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Article

Study of the Conformation of Bromated Guanidines with Potential Anti-Leishmania Activity

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Abstract: Leishmaniasis are neglected diseases that affect regions such as South Asia, South Africa and Latin America, less developed regions. Current treatment, for cutaneous, mucocutaneous and visceral leishmaniasis, which are the main forms of the disease, is based on the administration of pentavalent antimonial and amphotericin B. However, these drugs have low efficacy and high toxicity. For this reason, the search for new treatments for this disease is necessary, based on studies of new molecules and their conformations. The research proposed the conformational study of brominated guanidine compounds with potential antileishmanial activity using Nuclear Magnetic Resonance (NMR) and X-ray diffraction (XRD) techniques. The study involved two brominated molecules, LQOF-G35 and LQOF-G35-Br. The later was synthesized by the reaction of LQOF-G35 and NBS under IR irradiation at 120 Watts of potency and dichloromethane as solvent by 12-h of exposition. The obtained results demonstrated the efficiency of the bromination method, since two bromines atoms entered the molecule. Furthermore, NMR analysis show the occurrence of a *Z* to *E* conformational change. This feature was well confirmed by comparative DRX study of the LQOF-G35 and LQOF-G35-Br compounds.

Keywords: Guanidines compounds; Leishmaniasis; Bromination; NBS; X-ray and NMR conformational study

1. Introduction

The research and development of drugs to treat diseases, such as leishmaniasis, involves the creation of molecules through pharmacological design, synthesis, and biological and structural characterization. Leishmaniasis, prevalent in Southeast Asia, Sub-Saharan Africa and Latin America [1], encompasses several clinical-pathological manifestations, among them the most common known as cutaneous leishmaniasis, caused by *Leishmania amazonensis*, manifests itself through lesions and ulcers on the skin, and the most severe, visceral leishmaniasis, is characterized by anemia, significant weight loss, enlarged spleen, fever and often results in death, the latter being observed in 85-90% of untreated cases [1,2,3].

Current treatments, such as chemotherapy with pentavalent antimonials, paromomycin, amphotericin B, and miltefosine, present challenges, including low efficacy and severe side effects such as cardiotoxicity, pancreatitis, hepatotoxicity, parasite resistance, and long-term administration [2-5]. Therefore, the search for safer and more effective antileishmania compounds against leishmaniasis is essential [4,5,6].

Guanidine compounds are often found in nature and used in the synthesis of a variety of organic compounds, such as quinazolines, oxazolidinones, lactones, and carbonates [7-11]. Novel guanidine derivatives exhibit several biological effects, such as cardiovascular dilation, antihistamine properties, anti-inflammatory activity, antidiabetic effects, antibacterial, antifungal, antiprotozoal/antiparasitic properties, and antiviral activity [7,12-21]. Recently, some guanidine compounds have been studied in relation to leishmaniasis [22,23].

In 2019, Espírito Santo R. D. *et al.*, carried out the structural characterization by nuclear magnetic resonance (NMR) and the evaluation of the antiparasitic activity of a series of guanidine compounds [20], in this study some compounds proved to be highly effective and promising due to their low toxicity against mammals and high lethality for parasites. One such compound was LQOF-G2, which has the bromo atom in the "para" position of the aniline ring.

This compound stood out as the most effective candidate against leishmaniasis in the study, with a preferential 'Z' conformation being observed [22]. Another guanidine compound, LQOF-G35, which has a bromine atom as a substituent in the "ortho" position of the aniline ring, was studied in relation to *Leishmania amazonensis* and *Leishmania braziliensis* in the promastigote forms. This compound demonstrated an IC₅₀ value of 29.82 and 25.87 μM, respectively, standing out as an even more promising candidate when looking at the IC₅₀ values in amastigotes 7.63 and 4.62, respectively [24].

Based on these reported results, it was decided to examine the effects of incorporating additional bromine atoms into the structure of the guanidine compound LQOF-G35. Therefore, the bromination reaction with *N*-Bromosuccinimide was performed using irradiation with an IR lamp [25-30] and the results are described here.

2. Results and Discussion

The characterization of the compounds was carried out initially by electronic ionization mass spectrometry and melting point measurements (Table 1).

Table 1. Melting point and molecular ion data for guanidines.

Compound	Melting point (°C)	Molecular Ion [M] ⁺
LQOF-G35	103.0 – 104.0	406 <i>m/z</i>
LQOF-G35-Br	147.9 – 149.6	565 <i>m/z</i>

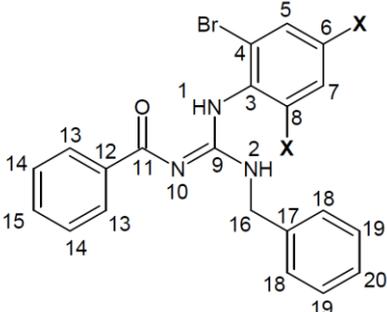
LQOF-G35-Br also was studied by high resolution mass spectrometry with electrospray ionization in the positive ion mode (ESI(+)-MS) and was identified by the detection of their intact protonated molecule, which were selected and further fragmented via ESI(+)-MS/MS experiments. Figure S3 shows the HRESI(+)-MS and ESI(+)-MS/MS spectra of the compound LQOF-G35-Br (*m/z* 565.8898 [M+H]⁺). The ESI(+)-MS/MS experiments yielded fragment ions at *m/z* 487 and 365 that are in agreement with formation of the desired product.

NMR studies were conducted at -10°C (263K) for LQOF-G35 and 10°C (283K) for LQOF-G35-Br. The NMR spectra can be found in the supplementary material and the most relevant data are highlighted in the Table 2.

The ¹H NMR of LQOF-G35 revealed the signals of the hydrogens NH 1 and 2, with chemical shifts of 5.19 and 12.33 ppm respectively, which were confirmed by HMBC ¹Hx¹⁵N (Figure S4). In contrast, the NH 1 and 2 hydrogens of LQOF-G35-Br were observed at 8.56 and 8.30 ppm respectively. This difference is due to the stabilization of both different conformations by the formation of a hydrogen bond between the hydrogen atom of the coplanar NH groups and the carbonyl oxygen atom, forming a pseudo six-member cycle. In this way, the NH hydrogen atom involves in the intramolecular hydrogen bond becomes less shielded. Therefore, it is possible to state that there was a conformational change from *Z* to *E* after or simultaneously bromination. This result was confirmed by XRD study (Figure 2).

Another important signal for LQOF-G35-Br was observed at 7.73 ppm, a singlet that integrates two protons, this signal represents the hydrogens H5/H7 (chemically equivalent). Which was also confirmed by HMBC ¹Hx¹³C due to its correlations with four ¹³C signals, three of which are associated with quaternary carbons (C4/C8 115.4 ppm, C3 119.3 ppm and C6 143.6 ppm), as well as the linker carbon C5/C7 itself, observed at 134.6 ppm.

Table 2. Chemical shifts (ppm) obtained by ¹³C NMR (500 MHz; CDCl₃) of guanidine compounds.



LQOF-G35. X = H

LQOF-G35-Br. X = Br

HYDROGENS											
	H1	H2	H5	H8	H10	H13	H14	H16	H18	H19	
LQOF-G35	5.19	12.33	7.68	7.31	-	8.30	7.43	4.76	7.38	7.38	
LQOF-G35-Br	8.56	8.30	7.73	-	7.56	7.56	7.49	4.72	7.48	7.37	

CARBONS												
	C3	C4	C6	C8	C9	C11	C12	C13	C14	C16	C17	C18
LQOF-G35	138.1	121.5	-	-	158.1	177.7	138.4	129.3	128.1	45.0	134.8	127.8
LQOF-G35-Br	119.3	115.4	143.6	115.4	144.4	166.7	132.8	127.1	127.7	45.1	138.1	129.2

Aliphatic hydrogens (H16) were identified at 4.7 ppm, corroborated by the HSQC $^1\text{H}\times^{13}\text{C}$ technique. Aromatic hydrogens were observed in the region between 7.0-8.5 ppm.

Through ^{13}C NMR it was possible to observe the C16 aliphatic signal for both guanidines. Between 115 and 166 ppm, all remaining carbons in the structures were observed. However, for LQOF-G35-Br there was the appearance of an additional quaternary carbon compared to LQOF-G35. Initially, this signal could indicate mono bromination in the structure. However, one of the bromines bonds (in ortho) makes carbons C4 and C8 chemically equivalent, resulting in only one signal for these and another for C6 (para position).

In conjunction with these analyses, a NOESY study was conducted, providing corroborating spatial information between hydrogens. Figure 1 illustrates the main spatial correlations for guanidines. We can observe intense correlations between H2-H16 and H18-H16 for LQOF-G35, with emphasis on the H13-H16 and H8-H2 correlations, what indicate the *Z* conformation. For LQOF-G35-Br, intense correlations were observed between H1-H16, H18-H16 and H19-H16, therefore, indicating the *E* conformation.

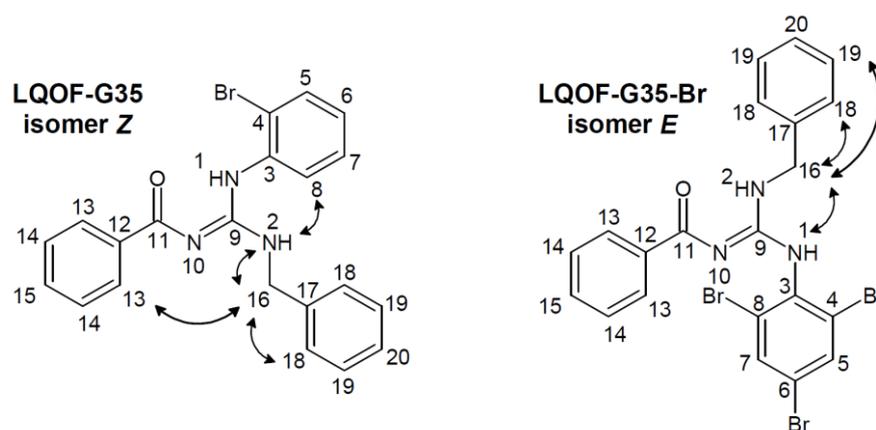


Figure 1. Conformational NOESY of guanidines LQOF-G35 and LQOF-G35-Br.

Single crystals of both compounds were obtained and analyzed using the SCXRD technique. The compound LQOF-G35 crystallizes in the non-centrosymmetric trigonal space group $P3_2$ and contains three independent molecules in the asymmetric unit, as shown in Figure 2. The SCXRD

study of LQOF-G35 show that the guanidine group present a resonant structure, with C–N bond distances ranging from 1.330 Å to 1.357 Å, formed by the contribution of the different resonance hybrids. As indicated by the NMR analyses, the LQOF-G35 has the *Z* conformation, with the amine group from the aniline stabilized by a six membered ring intramolecular hydrogen bond, N1H...O1. The presence of the hydrogen atoms bonded to the nitrogen atoms from the aniline and aminobenzyl groups was also indicated by the SCXRD analysis, with the evaluation of the electronic density maps.

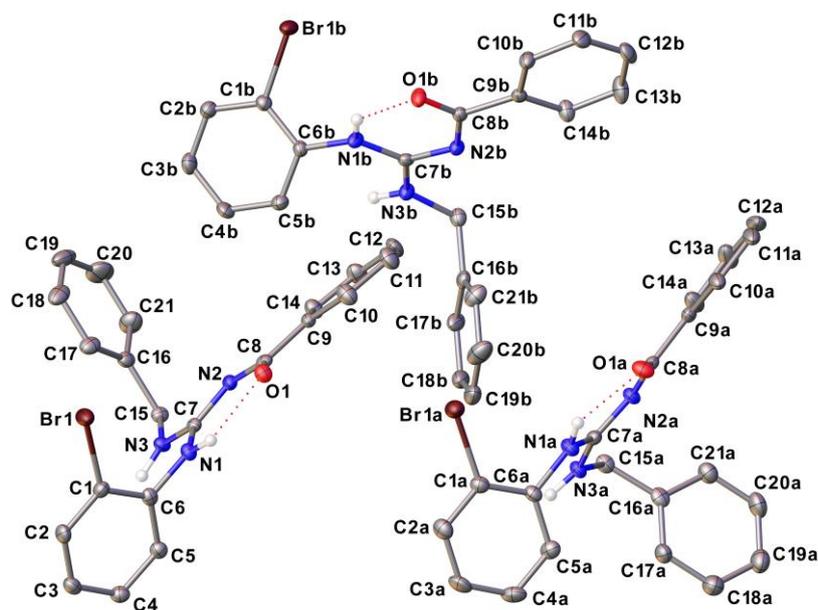


Figure 2. ORTEP type representation of the asymmetric unit of LQOF-G35, with thermal ellipsoids represented with 50% of probability. Hydrogen atoms bonded to the carbon atoms were omitted for clarity.

On the other hand, compound LQOF-G35-Br crystallized in the monoclinic space group $P2_1/c$, containing just one molecule per asymmetric unit (Figure 3). The analysis of the C–N bond lengths of the guanidine group, C7–N1 (1.290(3) Å), C7–N2 (1.409(3) Å) and C7–N3 (1.348(3) Å), indicate that the double bond is not in resonance in this case, just involving the atoms C7 and N1. An inversion in the position of the aniline and aminobenzyl groups was verified in the structure of LQOF-G35-Br, presenting an *E* conformation about the C7–N2, being now stabilized by another six membered ring intramolecular hydrogen bond, in this case involving N3H...O1. The electronic density maps analysis also confirmed the presence of the hydrogen atoms bonded to the nitrogen atoms N2 and N3.

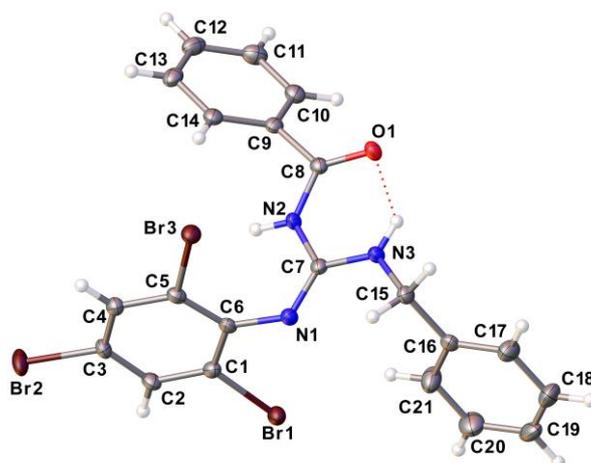
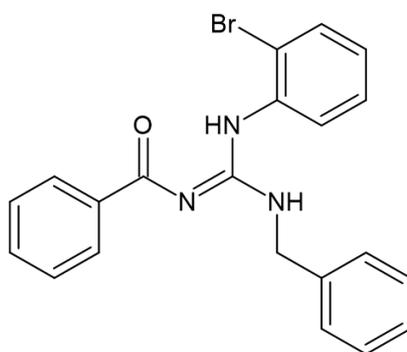


Figure 3. ORTEP type representation of the asymmetric unit of LQOF-G35-Br. Thermal ellipsoids are represented with 50% of probability.

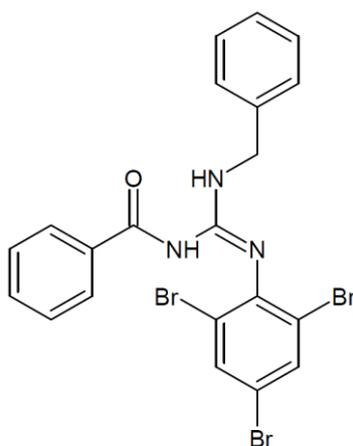
To better understanding the conformational change, LQOF-G35 was subjected to 12 hours of infrared irradiation and subsequently analyzed by ^1H NMR. This experiment revealed that without NBS the initial *Z* conformation of LQOF-G35 was maintained. Therefore, the polybromination was responsible for this conformational change in LQOF-G35-Br, more specifically the entry of the second atom of bromo, which was confirmed because the compound with two bromine atoms in the aniline ring was synthesized and it showed only the *Z* conformation.

The *Z/E* conformational ratio is directly related to the position of the double bond $\text{N10}=\text{C9}$ (Figure 1). The entry of the second atom of bromo in the aniline moiety, promote the increase in energy resulting in the opening of the *pi* bond and subsequent rotation around the $\text{N10}-\text{C9}$ sigma bond to reach the *E* conformation with the restored double bond localized between in $\text{N10}=\text{C9}$.

3.1. Structural data of compounds



(*Z*)-*N*-benzoyl-*N*-benzyl-*N*-(2-bromophenyl)guanidine (LQOF-G35). MM: 407.06 g.mol⁻¹. White solid. Melting point: 103-104°C. ^1H NMR 263K (500.16 MHz, CDCl_3) δ *Z* isomer = 12.33 (s, 1H), 8.30 (d, 2H), 7.69 (d, 1H), 7.50 (q, 2H), 7.43 (t, 3H), 7.41 – 7.34 (m, 4H), 7.31 (m, 1H), 7.16 (t, 1H), 5.19 (t, 1H), 4.84 (d, 2H). ^{13}C NMR (125.765 MHz, CDCl_3) δ = 177.7 (C=O), 158.1 (C=N), 138.2 (C), 137.8 (C), 134.6 (C), 134.0 (CH), 131.4 (CH), 129.1 (CH), 128.8 (CH), 128.7 (CH), 127.9 (CH), 127.6 (CH), 127.5 (CH), 127.4 (CH), 121.2 (C-Br), 45.0 (N-CH₂). GC-MS/EI (*m/z* 406). ESI(+)-MS *m/z* found 408.0705, *m/z* calculated for [$\text{C}_{21}\text{H}_{18}\text{BrN}_3\text{O} + \text{H}$]⁺: 408.0706; ESI(+)-MS/MS: $\text{M} + \text{H} - \text{C}_6\text{H}_5\text{CONH}_2$ ⁺ *m/z* 287.0175, [$\text{M} + \text{H} - \text{C}_{15}\text{H}_{12}\text{N}_2\text{O}$]⁺ *m/z* 171.9757, [$\text{M} + \text{H} - \text{C}_{13}\text{H}_{12}\text{BrN}_2$]⁺ *m/z* 122.0603;



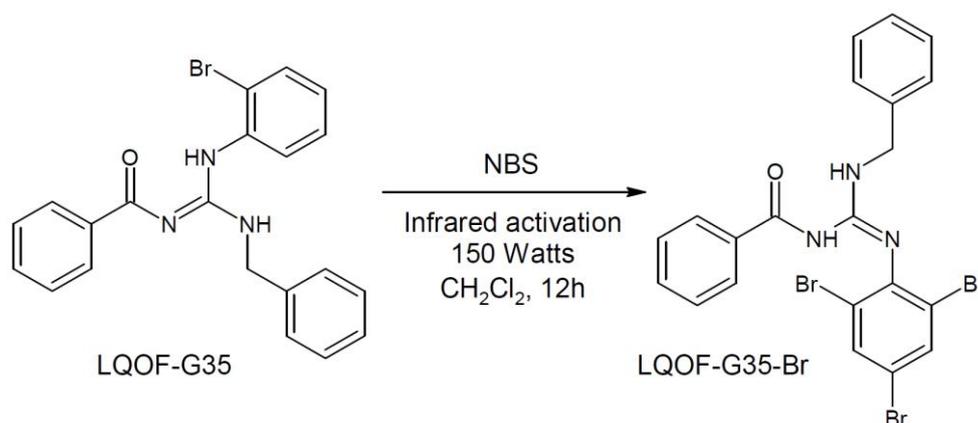
(*E*)-*N*-benzoyl-*N*-benzyl-*N*-(2,4,6-tri-bromophenyl)guanidine (LQOF-G35-Br). MM: 566.09 g.mol⁻¹. White solid. Melting point: 148-149°C. ^1H NMR 10 °C (500.16 MHz, CDCl_3) δ *E* isomer 8.86 (s, 1H), 8.22 (s, 1H), 7.73 (s, 2H), 7.62 – 7.53 (m, 3H), 7.51 – 7.43 (m, 6H), 7.37 (t, 3H), 7.32 – 7.27 (m, 2H), 4.72 (d, 2H). ^{13}C NMR (125.765 MHz, CDCl_3) δ = 166.6 (C=O), 144.4 (C=N), 143.6 (C-Br), 138.1 (C), 134.6

(CH), 133.2 (CH), 132.8 (C), 129.2 (CH), 128.6 (CH), 127.7 (CH), 127.3 (CH), 127.0 (CH), 119.3 (C), 115.4 (2C-Br), 45.0 (N-CH₂). GC-MS/EI (m/z 565): 77, 91, 105, 122, 285, 406, 486. HRESI-MS m/z 565.8898 [M+H]⁺ (calcd for C₂₁H₁₆Br₃N₃O₁, 566.091, Δ = -0.2012 ppm). Ionic fragment observed in ESI(+)-MS/MS: [M+H-C₁₄H₁₁BrN₂O]⁺ m/z 486.5555.

4. Materials and Methods

4.1. Guanidine bromination using NBS and IR irradiation with a IR lamp (150 Watts)

As a bromination method, N-Bromosuccinimide (NBS) catalyzed by 150-watt infrared light was used, under magnetic stirring for 12 hours (Scheme 1), a reflux system was added to avoid evaporating the solvent (Dichloromethane). This choice was motivated by the commercial availability of NBS and its good stability, which minimizes self-decomposition. Furthermore, NBS is compatible with the solubility of guanidines. The reaction was performed using a molar relationship guanidine: NBS of 1 mol of guanidine to 1.4 mol of NBS. After the reaction end, the reaction mixture was cooled to 10°C and washed with water, then the organic layer (dichloromethane) was separated and dried on magnesium sulfate. Subsequently, the reaction mixture was filtered, the solvent was evaporated, and the cru product was recrystallized from a mixture of petroleum ether and diethyl ether (9:1 v/v). Then, the suspension was filtered off and the solid was dried by reduced pressure and suspended in ethanol with ultrasound and recrystallized one more time.



Scheme 1. Bromination of the compound LQOF-G35 with formation of the polybrominated compound LQOF-G35-Br as confirmed by XRD.

4.2. Melting point

Melting point was obtained using a WRS-2 Micro Processor Melting-point apparatus. The samples were placed in a capillary tube, and pre-heating and final ramp temperatures were selected for 60°C and 250°C, respectively. The heating rate used was 2.0°C/min.

4.3. Electronic ionization mass spectrometry

The mass spectra of the guanidines were acquired by direct introduction (DI) in the Mass Spectrometry, with a Shimadzu QP-2010 Plus equipment with Electron Ionization (EI). The parameters used for the analysis were: Interface temperature: 240°C; Ionization chamber: 300°C; Time to solvent cut; 0.5 min; Initial time: 0.7 min; Final time: 25 min; DI temperature program: Initial temperature: 50°C, with heating from 20°C/min to 350°C and standby time 10min. The analysis was performed using acetonitrile and dichloromethane as a solvent. The energy used for ionization was 70eV.

4.4. RMN

¹H, ¹³C and other 2D NMR spectra were recorded on a Bruker Avance III-HD – 500 MHz. The resonance frequency for ¹H NMR was 500.16 MHz and for ¹³C NMR 125.765 MHz. Chemical shifts

for ^1H NMR and ^{13}C NMR were referenced to TMS, analysis was performed in CDCl_3 and all chemical shifts were reported in ppm. Data are presented as following: chemical shift, multiplicity (s = singlet, d = doublet, dd = doublet of doublet, t = triplet, qua = quadruplet, qu = quintuplet, m = multiplet, br s = broad singlet), integration, and coupling constants (in Hertz).

4.5. HRESIMS

The UPLC-ESI-QTOF-MS/MS analysis was performed on a Waters modular UHPLC consisting of a QSM Acquity® HClass quaternary pump, an Acquity® Sample Manager – FTM autosampler, and an Acquity® PDAe λ diode array detector, hyphenated to a Waters Xevo® G2-XS mass spectrometer, equipped with an electrospray ionization source (ESI), a quadrupole analyzer, and a collision cell configured in sequence with a time-of-flight (TOF) analyzer. The chromatographic separation was performed in gradient mode using a solution of formic acid at 0.1% v/v as mobile phase in ultrapure water as component A and acetonitrile acidified with formic acid at a concentration of 0.1% v/v as component B. Chromatographic runs were performed at a flow rate of 0.5 mL.min $^{-1}$, starting with 5% of component B in the mobile phase composition. The variation of the elution force was programmed so that in 5 min the %B increased until reaching the value of 100%. The analysis was performed on a Waters Acquity® UPLC HSS T3 reversed phase column (C18) containing 1.8 μm particles with 2.1 mm internal diameter and 100 mm length. The injection volume applied to the column was 0.2 μL . Full scan and MS/MS mass spectra were acquired in positive mode in the range of m/z 100 to 1500. Electrospray capillary and transfer cone voltages were maintained at (+) 2.5 kV and 20V respectively. The ESI source and desolvation gas temperatures were set to 120.0 and 350.0 $^\circ\text{C}$ respectively. The acquisition of MS/MS spectra was performed in Fast DDA mode (data dependent acquisition), in which only ions with intensity above 100 were fragmented. Ion fragmentation was performed by applying a voltage ramp so that for m/z 100 the energy ranged from 10 to 40 eV and for m/z 1500, from 40 to 70 eV.

4.6. Single crystal X-ray diffraction (SCXRD)

Single crystals of LQOF-G35 and LQOF-G35-Br were obtained and analyzed at 100 K on a Rigaku XtaLAB Synergy-S Dualflex diffractometer equipped with an HyPix 6000HE detector, using a Cu $K\alpha$ radiation (1.54184 \AA). Data collection, cell refinement, data reduction and absorption correction were performed using the CrysAlisPro software [31]. Intrinsic phasing method from the SHELXT-2018/2 program [32] was used to solve the structures, while the refinement of the non-hydrogen atoms was performed with non-linear the least-squares minimization on F^2 using the SHELXL-2019/2 program [33] and considering anisotropic displacement parameters. Hydrogen atoms were calculated at idealized positions using the riding model. The calculated positions of the hydrogen atoms bonded to the nitrogen atoms from the guanidine group were confirmed by the analyses of the electron density maps. Olex2 [34] was used for the solution and refinement of the structures, and to generate the graphical illustrations. Tables containing the information about the data collection, refinement and structural parameters are available in the Supplementary Materials.

The CIF file of LQOF-G35 and LQOF-G35-Br were deposited in the Cambridge Structural Data Base with CCDC number 2320314 and 23200315, respectively. Copies of the data can be obtained, free of charge, via www.ccdc.cam.ac.uk.

5. Conclusions

Two guanidine compounds have been unambiguously characterized. MS-EI and HRESI-MS analyzes showed one bromine atom in the compound LQOF-G35 and three Br in LQOF-G35-Br structures.

NMR and DRX have confirmed that all the bromine atoms are in the aniline ring. A conformational change from Z to E had also corroborated by these studies.

Finally, existence of the guanidine compound containing two-Br atoms on the same aniline ring, is strong evidence of the conformational change is occurring simultaneously to second bromine entrance.

Supplementary Materials: The following supporting information can be downloaded at: www.mdpi.com/xxx/s1, Figure S1: EI-MS (70 eV) spectrum of compound LQOF-G35 is displayed below; Figure S2: EI-MS (70 eV) spectrum of compound LQOF-G35-Br is displayed below; Figure S3: HRESI-MS Spectrum of LQOF-G35-Br; Figure S4. HPLC-UV analysis of LQOF-G35; Figure. S5. HRESI-(+) MS analysis of LQOF-G35; Figure. S6. HRESI-(+) MS/MS analysis of LQOF-G35. Figure S7: ^1H NMR spectrum of the aromatic region for LQOF-G35 at 263K in CDCl_3 ; Figure S8: ^{13}C NMR spectrum for LQOF-G35 at 263K in CDCl_3 ; Figure S9: HMBC $^1\text{Hx}^{15}\text{N}$ NMR spectrum for LQOF-G35 at 263K in CDCl_3 ; Figure S10: HSQC $^1\text{Hx}^{13}\text{C}$ NMR spectrum for LQOF-G35 at 263K in CDCl_3 ; Figure S11: HMBC $^1\text{Hx}^{13}\text{C}$ NMR spectrum for LQOF-G35 at 263K in CDCl_3 ; Figure S12: NOESY $^1\text{Hx}^1\text{H}$ NMR spectrum for LQOF-G35 at 263K in CDCl_3 ; Figure S13: ^1H NMR spectrum for LQOF-G35-Br at 283K in CDCl_3 ; Figure S14: ^{13}C DEPTQ-135 NMR spectrum for LQOF-G35-Br at 283K in CDCl_3 ; Figure S15: HMBC $^1\text{Hx}^{13}\text{C}$ NMR spectrum for LQOF-G35-Br at 283K in CDCl_3 ; Figure S16: HSQC $^1\text{Hx}^{13}\text{C}$ NMR spectrum for LQOF-G35-Br at 283K in CDCl_3 ; Figure S17: NOESY $^1\text{Hx}^1\text{H}$ NMR spectrum for LQOF-G35-Br at 283K in CDCl_3 ; Table S1: Crystal data and refinement parameters of LQOF-G35 and LQOF-G35-Br.; Table S2: Selected bond lengths and angles for LQOF-G35; Table S3: Selected bond lengths and angles for LQOF-G35-Br.

Author Contributions: E.H.Z. —organic synthesis and, NMR and DRX conformational study; L.R.d.A. — organic synthesis and NMR conformational study; P.H.d.O.S. — X-ray measurements and process for conformational study; T.R.d.S.N. — organic synthesis; J.A.E — X-ray measurements, process for conformational study and supervision; E.R.P.G.—general supervision. All authors have read and agreed to the published version of the manuscript.

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

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