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## Cardiotoxicity of Cancer Treatments: Focus on Anthracycline Cardiomyopathy

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### Abstract

Significant progress has been made in developing new treatments and refining the use of pre-existing ones against cancer. Their successful use and the longer survival of cancer patients have been associated with reports of new cardiotoxicities and the better characterization of the previously known cardiac complications. Immunotherapies with monoclonal antibodies against specific cancer-promoting genes, chimeric antigen receptor T cells (CAR-T-cells), and immune checkpoint inhibitors have been developed to fight cancer cells, but they can also show off-target effects on the heart. Some of these cardiotoxicities are thought to be due to non-specific immune activation and inflammatory damage. Unlike immunotherapy-associated cardiotoxicities which are relatively new entities, there is extensive literature on anthracycline-induced cardiomyopathy. Here, we provide a brief overview of the cardiotoxicities of immunotherapies for the purpose of distinguishing them from anthracycline cardiomyopathy. This is especially relevant as the expansion of oncologic treatments present greater diagnostic challenges in determining the etiology of cardiac dysfunction in cancer survivors with history of multiple cancer treatments including anthracyclines and immunotherapies administered concurrently or serially over time. We then provide a focused review of the mechanisms proposed to underlie the development of anthracycline cardiomyopathy based on experimental data mostly in mouse models. Insights into its pathogenesis may stimulate the development of new strategies to identify patients who are susceptible to anthracycline cardiomyopathy while permitting low cardiac risk patients to receive optimal treatment for their cancer.

### Keywords

anthracycline; cardiomyopathy; immunotherapy; mitochondria; mitochondrial genome; p53

## INTRODUCTION

Chemotherapy is the primary treatment for many types of cancers, and its improvement has contributed to the increased survival of cancer patients. Although chemotherapeutic agents generally target proliferating neoplastic cells, quiescent cardiomyocytes can also

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be damaged thereby leading to cardiomyopathy in cancer survivors. Treatment with one of the anthracycline class of chemotherapeutic agents such as daunorubicin, epirubicin, mitoxantrone, or doxorubicin can result in a progressive form of heart failure with a high mortality rate<sup>1-4</sup>. Among these anthracyclines, doxorubicin is an old but highly effective and commonly utilized agent. In a cumulative dose-dependent manner, it can cause myocardial dysfunction in a subset of patients for unclear reasons, and the onset of heart failure can occur many years after exposure to the chemotherapy. Despite extensive clinical characterization and a wealth of experimental data using model systems, there are still no preventive strategies or targeted treatment for anthracycline cardiomyopathy.

The introduction of immunotherapies against cancer has brought to the forefront the need to better understand the risk factors for anthracycline-induced cardiac damage. The use of trastuzumab, an anti-human epidermal growth factor receptor 2 (HER2) monoclonal antibody, in combination with an anthracycline resulted an unanticipated and significant increase in the risk of cardiac dysfunction<sup>5</sup>. More recently, immunotherapies using monoclonal antibodies against other cancer promoting genes and anti-cancer immune activating treatments such as immune checkpoint inhibitors (ICI) and chimeric antigen receptor T cells (CAR-T) have also turn out to have cardiovascular toxicities<sup>6, 7</sup>. Like the unexpected effect of anthracyclines on non-proliferating cardiomyocytes, eliciting cytotoxic effects using monoclonal antibodies for cancer cell recognition have also resulted in adverse side-effects. With these developments, the sub-specialty of cardio-oncology has coalesced with an emphasis on incorporating the advances in immune-oncology, such as CAR-T cell and ICI immunotherapy, for preventing and managing cancer treatment cardiotoxicities<sup>7</sup>.

Because over half of all cancer patients may be eligible for some form of immunotherapy<sup>78</sup>, it is useful to briefly summarize some of their cardiotoxicities in order to distinguish them from that of anthracyclines. Heart failure is the major cause of morbidity associated with anthracyclines, so we will review some of the immunotherapies and targeted therapies that have been associated primarily with adverse effects on myocardial contractility or the cardiovascular system. We will then focus on detailing the current understanding of the molecular mechanisms underlying anthracycline cardiotoxicity based mostly on *in vivo* mouse models with the goal of potentially translating these findings to assist in the management of cancer patients treated with anthracyclines. For the cardiotoxicities of chemotherapeutic agents such as proteasome inhibitors and tyrosine kinase inhibitors, the reader is referred to specialized reviews that more comprehensively discuss these and other chemotherapeutic agents.<sup>9,10,11, 12</sup>.

## **SOME CARDIOTOXICITIES ASSOCIATED WITH CANCER IMMUNOTHERAPIES AND TARGETED ANTI-ONCOGENIC SIGNALING ANTIBODIES**

### **Immune Checkpoint Inhibitors (ICIs)**

ICIs are therapeutic monoclonal antibodies that block T cell inhibitory regulators such as the cytotoxic T lymphocyte-associated protein-4 (CTLA-4), programmed cell death protein-1 (PD-1) and PD-1 ligand (PD-L1)<sup>6, 8</sup>. Disrupting these inhibitory activities with

ICI treatment results in varying degrees of uncontrolled T cell activation and subsequent off-target immune-related adverse events that affect different organs such as skin, thyroid, intestine, liver, lung and heart. Examining mouse models with ablation of the immune checkpoint inhibitor genes support the concept of cardiac damage from T cell overactivation and infiltration into the myocardium (Table 1)<sup>13, 14</sup>. The PD-1 signaling pathway appears to play an important role in restraining CD4+ and CD8+ T cell activities in the myocardium, but the specific molecular events leading to their myocardial infiltration and subsequent pathogenesis remain to be elucidated<sup>15, 16</sup>. Human pathological studies show PD-L1 activation with inflammatory cell infiltrates while there are macrophage polarization changes observed in mouse models after ICI treatment<sup>17,18</sup>. PD-L1-expressed endothelial cells other than cardiomyocytes may be also targets of ICI-related cardiac injury<sup>19</sup>. Besides a direct effect on the heart, such interactions of the vasculature, cytokines or autoantibodies may explain the complex mechanism of ICI-related systemic adverse events which are more comprehensively reviewed elsewhere<sup>13, 16</sup>.

Although the incidence of clinically significant cardiotoxicities with a single ICI is low (range of 1% or less), the combined use of two different ICIs significantly increases the risk<sup>20, 21</sup>. Also, inhibitors of protein kinases BRAF and MEK can cause cardiovascular toxicities<sup>4</sup>, and their combined therapy with ICIs have been reported to worsen ICI-related myocarditis<sup>22</sup>. Distinct from the late onset cardiomyopathy caused by anthracyclines, ICI cardiotoxicity can occur within days after treatment but the median time is about 4 wk with most occurring within the first 6 wk after treatment. Of the different cardiotoxicity manifestations which include arrhythmias and non-inflammatory myocardial dysfunction such as stress cardiomyopathy, myocarditis poses the greatest risk with ~50% of the severe cases being fatal in some studies<sup>8, 21</sup>. A combination of clinical presentation, cardiac injury biomarkers and cardiac imaging (echocardiogram or magnetic resonance imaging) are used to diagnose ICI associated myocarditis, but endomyocardial biopsy, if available, is considered the gold standard criterion. In addition to heart failure treatments, immune suppression with corticosteroids is the mainstay of ICI myocarditis management<sup>21</sup>. There is general agreement to permanently withhold ICI treatment if a patient develops moderate to severe cardiotoxicity.

### Chimeric Antigen Receptor T-cell Therapy (CAR-T)

CAR-T cell is another form of anti-cancer immunotherapy in which T cells are genetically modified to express a chimeric protein comprised of the cancer antigen binding receptor and T cell activating domains<sup>23</sup>. This directs both the innate and adaptive immune systems to target, for example, diffuse large B-cell lymphoma that express CD19 antigen using the axicabtagene ciloleucel or tisagenlecleucel CAR-T cells. The cardiac complications observed in patients undergoing CAR-T cell therapy include cardiac troponin elevation, heart failure, cardiogenic shock and arrhythmias which appear to stem from the phenomenon of cytokine release syndrome (CRS)<sup>24</sup>. CRS is characterized by a constellation of inflammatory signs and symptoms with hemodynamic instability that can be treated using tocilizumab, an anti-IL-6 receptor antagonist<sup>23, 25</sup>. Although some specific mechanisms for these side effects have been proposed such as the shared expression of B cell CD19 antigen by vascular mural cells in the brain, the full spectrum of CAR-T cell-induced

toxicities of the heart and other tissues remain to be further elucidated<sup>26</sup>. Unlike some other cancer treatment toxicities, the side effects of CAR-T therapy are actually “on-target” and may be reversible when the specific cancer cells are eliminated. The interaction between CAR-T and other anti-cancer treatments also continue to be clarified. Although prior treatment with anthracyclines was thought to increase the risk of cardiovascular events associated with CAR-T cell therapy, this does not appear to be significant according to a retrospective analysis<sup>25</sup>. On the other hand, CAR-T therapy could be directed towards providing beneficial cardiac effects. An innovative pre-clinical study has proposed the use of CAR-T cells directed against a cardiac fibroblast-specific protein (fibroblast activation protein, FAP) to prevent myocardial fibrosis, a characteristic finding of anthracycline cardiomyopathy<sup>27</sup>.

### Vascular Endothelial Growth Factor (VEGF) Inhibitors

Cancer cells require adequate blood supply for their growth, thus inhibiting the angiogenesis promoting activity of VEGF has been shown to have efficacy in cancer treatment, especially in suppressing metastatic cancer cell growth<sup>28</sup>. The monoclonal antibodies bevacizumab and ramucirumab can inactivate VEGF-A and block the signaling of its receptor VEGFR2, respectively. However, disrupting VEGF signaling can result in adverse cardiovascular effects such as systemic and pulmonary hypertension, cardiac ischemia, thromboembolism, and cardiac dysfunction<sup>29</sup>. Systemic hypertension is the most striking adverse effect of inhibiting VEGF while the least prominent cardiovascular effect is myocardial dysfunction, suggested to be in part secondary to hypertension exacerbation<sup>28, 29</sup>. VEGF receptors can also be blocked by small molecule kinase inhibitors such as sorafenib and sunitinib but their inhibition of other important receptor tyrosine kinases complicate the interpretation of their adverse cardiovascular effects<sup>28, 30</sup>.

### Human Epidermal Growth Factor Receptor 2 (HER2) Inhibitor

The overexpression of human epidermal growth factor receptor 2, associated with aggressive disease, is observed in a significant fraction of breast cancers, and its inhibition using the monoclonal antibody trastuzumab improves the prognosis of patients with HER2-positive breast cancer<sup>5</sup>. However, this therapeutic benefit can be associated with symptomatic heart failure and declines in left ventricular ejection fraction initially reported in 2–3% of patients, usually within weeks of starting the treatment although it generally recovers in the months following therapy interruption<sup>10, 31</sup>. In a more recent clinical trial, about 30% of patients exhibited cardiotoxicity after HER2 inhibitor treatment while those concurrently or serially treated with anthracyclines were at even higher risk<sup>32</sup>. Unlike anthracyclines, HER2 cardiotoxicity does not show a cumulative dose dependent increase in risk<sup>10</sup>. These clinical observations were confirmed in the laboratory by the cardiac specific deletion of *ErbB2*, the mouse homolog of the human *HER2* gene, which resulted in dilated cardiomyopathy and increased susceptibility to anthracycline cardiotoxicity while ErbB2 overexpression was protective<sup>33–35</sup>. These findings are also consistent with the involvement of HER2 in diverse cellular processes related to cell growth, survival, and energy metabolism<sup>10, 36</sup>.

## CARDIOTOXICITY ASSOCIATED WITH ANTHRACYCLINES

### Clinical Considerations of Anthracycline Cardiomyopathy

Although doxorubicin is the most commonly used anthracycline, there are 5 other class members available for the FDA- and non-FDA approved treatment of 14 and 7 different types of cancer, respectively<sup>37</sup>. Annually, an estimated one million patients received anthracyclines in North America<sup>38</sup> so even a relatively low incidence of severe anthracycline cardiotoxicity translates to a large number of affected individuals. The recommended lifetime cumulative dose of doxorubicin is 550 mg/m<sup>2</sup> or less, but the incidence of clinically significant decline (>10%) in left ventricular ejection fraction (LVEF) can be lowered further from ~65% at this dose to ~7% and ~18% at 150 mg/m<sup>2</sup> and 350 mg/m<sup>2</sup>, respectively. This risk continuum is also sensitive to patient factors, such as age (>65 or <18 yr), female gender, race, and pre-existing cardiac conditions<sup>3</sup>.

Acute cardiotoxicity (during or immediately after infusion) of anthracycline can manifest as transient arrhythmias and LVEF depression, but it is rare (<1%) and usually reversible<sup>11</sup>. The timing of the cardiotoxicity has been further divided into early ( < 1 yr) and late (median 7 yr after treatment) cardiotoxicity although the pathogenesis of anthracycline cardiomyopathy is most likely continuous, progressive, and irreversible unless detected early and intervened<sup>11</sup>. It is important to note here that the development of more sensitive imaging modalities such as cardiac magnetic resonance imaging has enabled the earlier detection of subclinical cardiac damage induced by anthracyclines<sup>39, 40</sup>. All 4 chambers of the heart may be dilated but not as severely as in some other forms of dilated cardiomyopathy, and the left ventricular wall may not be thinned significantly<sup>1</sup>. The affected left ventricle displays both systolic and diastolic contractile dysfunction. Other than careful risk stratification and monitoring of patients in order to intervene early with standard heart failure therapies if needed, the only approved preventive medication is dexrazoxane which appears to provide cardioprotective effects as measured by cardiac troponin and echocardiographic parameters<sup>3, 41, 42</sup>. Notably, the protective effects of dexrazoxane appear to be mediated in part by its activities on topoisomerases and mitochondrial iron homeostasis as discussed later in this review<sup>43, 44</sup>.

### Pathogenesis Considerations in Human Anthracycline Cardiomyopathy

Although a number of different molecular mechanisms have been proposed to mediate anthracycline cardiomyopathy, there is general consensus on the histopathologic changes observed upon endomyocardial biopsy of the right ventricle, considered to be the most sensitive and specific diagnostic test for this condition<sup>45</sup>. The biopsy may show patchy interstitial fibrosis, fibroblast proliferation, histiocytic (monocytic origin) infiltration, loss of myofibrils, distension of the sarcoplasmic reticulum, and vacuolated cardiomyocytes next to the areas of fibrosis<sup>1</sup>. These histopathologic features are used to grade the degree of anthracycline-induced damage, but the availability of this diagnostic test is limited to specialized medical centers.

## Relevance of Experimental Anthracycline Cardiotoxicity Models to Human Cardiomyopathy

From a mechanistic perspective, there have been many studies examining anthracycline cardiotoxicity using both *in vitro* and *in vivo* models which contrasts with the relative paucity of experimental data on immunotherapy associated cardiotoxicity albeit it is a newer treatment. There are many interesting *in vitro* studies, but in this review the mechanistic insights derived mostly from *in vivo* mouse models of anthracycline cardiotoxicity are discussed in order to associate them with cardiac function.

It is important to distinguish between the acute cardiotoxicity of doxorubicin, the most commonly used anthracycline in animal studies, and the late onset cardiomyopathy generally observed in patients. In fact, when apoptosis studies in doxorubicin cardiotoxicity were initially being performed, the important distinction between acute and chronic administration of doxorubicin in animal models and concerns about the relevance of supra-clinical doses and disease timing to the human condition were raised by some investigators<sup>46, 47</sup>. It is notable that in the representative anthracycline cardiotoxicity studies listed (Table 2), those that involve cell death mechanisms or early onset cardiac dysfunction tend to utilize an acute model and examine the cardiac response within days (versus weeks in chronic models) of anthracycline treatment (either bolus- or multi-injection dosing). In contrast, doxorubicin is usually given in low doses weekly in the clinics to prevent acute drug toxicity, now a rare complication in patients.

Generally, the acute cardiotoxicity models show cardiac dysfunction shortly after treatment with mortality rates sometimes exceeding 50%<sup>48, 49</sup>. This does not reflect anthracycline cardiomyopathy in the clinics where the heart failure presents only in a subset of patients and often occurs years after treatment exposure. Highlighting the importance of drug administration mode on cardiotoxicity outcome, the dosing of doxorubicin (2 mg/kg/wk over 7 wk) in a chronic cardiotoxicity model permitted the detection of rat heart mitochondrial dysfunction while this was not observed with an acute toxicity bolus dosing regimen (20 mg/kg once)<sup>50</sup>. Thus, in this review we have listed the available information on the dosing protocol of doxorubicin in various mouse models to provide context for interpreting the proposed mechanisms although it should be noted that in some instances both acute and chronic dosing schedules were used within the same study (Table 2).

### Anthracyclines Can Induce DNA Damage via Topoisomerases

Anthracyclines such as doxorubicin can induce DNA strand breaks by interacting with cleavable topoisomerase II $\alpha$  (TOP2A)-DNA complexes, thereby inhibiting the proliferation of cancer cells<sup>51</sup>. Non-proliferating cardiomyocytes express TOP2B rather than TOP2A (Figure 1)<sup>52</sup>, so an elegant study took advantage of this differential expression of topoisomerase II subtypes and showed that doxorubicin cardiotoxicity could be prevented by the cardiac-specific knockout of *TOP2B* gene in mice<sup>53</sup>. TOP2B involvement in mediating anthracycline cardiotoxicity was previously suggested by decreases in doxorubicin-induced DNA damage response upon disruption of *TOP2B* in mouse embryonic fibroblasts or by treatment with the cardio-protective agent dexrazoxane in rat H9C2 cardiomyoblasts<sup>43</sup>.

## Role of Oxidative Stress

In addition to the mechanisms by which it is known to damage DNA, doxorubicin accumulates in mitochondria to induce cytotoxic free radicals (Table 2)<sup>2</sup>. The redox metabolism of anthracyclines by the respiratory complex I NADH dehydrogenase can produce cytotoxic reactive oxygen species (ROS) in purified mitochondria<sup>54</sup>. Paralleling this *in vitro* study, mitochondrial manganese superoxide dismutase (MnSOD) overexpressing transgenic mice were protected from acute doxorubicin cardiotoxicity<sup>55</sup>. Other subcellular sources of ROS could also contribute to pathogenesis. For example, the disruption of the plasma membrane NAD(P)H oxidase gp91 has been shown to be protective against chronic doxorubicin cardiotoxicity in a mouse model<sup>56</sup>. This study also showed that specific polymorphisms of NAD(P)H oxidase subunits conferred differential sensitivity to anthracycline-induced cardiotoxicity in cancer patients.

Doxorubicin has also been suggested to cause myocardial injury through another free radical nitric oxide (NO) generated by inducible nitric oxide synthase (iNOS) enzyme in macrophages, but the role of NO in pathogenesis has been controversial<sup>57, 58</sup>. Another study which also examined acute toxicity within days after a bolus injection (20 mg/kg) of doxorubicin reported that iNOS deficient mice actually had worse cardiac outcome after treatment, suggesting an antioxidant role for NO<sup>57</sup>. Furthermore, the mitochondrial damage caused by doxorubicin in these iNOS deficient mice could be rescued by crossing them into transgenic mice overexpressing the antioxidant enzyme manganese superoxide dismutase (MnSOD)<sup>57</sup>. Another oxidative stress response enzyme NAD(P)H quinone oxidoreductase 1 (NQO1) has been shown to attenuate doxorubicin cardiotoxicity in association with reduced oxidative stress and inflammation via the NAD<sup>+</sup>/SIRT1 pathway<sup>59</sup>. Despite these and other pre-clinical studies showing their benefits in mouse models, the available clinical studies examining the protective effect of antioxidants against anthracycline toxicity in the heart have not been conclusive, except for dexrazoxane, an iron chelator which can mitigate mitochondrial oxidative stress<sup>42, 44, 60, 61</sup>.

## Cell Death and Other Cellular Processes in Anthracycline Cardiotoxicity

Various types of cell death such as apoptosis, necrosis, necroptosis, and ferroptosis with or without the involvement of p53, a well-established mediator of apoptosis<sup>62</sup>, have been reported to contribute to doxorubicin cardiotoxicity (Table 2). The apoptosis studies cited in Table 2 represent only a fraction of many that have been reported by various investigators over the years, but due to space limitations we provide only some representative *in vivo* studies. The apoptosis caused by oxidative DNA damage sensors ATM and p53 has been associated with doxorubicin cardiac injury and dysfunction, however in another study, p53 inhibition of mTOR signaling, not apoptosis, was proposed to be the major contributor of acute cardiotoxicity<sup>63–65</sup>. In certain cellular contexts, p53 can even prevent apoptosis via p21 mediated cell cycle arrest<sup>62</sup>, so perhaps it is not unexpected that the cardiac-specific knockout of p53 showed increased apoptosis with bolus dosing of doxorubicin albeit the homozygous cardiac-specific Cre mouse was embryonic lethal so that the heterozygous state had to be utilized<sup>67</sup>. The survival of cardiomyocytes in acute doxorubicin cardiotoxicity has also been shown to be promoted by tyrosine kinase FAK mediated upregulation of p21<sup>68</sup>. An additional factor to consider is the relative resistance of adult cardiomyocytes

to apoptosis compared with young cardiomyocytes due to decreased expression of the apoptotic machinery, suggesting potentially different mechanisms of anthracycline cardiotoxicity depending on age<sup>69</sup>. Both necrosis (unprogrammed inflammatory cell death) and necroptosis (programmed inflammatory cell death) through BNIP3 and RIP3-CaMKII, respectively, have also been reported to play a role in the single bolus dosing model of acute doxorubicin cardiotoxicity<sup>48, 70</sup>. The importance of these cell death mechanisms in doxorubicin cardiotoxicity have been supported by its prevention in BAX deficient mice as well as by the innovative use of a small molecule inhibitor of BAX<sup>71</sup>.

Consistent with the essential role of iron in oxidative stress homeostasis, doxorubicin cardiotoxicity has been associated with mitochondrial iron overload and cell death by ferroptosis<sup>44, 49, 72</sup>. Reducing mitochondrial iron stores either by promoting its efflux out of the mitochondria or decreasing the pool of iron available for uptake into the mitochondria was cardioprotective while higher cellular iron content in a hemochromatosis mouse model was associated with increased cardiac sensitivity to doxorubicin<sup>73</sup>. Intriguingly, the results of various iron chelator studies in anthracycline cardiotoxicity have indicated that iron binding capacity may not be the sole determinant of their cardioprotective activity although the beneficial effect of dexrazoxane does appear to be mediated mainly by decreasing mitochondrial iron levels<sup>44, 60</sup>. Further studies may help resolve some of these differences observed in model systems.

There is also evidence that autophagy and mitophagy are involved in anthracycline cardiotoxicity. Cytoplasmic p53 has been reported to interact with parkin and impair mitophagy causing cardiac dysfunction in doxorubicin-treated mice, and to inhibit autophagy via mTOR and decrease cell survival<sup>74, 75</sup>. In addition to these inhibitory activities in the cytoplasm through protein-protein interactions, an *in vitro* study showed that p53 overexpression can increase BNIP3 mRNA and protein expression in association with increased autophagy and cell death<sup>76</sup>. It should be noted that p53 can also induce apoptosis via its nuclear transactivation of autophagy regulator gene *DRAM*<sup>77</sup>. Further linking autophagy to doxorubicin cardiotoxicity<sup>78</sup>, another group used Beclin1 mouse models to show that doxorubicin blocks autophagic flux, resulting in the accumulation of autolysosomes which can increase ROS and cause cardiac damage<sup>79</sup>.

Inflammation has been implicated in anthracycline cardiomyopathy<sup>80</sup>, and this is evidenced by studies, for example, showing the attenuation of cardiac dysfunction with the inhibition of either cyclooxygenase 2 (COX-2) or toll-like receptor 2 (TLR 2)<sup>81, 82</sup>. Although the antibody-mediated neutralization of TLR2 ameliorated inflammation, fibrosis, and cardiac dysfunction caused by doxorubicin treatment, neutralizing TLR4 worsened these parameters and suppressed autophagy indicating the complementary and distinct roles of TLRs in pathogenesis<sup>82</sup>. Related to inflammation, aberrant coagulation caused by the activation of the G-protein coupled protease-activated receptor 1 (PAR1) has also been shown to contribute to doxorubicin cardiotoxicity<sup>83</sup>. Additionally, the inhibition of matrix metalloproteinase inhibitors have been shown to ameliorate cardiac dysfunction and prevent myocardial fibrosis in a mouse model with chronic weekly administration of doxorubicin<sup>84</sup>. These representative *in vivo* mouse model studies of anthracycline cardiotoxicity, although



not comprehensive, show mechanistic complexity and the need for further delineation as translation into the clinics is considered.

### **A Central Role for the Mitochondrion in Anthracycline Cardiomyopathy**

Many different lines of investigations have led to the conclusion that mitochondrial structure and function are altered in anthracycline cardiomyopathy as well detailed in a comprehensive review<sup>85</sup>. As a positively charge hydrophobic molecule at physiologic pH, doxorubicin is known to interact with the negatively charged inner mitochondrial membrane lipid cardiolipin, potentially disrupting cytochrome c and other essential components of the respiratory chain<sup>86–89</sup>. Early studies in rat models showed that there was cumulative and irreversible cardiac mitochondrial dysfunction including abnormalities of calcium homeostasis that may affect cell death signaling upon chronic treatment with doxorubicin<sup>90, 91</sup>. Subsequent studies have continued to support a central role of the mitochondria in anthracycline cardiotoxicity ranging from the finding of preventing BNIP3 mediated necrosis by mitochondrially localized SIRT3 to a study demonstrating impaired mitochondria function/content, calcium handling, and antioxidant activity in induced pluripotent stem cell-derived cardiomyocytes of patients with doxorubicin cardiotoxicity<sup>70, 92, 93</sup>.

Besides the various mechanisms of damage to the myocardium depending on doxorubicin dosing, the manifestation of slowly progressive heart failure well after the chemotherapeutic agent has been eliminated from the body suggested a more persistent alteration of the mitochondrion in anthracycline cardiotoxicity<sup>85</sup>. Changes to the cardiac mitochondrial genomic DNA (mtDNA) could explain such a permanent alteration, and indeed an early investigation reported that the impaired synthesis of mtDNA was slower to recover after doxorubicin treatment compared with that of nuclear DNA<sup>94</sup>. Notably, the higher sensitivity of the heart than the liver to doxorubicin damage could be due to its slower recovery of mtDNA synthesis after being impaired by treatment compared with that of the liver. Furthermore, mtDNA of heart has a faster turnover rate than that of liver ( $T_{1/2}$  ~7 d versus ~9 d, respectively; nuclear DNA  $T_{1/2}$  >30 d), suggesting greater susceptibility to DNA damaging events<sup>94</sup>.

A study using chronic low dose doxorubicin treatment of mice reported increased mtDNA deletions which could be prevented by the antioxidant coenzyme Q10, invoking oxidative stress as the cause of damage<sup>95</sup>. In other studies, acute doxorubicin treatment of rats (single injection of 15 mg/kg) resulted in higher levels of oxidative mtDNA modifications (8-hydroxydeoxyguanosine) compared with nuclear DNA, again with cardiac tissue showed greater mtDNA modifications than liver<sup>96</sup>. Sub-chronic doxorubicin treatment (2 mg/kg/wk over 6 wk) caused the greatest cumulative damage in cardiac mtDNA that persisted even after the treatment was complete, suggesting the temporal and genetic bases of anthracycline cardiomyopathy in cancer patients<sup>97</sup>. Strong evidence for the role of mtDNA in anthracycline cardiomyopathy was provided by the finding of increased mtDNA deletions, decreased mtDNA content, and decreased mtDNA-encoded, but not nuclear DNA-encoded, respiratory subunits in autopsy cardiac tissues of patients treated with doxorubicin<sup>98</sup>. Similar reductions in mtDNA content and mtDNA-encoded

respiratory subunit gene expression were reproduced in a rat model of chronic doxorubicin cardiomyopathy by these investigators<sup>99</sup>.

Mechanistically, a number of different genes have been linked to changes in mtDNA homeostasis after doxorubicin exposure. In addition to increased defects in cardiac cell myofibrils and mitochondrial ultrastructure by electron microscopy, mice that are deficient in mitochondrial topoisomerase I (TOP1MT) were observed to have greater loss of mtDNA content and respiratory complex subunit proteins upon doxorubicin exposure<sup>100</sup>. As mentioned earlier, doxorubicin cardiotoxicity was proposed to be mediated by TOP2B induced nuclear DNA strand breaks with resultant defective mitochondrial biogenesis<sup>53</sup>, but TOP2B is also known to be the only type II topoisomerase in the mitochondria and thus may affect mtDNA homeostasis as shown for TOP1MT<sup>100, 101</sup>. In fact, it has been suggested that TOP2B in the mitochondria may be the primary mediator of doxorubicin cardiotoxicity, because the gene program regulated by TOP2B in the nucleus is not typically related to mitochondrial biogenesis<sup>101, 102</sup>. Other genes that are well-established to promote mitochondrial biogenesis and activities such as SIRT1 and SIRT3 have been demonstrated to attenuate doxorubicin cardiotoxicity, further supporting the central role of the mitochondrion in its pathogenesis<sup>59, 103</sup>.

### **p53 as Guardian of mtDNA in Anthracycline Cardiomyopathy**

p53 is best known as a mediator of cell death for cancer prevention, but it has a dual nature and is equally important for promoting cell survival through its DNA repair, cell cycle regulatory, antioxidant, metabolic, and other activities depending on factors such as the cellular context and level of p53 induction<sup>104, 105</sup>. In fact, there are evidence that the translocation of p53 into the mitochondria under physiological conditions may serve to maintain the integrity of the mitochondrial genome<sup>106</sup>. An early study reported increased p53 immunoreactivity in cardiomyocyte mitochondria after doxorubicin treatment, and increased oxidative mtDNA damage was observed in *p53*<sup>-/-</sup> (null) mouse hearts (Table 2)<sup>107</sup>. A subsequent insightful study using mice with cardiac overexpression of a dominant-interfering mutant p53 showed protection from acute doxorubicin cardiotoxicity, presumably due to loss of wild-type p53 cell death activity, but the cardiac function of these mice deteriorated in the chronic phase indicating a protective role for p53<sup>108</sup>. The blunting of doxorubicin-induced STAT3 by mutant p53 was associated with increased expression of DNA damage markers, portending the worse late cardiac outcome.

We examined chronic doxorubicin cardiotoxicity in p53 null mice compared with mice that had knockin of the p53 R172H mutation, the mouse homolog of the human hotspot R175H mutation that causes the inherited early onset cancer disorder Li-Fraumeni syndrome<sup>109</sup>. This mutation had previously been shown to retain the mitochondrial biogenesis activity of wild-type p53 while losing its other activities such as apoptosis and cell cycle arrest<sup>110, 111</sup>. The mitochondrial activities of p53 could be mediated by its regulation of mitochondrial biogenesis genes such as *TFAM* and *p53R2* in the nucleus or by its interactions with other proteins such as POLG in the mitochondria<sup>106</sup>. As in wild-type p53 mice, the hearts of p53 R172H mutant knockin mice were relatively protected from loss of cardiac and mitochondrial function in contrast to that of p53 null mice after doxorubicin

treatment (Table 2)<sup>109</sup>. Furthermore, the selective decrease in the expression of mtDNA-encoded versus nuclear DNA-encoded mitochondrial respiratory subunit genes in p53 null hearts upon doxorubicin exposure was consistent with the previous reports of mtDNA depletion<sup>85, 107, 109</sup>. Taken together, the finding of p53-regulated mtDNA homeostasis playing a critical role in preventing anthracycline-induced cardiotoxicity correlates well the clinical observation of mtDNA damage observed in the heart samples of patients with anthracycline cardiomyopathy<sup>98</sup>. Thus, it could be interesting to examine whether the molecular components involved in regulating mtDNA in easily accessible patient samples such as blood provide diagnostic or prognostic information for the management of anthracycline treated patients.

## CONCLUSIONS

The various cardiotoxicities associated with different types of cancer treatments are increasing with the advancement of cancer therapies. With the improved survival of cancer patients from their primary disease, a new specialty of cardio-oncology has emerged for managing the cardiovascular complications of anti-cancer therapies<sup>112</sup>. While it is important to recognize and understand the molecular bases of the cardiotoxicities associated with the newer oncologic treatments, the cardiomyopathy caused by one of the oldest classes of chemotherapeutic agents, the anthracyclines, still does not have a widely accepted and targeted prevention treatment despite the effort of many investigators<sup>4, 40, 112</sup>.

Because of the different mechanisms by which anthracyclines can cause cardiotoxicities, one challenge is focusing on the ones that are most relevant to human cardiomyopathy pathogenesis. One criterion may be the dosing of the chemotherapy in the *in vivo* model system which is summarized in Table 2 for some representative studies. As a DNA damaging agent that concentrates in the mitochondria, various studies from different laboratories point to damaging of the mitochondrial genome as a major contributor<sup>85</sup>. Because newer anti-cancer treatments utilizing immune checkpoint inhibition are likely to be used in combination with anthracyclines, interaction or augmentation of their specific cardiotoxicity mechanisms could occur with potentially worse outcome, making it imperative to understand pathogenesis. Besides bioenergetic deficiency secondary to mitochondrial dysfunction, the intracellular release of mtDNA damaged by doxorubicin may activate innate immune signaling<sup>113</sup>, known to be stimulated by immunotherapy<sup>114</sup>, and synergistically increase myocardial fibrosis in the setting of ICI therapy<sup>115, 116</sup>. Although based on an anecdotal case report, it is also tempting to consider underlying subclinical mitochondrial genomic DNA disease as a susceptibility factor for anthracycline cardiotoxicity<sup>117</sup>. In parallel, the development of a doxorubicin analog that uncouples its cardiotoxic DNA damaging effect from its cancer cell destroying chromatin modification activity would be of great benefit and timely needed for the care of cancer patients<sup>118</sup>.

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### Nonstandard Abbreviations and Acronyms

<b>CAR-T</b>	chimeric antigen receptor T-cell
<b>CRS</b>	cytokine release syndrome
<b>HER2</b>	human epidermal growth factor receptor 2
<b>ICI</b>	immune checkpoint inhibitors
<b>mtDNA</b>	mitochondrial DNA
<b>nDNA</b>	nuclear DNA
<b>PD-1</b>	programmed cell death protein-1
<b>PD-L1</b>	PD-1 ligand
<b>VEGF</b>	vascular endothelial growth factor

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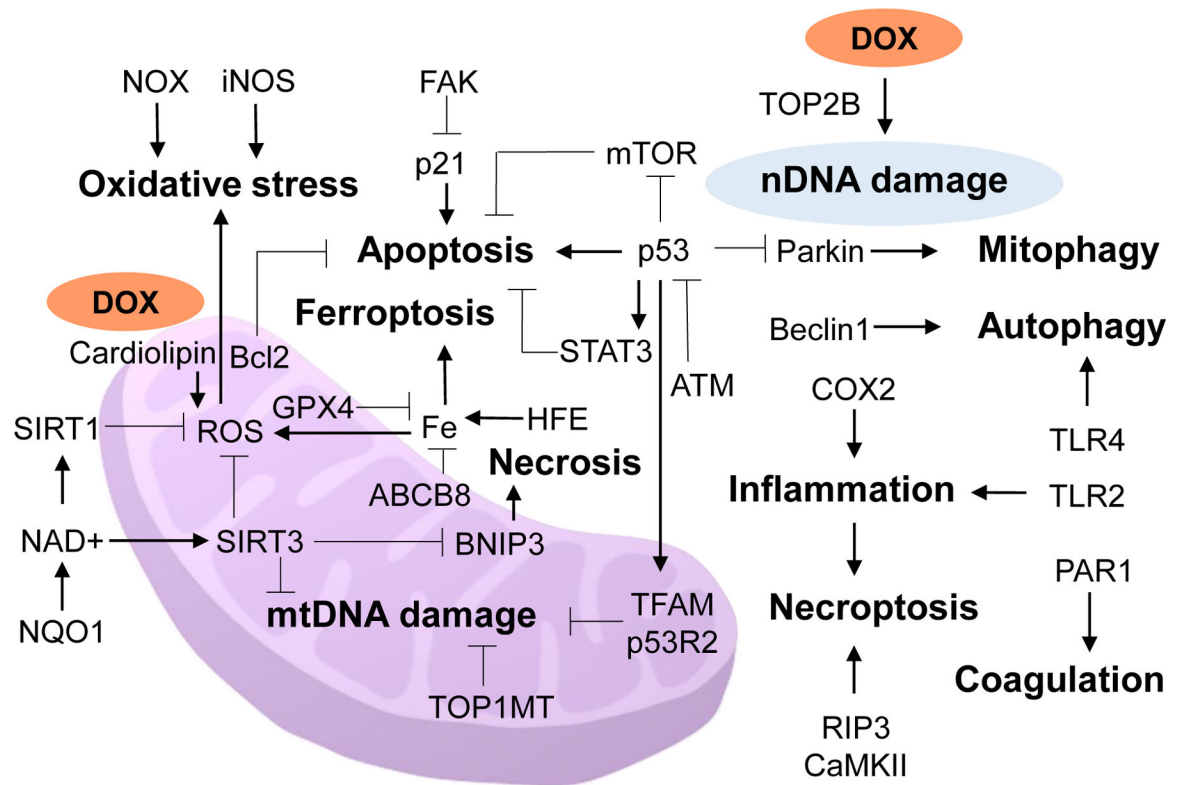
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### Highlights

- With recent advances in oncology and improved survival of cancer patients, the cardiotoxicities of new treatments such as immunotherapy are becoming more evident and require further understanding for their management and prevention.
- The cardiotoxicity of anthracyclines, one of the oldest but most effective class of chemotherapeutic agents, is the major limiting factor in its use, and various molecular mechanisms of this side effect have been proposed using experimental models.
- Determining the mechanisms most relevant to human anthracycline cardiomyopathy pathogenesis remains a challenge, but one criterion may be the dosing protocol of the chemotherapy in the *in vivo* model.
- Accumulation evidence points to damage of the cardiomyocyte mitochondrial genome as a major determinant of anthracycline cardiomyopathy, and its protection during treatment or better risk stratification based on a firm molecular understanding of pathogenesis may be key to minimizing this side effect while optimizing treatment of the cancer.



**Figure 1.**

Schematic diagram of the major cellular processes (bold print) and their mediators/pathways of anthracycline cardiotoxicity. ABCB8, ATP binding cassette subfamily B member 8; ATM, ataxia telangiectasia mutated kinase; Bcl2, B-cell lymphoma 2; BNIP3, Bcl2 interacting protein 3; CaMKII, calcium/calmodulin-dependent protein kinase II; COX-2, cyclooxygenase 2; DOX, doxorubicin; FAK, focal adhesion kinase; GPX4, glutathione peroxidase 4; HFE, human homeostatic iron regulator protein; iNOS, inducible nitric oxide synthase; mtDNA, mitochondrial DNA; mTOR, mammalian target of rapamycin; NOX, NAD(P)H oxidase; NQO1, NAD(P)H quinone oxidoreductase 1; nDNA, nuclear DNA; PAR1, protease-activated receptor 1; p53R2, p53-inducible ribonucleotide reductase; RIP3, receptor-interacting protein 3; SIRT, sirtuin; STAT3, signal transducer and activator of transcription 3; TFAM, transcription factor A, mitochondrial; TLR, toll-like receptor; TOP1MT, topoisomerase I mitochondrial; TOP2B, topoisomerase 2-beta.

**Table 1.**

Cardiotoxicity mechanisms of some anti-cancer immunotherapies and targeted treatments

Therapeutic agent	Molecular target	Proposed cardiotoxicity mechanism	References
<b>Immune Checkpoint Inhibitors</b>			
Lipilimumab	CTLA-4	Activation of CD4+ and CD8+ T cells with marked myocardial infiltration in mice	8, 14, 119–122
Pembrolizumab	PD-1	Activated T cells in myocardium; cardiac troponin I autoantibodies and dilated cardiomyopathy observed in PD-1-deficient mice	
Nivolumab			
Cemiplimab			
Atezolizumab	PD-L1	Activated T cells in myocardium	
Avelumab			
Durvalumab			
<b>CAR-T cells</b>			
Axicabtagene ciloleucel	CD19	Cytokine release syndrome; IL-6 implicated as key mediator	24, 25
Tisagenlecleucel			
<b>HER2 inhibitors</b>			
Trastuzumab	HER2	Inhibition of HER2 signaling pathway involved in cell growth, survival, angiogenesis and migration	33–36
Pertuzumab			
<b>VEGF inhibitors</b>			
Bevacizumab	VEGF	VEGF inhibition alters blood vessel homeostasis and nitric oxide signaling which can both affect blood pressure and thrombosis	28, 29
Ramucirumab	VEGF receptor		

Cardiotoxicity mechanisms of some anti-cancer immunotherapies and targeted treatments are summarized and referenced. CAR-T, chimeric antigen receptor T-cell; CTLA-4, cytotoxic T lymphocyte-associated protein-4; HER2, human epidermal growth factor receptor 2; PD-1, programmed cell death protein-1; PD-L1, PD-1 ligand; VEGF, vascular endothelial growth factor.

**Table 2.**

Mechanisms of doxorubicin-induced cardiotoxicity in mouse models

Mechanisms	Pathway/Genes	Mouse genotypes	Dosage	Ref
DNA damage	TOP2B	Cardiac specific- <i>TOP2B</i> <sup>-/-</sup>	Acute: 25 mg/kg once Chronic: 5 mg/kg/wk for 5 wk	53
Oxidative stress	MnSOD	MnSOD Tg	10, 20 or 25 mg/kg once	55
	NOX subunit	<i>gp91</i> <sup>-/-</sup>	3 mg/kg/wk for 3 wk then once at wk 5	56
	iNOS	<i>iNOS</i> <sup>-/-</sup> ; MnSOD Tg	20 mg/kg once	57, 58
	NQO1 SIRT1	<i>NQO1</i> <sup>-/-</sup>	4 mg/kg/d × 3 d	59
Apoptosis	p53	<i>p53</i> <sup>-/-</sup>	20 mg/kg once	64
	p53 ATM Bcl2	<i>p53</i> <sup>+/-</sup> ; Bcl2 Tg	6 mg/kg/wk for 4 wk	65
	p53	Cardiac specific- <i>p53</i> <sup>-/-</sup>	20 mg/kg once	67
	FAK p21	Muscle specific- <i>FAK</i> <sup>-/-</sup> ; cardiac specific- superFAK Tg	20 mg/kg once	68
	BAX	<i>BAX</i> <sup>-/-</sup>	Acute: 20 mg/kg once Chronic: 3 mg/kg × 8 over 2 wk	71
Necrosis	BNIP3	<i>BNIP3</i> <sup>-/-</sup>	20 mg/kg once	70
Necroptosis	RIP3 CaMKII	<i>RIP3</i> <sup>-/-</sup>	Acute: 20 mg/kg once Chronic: 5 mg/kg/wk for 4 wk	48
Ferroptosis	HFE	<i>HFE</i> <sup>-/-</sup>	Acute: 20 mg/kg once Chronic: 5 mg/kg × 6 at 2 wk intervals	73
	ABCB8	Cardiac specific- <i>ABCB8</i> <sup>-/-</sup> ; cardiac specific-ABCB8 Tg	10 mg/kg × 3 over 5 d; 6 mg/kg × 4 over 10 d	44
	NRF2 HO1	<i>NRF2</i> <sup>-/-</sup>	20 mg/kg once	49
	GPX4	<i>GPX4</i> <sup>+/-</sup> ; GPX4 Tg	6 mg/kg × 3 at 2 d intervals	72
Autophagy	p53 mTOR	Cardiac specific-CB7 (dominant interfering p53) Tg; cardiac specific-mTOR Tg	10 mg/kg × 2 at 3 d interval	66
	p53 Parkin	<i>p53</i> <sup>-/-</sup> ; <i>Parkin</i> <sup>-/-</sup> ; Parkin Tg	2.5 mg/kg × 5 over 2 wk	75
	Beclin1	<i>Beclin1</i> <sup>+/-</sup> ; cardiac specific-Beclin1 Tg	5 mg/kg/wk for 4 wk	79
Inflammation, Coagulation	COX-2	Mice treated with COX-2 inhibitor	4 mg/kg/wk for 6 wk	81
	TLR2 TLR4	Mice treated with TLR2 or TLR4 neutralizing antibody	Acute: 10 mg/kg once Chronic: 3.5 mg/kg/wk for 8 wk	82
	PAR1	<i>PAR1</i> <sup>-/-</sup>	Acute: 20 mg/kg once Chronic: 5 mg/kg/wk for 5 wk	83
Mitochondrial damage	p53 mtDNA	<i>p53</i> <sup>-/-</sup>	20 mg/kg once	107
	TOP1MT	<i>TOP1MT</i> <sup>-/-</sup>	4 mg/kg/wk for 8 wk	100
	p53 STAT3	Cardiac specific-CB7 (dominant interfering p53); cardiac specific- <i>STAT3</i> <sup>-/-</sup>	5 mg/kg/wk for 5 wk	108
	SIRT3 OGG1	<i>SIRT3</i> <sup>-/-</sup> ; SIRT3 Tg	5 mg/kg every 15 d × 3	103

Mechanisms	Pathway/Genes	Mouse genotypes	Dosage	Ref
	p53 TFAM p53R2	<i>p53</i> <sup>-/-</sup> ; <i>p53</i> R172H knockin	5 mg/kg/wk for 5 wk	109

The genes/pathways involved in mediating anthracycline cardiotoxicity, the mouse model genotypes, and the corresponding references are shown. The dose and schedule of doxorubicin treatment used in each of the studies are also shown to better assess the chronicity of the cardiotoxicity model. ABCB8, ATP binding cassette subfamily B member 8; ATM, ataxia telangiectasia mutated kinase; BAX, Bcl2 associated x protein; Bcl2, B-cell lymphoma 2; BNIP3, Bcl2 interacting protein 3; CaMKII, calcium/calmodulin-dependent protein kinase II; COX-2, cyclooxygenase 2; FAK, focal adhesion kinase; GPX4, glutathione peroxidase 4; HFE, homeostatic iron regulator protein; HO1, heme oxygenase 1; iNOS, inducible nitric oxide synthase; MnSOD, manganese superoxide dismutase; mTOR, mammalian target of rapamycin; NOX, NAD(P)H oxidase; NRF2, nuclear factor erythroid 2-related factor 2; NQO1, NAD(P)H quinone oxidoreductase 1; OGG1, 8-oxoguanine DNA glycosylase; PAR1, protease-activated receptor 1; p53R2, p53-inducible ribonucleotide reductase; RIP3, receptor-interacting protein 3; SIRT, sirtuin; STAT3, signal transducer and activator of transcription 3; TFAM, transcription factor A, mitochondrial; TLR, toll-like receptor; TOP1MT, topoisomerase I mitochondrial; TOP2B, topoisomerase II $\beta$ .