to the negative control. MT shows a significant dose dependent reduction of the effect of the sample on PCV, though in a negative manner. The effect of MT on Hb concentration shows the same effect as in PCV.

The results in Table 8 showed a general dose dependent significant parasitemia inhibition compared with the negative control with MT inhibition of 84.02% at 50 mg/kg close to the value (87.22%) of the standard drug—Artesunate at 5 mg/kg. At 25 mg/kg, MT gave 65.81%. This implies that MT can be used as an antimalarial drug after further tests.

2.7. In Silico Studies

COVID-19 is a deadly disease that has led to the loss of so many lives all over the world. This disease is still ravaging the human race irrespective of the fact that some vaccines have been discovered. There is need for discovery of more drugs. It is therefore of paramount importance that drug-like molecules should target the SARS-CoV-2 6Y84 so as to identify compounds with high binding affinity, hence they could effectively combat COVID-19.

Table 9 shows the binding free energies of the ligand and the complexes for both SARS-CoV-2 and antibacterial in silico studies. The binding energy reveals a strong binding affinity the ligand and the complexes have with the receptors. MT showed the highest binding affinity with *E. coli* DNA gyrase while the complex, Er(III)MT demonstrated the highest binding affinity to the SARS-CoV-2 receptor (Figure 1). To gain further insights into the nature and type of chemical interactions involved, the binding poses of these compounds in the active binding sites of the receptors were analyzed and are shown in Figures 2–4. The atoms of the ligand, MT interacted with various amino acid residues of 5MMN. Significant interactions include H-bonding interaction between the O-atom of MT and VAL 160 through a distance of 3.02 Å. Pi-cation was observed between the pi-electrons of the phenyl group and ARG 76 (4.03 Å). Other amino acid residues involved include ALA 47, ILE 78 ILE 94, and PRO 79. Figure 3 depicts a well-fitted complex, Er(III)MT in the active binding site of 6Y84. The atoms of the complex interacted mostly with the amino acid residues through various hydrophobic interactions.

Compound	SARS-CoV-2. (6Y84)	E. coli DNA Gyrase (5MMN)
MT	-5.69	-7.03
Dy(III)MT	-7.30	-5.82
Er(III)MT	-7.98	-7.03
Gd(III)MT	-7.55	-6.59
CQ	-6.31	-
HCQ	-6.30	-
Cipro	-	-5.40

Table 9. Free binding energy of compounds against SARS-CoV-2 (6Y84) and *E. coli* DNA Gyrase (5MMN).



Figure 1. (a) *E. coli* DNA Gyrase B 24 kDa ATPase domain in complex with 1-ethyl-3-[8-methyl-5-(2-methyl-pyridin-4-yl)-isoquinolin-3-yl]-urea. (b) SARS-CoV-2 main protease with unliganded active site.



Figure 2. Binding interactions of MT with amino acid residues of *E. coli* DNA gyrase (5MMN). Key to bond type: orange dotted line = pi-cation bond; green dotted lines = H-bond; purple dotted line = pi-alkyl.



Figure 3. Binding pose of Er(III)MT in the active binding site of 6Y84.



Figure 4. 2D interaction of MT with the amino acid residues of E. coli DNA gyrase (5MMN).

3. Materials and Methods

3.1. Materials

All the chemicals used were of analytical reagent grade, purchased from Zayo–Sigma and were used as supplied without further purification. The melting points of the compounds were determined using Fischer Jones melting point apparatus and were uncorrected. Molar conductance measurements were carried out using 10^{-4} mol/L solutions of the complexes in methanol at room temperature using a WTW-LF 90 conductivity meter. Electronic spectra (in methanol) were recorded on a UV-Vis 1800 SHIMADZU spectrophotometer. Infrared spectra of the compounds were performed using KBr discs on a Perkin–Elmer (Waltham, MA, USA) 100 series version 10.03.08 FTIR spectrophotometer. The ¹H and ¹³C-NMR spectra of the compounds were recorded on a Bruker (Billerica, MA, USA) DPX 300 spectrometer in DMSO-d₆ at 300.13 MHz and 75.47 MHz respectively. Elemental analysis for C, H, and N were carried out using an LECO–CHN–932 analyzer.

3.2. Synthesis of 2,4,6-Tris(4-carboxybenzimino)-1,3,5-triazine(MT)

The method reported by Uysal and Ucan (2009) [39] was adopted. Melamine (0.63 g, 0.005 mol) was dissolved in benzene (5 cm³) stirred for 1 h, then 4-carboxybenzaldehyde (2.25 g, 0.015 mole) was added and refluxed for 4 h. A white precipitate was obtained, filtered, and recrystallized from a mixture of methanol and water, dried and stored over CaCl₂.

3.3. Synthesis of Ligand Complexes

The ligand complexes were prepared by addition of concentrated ammonia solution to a solution of $[Dy/Er/Gd(salen)]_2O$ in absolute ethanol, stirred at 50 °C until a pH of 12 was achieved [50,51].

3.4. Synthesis of [{Dy/Er/Gd(salen)}₃(MT)].3H₂O

 $[Dy/Er/Gd(salen)]_2O(0.00037 \text{ mol})$ was suspended in hot absolute ethanol (25 cm³) and a solution of MT (0.13 g, 0.00025 mol) in absolute ethanol was added while stirring. The reaction mixture was boiled under reflux for 4 h. The light yellow solid formed was washed with water and dried over CaCl₂ (Scheme 2).

3.5. In Vitro Antimicrobial Activity

The tripodal ligand and its trinuclear complexes were tested in vitro for their antimicrobial activities against American Type Culture Collection (ATCC) bacteria strains obtained from Rockville, MD, USA, by the Department of Microbiology, University of Nigeria, Nsukka while the fungi strains were isolated under clinical conditions. The typed bacteria culture was comprised of Gram-positive bacteria: *Staphylococcus aureus* (ATCC 6538P) and *Bacillus cereus* (ATCC 14579); Gram-negative bacteria: *Escherichia coli* (ATCC 6749) and *Pseudomonas aeruginosa* (ATCC 9027). The fungi strains used were *Candida albicans* and *Aspergillus niger*. The bacteria strains were tested for sterility on nutrient agar and then grown in nutrient broth at 37 °C for 24 h while the fungal strains were tested on Sabourand Dextrose Agar (SDA) and cultured in Sabourand Dextrose Liquid medium at 25 °C for 24 h. The overnight cultures were subsequently diluted and suspensions were made in normal saline and adjusted to 0.5 McFarland standards [52].

3.6. Antimicrobial Assay

The antimicrobial activities of all the synthesized compounds were determined by the agar cup diffusion technique [53]. The nutrient agar and SDA plates were inoculated with 0.1 mL broth culture of the test bacteria or fungi. Using a sterile cork borer, wells (5 mm in diameter and 2.5 mm deep) were bored into the inoculated agar. Fresh stock solutions (1000 μ g/mL) of the synthesized compounds were prepared in DMSO. The stock solution was further diluted with sterilized distilled water to 12.5, 25, and 50 μ g/mL for

antimicrobial evaluation. The wells were filled with 100 μ L of the test compounds by means of a sterile micropipette. Standard antibiotics, namely Ciprofloxacin, Tetracycline, Gentamycin, and Fluconazole were used as positive control while sterile DMSO served as negative control. Subsequently, 12.5, 6.25, and 3.125 μ g/mL of each positive control were prepared in DMSO. The bacteria plates were incubated at 37 °C for 24 h while fungal plates were incubated at 25 °C for 24 h. Inhibition zone diameter (IZD) around each well was measured in millimeters and recorded. The graph of IZD² against the log of concentration was plotted for each plate containing a specific compound and a micro-organism. The anti-log of the intercept on *x*-axis is the MIC.

3.7. Determination of Acute Toxicity (LD₅₀)

All experiments involving the use of mice were conducted in compliance with NIH guidelines for care and use of animals [54]. MT was used for the test because it showed higher sensitivity against the tested micro-organisms. The oral acute toxicity of the ethanolic solutions of the samples were estimated in albino mice (100-250 g) by upper-level lethal dose (LD_{50}) described by Lorke's method [55]. A total of four mice of both sexes were employed and an acclimatization period of 24 h was allowed. The samples were weighed and dissolved in 3% ethanol. The ethanolic solutions of the samples were administered orally at doses of 1000, 1600, 2900, and 5000 mg/kg. The animals were monitored for 24 h and the number of deaths per group recorded. Then, the mice were observed continuously for one hour after treatment, intermittently for three hours, and thereafter over a period of 24 h. The mice were observed for gross behavioral changes such as feeding, hair erection, lacrimation, mortality, and other signs of toxicity manifestation. The mice were given access to food and clean water during the study.

3.8. In Vivo Studies

The in vivo antimalarial assay was done based on a 4-day suppressive test using mice. The evaluation of the antimalarial activity against the methanolic solutions of the samples and Artesunate sensitive Plasmodium berghei (NK 65) was carried out according to a standard protocol of Peter's 4-day suppressive test [56]. Each of the 12 healthy experimental mice was inoculated intraperitoneally on the first day (Day 1). The infected mice were weighed and randomly divided into four groups of three mice each and four hours post inoculation were treated orally thereafter 24, 48, and 72 h post inoculation. For groups 1 and 2, 25 mg of sample/kg of mouse and 50 mg of sample/kg of mouse was administered orally for four consecutive days. For group 3, 5 mg of Artesunate/kg of mouse was administered while group 4 was given 5 mL of distilled water/kg of mouse for four consecutive days. On day 4 post inoculation, one drop of blood was taken from the tail of each experimental mouse and smeared on a microscope slide to make a thin film [57]. The thin films were fixed with methanol, stained with 10% Giemsa solution at pH 7.2 for 10 min, and examined microscopically. Parasitemia level was determined by counting the number of parasitized erythrocytes out of 100 erythrocytes per field in 4 random fields under a light microscope at magnification (\times 100) while average percentage parasitemia suppression was determined by comparing the parasitemia in the control group with the treated group.

Average percentage suppression
$$= pc - \frac{pt}{pc} \times 100$$

where pc = average parasitema in the control group, pt = average parasitemia in the treated group [58].

3.9. In Silico Study

The 3D structures of the receptors of SARS-CoV-2 (6Y84) and *E. coli* DNA gyrase (5MMN) as shown in Figure 1 were retrieved from the Protein Data Bank. The 3D structures of the ligand and complexes drugs drawn using MarvinSketch 17.2.6.0 were energy minimized and docked into the active binding sites of these receptors. Docking protocols

were validated by reproducing the PDB crystal structures in silico. The various binding poses produced were analyzed with Discovery Studio vs16.1.0.15350.

DNA gyrase, a type II topoisomerase, is found in all bacteria. They consist of two subunits: GyrA and GryB of *E. coli* gyrase. These enzymes are responsible for catalyzing topological changes in DNA and have proved to be drug targets for therapeutic agents.

4. Conclusions

A tripodal Schiff base ligand derived from 2,4,6-triamino-1,3,5-triazine and its novel trinuclear Dy(III), Er(III), and Gd(III) salen capped complexes were synthesized and characterized based on various physico-chemical and spectral studies. The ligand was found to be tripodal and to coordinate to the ligand complexes via the carboxylic group. In vitro antimicrobial tests indicated that the tripodal ligand, MT is more active against the test micro-organisms relative to the trinuclear complex. Acute toxicity studies reveal that MT is safe and has a wide range of effective doses (ED_{50}). In vivo antimalarial studies indicate that MT could serve as an effective antimalarial agent since it has parasitemia inhibition of 84.02% at 50 mg/kg and 65.81% at 25 mg/kg close to the value (87.22%) of the standard drug—Artesunate at 5 mg/kg. The chemical interactions of our synthesized ligand and complexes with SARS-CoV-2 and DNA gyrase revealed multiple effective bonding interactions. Through docking simulation, the binding energy calculated for the two drug targets showed active molecules with the ability to inhibit SARS-CoV-2 (COVID-19) and DNA gyrase.

Supplementary Materials: Figure S1: Electronic absorption spectrum of MT, Figure S2: Electronic absorption spectrum of Dy(III)MT, Figure S3: Electronic absorption spectrum of Er(III)MT, Figure S4: Electronic absorption spectrum of Gd(III)MT, Figure S5: Infrared Spectrum of MT, Figure S6: Infrared Spectrum of Dy(III)MT, Figure S7: Infrared Spectrum of Er(III) MT, Figure S8: Infrared Spectrum of Gd(III)MT, Figure S9: ¹H-NMR Spectrum of MT, Figure S10: ¹H-NMR Spectrum of Dy(III)MT, Figure S11: ¹³C-NMR Spectrum of MT, Figure S12: ¹³C-NMR Spectrum of Dy(III)MT, Figure S13: Structures of the drugs used as standard, Figure S14: Structure of Artesunate, Table S1: Inhibition Zone Diameter (IZD in mm) of the Controls, Table S2: Determination of Acute Toxicity (LD50), Table S3: Anti-plasmodic Effect of Samples on Albino mice.

Author Contributions: Conceptualization, U.S.O. and P.O.U.; methodology, U.S.O. and P.O.U.; software, C.J.E., U.S.O., J.C.N., and F.U.E.; validation, C.J.E., U.S.O., T.E.E., and C.P.U.; formal analysis, U.S.O. and P.O.U.; investigation, U.S.O., P.O.U., C.J.E., T.E.E., and N.N.U.; resources, U.S.O. and P.O.U.; data curation, U.S.O., F.U.E., J.C.N., and U.V.C.; writing—original draft preparation, U.S.O. and S.N.O.; writing—review and editing, C.J.E., L.C.E., N.N.U., and S.C.U.; visualization, U.S.O., S.C.U., and C.P.U.; supervision, P.O.U.; project administration, U.S.O. and P.O.U. All authors have read and agreed to the published version of the manuscript.

Funding: This research received no external funding.

Institutional Review Board Statement: Not applicable.

Informed Consent Statement: Not applicable.

Data Availability Statement: Not applicable.

Conflicts of Interest: The authors declare no conflict of interest.

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