

Cardiac Dysfunction Occurs in the HIV-1 Transgenic Mouse Treated with Zidovudine

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SUMMARY: Cardiomyopathy in AIDS is an increasingly important clinical problem. Mechanisms of AIDS cardiomyopathy were explored using AIDS transgenic mice that express replication-incompetent HIV-1 (NL4-3Δ *gag/pol*). Transgenic and FVB/n mice ($n = 3$ to 6 per cohort) received water ad libitum with and without zidovudine (3'-azido-2',3'-deoxythymidine; AZT; 0.7 mg/ml) for 21 or 35 days. After 21 days, echocardiographic studies were performed and abundance of mRNA for cardiac sarcoplasmic reticulum calcium ATPase (SERCA2), sodium calcium exchanger (NCX1), and atrial natriuretic factor were determined individually using Northern analysis of extracts of left ventricles. After 35 days, contractile function and relaxation were analyzed in isolated work-performing hearts. Histopathological and ultrastructural (transmission electron microscopy) changes were identified. After 21 days, molecular indicators of cardiac dysfunction were found. Depressed SERCA2 and increased atrial natriuretic factor mRNA abundance occurred in left ventricles from AZT-treated transgenic mice. NCX1 abundance was unchanged. Eccentric left ventricle hypertrophy was determined echocardiographically. After 35 days, cardiac dysfunction was worst in AZT-treated and AZT-untreated transgenic mice. Decreases in the first derivative of the maximal change in left ventricle systolic pressure with respect to time (+dP/dt) occurred in transgenic mice with and without AZT. Increased half-time of relaxation and ventricular relaxation (-dP/dt) occurred in AZT-treated and -untreated transgenic mice. Increased time to peak pressure was found only in AZT-treated transgenic mice. In AZT-treated FVB/n mice, -dP/dt was decreased. Ultrastructurally, mitochondrial destruction was most pronounced in AZT-treated transgenic mice, but also was found in AZT-treated FVB/n mice. Transgenic mice that express HIV-1 demonstrate cardiac dysfunction. AZT treatment of FVB/n mice causes mitochondrial ultrastructural alterations that are similar to those in other species. In transgenic mice, AZT treatment worsens molecular and ultrastructural features of cardiomyopathy. HIV-1 constructs and AZT each contribute to cardiac dysfunction in this murine model of AIDS cardiomyopathy. (*Lab Invest* 2000, 80:187-197).

AIDS cardiomyopathy (CM) was described more than a decade ago (Cohen et al, 1986), and is gaining clinical importance (Barbaro et al, 1998). In patients with advanced AIDS, 8% had CM (De Palo et al, 1995). A similar proportion was found in hemophiliacs with AIDS (Jacob et al, 1993). With the advent of effective treatment of AIDS complications and increasing survival with AIDS, it seems logical that CM will become a prevalent clinical problem (Currie and Boon, 1983; Hsia et al, 1993).

To date, clinical studies fail to elucidate the etiology or pathogenesis of AIDS CM and thus mechanisms of AIDS CM remain a mystery. Myocardial filling and relaxation abnormalities were found in AIDS patients' echocardiographs (Barbaro et al, 1996; Cardoso et al, 1998; Coudray et al, 1995) and became early indicators of CM.

Many etiologies and pathogeneses are postulated for AIDS CM (Lewis, 1998; Michael et al, 1997) and it is possible that more than one etiology may contribute to CM development in the same AIDS patient. Potential etiologies include direct HIV-1 infection of the heart (Barbaro et al, 1998; Calabrese et al, 1987; Grody et al, 1990; Lipshultz et al, 1990; Wu et al, 1990), myocarditis caused by HIV-1 or by concomitant (or opportunistic) myocardial infection (Anderson et al, 1988; Barbaro et al, 1998; Reilly et al, 1988), cardiotoxicity of prolonged administration of antiviral therapeutics including zidovudine (AZT; 3'-azido-2',3'-deoxythymidine; Dalakas et al, 1991; Herskowitz et al, 1992; Lewis et al, 1991) or other agents, cardiodepressant effects from localized or systemic cytokine expression, cardiotoxicity of illicit or self-prescribed drugs (particularly alcohol, cocaine, or heroin) or pharmaceuticals, and toxicity of home remedies (reviewed in Lewis, 1998; Lewis and Dalakas, 1995).

The hemizygous NL4-3Δ *gag/pol* transgenic mouse (TG) contains an internal deletion of the *gag/pol* coding sequence that renders the viral construct replication-incompetent, non-infectious, relatively easy to handle, and useful to study AIDS CM (Bruggeman et al, 1997; Klotman and Notkins, 1996; Kopp et al, 1992).

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Phenotypic changes of AIDS in the TG attest to the authenticity of the model and include AIDS nephropathy (Bruggeman et al, 1997), failure to thrive and wasting (of pups; Kopp et al, 1992), and dermatological lesions (Kopp et al, 1993). Because the TG genetic background is FVB/n, FVB/n wild type mice served as controls (WT).

Using a mouse model system, experiments here dissect the role of a transgenically inserted, replication incompetent HIV-1 cDNA construct on the development of CM in AIDS. Treatment of these mice with AZT, the principal antiretroviral agent used in AIDS, was initiated to mimic therapeutic intervention and to define cardiotoxicity.

Results

General. All groups of mice survived for the duration of the 21-day and 35-day experiments. Heart weights (mg; mean ± standard deviation sd) were determined after 35 days. The heart rates ranged from 161 ± 30 for untreated WT to 221 ± 36 in AZT-treated TG (37% increase above WT; *p* < 0.01; Table 1).

AZT Consumption Data. TG and WT each consumed AZT in drinking water. Consumption of AZT was 182 to 215 mg/kg/day.

Physiological Changes in the Isolated Work-Performing Hearts of AIDS Transgenic Mice at 35 Days. Contractile dysfunction was observed in the TG. Figure 1 is a composite of histograms depicting cardiac performance in all of the groups used in these experiments. Parameters evaluated included the first derivatives of maximal changes, with respect to time, in LV systolic pressure (+dP/dt) and LV relaxation (-dP/dt), time to peak pressure (TPP/mmHg), and half time of relaxation (RT_{1/2}/mmHg). All groups were initially compared at identical preloads and afterloads and similar heart rates (approximately 320 beats/minute).

+dP/dt showed a small but significant decrease (*p* < 0.01) in untreated TG, compared with the untreated WT. Ventricular relaxation (-dP/dt) revealed a significant decrease in rate in AZT-treated WT (*p* < 0.01) and AZT-treated TG, but untreated TG also demonstrated defective relaxation (*p* < 0.001 respectively). RT_{1/2}/mmHg increased significantly in AZT-treated and -untreated TG (*p* < 0.05) and a small increase in TPP/mmHg was found in AZT-treated TG. Tables 2 through 5 present the numerical summaries of all measured and derived physiological measure-

Table 1. Heart Weight and Body Weight of TG and WT

Group	Heart weight (mg)	Body weight (g)	Heart/Body weight (mg/g)
WT (n = 7)	161 ± 30	24.7	6.5
WT + AZT (n = 4)	201 ± 46	26.6	7.6
TG (n = 5)	198 ± 31	23.2	8.5
TG + AZT (n = 5)	221 ± 36*	24.3	9.1*

* *p* < 0.01 compared with WT.
TG, transgenic mouse; WT, wild type mouse.

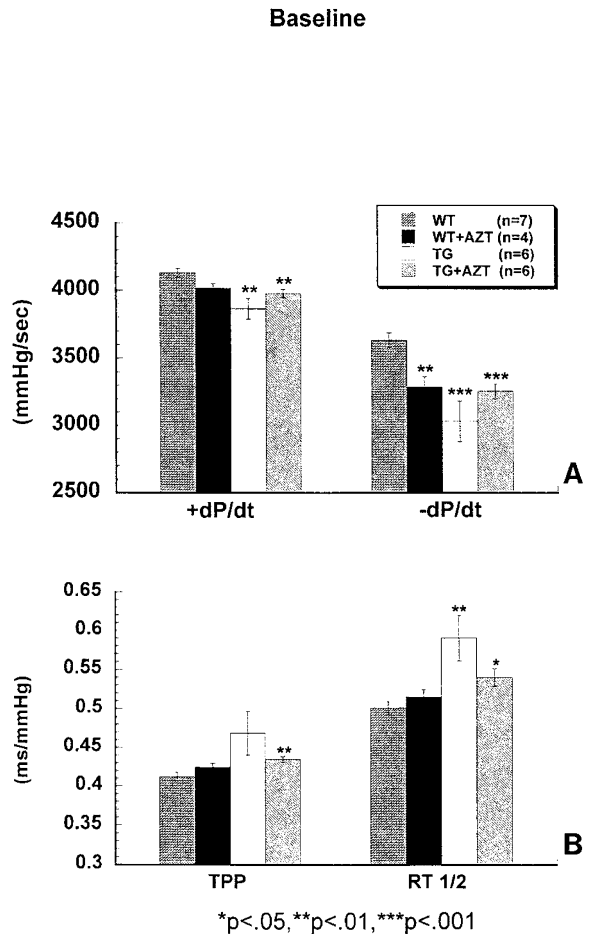


Figure 1.

Myocardial contractile parameters (+dP/dt in mmHg/seconds and TPP/mmHg in ms/mmHg of intraventricular pressure) and relaxation parameters (-dP/dt in mmHg/second and RT_{1/2}/mmHg in ms/mmHg of intraventricular pressure) of NL4-3Δ gag/pol wildtype mice (WT), WT treated with zidovudine (3'-azido-2',3'-deoxythymidine; AZT), transgenic mice (TG), and TG treated with AZT, tested in work-performing heart preparations under identical loading conditions. Basal loading was 5 ml/minute cardiac output and 50 mmHg mean aortic pressure. Heart rates were approximately 315 beats/minute and work load was 250 (mmHg × ml/minute). A, A decrease in the rate of pressure development (+dP/dt) is present in both AZT-treated and untreated TG. Changes in the rate of relaxation (-dP/dt) are decreased in AZT-treated TG and WT and in untreated TG. B, TPP/mmHg is increased in AZT-treated TG. RT_{1/2}/mmHg increased dramatically in TG with and without AZT treatment.

ments in the histograms. Physiological data above and pathological data below indicate that both AZT and HIV-1 each play a role in the development of CM, although possibly through different pathophysiological mechanisms.

It was of great importance to evaluate the performance of all hearts in all groups during decreases and increases of afterload and preload. Changes in afterload were achieved by decreasing and increasing aortic pressure. Changes in preload were accomplished by decreasing and increasing venous return. Stepwise decreases of pressure load were tolerated in all groups with respect to contraction (+dP/dt and TPP/mmHg of intraventricular pressure [IVP]). However, the relaxation parameters (-dP/dt and RT_{1/2}/mmHg of IVP) showed significant impairment in both

Table 2. Minimum Pressure Loading

Parameter	WT (n = 7)	WT + AZT (n = 4)	TG (n = 5)	TG + AZT (n = 5)
Heart rate (Bpm)	307.7 ± 19.1	328.0 ± 20.7	311.0 ± 20.3	300.0 ± 23.5
MAP (mmHg)	30.4 ± 0.3	30.3 ± 0.3	32.3 ± 1.7	32.2 ± 2.0
EDP (mmHg)	4.86 ± 0.8	5.3 ± 0.9	10.5 ± 4.2	7.0 ± 2.2
+dP/dt (mmHg/sec)	3404.6 ± 106.8	3136.0 ± 92.3	3188.0 ± 164.7	3096.0 ± 107.9
-dP/dt (mmHg/sec)	-2965.7 ± 115.1	-2472.0 ± 216.4	-2380.0 ± 178.0*	-2260.8 ± 119.4**
TPP/mmHg (msec/mmHg)	0.505 ± 0.02	0.533 ± 0.02	0.546 ± 0.04	0.542 ± 0.02
RT _{1/2} (msec/mmHg)	0.554 ± 0.02	0.617 ± 0.04	0.652 ± 0.04*	0.702 ± 0.06*
CO (ml/min)	5.17 ± 0.04	5.2 ± 0.06	5.1 ± 0.04	5.12 ± 0.04
Work	156.3 ± 1.3	156.7 ± 3.3	163.8 ± 9.1	163.8 ± 10.0
Coronary flow (ml/min)	1.27 ± 0.3	1.2 ± 0.2	1.2 ± 0.2	1.93 ± 0.5
Aortic flow (ml/min)	3.89 ± 0.3	4.0 ± 0.1	3.88 ± 0.2	3.18 ± 0.5

* p < 0.05, ** p < 0.01, *** p < 0.001

Table 3. Maximum Pressure Loading

Parameter	WT (n = 7)	WT + AZT (n = 4)	TG (n = 5)	TG + AZT (n = 5)
Heart rate (Bpm)	325.0 ± 8.8	356.8 ± 14.9	377.8 ± 13.9	346.3 ± 15.7
MAP (mmHg)	86.7 ± 1.8	85.3 ± 3.0	84.3 ± 3.5	81.17 ± 4.5
EDP (mmHg)	8.9 ± 1.6	8.5 ± 3.3	13.3 ± 3.1	7.8 ± 1.5
+dP/dt (mmHg/sec)	5640.0 ± 141.6	5520.0 ± 319.8	5632.0 ± 222.0	5556.0 ± 312.1
-dP/dt (mmHg/sec)	-4738.3 ± 214.0	-4296.0 ± 402.4	-4576.0 ± 284.8	-4464.0 ± 121.8
TPP/mmHg (msec/mmHg)	0.3064 ± 0.01	0.3185 ± 0.02	0.3141 ± 0.01	0.3158 ± 0.02
RT _{1/2} (msec/mmHg)	0.4513 ± 0.02	0.4495 ± 0.01	0.4376 ± 0.02	0.445 ± 0.03
CO (ml/min)	5.057 ± 0.04	5.2 ± 0.00	5.2 ± 0.03	5.13 ± 0.05
Work	436.3 ± 9.9	438.8 ± 14.5	432.2 ± 19.2	414.7 ± 25.0
Coronary flow (ml/min)	3.1 ± 0.4	3.9 ± 0.5	3.5 ± 0.3	3.8 ± 0.4
Aortic flow (ml/min)	1.9 ± 0.4	1.3 ± 0.5	1.6 ± 0.4	1.38 ± 0.4

* p < 0.05, ** p < 0.01, *** p < 0.001

Table 4. Minimum Volume Loading

Parameter	WT (n = 7)	WT + AZT (n = 4)	TG (n = 5)	TG + AZT (n = 5)
Heart rate (Bpm)	299.1 ± 20.2	286.3 ± 27.9	293.0 ± 15.6	314.67 ± 13.3
MAP (mmHg)	34.57 ± 3.3	34.7 ± 1.5	31.3 ± 2.0	37.0 ± 2.6
EDP (mmHg)	4.14 ± 0.4	3.7 ± 0.9	6.8 ± 2.3	5.7 ± 1.0
+dP/dt (mmHg/sec)	2890.3 ± 178.3	2880 ± 104.6	2708.0 ± 91.2	2804.0 ± 120.2
-dP/dt (mmHg/sec)	-2149.7 ± 120.0	-1848.0 ± 183.3	-1780.0 ± 76.3*	-1804.0 ± 94.3*
TPP/mmHg (msec/mmHg)	0.5913 ± 0.04	0.589 ± 0.03	0.6376 ± 0.02	0.6133 ± 0.03
RT _{1/2} (msec/mmHg)	0.7177 ± 0.041	0.759 ± 0.071	0.8215 ± 0.037	0.8588 ± 0.087
CO (ml/min)	3.11 ± 0.03	3.1 ± 0.06	3.3 ± 0.1	3.233 ± 0.16
Work	108.0 ± 10.1	108.0 ± 5.7	103.8 ± 12.2	121.7 ± 15.4
Coronary flow (ml/min)	1.11 ± 0.22	1.10 ± 0.06	1.15 ± 0.14	1.88 ± 0.36
Aortic flow (ml/min)	2.06 ± 0.20	2.07 ± 0.07	2.15 ± 0.07	7.52 ± 6.0

* p < 0.05, ** p < 0.01, *** p < 0.001

groups of TG and with or without AZT. In contrast, increasing pressure loads resulted in good Starling responses in all parameters for all groups (Fig. 2).

The physiological effects on TG were further analyzed by decreasing preload. Stepwise decreases of preload (volume reduction) to minimum venous return (3 ml/min) resulted in reduced relaxation rates (-dP/dt) in AZT-treated and AZT-untreated TG (p < 0.05).

Other parameters were not significantly affected (Fig. 3). Increasing preload (increasing venous return) showed only prolongation of contraction time (TPP/mmHg). All other parameters showed no significant change compared with the WT (Fig. 3).

Pathological Changes in Control and Transgenic Mice at 35 Days. Necropsies were performed on mice. Hearts revealed globular configuration in AZT-treated

Table 5. Maximum Volume Loading

Parameters	WT (n = 7)	WT + AZT (n = 4)	TG (n = 5)	TG + AZT (n = 5)
Heart rate (Bpm)	342.0 ± 14	348.0 ± 16	360.0 ± 6.43	354.3 ± 12.17
MAP (mmHg)	64.1 ± 3.5	61.8 ± 2.3	64.0 ± 2.30	64.2 ± 2.47
EDP (mmHg)	9.71 ± 1.6	12.3 ± 3.8	16.8 ± 4.43	9.833 ± 2.21
+dP/dt (mmHg/sec)	5218.3 ± 204.4	5250.0 ± 641.3	4627.2 ± 359.79	5108.0 ± 243.76
-dP/dt (mmHg/sec)	-4433.1 ± 281.2	-4284.0 ± 640.0	-3907.2 ± 399.83	-4344.0 ± 282.62
TPP/mmHg (msec/mmHg)	0.3351 ± 0.009	0.3475 ± 0.045	0.3744 ± 0.0180*	0.3407 ± 0.0189
RT _{1/2} (msec/mmHg)	0.465 ± 0.03	0.4435 ± 0.06	0.5266 ± 0.038	0.4657 ± 0.039
CO (mL/min)	6.8 ± 0.2	7.4 ± 0.2	7.1 ± 0.3	7.067 ± 0.056
Work	439.9 ± 34.9	458.3 ± 26.3	451.8 ± 22.96	455.33 ± 19.548
Coronary flow (mL/min)	3.1 ± 0.4	3.9 ± 0.4	3.76 ± 0.607	4.06 ± 0.361
Aortic flow (mL/min)	3.7 ± 0.5	3.5 ± 0.5	3.28 ± 0.441	3.04 ± 0.376

* p < 0.05, ** p < 0.01, *** p < 0.001

TG. Cross-sections of myocardium taken at the base of the left ventricle (LV) papillary muscles from perfusion-fixed samples permitted visual comparison of the LV cavities in similar planes. LV cavity configurations were unremarkable in untreated WT and TG (Fig. 4, A and B). Hearts from WT and TG treated with AZT revealed increased LV cavity size (Fig. 4, C and D). Neither interstitial inflammation nor repair fibrosis was found.

Ultrastructural changes in cardiac myocytes were examined quantitatively in samples from treatment groups and their respective controls (Fig. 5, A and B compared with 5, C and D). Transmission electron microscope (TEM) photomicrographs of hearts of mice treated with AZT revealed ultrastructural changes in cardiac myocyte mitochondria that resembled those observed previously by us and by others. The characteristic features of damage from AZT included fragmentation of mitochondrial cristae and swelling (Fig. 5, C and D).

Morphometric analysis revealed cardiac mitochondrial damage that was greater in the AZT-treated TG than in untreated TG or AZT-treated WT (Table 6). Comparing AZT-treated WT with untreated WT, structural damage to mitochondria increased 5-fold. AZT-treated TG demonstrated a 19-fold increase in the abundance of damaged mitochondria over untreated WT.

Echocardiographic Findings. In WT, after 3 weeks of AZT treatment, the AZT treatment tended to increase LV end-diastolic dimension (EDD) ($p = 0.058$), but did not reach a level of significance. However, the effects of AZT on LV end-systolic dimension (ESD) did reveal a significant increase. As a result, there was a significant (30%) reduction in LV fractional shortening (Table 7). Accordingly, LV mass increased significantly without a change in geometry (ie, the ratio of wall thickness to cavity radius), indicating eccentric LV hypertrophy and cardiac remodeling.

Steady-State Abundance of Cardiac mRNA in TG and WT Treated With and Without AZT, at 21 Days. Figure 6A is a composite histogram indicating the relative signal of atrial natriuretic factor (ANF) normalized to glyceraldehyde-3-phosphate dehydrogenase

(GAPDH) and compared with the control. Untreated TG females demonstrated a 1.8-fold increase in steady-state abundance of ANF mRNA compared with WT, a 22% decrease in SerCa ATPase (SERCA2) mRNA, and essentially no change in Na/Ca Exchanger (NCX1) mRNA. In contrast, female WT+AZT revealed no change in ANF, a 24% decrease in SERCA2 mRNA, and no change in NCX1 (Fig. 6, A to C).

SERCA2 mRNA abundance decreased significantly in both female AZT-treated WT and female TG compared with untreated, gender-matched WT. Untreated female TG demonstrated a 20% decrease in myocardial SERCA2 mRNA. The most profound effect on SERCA2 was found in AZT-treated TG of both genders; the observed decrease in SERCA2 mRNA exceeded 35% in each (Fig. 6B).

Overall, the TG+AZT cohort demonstrated molecular features of cardiac dysfunction based on altered expression of these mRNA. Changes included 2- to 4-fold increase in ANF expression (both genders), approximately 35% decrease in SERCA2 (both genders), and no changes in NCX1 compared with controls.

Discussion

Using a transgenic AIDS model, this study addresses the contribution of HIV-1 (as expressed in a transgenic HIV-1 construct) and of AZT treatment on CM in AIDS. With increased survival of AIDS patients, an increase in the prevalence of AIDS CM may occur. Accordingly, the clinical importance of AIDS CM may rise.

Our data demonstrate that cardiac dysfunction is present in an established murine transgenic AIDS line (containing an HIV-1 construct). Although the TG construct alone was associated with cardiac dysfunction, and AZT treatment caused structural and functional changes in WT, antiretroviral treatment with AZT amplified cardiac dysfunction and worsened ultrastructural features of AIDS CM in TG. Physiological changes in contraction (+dP/dt or TPP/mmHg) and relaxation (-dP/dt and RT_{1/2}/mmHg) were found in AZT-treated and untreated TG. Alterations in relax-

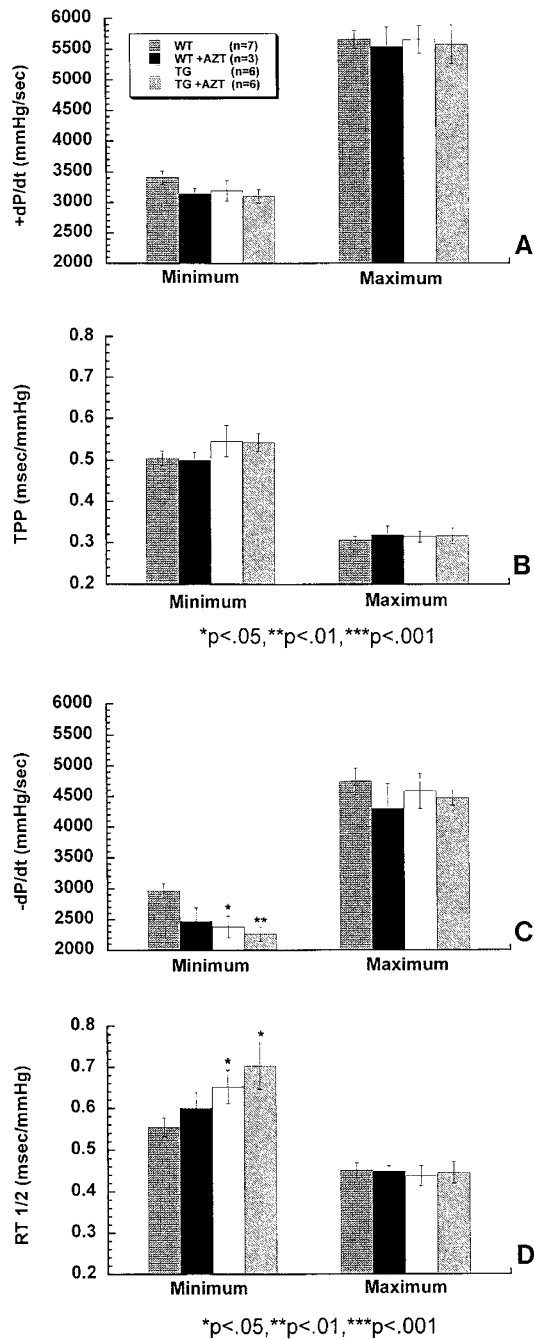


Figure 2.

A and B, Comparison of contractile parameters during pressure loading in TG and WT with and without AZT treatment. Venous return was kept constant at 5 ml/minute. Minimum ranges of aortic pressure were approximately 30 mmHg and maximum ranges approximately 80 mmHg. The resulting work load was approximately 160 mmHg × ml/minute during minimal loading, and 414 to 438 mmHg × ml/minute during maximal loading. C and D, Comparison of relaxation parameters during pressure loading.

ation ($-dp/dt$) were found in AZT-treated WT. Molecular and echocardiographic changes early in treatment were consistent with a pathophysiologic picture of cardiac remodeling and dysfunction, which could lead to CM.

The overall response to loading of the hearts (pressure or volume, respectively) after 35 days AZT treat-

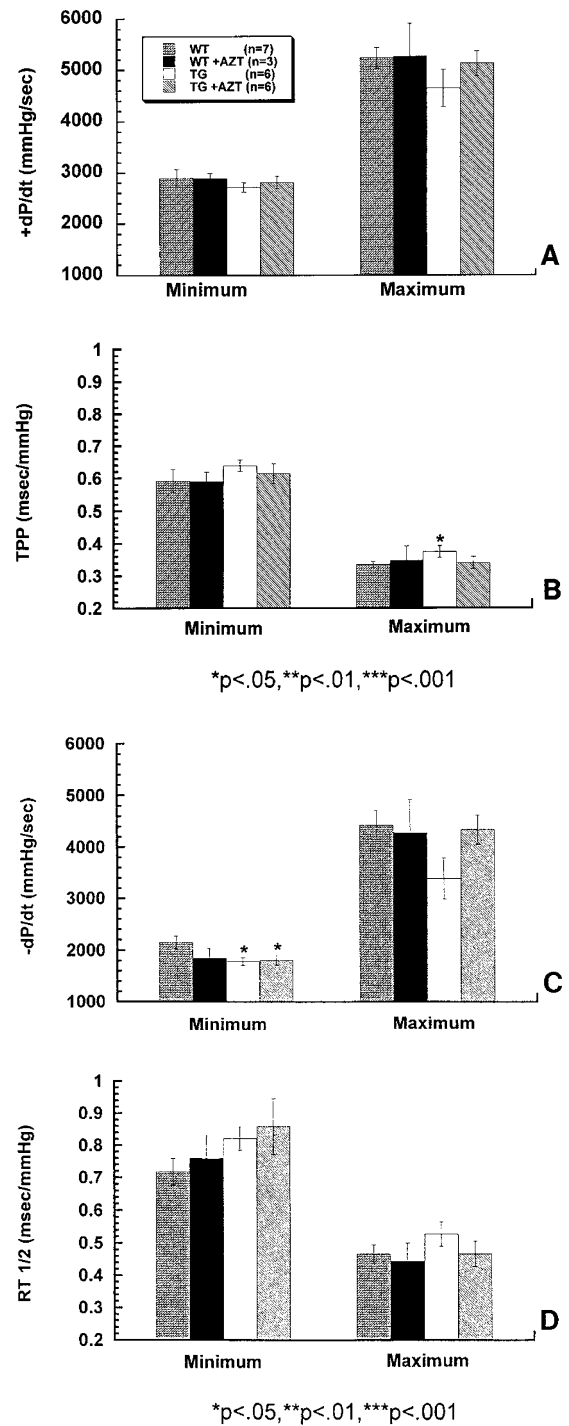


Figure 3.

A to D, Comparison of contractile and relaxation parameters of AZT-treated and untreated WT and TG during volume loading (minimum work load: 108 to 121 mmHg × ml/minute; maximum work load, 440 to 458 mmHg × ml/minute).

ment indicated that both murine LV systolic and diastolic function was sensitive to the HIV-1 TG; this was particularly true with AZT treatment. Dose-response curves indicated no change in response to isoproterenol amongst the groups (data not shown). This lack of response suggested that β -adrenergic receptor regulation may not be involved directly.

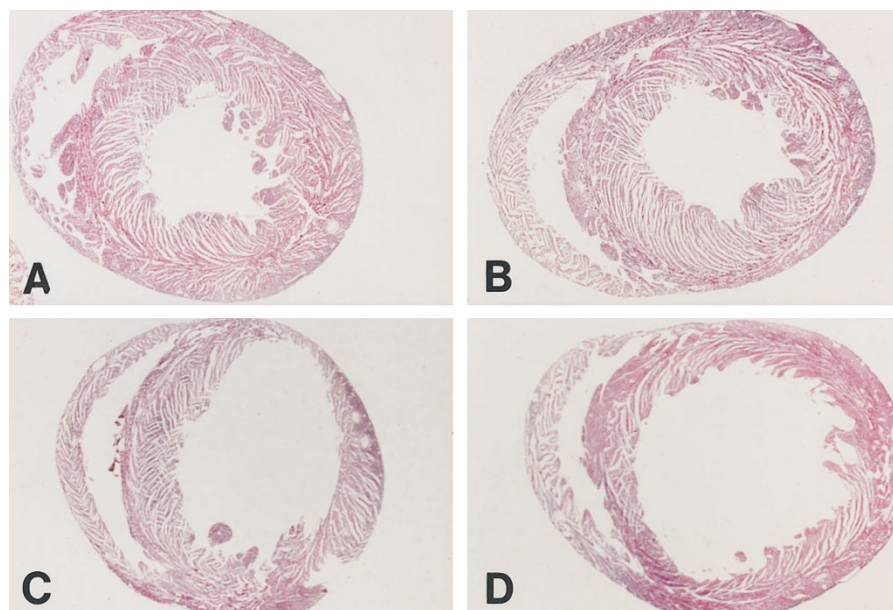


Figure 4.

Representative light photomicrographs from AZT-treated and untreated WT and TG. Upon completion of cardiac performance measurements, hearts were perfusion fixed in 10% neutral buffered formalin, serially sectioned from apex to base, and processed for routine histopathology. For ventricular chamber dimensions, sections were uniformly taken 3 mm from the cardiac apex. Both untreated WT (A) and untreated TG (B) mouse hearts reveal preservation of native left ventricular (LV) and right ventricular (RV) configuration. With AZT treatment (lower two panels), hearts from both WT (C) and TG (D) mice reveal dilation of the LV chamber. More dilation seems to be present in the AZT-treated TG than in the AZT-treated WT. (Original magnification, $\times 7$.)

Pathological changes were found in the hearts of TG and WT treated with AZT for 35 days. Cardiomegaly in AZT-treated TG correlated with increased LV cavity size. Characteristic mitochondrial ultrastructural changes of AZT CM were found in AZT-treated TG and WT. The pathological changes resembled findings with AZT treatment of other species (Cocuera-Pindado et al, 1994; Lamperth et al, 1991; Lewis et al, 1991; McCurdy and Kennedy, 1998).

AZT damaged cardiac mitochondria in WT, with destruction, swelling, cristae dissolution, and fragmentation. Similarly, hearts from AZT-treated TG showed increased mitochondrial damage, but with greater intensity. Based on the ultrastructural findings, the HIV-1 transgene and AZT acted cooperatively to worsen mitochondrial damage.

On a clinical basis, the role of cardiac HIV-1 infection in AIDS CM remains controversial. HIV-1 was isolated and cultivated from an endomyocardial biopsy obtained from a patient with AIDS CM (Calabrese et al, 1987). Our group and others identified HIV-1 in human myocardial tissues by *in situ* hybridization with radiolabeled riboprobes (Grody et al, 1990; Lipshultz et al, 1990; Wu et al, 1990); but the overall pathogenic impact of HIV-1 in CM is not completely elucidated.

High-dose AZT caused mitochondrial CM, decreased cardiac mtDNA, and caused skeletal mitochondrial myopathy in rats (Cocuera-Pindado et al, 1994; Lamperth et al, 1991; Lewis et al, 1991; 1993; McCurdy and Kennedy, 1998, 1996). Those pathological observations correlated with inhibition of DNA polymerase- γ *in vitro* by AZT triphosphate (Konig et al, 1989). Mixed inhibition kinetics were found with DNA

polymerase- γ extracted from the bovine heart (Lewis et al, 1994a) and DNA polymerase- γ was among the most sensitive eukaryotic DNA polymerases to be inhibited by triphosphates of nucleoside analogs. Other thymidine analogs caused similar biochemical pathological changes (Lewis et al, 1994b, 1996). A proposed mechanism for mitochondrial toxicity from antiviral nucleoside analogs suggested altered mitochondrial biogenesis was crucial (Lewis and Dalakas, 1995).

Despite intriguing data *in vivo* and *in vitro*, the impact of antiretroviral therapeutics on the development of clinical AIDS CM remains unclear. In children, one group found progressive LV dilation in HIV-1-infected pediatric patients irrespective of AZT therapy (Lipshultz et al, 1992). In contrast to the former findings, another report suggested AZT caused CM in HIV-1-infected children, whereas didanosine did not (Domanski et al, 1995). In primates, low-dose AZT treatment of the dam resulted in changes of mitochondrial structure and function in the heart and skeletal muscle of the fetus (Gerschenson et al, 1999). Previously, high-dose didanosine administration to Sprague-Dawley rats caused no mitochondrial lesions in the heart (Lewis et al, 1992). The relationship of AZT pharmacokinetic events in the cardiac myocyte (particularly nucleoside phosphorylation), tissue distribution and compartmentalization, and a "threshold effect" of AZT on mitochondrial dysfunction may all affect the development of CM in AIDS (Lewis and Dalakas, 1995; Wallace, 1999).

Recently, support emerged for the concept that altered mitochondrial DNA replication is linked to

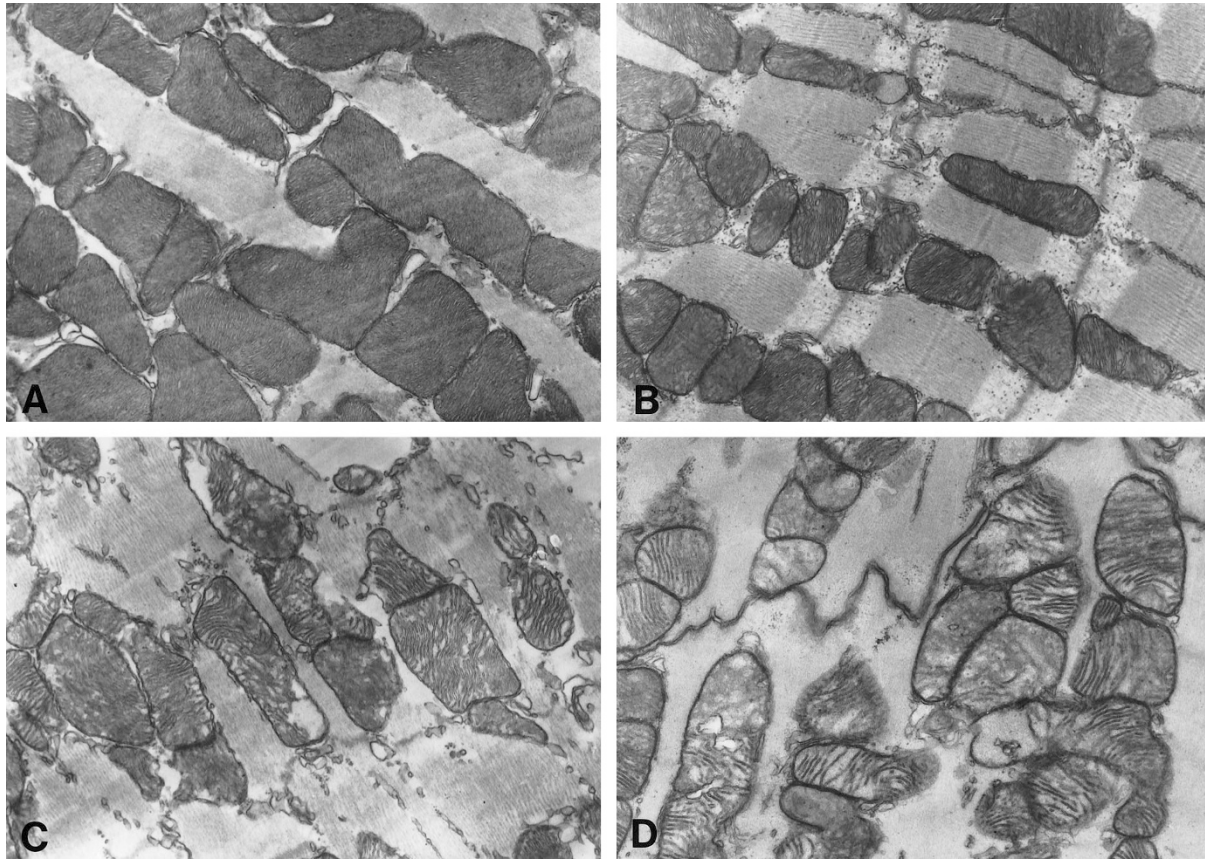


Figure 5.

Ultrastructural changes in cardiac myocytes: With AZT treatment in both WT (C) and TG (D), mitochondrial changes of cristae dissolution and disruption were abundant compared with findings in hearts from WT (A). TG without AZT demonstrated minimal destruction of mitochondria (B). In AZT-treated WT (C) or AZT-treated TG (D), mitochondrial destructive change was increased compared with WT or TG. (Original magnification, $\times 14,000$ for each.)

Table 6. Fraction of Damaged Cardiac Mitochondria

Group	Mean % Mitochondrial Damage	Total Number of Mitochondria Viewed	Total Number of Damaged Mitochondria
WT (n = 7)	2.3	925	21
WT + AZT (n = 4)	11.4	1538	175*
TG (n = 5)	9.6	939	90
TG + AZT (n = 5)	42.4	1674	710†

* $p < 0.03$ compared with WT; † $p < 0.00001$ compared with WT.

mitochondrial toxicity of nucleoside agents (Swartz, 1995; Wright and Brown, 1990). As a cautionary note, current doses of AZT in clinical use are not similar to those employed in experiments here on a mg/kg/day basis. Nonetheless, AZT doses used in the present study resemble those of other studies in different species (Lewis et al, 1991; 1992; McCurdy and Kennedy, 1996, 1998; Ruprecht et al, 1986) and similar findings occurred with low-dose AZT treatment of *E. patas* (Gerschenson, et al, 1999).

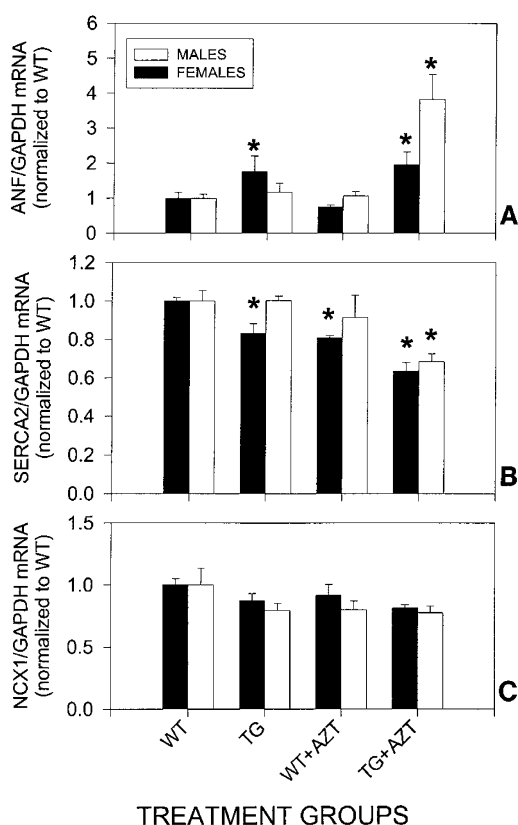
In summary, findings from this AIDS TG model implicate a transgenic HIV-1 construct in the mouse as

a contributor to the development of CM. The precise role of HIV-1 in cardiac dysfunction requires further exploration. AZT also contributes to structural and functional changes of mitochondrial CM in WT and TG mice. Pathophysiologic changes in WT resemble those found in other species. Contractility and relaxation were decreased significantly in TG. Mitochondrial damage observed in AZT-treated TG was greater than that found in untreated WT, untreated TG, or AZT-treated WT. Molecular correlates of CM were most striking in TG+AZT. They included increased steady-state abundance of mRNA encoding ANF and

Table 7. Echocardiographic Data from TG and WT Treated with AZT (21 days)

	WT (n = 3)	WT ± AZT (n = 3)	TG (n = 3)	TG ± AZT (n = 3)
EDD	3.21 ± 0.18	3.71 ± 0.06	3.09 ± 0.19	3.36 ± 0.04
ESD	1.98 ± 0.11	2.77 ± 0.07*	2.07 ± 0.14	2.42 ± 0.24
FS	38.2 ± 0.4	26.6 ± 3.8*	34.0 ± 1.4	28.1 ± 6.4
PWT _{ed}	0.32 ± 0.02	0.36 ± 0.01	0.43 ± 0.01	0.37 ± 0.05
SWT _{ed}	0.32 ± 0.02	0.35 ± 0.01	0.46 ± 0.02	0.38 ± 0.02
h/r	0.20 ± 0.01	0.19 ± 0.01	0.29 ± 0.02	0.22 ± 0.23
LV mass	20.7 ± 2.3	29.9 ± 1.4*	28.7 ± 3.3	26.4 ± 2.2

Mean ± SEM.

* $p < 0.05$ vs WT.EDD, end diastolic dimension; ESD, end systolic dimension; FS, fractional shortening; PWT_{ed}, posterior wall thickness; SWT_{ed}, systolic wall thickness; h/r, wall thickness/cavity radius; LV mass, left ventricular mass.**Figure 6.**

Alterations in atrial natriuretic factor and Ca transporter gene expression in transgenic and AZT-treated mice. Total RNA was isolated from the LV myocardium of male (open bars) and female (filled bars) WT and NL4-3Δ *gag/pol* TG mice. RNA was also isolated from male and female WT and TG animals that also received AZT. The RNA (15 μg) was size-fractionated by denaturing agarose gel electrophoresis, transferred to nitrocellulose, and the resulting Northern blots were sequentially probed with ³²P-labeled cDNA specific for atrial natriuretic factor (ANF), SR Ca ATPase (SERCA2), Na-Ca Exchanger (NCX1), and glyceraldehyde-3-phosphate dehydrogenase (GAPDH). The amounts of ANF and Ca transporter mRNA were quantified by scintillation spectroscopy, and are expressed relative to the amount of GAPDH mRNA. Data were further normalized to mRNA levels of the WT male or female animals. Data were compared by one-way ANOVA or Kruskal-Wallis one-way ANOVA by ranks, as appropriate. * $p < 0.05$ versus respective male or female WT animals; $n = 4$ in each group.

decreased mRNA encoding SERCA2 in LV extracts. Echocardiographic correlates of CM were present after 21 days.

Together these data indicate that the TG and AZT treatment each contribute to cardiac dysfunction. The present studies indicate that transgenic mouse models of AIDS serve as useful tools to recapitulate and explore some features of human AIDS CM.

Materials and Methods

Animals and Retrieval of Hearts

Hemizygous TG employed for the study came from the colony established by Klotman and colleagues and were brought to the specific pathogen free barrier facility at our institution. Tg26 is a well-described murine line of NL4-3Δ *gag/pol* that develops nephropathy (Bruggeman et al, 1997; Kopp et al, 1992) skin lesions in hemizygotes and severe wasting in homozygotes (Kopp et al, 1993). For this study, a 2 × 2 protocol included both hemizygous TG and FVB/n (WT) treated with and without AZT for either 21 days (for echocardiographic and molecular studies) or 35 days (for isolated heart and pathological studies).

For both studies, each mouse received either water without drug or water containing AZT (courtesy of Glaxo Wellcome, Research Triangle, NC; 0.7 mg/ml) ad libitum. Volume consumed was determined in individually housed mice, and amount of AZT (in mg/kg/day) was derived. Fresh water was replenished on alternate days. Total AZT consumption and fractional consumption was determined for each mouse.

Preparation of the Heart for Physiological Measurements and Cardiac Performance Ex Vivo. The work-performing heart preparation was as described (Grupp et al, 1993; Jones et al, 1996). Mice were age-matched littermates (12 to 16 weeks after birth). For retrieval of the hearts, TG ($n = 10$) and WT ($n = 11$) were anesthetized with 30 mg/kg pentobarbital sodium injected ip with 500 U/kg of heparin sodium (Elkins-Sinn, Cherry Hill, New Jersey) to prevent intracardiac blood coagulation. A 12-lead electrocardiogram was obtained and computation of heart rates was performed. For work-performing heart preparations, the hearts were removed through a midsternal incision, weighed, and immediately suspended on a 20-gauge stainless steel cannula and connected to the perfusion apparatus. Retrograde perfusion with heated (37.4°C) and oxygenated Krebs-Henseleit buffer (KHS; con-

taining, in mM: 118 NaCl, 4.7 KCl, 2.5 CaCl₂, 1.2 MgSO₄, 1.2 KH₂PO₄, 0.5 Na-EDTA, 25 NaHCO₃, and 11 glucose) cleared the coronary vasculature of remaining blood and served to insert the left intraventricular catheter for pressure measurements. After a short stabilization period, the pulmonary vein was cannulated. Perfusion of the heart then was switched to antegrade. Venous return (preload) was adjusted and regulated with a micrometer and continuously measured using a dual channel Transonic flow meter (model T206; Transonic Systems, Ithaca, New York). Left ventricular pressure, aortic pressure, and left atrial pressure were measured with COBE pressure transducers.

Aortic flow was monitored on the second channel of the flow meter. Coronary flow was calculated. Aortic pressure (afterload) was adjusted with a second micrometer. Heart rate was computed with a Tachograph. The recording, amplification, and differentiation system was from Grass Instruments (models 7DA, 7PHA, 7P1, 720; Quincy, Massachusetts) and all parameters were analyzed and calculated with a custom-made program. The fluid-filled catheter system responded well within experimental requirements, without distortion up to a frequency of 600 bpm. Heart rate was calculated from an ECG R-R interval. Values measured included: mean aortic pressure, IVP, peak systolic pressure, TPP/mmHg, RT_{1/2}/mmHg, +dP/dt, aortic and coronary flow, venous return (cardiac output), left ventricular minute work (mean aortic pressure × cardiac output), stroke volume (cardiac output ÷ heart rate), stroke work (stroke volume × mean aortic pressure), left atrial pressure, and perfusate temperature. The arterial PO₂ was 650 mmHg, and the PCO₂ ~ 30 mmHg, pH 7.4. The data from the working heart preparations were expressed as mean ± SEM. Starling curves were generated by linear regression using Statview, version 4.01 (Abacus Concepts, Berkeley, California). Regression lines were compared using an F test.

Flow was adjusted to deliver an initial "venous return" of ~5 ml/minute and aortic pressure of 50 mmHg. Cardiac function curves (Starling loading curves) were performed after stabilization of the heart in increasing and decreasing aortic pressure (pressure loading) followed by increasing and decreasing venous return (volume loading). Ventricular end-diastolic and left atrial pressures were monitored continuously.

Preparation for Histologic and Ultrastructural Examination. After physiological measurements were completed, each heart was placed on a Langendorff apparatus and fixed by perfusion at 50 mmHg constant pressure (monitored) with 10% neutral buffered formalin (25 ml) for 5 minutes. The formalin-fixed heart was serially sectioned (2 mm) from apex to base. The second section (2 to 4 mm from apex) was further fixed in diluted Karnovsky's fixative and used for TEM (see below).

For histopathological evaluation, each heart slice was dehydrated through a graded series of alcohols, and paraffin embedded (uniformly oriented apex down). Serial sections from the paraffin-embedded

tissue slice were stained with hematoxylin and eosin, and Masson's trichrome stain. Each tissue slice was examined microscopically. Photomicrography of the stained cross-section was performed with an Olympus photomicroscope (Olympus Optical, Tokyo, Japan).

Transmission Electron Microscopy. The LV free wall was identified in the perfusion-fixed 2-mm slices of heart. Slices of myocardium were sectioned longitudinally and transversely to generate 1-mm cubes that were further fixed in 2% glutaraldehyde-cacodylate for 16 to 24 hours at 4° C for TEM. Each heart provided approximately 20 samples for embedding. Myocardium was rinsed in cold Ringer's solution and post-fixed in 1% OsO₄ (Sigma, St. Louis, Missouri) in PBS, pH 7.4, for 2 to 3 hours. After osmication and rinses, tissue was dehydrated with graded ethanols and embedded in Spurr (Spurr, 1969). Myocardial samples were sectioned (100 nm), stained with uranyl acetate, and examined on a JEOL-JEM-100CX electron microscope. (JEOL, Tokyo, Japan). For each embedded sample of myocardium, 10 random sections were selected and photographed at a uniform magnification (14,000x) by a single investigator.

Photomicrographs were enlarged to 8×10-inch prints and reviewed independently by two investigators for the presence of structurally abnormal mitochondria, for presence of intramitochondrial lamellar bodies, and for intramitochondrial paracrystals (Dalkas et al, 1990). Structurally abnormal mitochondria were operationally defined as mitochondria that had loss or dissolution of ≥ 25% of cristae by area. Damaged mitochondria were scored, and the fraction of damaged mitochondria was defined in the profiles. In each cohort, at least 60 myocyte profiles were examined.

Echocardiography of TG and WT with AZT Administration. Echocardiographic studies were performed in 10-week-old gender-matched (littermate) WT and TG that were treated with or without AZT for 3 weeks (*n* = 3 per cohort). Echocardiographic methods were described previously (Hoit et al, 1995, 1997; Kadambi et al, 1996). Briefly, mice were anesthetized with avertin (0.25 mg/g of body weight) and two-dimensionally targeted M-mode studies were performed with a 9-MHz imaging and a 5 to 7.75-MHz Doppler transducer (Apogee X-200).

M-mode measurements of EDD, ESD, end-diastolic posterior wall thickness (EDPWTH), and end-diastolic septal wall thickness (EDSWTH) were made from original tracings. Calculated variables included: left ventricular fractional shortening (FS = [(EDD - ESD)/EDD]), and LV mass (1.06 × [(EDD + EDPWTH + EDSWTH)]³ - (EDD)³), and the ratio of wall thickness to cavity dimension (h/r).

RNA Extraction from Hearts. For mRNA studies in the heart, treatment protocols were identical to those used above with AZT treatment of 21 days. Total RNA was isolated from LV tissue samples using the acid guanidinium thiocyanate-phenol-chloroform method (Chomczynski and Sacchi, 1987). RNA integrity was examined by electrophoresis and quantified as 260/

280 nm absorbance ratio. Total RNA (10 to 20 μg) was size-fractionated by denaturing agarose gel electrophoresis and transferred by capillary action to nylon membranes for Northern blotting. Northern blots were sequentially probed with ^{32}P -labeled cDNA probes specific for the SERCA2 (a 2.3-kb cDNA fragment of the rat cardiac SERCA2, provided by Dr. W. Dillmann, University of California, San Diego, California), NCX1 (a 1.5-kb cDNA fragment of guinea-pig cardiac sodium calcium exchanger from Dr. K. Philipson, University of California, Los Angeles, California), and ANF (a 0.8-kb cDNA fragment of rat ANF courtesy of T. Inagama, Vanderbilt University, Nashville, Tennessee).

To ensure equal loading conditions and uniformity of transfer, hybridization signal intensity was normalized to the intensity of GAPDH mRNA. The cDNA probes were radiolabeled and hybridized as previously described (Eleftheriades et al, 1993). The amount of each respective mRNA relative to the amount of GAPDH mRNA was then quantified by autoradiography at -80°C and laser densitometry of the signals and normalized to those from untreated controls.

Statistical Analysis. Steady-state abundance of mRNA, heart weights, and mitochondrial damage were compared by one-way ANOVA. Student *t* tests were employed to compare the physiological data in isolated work-performing hearts. $p < 0.05$ was used to establish significance.

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