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Impaired mitochondrial function is abrogated by dexrazoxane in doxorubicin-treated childhood acute lymphoblastic leukemia survivors

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Abstract

BACKGROUND—Impaired cardiac function in doxorubicin-treated childhood cancer survivors is partly mediated by disruption of mitochondrial energy production. Doxorubicin intercalates into mitochondrial DNA (mtDNA) disrupting genes encoding for polypeptides that make ATP.

METHODS—This cross-sectional study examined mtDNA copy numbers/cell and oxidative phosphorylation (OXPHOS) in peripheral blood mononuclear cells (PBMCs) in 64 childhood survivors of high-risk acute lymphoblastic leukemia (ALL) treated on Dana-Farber Cancer Institute Childhood ALL protocols who had received doxorubicin alone (42%) or with dexrazoxane (58%), a cardioprotectant. Mitochondrial DNA copies per cell and OXPHOS enzyme activities of nicotinamide adenine dinucleotide (NADH) dehydrogenase (Complex I, CI) and cytochrome *c* oxidase (Complex IV, CIV) were measured by quantitative real time-polymerase chain reaction (qRT-PCR) immunoassay and thin layer chromatography, respectively.

RESULTS—At a median follow-up of 7.8 years after treatment, the median number of mtDNA copies per cell for patients treated with doxorubicin alone was significantly higher than for those who also received dexrazoxane (medians, 1106.3 and 310.5; $P=0.001$). No significant differences were detected between groups for CI or CIV activities.

CONCLUSIONS—Doxorubicin-treated survivors had increased PBMC mtDNA copies/cell and concomitant use of dexrazoxane was associated with lower mtDNA copies/cell. Due to a possible

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compensatory increase in mtDNA copies/cell to maintain mitochondrial function in the setting of mitochondrial dysfunction, overall OXPHOS activity was not different between groups. The long-term sustainability of this compensatory response in these survivors at risk for cardiac dysfunction over their lifespan is concerning.

Keywords

Childhood cancer survivors; dexrazoxane; doxorubicin; mitochondrial function; mitochondrial DNA

INTRODUCTION

Childhood cancer survival rates have improved dramatically over recent decades. The five-year event-free survival rate for childhood acute lymphoblastic leukemia (ALL) in 2011 was 82%, a significant improvement from less than 50% in the 1970s (1). Much of this success can be attributed to the widely used and highly effective anthracycline, doxorubicin. However, long-term follow-up studies of this growing population have revealed that doxorubicin treatment is associated with cardiac late-effects years later (2-10).

Doxorubicin-associated cardiotoxicity can occur both during and years after therapy, progressing from asymptomatic left ventricular (LV) dysfunction, to cardiomyopathy, heart failure, and in some cases, the need for heart transplantation or cardiac death (7,9,11,12). Identifying patient risk factors such as female sex, anthracycline cumulative dose and dose-rate, trisomy 21, younger age at diagnosis, black race, prior and concomitant cardiotoxic therapy, preexisting risk for developing cardiovascular disease or identified cardiovascular disease, and increasing time since cancer treatment has helped characterize those at higher risk for these anthracycline-associated cardiac late-effects (5,13-17), although predisposition to cardiac damage varies substantially among individuals.

Recent evidence suggests that doxorubicin-associated cardiotoxicity is at least partially due to the generation of semiquinone free radicals by doxorubicin-iron complexes in the myocardium. Furthermore, one animal study found a dose-dependent increase of doxorubicin in mitochondria, indicating that doxorubicin accumulates in the mitochondria and leads to an increase in mitochondrial iron levels (18). Reaction of these free radicals with oxygen (reactive oxygen species) leads to lipid peroxidation and ultimately DNA damage (19-22). Dexrazoxane, a cardioprotectant, chelates iron, thereby reducing iron from forming doxorubicin-iron complexes and, as a result, lowering free radical DNA damage (23).

Mitochondria contain mitochondrial DNA (mtDNA) that is separate from nuclear DNA and encodes 13 polypeptides of oxidative phosphorylation (OXPHOS). Preclinical studies have associated doxorubicin exposure with irreversible cardiac mitochondrial dysfunction (24-26); clinical histological studies have observed doxorubicin-related structural abnormalities; and one recent study of childhood ALL survivors found that gene polymorphisms of mitochondrial expression may affect mtDNA replication and possibly indicate susceptibility to doxorubicin (27). We have found that doxorubicin-treated long-term survivors of childhood ALL had significantly more mitochondrial DNA mutations or

polymorphisms than healthy children (28). Our findings were similar to what has been observed in murine late doxorubicin cardiotoxicity where increased peripheral blood lymphocytes contained mitochondrial mutations that were related to respiratory chain defects and late onset cardiomyopathy (29). These studies demonstrate that doxorubicin has deleterious effects on mitochondrial structure and DNA, which could impact mtDNA copy numbers and OXPHOS.

Further, preclinical studies on anthracycline exposure in mitochondria have shown disrupted OXPHOS, particularly in OXPHOS enzyme activities of NADH dehydrogenase (complex I (CI) and cytochrome *c* oxidase (complex IV (CIV) activity (30,31), as well as changes in the mtDNA in the form of mutations, deletions, and reduced copy numbers per cell in heart tissue (32,33). If the mitochondria of doxorubicin-exposed cancer survivors are indeed impaired, either through DNA intercalation or oxidative stress, they may undergo clonal expansion of their mtDNA in order to compensate for mutations or deletions that could lead to normal OXPHOS and ATP production (34) in the myocardium. Studies in rats treated with dexrazoxane and doxorubicin demonstrated that concomitant dexrazoxane treatment prevented doxorubicin-induced cardiac mitochondrial dysfunction by maintaining OXPHOS activities and mtDNA integrity (32,35).

Therefore, we examined mtDNA and OXPHOS enzyme activities in peripheral blood mononuclear cells (PBMCs) in childhood survivors of high-risk ALL who received doxorubicin alone or dexrazoxane with doxorubicin.

MATERIALS AND METHODS

Participants

Following institutional review board approval, patients were enrolled on DFCI ALL Consortium Protocol 05-336 if they met all of the following eligibility criteria: were previously diagnosed with ALL and treated on DFCI Childhood ALL protocols, had no prior relapse (in first complete remission), and had no prior allogeneic stem cell transplant. For this cross-sectional study only high-risk ALL patients were included and additional inclusion criteria were established: patients had to have been diagnosed with high-risk ALL at least 4 years prior and did not have a history of secondary malignancies treated with chemotherapy or radiation. Informed consent was obtained from patients who were >18 years old or from parents of patients 18 years old. The total planned cumulative doxorubicin dosage for high-risk ALL patients was 300-360 mg/m². All patients received doxorubicin alone or doxorubicin plus dexrazoxane.

Sample collection

Peripheral blood was collected at the time of enrollment. All samples were shipped overnight at room temperature to a central processing laboratory. Whole blood was diluted 1:1 with basic medium saline/PBS. Samples were centrifuged on lymphocyte separation solution (LSS, Organon Biosciences, Kenilworth, NJ) and the mononuclear cell layer was isolated and washed twice with two volumes of PBS. Cell samples were enumerated and adjusted with PBS to achieve a concentration of 1×10^6 cells/mL and again centrifuged and

aspirated. Resulting mononuclear cells constituting primarily lymphocytes were cryopreserved by freezing at -70 °C until mtDNA isolation and analysis. All samples were processed within 30 hours of collection.

PBMC isolation

Blood samples were drawn into ethylenediaminetetraacetic acid and heparin containing vacutainers at each study site. PBMCs were isolated within 8 hours via a Ficoll gradient (GE Healthcare, Waukesha, WI) and platelets were removed by two phosphate-buffered saline washes and centrifugation at 300 relative centrifugal force. The PBMCs were then pelleted and frozen at -80°C.

Acquisition of mtDNA copy numbers and OXPHOS activity

PBMC mtDNA copies/cell were measured by quantitative real time-polymerase chain reaction. DNA was extracted from frozen PBMC using a Qiagen DNA kit (Qiagen, Valencia, CA). Standardization of real-time PCR was performed using LightCycler FastStart DNA Master SYBR Green I with the Roche LightCycler instrument (Roche, Indianapolis, IN). A dilution series of the control plasmid containing the 90 base-pair mtDNA NADH dehydrogenase, subunit 2 and the 98 base-pair Fas Ligand gene was prepared to set up the standard. Each sample and standard was run in a duplicate 20 µl reaction mixture. At the conclusion of the PCR, a melt curve analysis was started at 65°C, and the temperature was increased half a degree every 30 sec for 60 cycles. The results were analyzed with Version 4.0 LightCycler software (36).

CI and CIV were measured by immunoassay and thin layer chromatography, respectively. Each vial of viable PBMC was thawed and washed in 0.5 ml of PBS twice before addition of 0.5 ml of ice-cold extraction buffer [1.5% lauryl maltoside, 25mM Hepes (pH 7.4), 100 mM NaCl, plus protease inhibitors (Sigma, P-8340)]. Samples were mixed gently and kept on ice for 20 minutes, and then they were spun in a micro-centrifuge at 16,400 rpm at 4°C for 20 minutes to remove insoluble cell debris. The supernatant, an extract of detergent-solubilized cellular proteins, was then assayed with the OXPHOS dipstick assays. All samples were loaded on dipsticks with equal amounts of total cell protein or enzyme activity using an amount previously established with control samples to generate signals within the linear range of the assay. Therefore, the resulting dipstick signals were directly proportional to the amount of OXPHOS protein or enzyme activity in the sample. Quantitation of dipstick signals was done by densitometric scanning (36). OXPHOS CI and CIV were specifically measured, as they are the beginning and end of the electron transport chain and are partially encoded by mtDNA. Furthermore, preclinical studies on anthracycline exposure observed disrupted OXPHOS, particularly in CI and CIV, in heart tissue (30,31). We expect that any disturbances to OXPHOS would be measurable at these complexes.

Statistical analysis

A Wilcoxon rank-sum test was used to compare continuous measures between treatment groups. A multiple linear regression model for each continuous outcome measure (PBMC mtDNA copy number and CI and CIV enzyme activity measures) was constructed to test for differences between treatment groups while adjusting for patient and sample characteristics

including age at study registration (<10 years versus ≥10 years), sex, T-cell immunophenotype, initial white blood cell count (WBC), and red blood cell (RBC) contamination. The cumulative doxorubicin dose was also included in the modeling on a subset of the data, excluding those patients dosed as mg/kg. Due to the skewness of the distributions, the values for mtDNA copy number and CI and CIV enzyme activities were log₁₀ transformed prior to modeling. All analyses were performed using SAS 9.2. *P*-values were two-sided and considered significant at the 0.05 level.

RESULTS

Sixty-four patients provided samples at a median of 7.8 years (range, 2.9-30.2 years) after doxorubicin treatment (Table 1). The median cumulative dose of doxorubicin was 300 mg/m² (120-382 mg/m²) and 37 patients (58%) received concomitant dexrazoxane.

In a univariate analysis, a significant difference was detected for mtDNA copy number between patients who received doxorubicin only (median, 1106.3 copies/cell; range, 144.2-6746.8 copies/cell) and those who also received dexrazoxane (310.5 copies/cell; range, 15.3-1859.2 copies/cell) (*P*=0.001; Figure 1). No significant differences were found between treatment groups in CI or CIV enzyme activities (Table 2). However, the median CI activity was significantly higher for males (12.6 OD/μg × 10³; n=31; range, 5-41.4) compared to females (6.7 OD/μg × 10³; n=27; range, 5.0-27.7) (*P*=0.037), but not for CIV. Similarly, there was no significant difference in the mtDNA copies/cell when comparing girls to boys.

In a regression analysis, treatment remained associated with mtDNA copy numbers after adjusting for age, sex, T-cell immunophenotype, RBC contamination, and initial WBC cell count (*P*=0.001, Table 3). No other covariates were significant in the modeling. When considering a model including the cumulative dose of doxorubicin for the subset of patients dosed as mg/m² (n=48) adjusting for the same covariates, the effect of treatment remained significant (*P*=0.007; model not shown). Drug treatment was not associated with CI or CIV enzyme activities. However, sex remained associated with CI enzyme activity (*P*=0.03; model not shown).

In subset analysis of each treatment group the sex difference was only present in the doxorubicin plus dexrazoxane group (male versus female median 16.4 OD/μg × 10³ versus 5.3 OD/μg × 10³; *P*=0.007), not in the doxorubicin alone group (male versus female median 9.8 versus 12.3; *P*=0.85).

DISCUSSION

In this study, after a median of seven years since completing doxorubicin treatment, survivors of childhood high-risk ALL treated with doxorubicin alone had three times the number of mtDNA copies/cell in PBMCs than did children treated with doxorubicin plus dexrazoxane, but OXPHOS enzyme activities did not differ between groups. In addition, mtDNA copies/cell in those who received dexrazoxane were within the normal ranges observed in our previous studies (100-350 copies/cell) (36,37).

As described earlier, in the myocardium, doxorubicin accumulates in the mitochondria, forms complexes with iron, inducing oxidative stress, which then leads to lipid peroxidation and DNA damage (18-22). Additionally, elevated oxidative stress is associated with higher mtDNA copies per cell in aging tissue (34). Dexrazoxane protects the mitochondria by chelating the iron and preventing the formation of the doxorubicin-iron complex (38). Also, *in vitro* dexrazoxane antagonizes doxorubicin-induced DNA damage in H9C2 cardiomyocytes through its interference with DNA topoisomerase II, which could implicate this topoisomerase in doxorubicin cardiotoxicity (39). *In vivo* mouse studies with a cardiomyocyte-specific deletion of DNA topoisomerase II prevented mice from developing doxorubicin progressive heart failure whereas in control mice treated with doxorubicin, the genes involved in the regulation of mitochondrial biogenesis and function were decreased (40). Together, this might explain why those who were treated with doxorubicin alone had higher mtDNA copies/cell than those who received dexrazoxane before each doxorubicin dose. This may reflect doxorubicin-associated accelerated mitochondrial aging.

Although a decrease in OXPHOS enzyme activity might be expected, due to impaired mitochondrial function, which was not the case in our study. As with aging tissue, we suspect that doxorubicin-exposed mitochondria may undergo clonal expansion to compensate for doxorubicin-induced mtDNA deletions or mutations to maintain OXPHOS and ATP production (34). A similar compensatory mechanism was seen in a study of adults with Leber's hereditary optic neuropathy, an inherited blinding disease caused by homoplasmic point mutations in the CI subunit genes of mitochondrial DNA, which are part of the same NADH dehydrogenase subunit examined in our study. This study showed that unaffected carriers of the mutation had a higher mtDNA copy number compared with their affected relatives and controls (41). This would result in normal transcription levels of the NADH dehydrogenase subunit, resulting in normal OXPHOS enzyme activity for this enzyme. The authors postulated that a higher mtDNA copy number per cell in carriers might override some of the pathogenic effects of mitochondrial DNA mutations (41).

Lebrecht et al. found that doxorubicin-exposed rats had higher mtDNA mutations and deletions, and lower mtDNA copies/cell in cardiac tissue than controls after only a few weeks after the last doxorubicin exposure (29). It is possible that mtDNA copies/cell may have been lower in our patients initially after doxorubicin treatment and then later increased through clonal expansion to maintain steady OXPHOS levels. However, this hypothesis cannot be confirmed, as we did not measure mtDNA mutations, deletions, or copies per cell immediately after treatment or serially after completing doxorubicin therapy.

This study has limitations. First, studies examining doxorubicin-induced cardiotoxicity typically use cardiac tissue to measure mitochondrial function, however, we examined systemic changes in PBMC mitochondria, as acquiring cardiac mitochondria is invasive and unethical. Yet, systemic mitochondrial pathologies have been shown to correlate in PBMCs and in the tissues of interest (37,42), therefore it is possible that what we observed in PBMC mitochondria may reflect what is occurring in cardiomyocyte mitochondria. Furthermore, the transcriptome in doxorubicin-exposed cardiac cells have been shown to have a high similarity in the gene expression profiles to those in PBMCs, suggesting that the PBMC transcriptome might serve as a surrogate marker of doxorubicin-induced cardiotoxicity (43).

However, further study is warranted in this area. Second, it is possible that the high number of mtDNA copies in the doxorubicin group was due to greater numbers of mitochondria present in the PBMCs. Lebrecht et al. (29) found higher citrate synthase activity, a marker for mitochondrial numbers, in cardiac cells of doxorubicin treated rats as compared to controls. An assay measuring the mitochondrial proteins citrate synthase or porin would have indicated if there were increased numbers of mitochondria in the PBMCs of our patients, but further samples were not available for testing. Future research should examine the number of mitochondria as well as the number of mtDNA copies. Lastly, we did not have sufficient echocardiograms taken at the same time as the samples to determine LV function. However, the patients enrolled in this study were sampled from a larger cohort where LV function was examined after doxorubicin alone or dexrazoxane with doxorubicin treatment [44]. While effects of dexrazoxane on LV function were mild and more significant in the females, generally there was an improvement in cardiac outcomes [44].

A recent study supports the importance of the mitochondrial respiratory chain as a target for doxorubicin cardiotoxicity. This study described a previously unidentified cardiomyocyte-signaling pathway that couples doxorubicin-induced mitochondrial respiratory chain defects and necrotic cell death in a mutually dependent and obligatory link to the BH3-only protein Bcl-2-like 19KDa-interacting protein 3 (Bnip3). Mitochondrial localization of Bnip3, increased reactive oxygen species production, loss of the mitochondrial membrane potential, and mitochondrial permeability transition pore opening resulting in doxorubicin-induced contractile failure and necrotic cell death were associated with cardiomyocyte vacuolization and disrupted mitochondria in an acute injury model that did not examine late-effects. Interventions that antagonize Bnip3 appear beneficial in preventing doxorubicin-induced mitochondrial injury and heart failure (45).

Overall, this is the first long-term study in pediatric high-risk ALL survivors to show that doxorubicin therapy affects mtDNA numbers and that concomitant dexrazoxane therapy improved mtDNA copies/cell; this was at a median of 7 years of completing treatment of doxorubicin using a median cumulative dose of 300 mg/m². This supports the evidence that dexrazoxane adjuvant therapy in pediatric high-risk ALL patients offer systemic mitochondrial protection as observed in PBMCs, in addition to the already documented cardioprotection seen in this same patient population by measuring cardiac biomarkers of myocardial injury during doxorubicin therapy (serum cardiac troponin T) and cardiomyocyte stress (N-terminal pro brain natriuretic peptide) during therapy (23,46) and by assessment of LV structure and function by echocardiography in long-term survivors (44). The increase in mtDNA copies/cell to maintain mitochondrial function is likely compensatory in the setting of mitochondrial dysfunction. Although, overall OXPHOS activity was not different between groups. The sex differences in CI OXPHOS activity observed in our study may help explain how dexrazoxane affects girls and boys differently. Girls have greater doxorubicin cardiotoxicity than boys as long-term survivors of childhood cancer (15). Girls also derive greater doxorubicin cardioprotection than boys when treated with dexrazoxane for ALL (44). These sex differences suggest that there are differences in both mitochondrial and free radical injury protective mechanisms, similar to what is found in preclinical doxorubicin studies where these pathways have been found to be important targets for cardiotoxicity (21,25,26). The importance of ATP-binding cassette transporter

and endothelial nitric oxide synthase gene mutations on the development of late cardiotoxicity in long-term survivors of doxorubicin-treated ALL was recently noted (47) and further supports the importance of mitochondrial and oxidative stress as targets. In that study, abnormal mitochondrial function after doxorubicin therapy might relate to late depressed LV contractility and in disturbed cardiac growth (47). This suggests the importance of both mitochondrial function and free radical pathways in the development of doxorubicin cardiotoxicity. Similarly, the doxorubicin cardiotoxicity free radical damage hypothesis was supported by the presence of hemochromatosis gene mutations, resulting in more iron and free radical formation by the coupling of iron with doxorubicin, are associated with more dead and dying cardiomyocytes during doxorubicin therapy for childhood ALL and in more persistent echocardiographic abnormalities years later (48). These findings support the possible role of dexrazoxane prior to anthracycline therapy to protect mitochondrial function in other healthy tissues such as the ovaries (49). We have further reported that the use of dexrazoxane before every doxorubicin dose resulted in less late cardiotoxicity for children treated for T-cell ALL (50), that its use in pediatric patients with osteosarcoma allowed dose escalation of doxorubicin to 600 mg/m² without evidence of cardiotoxicity (51), and that its use in osteosarcoma prevented the additive cardiotoxicity associated with the concurrent use of doxorubicin and trastuzumab (52).

Additional follow-up is needed to determine if this compensatory mechanism could eventually become insufficient for maintaining OXPHOS activity in the longer term.

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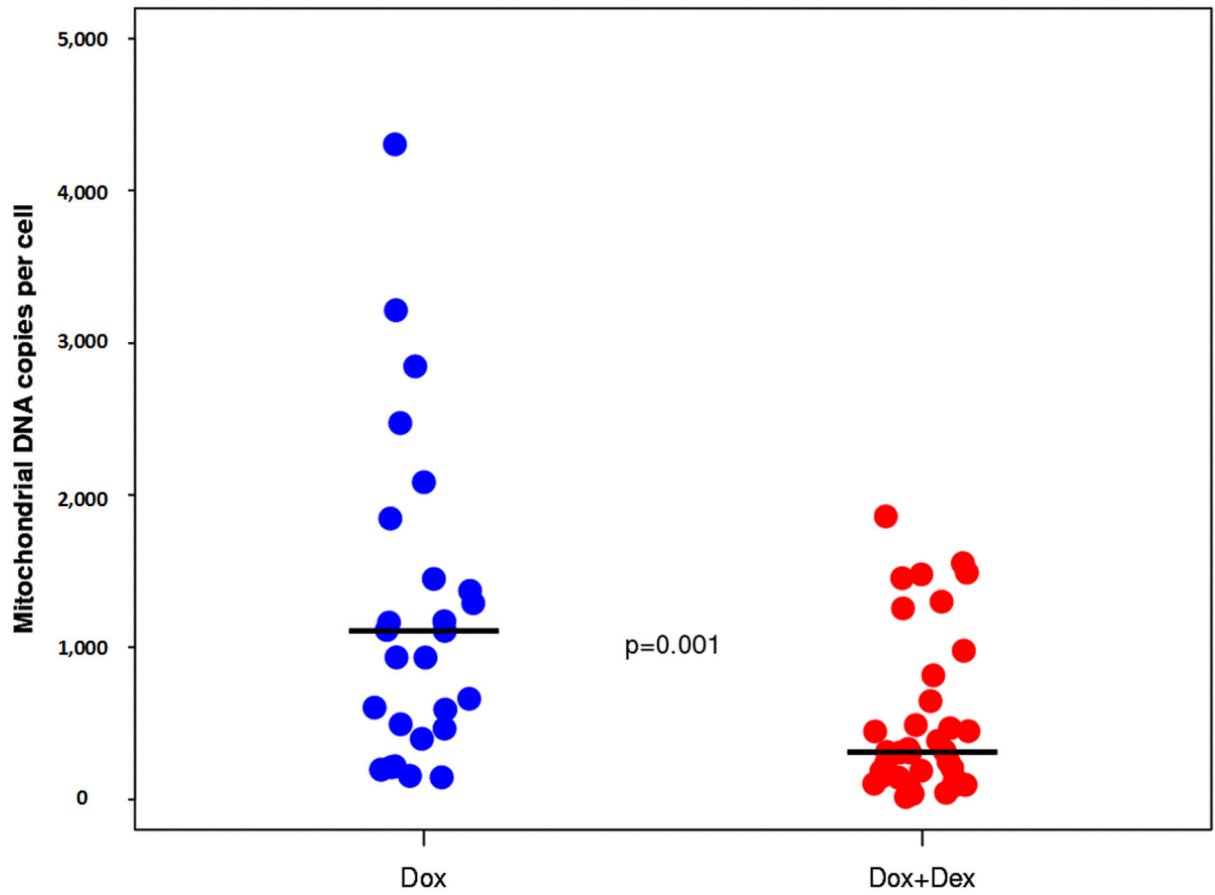


Figure 1. MtDNA levels for doxorubicin alone without dexrazoxane versus dexrazoxane administered before each dose of doxorubicin. DEX, dexrazoxane; DOX, doxorubicin; DOX+DEX, administration of dexrazoxane prior to every doxorubicin treatment.
*Horizontal black bars indicate median.

Table 1

Patient characteristics of all high-risk ALL patients (n=64).

	n (%)
Time to sample from DOX or DEX+DOX treatment years, median (range)	7.8 (2.9, 30.2)
Protocol	
77-01	1 (1)
81-01	1 (1)
87-01	4 (6)
91-01	13 (20)
95-01	14 (22)
00-01	31 (48)
Sex	
Male	33 (52)
Female	31 (48)
Age ^a , median years (range)	5.1 (1.4, 17.1)
<10 years	41 (64)
10 years	23 (36)
T-cell immunophenotype	
Yes	7 (11)
No	57 (89)
Initial WBC ($\times 10^3/\mu\text{L}$), median (range)	32.2 (1.3, 354.0)
Treatment	
DOX	27 (42)
DEX+DOX	37 (58)
Cumulative dose of DOX (mg/m^2) ^b , median (range)	300 (120, 382)

^a Age at the time of protocol registration.^b n=50 reported.

DOX, doxorubicin; DEX, dexrazoxane; DEX+DOX, administration of dexrazoxane prior to every doxorubicin treatment; WBC, white blood cell count.

Table 2

Mitochondrial Parameters and Treatment

	n	DOX Median (range)	n	DEX+DOX Median (range)	P-value
MtDNA copies/cell	27	1106.3 (144.2-6746.8)	35	310.5 (15.3-1859.2)	0.001
CI activity (OD/ $\mu\text{g} \times 10^3$)	25	10.5 (5.0-31.3)	33	11.7 (5.0-41.4)	0.97
CIV activity (OD/ $\mu\text{g} \times 10^3$)	25	9.8 (5.0-20.1)	34	8.1 (4.7-23.4)	0.40

CI, Complex I; CIV, Complex IV; DOX, doxorubicin; DEX, dexrazoxane; DEX+DOX, administration of dexrazoxane prior to every doxorubicin treatment; OD, optical density.

Table 3

Multiple regression model for mtDNA copy number for DOX alone versus DEX+DOX

Variable	Coefficient (β)	SE	P-value
Intercept	2.60	0.28	<0.001
Treatment (DOX alone versus DEX+DOX)	0.46	0.13	0.001
Sex (male versus female)	0.11	0.14	0.45
Age ^a (<10 versus 10 years)	0.02	0.14	0.89
Immunophenotype (B-cell versus T-cell)	-0.18	0.21	0.41
RBC contamination (no versus yes)	0.02	0.14	0.89
Initial WBC	-0.0006	0.001	0.60

^aAge at the time of protocol registration.

DOX, doxorubicin; DEX, dexrazoxane; DEX+DOX, administration of dexrazoxane prior to every doxorubicin treatment; RBC, red blood cell; SE, standard error; WBC, white blood cell count.

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