Contribution from the Department of Chemistry, Texas Christian University, Fort Worth, Texas 76129 USA and University of Belgrade, Belgrade, Yugoslavia

The Structure of (+)-dinitrobis-(L-arginato)cobalt(III) Nitrate Dihydrate*

W. H. Watson, D. R. Johnson, M. B. Celap, and B. Kamberi

Received February 8,1972

The structure and absolute configuration of (+)-dinitrobis-(L-arginato)cobalt(III) nitrate dihydrate was determined by single crystal X-ray diffraction techniques. The crystals belong to the triclinic space group P1 with unit cell dimensions of a = 11.250(5), b =9.867(5), c = 5.701(8) Å, $\alpha = 89^{\circ}(29)(5)'$, $\beta = 99^{\circ}26(5)'$ and $\gamma = 93^{\circ}13(15)'$. The observed density of 1.588 g. cm^{-3} is consistent with one formula unit per cell, $d_c = 1.594$ g. cm^{-3} . The anomalous dispersion of the cobalt(III) ion permitted the use of the Ps function to solve the structure directly. A total of 2044 independent reflections was collected by counter techniques and the model was refined to a conventional Rindex of 0.075 using the 1889 nonzero reflections. The crystal structure agrees with the R-configuration assigned by circular dichroism studies. The coordination around the cobalt(III) ion is distorted octahedral with the two nitro groups in a cis arrangement. The two amino acid ligands form five-membered chelate rings through use of the amino nitrogen and the carboxylate oxygen atoms; however, the conformations of the side chains differ. An extensive hydrogen bonding network exists in the solid state.

Introduction

Transition metal ions complexed with amino acids may serve as models for metal-protein interactions, and their structures may provide insight into the factors determining amino acid conformation. Generally, a-amino acids form five-membered chelate rings with the amino group and the carboxylate group acting as metal binding sites;¹ however, a number of amino acid side chain conformations have been observed. These variations may be attributed to extensive hydrogen bonding and packing interactions which exist in the solid state.

The structures of L-arginine dihydrate,² L-arginine monohydrobromide monohydrate,3 L-arginine monohydrochloride monohydrate.⁴ L-arginine monohydrochloride,⁴ and L-arginine phosphate monohydrate⁵ have been reported. The conformations of the side chains as well as bonding parameters for these

(1964).

molecules have been summarized in several reports.4-7 The bond lengths and angles are consistent among each of the reported structures.

The chemical and physical properties of (+)-dinitrobis-(L-arginato)cobalt(III) nitrate have been reported³ and solution circular dichroism studies were used to assign the configuration as R-. The crystal structure analysis of this complex was undertaken to confirm the absolute configuration, to determine the conformations of the extended carbon side chains and to determine if the guanidyl group also interacted with the metal ion. Interactions with the nitrogen of a folded side chain have been observed for a Cobalt(II) histidine complex,⁹ and a bis-glycylglycinate-cobalt(III) complex.10

Experimental Section

Reddish-brown crystals suitable for X-ray structural work were supplied by M. B. Celap and B. Kamberi. The following reaction describes the formation of the product:

Co(NO₃)₂+2 L-argHCl (25% KOH)+2 KNO₂---- $[Co(L-arg)_2(NO_2)_2]NO_3 \cdot 2H_2O + other products$

The cobalt complex was purified by recrystallization from water. Carbon, nitrogen, hydrogen, and metal analyses were consistent with the formula $[Co(C_5N_4 O_2H_{14})_2(NO_2)_2$ NO₃ · 2H₂O.

A crystal of dimensions 0.34 X 0.25 X 0.47 mm was used to collect all intensity data. The crystal was mounted with the long axis coincident with the spindle axis. The dimensions of the triclinic cell were determined at room temperature from c axis rotation and hk0 Weissenberg photographs calibrated with NaCl powder lines using zirconium-filtered MoKa radiation, ($\lambda = 0.71069$ Å).

Watson, fohnson, Celap, Kamberi | Structure of (+)-dinitrobis-(L-arginato)cobalt(III) Nitrate Dihydrate

^(*) Contribution Number 8 from the FASTBIOS Labratory.
(1) H. C. Freeman, « Advances in Protein Chemistry », Academic Press, New York, p. 257, 1967.
(2) J. L. Karle and J. Karle, Acta Cryst. 17, 835 (1964).
(3) S. K. Mazumdar and R. Srinivasan, Current Science, 33, 573

⁽⁴⁾ G. N. Ramachandran, S. K. Mazumdar, K. Venkatesan and
A. V. Lakshminarayanan, J. Mol. Biol., 15, 232 (1966).
(5) K. Aoki, K. Nagano, and Y. Iitaka, Acta Cryst., B27, 11 (1971).

<sup>(1971).
(6)</sup> A. V. Lakshminarayanan, V. Sasisekharan, and G. N. Rama-chandran, «Confirmation of Biopolymers», Academic Press, London, p. 61, 1967.
(7) R. E. Marsh and J. Donohue, Advances in Protein Chemistry, Academic Press, New York, p. 255, 1967.
(8) R. G. Denning, M. B. Celap and D. J. Radanovic, Inorg. Chim. Acta, 2, 58 (1968). M.B. Celap, B.A. Kamberi, and T.J. Janle, Proc. of 15th Annual Meeting of the Serbian Chemical Society, Novi Sad (Yuroslavia) 1920. Proc. of 15th Annuai Sad (Yugoslavia) 1920.

Cristal data

M = 597.4Co(C6N4O2H14)2(NO2)2NO3 . 2H2O a = 11.250(5) Å, b = 9.867(5) Å, c = 5.701(8) Å $\alpha = 89^{\circ}29(5)', \beta = 99^{\circ}26(5)', \gamma = 93^{\circ}13(5)'$

Space group: Pl (No systematic absences)

Z = 1; $V = 623.3 \text{ Å}^3$; $\mu = 60.4 \text{ cm}^{-1}(\text{CuK}\alpha)$; $d_o = 1.588$ g. cm⁻³; $d_c = 1.594$ g. cm⁻³

The errors in cell dimensions are standard deviations derived from the calculation of cell edges for a number of high angle reflections or layer lines. The errors in the angles are the average deviations obtained from several independent measurements of the same angle. The density was determined by the flotation method with a mixture of carbon tetrachloride and benzene.

Three-dimensional intensity data through the fourth level (hk4) were collected with a Phillips PAILRED diffractometer using equi-inclination geometry and the continuous ω -scan technique. The scan range varied from 3.4° for the zero level to 5.4° for the fourth level with a constant scan rate of 1°/min. Twenty second background counts were taken on either side of the peaks. CuKa radiation ($\lambda =$ 1.5418 Å) and a graphite monochromator crystal [d(002) = 3.354 Å] were used to collect all the data. CuKa radiation was used because of the large correction to the imaginary part of the anomalous dispersion for cobalt (Δ f'' = 3.95).¹¹ This was used to establish the absolute configuration. Several standard reflections were monitored throughout the data collection period. No statistically significant change in intensities was observed. The monitored reflections were used to scale the levels together.

In general, all of the nonequivalent reflections of the type hkl, hkl, hkl and $h\bar{k}l$ were collected up to a 20 limit of 135°. A total of 2044 nonequivalent intensities was measured, and 1889 were tabulated as being observed, *i.e.* $I > 2\sigma(I)$. Friedel pairs were collected over a more limited range.

Lorentz and polarization corrections were applied to the data.¹² An absorption correction was made using a program for Weissenberg geometry.¹³ The program is general and depends upon the equations of the crystal faces and not on the crystal class. The maximum and minimum transmission factors were calculated to be 0.344 and 0.166, respectively. Standard deviations were assigned on the basis of counting statics according to the equation

$$\sigma(F_{o})^{2} = \frac{A(N_{s} + RN_{B} + (0.031_{o})^{2})}{4L_{p}I_{o}}$$

$$I_o = (N_s - RN_B)$$

(13) J. A. Ibers, Northwestern University, Absorption Correction Program for Weissenberg Geometry, Modified for IBM 1800 by N. R. Stemple, Texas Christian University, 1969. (13) J.

where N_s is the total count obtained during a scan cycle, N_B is the total background count, R is a constant which relates the background count time to the total scan time, I_o is the number of counts associated with each reflection, and A is the absorption correction.

The scattering factors of Cromer and Waber¹⁴ were used for all nonhydrogen atoms. The scattering factor for cobalt(III) was corrected for the real and imaginary part of anomalous dispersion using the values of Cromer.¹¹ The hydrogen scattering factors were those calculated by Stewart, Davidson, and Simpson.15

Structure Determination

There were large differences in the magnitudes of the Friedel pairs and the P_s Patterson function of Pepinsky and Okaya¹⁶ was used to solve the structure. Since one anomalous scatter exists per cell, the P_s function yields the entire non-centrosymmetric distribution around the cobalt atom. All atoms in the two arginine ligands, the nitrate group and one of the NO₂ groups were located in the P₅ map. A structure factor calculation using 866 of the strongest reflections yielded a conventional R of 0.34 were R = $\Sigma ||F_0|$ - $|F_c|/\Sigma|F_o|$. Three cycles of block-diagonal least-squares using isotropic thermal parameters reduced the R value to 0.28. A three-dimensional Fourier map computed with the phases from the last least-squares calculation yielded all of the remaining nonhydrogen atoms. Reflections having $|F_o| > 2|F_c|$ were not used in the calculation.

All nonhydrogen atoms in the model were refined isotropically for 4 cycles using a block-diagonal leastsquares program. The cobalt atom position was fixed at (1/2, 1/2, 1/2). The agreement factors for 866 reflections were R = .099 and R(wtd) = .099 where $R(wtd) = \Sigma w ||F_o| - |F_c|| / \Sigma w |F_o|$. The function minimized in the refinement is $\Sigma w(kF_o - F_c)^2$ where $w = [1/\sigma(F_0)]^2$.

Anisotropic thermal parameters were assigned to cobalt, the six atoms coordinated to the cobalt ion and the four oxygen atoms of the two coordinated nitro groups. Limited computer memory and expenses prevented a complete anisotropic refinement. Eight cycles of block-diagonal refinement reduced R to .057 and R(wtd) to .055 for the 866 reflections. The inclusion of anisotropic thermal parameters in the model was found to be significant at the $\alpha = 0.01$ level based on Hamilton's significance test.¹⁷ Three cycles of full-matrix refinement were computed using 1299 reflections, and a three-dimensional difference Fourier map was calculated from the output. Most hydrogen atom positions could be assigned from The positions of the hydrogen atoms the map. associated with the water molecules are known with the least certainty. Three cycles of block-diagonal refinement yielded R = 0.0750 and R(wtd) =0.0752 for all 1889 nonzero reflections. The contri-

(17) W. C. Hamilton, Acta Cryst., 18, 502 (1956).

⁽⁹⁾ M. M. Harding and H. A. Long, J. Chem. Soc. A. 2554 (1968).
(10) R. D. Gillard, E. D. McKenzie, R. Mason and G. B. Robertson, Nature, 209, 1347 (1966).
(11) D. T. Cromer, Acta Cryst., 18, 17 (1965).
(12) A complete listing and description of all computer programs may be found in the Ph. D. dissertation of N. R. Stemple, Texas Christian University, 1970. The following programs written by N. R. Stemple were used in this investigation: Data Reduction, CDTRO; Fast Fourier (integer) FFOUR; Fourier, FRIER; Full-Matrix Least-Squares (modification of a program written by Y. Okaya), CLSQ; Interatomic Distances and Angles, CDIST; and Structure Drawing, CDRAW.
(13) I. A. Ibers. Northwestern University Absorption Correction

⁽¹⁴⁾ D. T. Cromer and J. T. Waber, Acta Cryst., 18, 104 (1965).
(15) R. F. Stewart, E. R. Davidson and W. T. Simpson, J. Chem. Phys., 42, 3175 (1965).
(16) Y. Okaya and R. Pepinsky, «Computing Methods and the Phase Problem in X-ray Analysis», cd. by R. Pepinsky, J. M. Ro-bertson and J. C. Speakman, Pergamon Press, New York, p. 271, 1962 1962

Table	Ι.	Squares	of the	observed	and	calculated	structure	factors fo	or (+)-dinitrobis-(L-arginato)co	balt(III)	nitrate	dihydrate

ノフエ

Atom	Х	Y	Z	B11 a	B ₂₂	B ₃₃	B ₁₂	B ₁₃	B ₂₃
Co	.5000	.5000	.5000	3.5	1.7	3.4	0.0	2.0	0.3
O(1)	.3983(4)	.4925(4)	.1969(5)	3.7	2.9	3.6	0.0	0.7	0.1
N(1)	.3730(5)	.5959(6)	.6059(14)	4.1	2.2	3.7	0.5	0.8	0.2
O(1')	.4191(3)	.3385(4)	.5963(5)	3.6	2.7	2.9	0.0	1.9	0.2
N(1')	.6104(5)	.3742(6)	.3910(10)	2.9	2.4	2.6	0.8	1.3	0.1
N(6)	.5868(7)	.6566(7)	.3422(10)	3.7	3.1	3.1	0.4	0.1	0.0
O(6)	.5307(5)	.7585(6)	.3442(11)	2.9	3.9	6.9	0.4	3.1	1.2
O(7)	.6945(4)	.6594(5)	.4211(9)	3.7	4.6	5.4	1.5	0.1	1.4
N(6′)	.5967(7)	.5141(10)	.8145(12)	4.3	5.7	1.8	-1.4	1.0	-1.1
O(6')	.6589(6)	.4232(8)	.8925(7)	7.2	6.7	3.5	2.5	0.1	0.9
O(7')	.5891(5)	.6163(8)	.9201(9)	6.3	7.7	3.5	-1.7	1.5	-2.2
C(1)	.2894(5)	.5169(7)	.2043(14)	2.6					
O(2)	.2090(4)	.4998(5)	.0206(7)	3.8					
C(2)	.2580(6)	.5667(7)	.4357(14)	2.9					
C(3)	.1770(7)	.6927(9)	.4022(20)	3.9					
C(4)	.2278(7)	.7973(10)	.2436(20)	4.4					
C(5)	.1690(6)	.9338(8)	.2478(16)	3.2					
N(2)	.2176(5)	.0189(6)	.0718(7)	3.5					
C(6)	.1947(6)	.1500(7)	.0388(17)	3.0					
N(3)	.2469(4)	.2213(6)	1148(10)	3.5					
N(4)	.1235(5)	.2093(7)	.1679(11)	3.8					
C(1')	.4502(5)	.2322(7)	.5182(17)	2.6					
O(2')	.4092(3)	.1184(5)	.5717(8)	4.0					
C(2')	.5398(6)	.2438(7)	.3380(15)	2.8					
C(3')	.6147(6)	.1207(8)	.3448(17)	3.3					
C(4')	.6868(7)	.1163(8)	.1379(17)	3.6					
C(5')	.7445(7)	0209(9)	.1458(17)	3.9					
N(2')	.8202(4)	0305(6)	0341(8)	2.9					
C(6')	.8651(6)	1442(8)	0850(14)	3.0					
N(3')	.9366(5)	—.1453(6)	2543(10)	3.7					
N(4′)	.8430(5)	2606(7)	.0290(15)	4.2					
N(5)	.9460(5)	.2253(7)	.5832(9)	4.0					
O(3)	.9841(3)	.1166(5)	.5292(10)	4.5					
O(4)	.8742(5)	.2267(7)	.7252(13)	6.6					
O(5)	.9789(6)	.3310(8)	.4963(13)	6.9					
O(8)	.9811(6)	.5645(7)	.7873(11)	6.2					
O(9)	.4064(4)	.8819(6)	.8417(9)	5.4					
Hydrogen	atoms parameters.	Temperature	factor of 3.0 assig	gned to all a	toms.				
H (1)	.394	.695	.608	H(1')	.680	.36		.521
H(2)	.362	.564	.767	H(2')	.637	.41	l	.243
H(3)	.208	.482	.499	H(3')	.484	.250)	.168
H(4)	.175	.737	.572	H(4')	.675	.118	3	.510
H(5)	.087	.660	.324	H(5′)	.557	.030)	.343
H(6)	.323	.812	.298	H(6')	.756	.196	5	.158
H(7)	.214	.760	064	H(7')	.627	.126	5	027
H(8)	.194	.979	.420	H(8')	.797	035	5	.318
H(9)	.073	.919	.197	H(9')	.675	101		.121
H(10)	.273	.976	033	H(10')	.840	.056	5	127
H(11)	.238	.320	129	H(11')	.953	061	l	342
H(12)	.296	.175	—.217	H(12')	.973	230)	290
H(13)	.114	.308	.155	H(13')	.790	262	2	.151
H(14)	.080	.154	.276	H(14')	.880	345	5	009
									0.0
H(15)	.407	.959	.754	H(17)	.055	.544	-	.863

Thermal parameters are of the form exp $-0.25(B_{11}h^2a^{*2}+B_{22}k^2b^{*2}+B_{33}l^2c^{*2}+2B_{12}hka^*b^*+2B_{13}hla^*c^*+2B_{13}hla^*c^*)$

^a The average standard deviation for the thermal parameters is approximately 0.3.

butions of the hydrogen atoms were calculated, but the parameters were not refined. Generally, those reflections assigned a zero magnitude have a calculated magnitude less than the observed minimum. The final observed and calculated structure factors for all 2044 reflections are included in Table I. sional difference Fourier map was calculated and no peak was larger than those associated with hydrogen atoms. Structure factor calculations for the Friedel pairs indicated the correct configuration has been assigned.

All shifts of the parameters for nonhydrogen atoms during the final cycle of refinement were less than one-seventh of the estimated standard deviations of the parameters. The final positional and thermal parameters along with their estimated standard deviations are given in Table II. A final three-dimen-

Discussion

Figure 1 shows a projection of the contents of the unit cell onto the *ab* plane. Both arginine ligands are bidentate and coordinate to the cobalt through the

 Table III. Interatomic distances and bond angles for (+)dinitrobis-(L-arginato)cobalt(III)nitrate dihydrate.

Co-O(1)	1 909(3) &	Co-O(1')	1 912(4)
Co-N(1)	1 935(6)	$C_0 N(1')$	1 977(6)
O(1)-C(1)	1 270(8)	O(1)- $C(1')$	1 234(8)
C(1)-O(2)	1.272(9)	C(1') - O(2')	1 244(8)
C(1)-C(2)	1.513(10)	C(1') - C(2')	1.244(0) 1.552(11)
C(2)-N(1)	1.501(10)	$C(1') \cdot C(2')$	1 480(9)
C(2)-C(3)	1.574(11)	C(2')-C(3')	1.100(3)
C(3)-C(4)	1.519(13)	C(3')-C(4')	1.539(13)
C(4)-C(5)	1.535(11)	C(4')-C5')	1.531(12)
C(5)-N(2)	1.458(9)	C(5')-N(2')	1.444(10)
N(2)-C(6)	1.339(9)	N(2')-C(6')	1.310(9)
C(6)-N(3)	1.311(10)	C(6')-N(3')	1.354(9)
C(6)-N(4)	1.333(9)	C(6')-N(4')	1.355(10)
Co-N(6)	1.894(7)	Co-N(6')	1.941(7)
N(6)-O(6)	1.240(9)	N(6')-O(6')	1.205(11)
N(6)-O(7)	1.201(9)	N(6')-O(7')	1.194(12)
N(5)-O(3)	1.236(8)	N(5)-O(4)	1.235(9)
N(5)-O(6)	1.216(10)		. ,
O(1)CoN(1)	85.8(2)°	N(1)CoN(6)	96.0(3)
O(1)CoN(1')	91.1(2)	O(1')CoN(1')	84.7(2)
O(1)CoO(1')	90.4(2)	O(1')CoN(6')	89.7(3)
O(1)CoN(6)	91.4(2)	N(1)'CoN(6')	92.2(3)
N(1)CoO(1')	86.0(2)	N(1')CoN(6)	93.5(3)
N(1)CoN(6')	90.9(3)	N(6')CoN(6)	88.6(3)
$C_0O(1)C(1)$	113.8(4)	CoO(1')C(1')	114.8(4)
O(1)C(1)O(2)	120.4(6)	O(1')C(1')O(2')	122.5(6)
O(1)C(1)C(2)	118.5(6)	O(1')C(1')C(2')	117.7(6)
C(1)C(2)N(1)	108.4(5)	C(1')C(2')N(1')	106.0(6)
CoN(1)C(2)	108.6(5)	CoN(1')C(2')	106.4(4)
N(1)C(2)C(3)	112.0(6)	N(1')C(2')C(3')	114.8(6)
C(1)C(2)C(3)	113.0(7)	C(1')C(2')C(3')	112.0(7)
C(2)C(3)C(4)	110.0(6)	C(2')C(3')C(4')	113.2(7)
C(3)C(4)C(5)	111.9(7)	C(3')C(4')C(5')	108.0(7)
C(4)C(5)N(2)	105.7(6)	C(4')C(5')N(2')	111.7(7)
V(5)N(2)C(6)	123.2(6)	C(5')N(2')C(6')	122.8(7)
N(2)C(0)N(3)	119.3(6)	N(2')C(6')N(3')	119.6(6)
N(2)C(0)N(4)	120.5(7)	N(2')C(6')N(4')	121.6(6)
N(3)C(0)N(4)	120.1(6)	N(3')C(6')N(4')	118.8(7)
CoN(6)O(6)	118.7(5)	CoN(6')O(6')	120.4(6)
O(6)N(6)O(7)	121.2(5)	O(6)N(6)O(7)	115.6(6)
O(3)N(5)O(4)	120.1(7)	O(0) IN(0) O(7)	124.0(7)
O(3)N(3)O(4)	120.1(7)	O(4) IN(5) O(5)	119.8(7)
O(3)N(3)O(5)	120.2(6)		

of rotation of the torsion angles is consistent with the rules proposed by Edsall, *et al.*¹⁸ and Klyne and Prelog.¹⁹ Table V gives the least-squares equations for planes passed through significant structural groups and the distance of each atom from the plane.



Figure 1. Projection of the unit cell contents onto the ab plane for (+)-dinitrobis-(L-arginato)cobalt(III) nitrate dihydrate.

The coordination geometry about the cobalt is that of a distorted octahedron with the two nitro groups in a *cis* configuration. Celap, *et al.*¹⁸ have shown that the nitro groups are in the *cis* configuration for a series of dinitrobis-(amino acidato)cobalt(III) ions. The distortions of the angles about the cobalt ion are associated primarily with the restrictions im-

Table IV. Summary of the torsion angles in Cobalt(L-arginine) and other reported arginine compounds.

Internal Rotation Angles	Unprimed Ligand	Primed Ligand	Arg.H ₁ PO ₁ . H ₂ O	Arg.2H ₂ O	Arg.I Mol I	HC1 Mol 2	Arg.HC Mol 1	1.H2O Mol 2	Arg.H Mol 1	Br.H₂O Mol2
(1) = O(1)C(1)C(2)N(1) O(1)C(1)C(2)C(3)	9	27	34 84		51 71	41 79	6 116	26 97	6 115	
(2) = O(2)C(1)C(2)N(1) O(2)C(1)C(2)C(3)			148 94	168 70	135 104	138 	178 —60	157 80	177 —61	156 80
	77 47	70 170	164 78	62 —60	171 49	168 49	63 178	54 175	61 178	
(2) = C(2)C(3)C(4)C(5) (3) = C(3)C(4)C(5)N(2) (4) = C(4)C(5)N(2)C(6)			175 64 113	151		175				
(5) = C(5)N(2)C(6)N(3) (6) = C(5)N(2)C(6)N(4)	176	-179 0		172	175	-179 6		165 —19		165 —18

amino nitrogens N(1) and N(1') and the carboxylate oxygens O(1) and O(1'). The guanidyl groups of the amino acids do not coordinate to the cobalt. Bond distances and bond angles are given in Table III along with several intermolecular distances which influence the packing in the unit cell. Torsion angles for the arginine ligands in this structure and in other reported structures are given in Table IV. The sense posed by the five-membered chelate rings, O(1)-Co-N-(1)=85.8° and O(1')-Co-N(1')=84.7°. These angles are consistent with those observed for other amino acid complexes. The average of seventeen O-Cu^{II}-N

(18) J. T. Edsall, P. T. Flory, J. C. Kendrew, A. M. Liquori,
G. Nemethy, G. N. Ramachandran, and H. A. Scheraga, J. Molec. *ibiol.*, 15, 399 (1966).
(19) W. Klyne and V. Prelog, *Experientia*, 16, 521 (1960).

angles in five-membered chelate rings involving amino acids is 83.5°.1 The absolute configuration about the cobalt ion is R^{-20} which is equivalent to the $P(C_2)$ notation pro-

Table V. Least-squares equations for groups of atoms and the deviations of the atoms from each plane. The equation of each plane is defined by Ax+By+Cz = D where y is parallel to b, x is parallel to the projection of a on a plane perpendicular to b and z is perpendicular to x and y^b

(1) Chelate Rin	g Plane Unprimed Ligand	Primed Ligand	(2) Carboxyl	ate Group Plane Unprimed Ligand	Primed Ligand
0(1)	-0.009	0.087	Q(1)	0.000	0.005
O(1)	0.049	0.130	O(2)	0.000	0.005
C(1)	-0.013	0.030	C(1)	0.000	0.013
C(1)	-0.072		C(2)	0.000	0.004
N(1)	0.063	0.203	$N(1)^{a}$	0.202	0.609
	-0.376	-0.256	Co ^a	0.251	0.112
A	0.209	0.545	Α	0.276	0.645
B	0.923	-0.175	В	0.934	0.075
Č	-0.324	0.820	С	0.276	0.761
D	4.831	4.557	D	4.980	4.999
(3) Guanidyl G	roup Plane		(4) Carbon	Side Chain Plane	
(0)	Unprimed	Primed		Unprimed	Primed
	Ligand	Ligand		Ligand	Ligand
N(2)	0.005	0.001	C(2)	0.069	0.045
C(6)	0.014	0.004	C(3)	0.091	0.055
N(3)	0.005	0.001	C(4)	0.059	0.033
N(4)	0.005	0.001	C(5)	0.094	0.051
A	0.678	0.702	N(2) ^a	0.017	0.007
В	0.204	0.168	Α	0.529	0.701
С	0.706	0.692	В	0.348	0.403
D	3.193	6.219	С	0.774	0.588
			D	5 044	6 0 3 4

(5) Nitrate Group Plane

N(5)	0.001
O(3)	0.000
O(4)	0.000
O(5)	0.000
Α	0.650
B	0.075
С	0.756
D	9.155

^a These atoms were not included in the least-squares plane calculation. ^b The transformation from the triclinic cell (X,Y,Z) to the orthogonal cell is given by (x) = (11232 - 0) = (207) (X)

(X)		[11.232	0	907 \	(X)
ĺv	=	631	9.867	.051	$[\mathbf{Y}]$
\ z /		0	0	5.628 /	$\langle z /$

Table	VI.	Intermolecular	distances	involving	hydrogen	atoms.
-------	-----	----------------	-----------	-----------	----------	--------

Donor (D)	Acceptor (A)	Distance (D)(A)	Distance H(A)	Angle (D)-H (A)
N(1)	O(W)	3.112(Å)	2.27(Å)	142°
O'(W)	O(5)	2.848	1.88	161°
O(W)	O'(2)	2.785	1.88	176°
O(W)	Q(6)	3.234	2.14	149°
N(2)	O(W)	3.054	2.04	165°
N'(3)	O'(W)	2.935	2.07	145°
N'(4)	O'(W)	2.887	2.00	148°
O'(W)	O(2)	2.794	2.10	133°
N(3)	O(2)	2.934	2.04	149°
N(4)	O(2)	3.124	2.31	139°
N'(4)	$\vec{O}(\vec{7})$	3.074	2.13	157°
N(4)	O(3)	2.899	1.96	15 7 °
N(4)	O(5)	2.976	2.67	98°
N'(3)	O(3)	2.914	1.93	169°
N'(2)	O(4)	2.954	1.92	153°
N(3)	O '(1)	2.926	2.65	117°
N(3)	O'(2)	2.978	2.01	165°

The cobalt(III)-oxygen bond lengths of 1.909 and 1.911 Å may be compared with the value of 1.91 Å found in (-)-sarcosinatobis(ethylene-diamine)cobalt-(III) diiodide dihydrate²² and 1.85 Å reported for a bis-glycylglycinate-cobalt(III) complex.¹⁰ The Co^{III}-nitrogen bond lengths of 1.935, 1.977, 1.894 and 1.941 Å may be compared with the values of 1.97 Å and 1.92 Å reported for the above two complexes.

Least-squares planes fitted to the five-membered chelate rings of the arginine ligands indicate the rings are not planar. Planes fitted to the carboxylate groups and the adjqacent carbon atoms show N(1) to be 0.20 Å out of the plane while N(1') is 0.61 Å from the plane. The torsion angle about C(1)-C(2) is $+9^{\circ}$ while that about C(1')-C(2') is $+27^{\circ}$. These torsion angles are consistent with the magnitudes observed for several α -amino acids, but they are of opposite signs.

Least-squares planes fitted to the guanidyl groups indicate they are planar within experimental error and normals to the carboxylate planes and the guanidyl planes of the two ligands make angles of 82° and 15°, respectively. The conformations of the two arginine ligands are indicated by the torsion angles given in Table IV. One of the major differences between the two ligands is the torsion angle N(1)-C(2)-C(3)-C(4). In the unprimed ligand this angle is 77° leaving C(4) staggered betweeen the NH₂ and COO groups. This is not the sterically favored orientation, but has been observed in the L-arginine dihydrate

(20) R. S. Cahn, C. Ingold and V. Prelog, Angew. Chem. Internat. Edit., 5, 385 1966).
(21) W. J. Mc Caffery, S.F. Mason and B. J. Norman, J. Chem. Soc., 5094 (1965).
(22) J. F. Blount, H. C. Freeman, A. M. Sargeson and K. R. Turnbull, Chem. Comm. 324 (1967).

structure.² The torsion angle for the primed ligand is -70° which leaves C(4') staggered between the NH₂ group and a hydrogen on C(2'). Packing interactions appear to be responsible for the orientation of the side chains.

The bond lengths and angles for each of the arginine ligands are consistent with those reported for other arginine structures.⁷ The average carbon-carbon distance is 1.54 Å and the average C-N distance is 1.33 Å. The C-N distances indicate some delocalization in the protonated guanidyl groups.

The N-O distances in the nitro groups average 1.22 Å while the N-O distances in the nitrate group average 1.23 Å. The bond angles are consistent with sp² hybridization. The cobalt(III) ion lies in the plane of each nitro group. The nitrate group is planar.

Table VI lists distances between potential hydrogen bonded atoms. Contact distances between nitrogen and oxygen atoms of 2.75 to 3.15 Å have been associated with hydrogen bonding in many cases.²³ The oxygen atoms of the water molecules act as both acceptors and donors in the formation of hydrogen bonds while the guanidyl groups probably are involved as hydrogen bond donors. The carboxylate oxygen atoms of both ligands are involved as hydrogen bond acceptors as probably are oxygen's O(7) and O(5) of the nitro and nitrate groups. Each group in the structure appears to assume an orientation in the unit cell which permits the most extensive hydrogen bonding network.

Acknowledgment. We would like to acknowledge the Robert A. Welch Foundation for their financial support and the TCU Research Foundation for the purchase of equipment.

(23) G. C. Pimentel and A. L. McCellan, «The Hydrogen Bond», Freeman, San Francisco and London, 1960.