03.1-5 STRUCTURE-ACTIVITY RELATIONSHIPS OF BIPYRI-DINE INOTROPIC CARDIAC AGENTS. <u>Vivian Cody</u> and Joe Luft, Medical Foundation of Buffalo, Inc., 73 High St., Buf-falo, NY 14203; Faith B. Davis and Paul J. Davis, VA Med-ical Center, Buffalo, NY, 14215, USA.

Hilrinone [2-methyl-5-amino-(3,4'-bipyridin)-6(1H)-one] (A) and amrinone (B) are members of a new class of oral nonglycosidic, noncatecholamine cardiac positiveinotropic agents developed for the treatment of conges-tive heart failure. These bipyridine inotropic agents strengthen myocardial contraction by increasing the availability of intracellular calcium. We have reported that militinone, but not amrinone, shares structural homo-logy with thyroid hormone and like thyroid hormone, stim-ulates rabbit myocardial membrane Ca^{2+} -ATPase activity as shown by Ca^{2+} -dependent ATP hydrolysis in vitro (Mylotte et al, Proc. Natl. Acad. Sci. USA, 82, 7974 (85)). Com-parison of these molecular structures showed that there are structural homologies between the phenolic ring of thyroxine and the substituted ring of milrinone, not shared by amrinone. To further delineate the structural features required for this activity, the crystal struc-tures of a number of milrinone analogues were investi-gated and are commerced with the reserved structures f The status is activity, the crystal status tures of a number of milrinone analogues were investi-gated and are compared with the reported structures of amrinone and milrinone (Cody, Acta Cryst., in press; Robertson et al, J. Med. Chem. 29, 635 (86)). The fol-lowing inotropic agents (C-F) have been studied to deter-mine the conformational aspects of drug specificity. (C) 2-methyl-(3,4'-bipyridin)-6(1H)-one (C_{1,H_10,Y_0}); P2/(c, z = 4, a = 7.574 (1), b = 11.132(2), c = 11.437(2)Å, β = 94.90(1)°), (D) 2-methyl-5-bromo-(3,4'-bipyridin)-6(1H)-one (C, 1B₀N₀Br; P-1, z = 2, a = 6.813(2), b = 7.269(2), c = 12.787(4)Å, α = 97.57(3), β = 92.61(2), γ = 113.74-(2)°), (E) 2,3-dimethyl-5-cyano-(3,4'-bipyridin)-6(1H)-one HBr (C₁₃H₁₁N₃0 HBr; Pbca, z = 8, a = 7.2993(8), b = 18.955(3), c = 19.814(3)Å), (F) 2,3-dimethyl-5-cyano-(3,4'-bipyridin)-6(1H)-one (C, 1H₁N₃0; P2₁/c, z = 4, a = 11.713(4), b = 7.892(3), c = 12.154(4)Å, β = 92.27(4)°). The biochemical data for these compounds show that only the bromo analogue (D) and a 2-CH₃-5-NH₂ analogue are active in this myocardial system. Structural analysis of these data show that all of the bipyridine rings are active in this myocardial system. Structure characteristic these data show that all of the bipyridine rings are twisted as observed in the structure of milrinone. The two determinations of the N-methyl analogue (E,F) show two determinations of the N-methyl analogue (L,r) show the largest variation in the twist angle. There is an N- $H_{\dots,0}$ hydrogen bond in C and D. These data show that the presence of the 2-methyl substituent forces the two pyri-dine rings to adopt a twist conformation which is favor-able for Ca²⁺-ATPase enzyme activity. The nature of the 5-substituent is less critical although these results suggest that a functional group that can mimic iodine in either size or electrochemical properties is preferred. Thus, prerequisites for enzyme stimulation include a 2-methyl for a twist conformation, a free 3-N position, and a 5-substituent.



A* Milrinone, Cody, Acta Cryst.; Milrinone HCl, Robertson et.al., J. Med. Chem. B* Amrinone H₂0, Robertson et. al.; Amrinone, 4 independent molecules, Cody. 03.1-6 THE SPATIAL ARRANGEMENT OF IMPORTANT MOLE-CULAR FEATURES IN CLASS I ANTIARRHYTHMIC AGENTS. By Marek L. Glowka and Penelope W. Codding, Departments of Chemistry and of Pharmacology and Therapeutics, University of Calgary, Calgary, Alberta, Canada.

Class I antiarrhythmic agents control irregular heart beat in cardiac disorders by binding to sodium channels and slowing conduction velocity. Class I agents are and slowing conduction velocity. Class I agents are characterized by three structural units: (a) an aro-matic ring that can intercalate between the alkyl matic ring that can intercalate between the alkyl chains of phospholipids, (b) an amino group that can ionize under biological conditions, and (c) a linkage region between (a) and (b) that contains a hydrogen bonding group. In this work, the optimal three dimen-sional arrangement of these crucial features has been sought through X-ray crystallography, molecular mechan-ics calculations and structural correlations. Two series of synthetic antiarrhythmic agents, one based on the 2,6-bis-(1-pyrrolidinylmethyl)phenol moiety (A) and one based on 2,2,5,5-tetramethyl-3-pyrolino-3-carboxyamide (B) are being characterized. These struc-tures have been compared to those of the traditional local anesthetics like disopyramide (C), quinidine, procainamide, lidocaine and mexiletine to identify the stereochemical requirements for arrhythmia control. These comparisons have been extended by calculating the These comparisons have been extended by calculating the potential energy surface for both anesthetics (mexiletine) and synthetic compounds (A). Comparisons of the low energy conformers of the active agents thus described indicate that, while flexibility is an impor-tant feature of these drugs, there are specific ranges for the distances between the three crucial groups mentioned above. Class I antiarrhythmic drugs have been divided into three subclasses according to the rate of recovery from channel block. We find that the separations between the three molecular features are different for each of these subclasses. Thus, Class I antiarrhythmic activity may be dependent on an induced fit of these flexible agents to a receptor. The simi-larity in the general three dimensional shape and functionality of these agents suggests that a common receptor mediates the antiarrhythmic action; these findings will be used to propose a model for this receptor site. Crystal data: A: N-[4-hydroxy-3,5-bis(1-pyrrolidiny]methy1)pheny1]-2-methy1-benzamide, C23H31N30.HC1, orthorhombic, Aba2, a=13.882(2), b=29.115(3), c=11.092(1)A, T=223K, Z=8, R=0.042, R_{w}=0.043. B: N-(ω -N-phthalimido-propyl)-2,2,5,5-tetramethyl-2-pyrrolino-3-carboxyamide, C₂₀H₂₅N₃0₃·HCI·1/2CH₃0H, triclinic, PI, a=8.995(1),

