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Prognostic Factors for Cardiotoxicity among Children with Cancer: Definition, Causes and Diagnosis with Omics' Technologies'

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Abstract: Improvements in the treatment of childhood cancer have considerably enhanced survival rates over the last decades, to over 80% as per today. However, this great achievement has been accompanied by the occurrence of several early and long-term treatment-related complications major of which is the cardiotoxicity. The chemotherapeutic agents that have been implicated as a cause of cardiotoxicity, are mostly anthracyclines and mitoxantrone, alkylating agents, proteasome inhibitors and to a lesser extent anti-microtubule agents, cisplatin, monoclonal antibodies, small molecular tyrosine kinase inhibitors, and nucleotide synthesis inhibitors. Routine diagnosis and monitoring of cardiotoxicity rely on electrocardiography and echocardiography. For the early detection of cardiotoxicity, in recent years, major studies have been conducted using biomarkers such as troponin, N-terminal pro b-natriuretic peptide, etc. Despite refinements in diagnostics still severe limitations exist, due to the increase of the above-mentioned biomarkers, only after significant cardiac damage has occurred. Lately, research has been expanded by introducing new technologies and finding new markers by omics approach. These new markers could be used not only for early detection, but also for early prevention of cardiotoxicity. The omics science, which includes genomics, transcriptomics, proteomics, and metabolomics, offers new opportunities for biomarker discovery in cardiotoxicity and may provide understanding of mechanisms of cardiotoxicity, beyond traditional technologies. This article reviews the contemporary definition of cardiotoxicity, older and newer chemotherapeutic agents mainly involved in cardiotoxicity and the methods of early and preventive diagnosis, using omics technology.

Keywords: cardiotoxicity; childhood cancer; chemotherapeutics agents; biomarkers; omics technology

Introduction

With the induction of new chemotherapeutic agents over the years, 5-year survival rate for childhood malignancies exceeds 80%. Caring for these patients includes not only early survival but also later outcomes. Chronic health conditions and health-related quality of life are noted among them as many long-term treatment related complications have resulted in increasing morbidity and mortality rates. Cardiotoxicity represents the most serious non-hematological toxicity of chemotherapeutic drugs. It is noted that childhood cancer survivors have an 8x higher risk in mortality due to cardiovascular disease and 6x increased risk of congestive heart failure. Most importantly early cardiotoxicity may affect the design of chemotherapy and the omission of radiotherapy resulting in incomplete cancer treatment and consequently inferior outcome. The study of cardiotoxicity among the pediatric population is, as expected, of particular importance due to the long-life expectancy. [1-6]

Cardiotoxicity

Definition

As a term, cardiotoxicity was first described in 1946 as the damage on the heart caused from local anesthetics, mercurial diuretics, and digitalis. Later in the 1970s, the term broadened to encompass cardiac complications related with anthracyclines (doxorubicin and daunorubicin), combination therapies like doxorubicin and radiation and drugs like for 5-fluorouracil. Presently there is an increased research interest both basic and clinical aiming in detecting and managing cardiotoxicity as early as possible.

The definition of cardiotoxicity has a great significance for patient's management. Based on International Cardio-Oncology Society (IC-OS) the cardiovascular complications of chemotherapy can be separated into the following clinical entities and/or categories: i) cardiac dysfunction: cardiomyopathy/ heart failure (HF), ii) vascular toxicity, iii) myocarditis, iv) arterial hypertension and v) arrhythmias and QT prolongation. [7,8]

The most preponderant diagnosis of cardiotoxicity is based on the changes found in left ventricular (LV) systolic function measured by left ventricular ejection fraction (LVEF). Different organizations have defined cardiotoxicity in several ways using different threshold changes in LVEF [8]. The need to harmonize all these definitions satisfied by the International Cardio-Oncology Society (IC-OS) and supported by 2022 ESC Guidelines. [7,8]

Cardiotoxicity can be categorized according to the time of presentation as acute, early onset and late onset. Cardiotoxicity could be reversible if addressed while in its early stages, [9] Acute (<1%) toxicity could occur either after administering a single dose, or after a course of chemotherapeutic agents, as long as the onset of clinical manifestations is within the first two weeks following the end of the administration. If presented within the first year of treatment, it is characterized as Early onset (1-18%). Late or chronic onset is manifested years or even decades following the treatment. [9] The percentage of late onset cardiotoxicity varies in literature mainly due to the different definitions used, the detection methods of cardiotoxicity, the population monitored, and also due to the study design. It seems that over 50% of pediatric cancer survivors show subclinical decline in myocardial function and over 16% show symptoms of clinical HF, especially those who have been exposed to anthracyclines. [9]

Abnormalities of ventricular repolarization and electrocardiographic QT-interval alterations, supraventricular and ventricular arrhythmias, acute coronary syndromes, and pericarditis and/or myocarditis-like syndromes are the hallmarks of acute or early onset cardiotoxicity. [9] In contrast, asymptomatic systolic and/or diastolic LVD, which can result in dilated cardiomyopathy, is the most typical indicator of chronic cardiotoxicity. [9,10] Clinical and sub-clinical cardiovascular damage, coronary artery disease, and cerebrovascular events are other conditions linked to treatment-related complications. Survivors had an almost six-fold higher risk of heart failure, a five-fold higher risk of myocardial infarction, a six-fold higher risk of pericardial disease, and an almost five-fold higher risk of valvular abnormalities compared to their siblings. [11-13]

Chemotherapeutic drugs

Anthracyclines

Anthracyclines, primarily doxorubicin but also daunomycin, epirubicin and idarubicin, are some of the commonest used agents for both hematologic and solid tumors. Acute cardiotoxicity due to anthracyclines may present as hypotension, tachycardia, arrhythmia, transient depression of left ventricular function, myocarditis, pericarditis, and acute coronary syndrome. The late onset cardiotoxicity caused by high cumulative dose of anthracyclines, mainly includes signs and symptoms of cardiomyopathy and chronic heart failure. [9]

Mitoxantrone is an anthracenedione, or anthracycline analogue and has similar anthracycline mechanisms of action. Mitoxantrone might cause a wide variety of heart conditions, such as disturbances of cardiac rhythm, chronic heart failure, and persistent diastolic dysfunction in the absence of an impairment of left ventricular ejection fraction. [10]

The prevalent concept of how anthracycline action may cause heart damage, involves the production of oxygen radicals, which in turn damage the DNA, proteins, and lipids and lead to cellular dysfunction and myocyte death. [14,15-16]

Cardiolipins are abundantly found on the inner mitochondrial cell membrane. By having an increased affinity for anthracyclines, they in turn allow for their increased cell entry. Upon cell entry by passive diffusion, they can reach much higher intracellular concentrations compared to extracellular compartments. Within the cell, they form complexes by binding to iron thus producing free radicals and reactive oxygen species, which in turn cause cell damage and death. By peroxidizing lipids of the cell membrane, those elements may also damage the cell membrane. As cardiomyocytes, contain an abundance of mitochondria, they are more susceptible to anthracycline damage because of the depletion of glutathione peroxidase (antioxidant). [15]

Other mechanisms of cardiotoxicity include alterations to gene expression and nitric oxide synthase activity, which lead to reduced creatine kinase activity and function in mitochondria and ultimately cell death. [15] After exposure to anthracyclines, many of these subcellular sequelae continue to develop for weeks, shedding light on the mechanisms of chronic cardiomyopathy.[14]

Another identified mechanism of doxorubicin-mediated cardiotoxicity is changes in topoisomerase-II (Top2). Topoisomerase II (TOP2) is a molecule that anthracyclines bind to and inhibit, preventing the growth of tumors. DNA's phosphate backbone is broken, twisted, and then resealed by topoisomerases, allowing the double helix's tension to be changed during transcription and replication. Anthracyclines intercalate into DNA, forming complexes with TOP2 that halted the enzyme's activity and trigger a DNA-damage reaction that results in cell death. [14,15,17].

The mechanisms of mitoxantrone associated cardiotoxicity remain yet to be completely understood, the formation of reactive oxygen species in myocardial cells is thought to lead to tissue damage through interactions with cellular iron metabolism, [10].

Nucleotide synthesis inhibitors

The clinical presentation of methotrexate and Fluorouracil (5-FU)-induced cardiotoxicity includes myocardial ischemia, cardiogenic shock, heart failure and cardiomyopathy [10]. Coronary spasm is the most frequently reported mechanism of 5-FU-induced cardiotoxicity [10]. Data derived from animal models indicate that these chemotherapeutic agents induce oxidative stress and subsequent apoptosis of cardiomyocytes and endothelial cells. [10,17]

Alkylating agents

Adjuvant DNA alkylating agents, such as cyclophosphamide (CP) and ifosfamide (IFO), suspend DNA synthesis in cancer cells. These two agents are similar in structure and engender a similar pattern of cardiotoxic effects causing acute heart failure, hemorrhagic myopericarditis and arrhythmia. [10,18] CP- and IFO-induced acute cardiotoxicity is attributed mainly due to a raise in free oxygen radicals and the lower antioxidant defense mechanism in the myocardium. A recent study by Sayed-Ahmed MM et al demonstrated that CP- and IFO-induced cardiotoxicity is due to the inhibition of long-chain fatty acid oxidation via repression of carnitine palmitoyl transferase I and fatty acid binding protein. [19]

Tyrosine kinase inhibitors

Dasatinib, imatinib, lapatinib, sorafenib, nilotinib, and sunitinib are examples of small molecule tyrosine kinase inhibitors (TKIs) that suppress cancer cell proliferation and induce apoptosis of cancer cells. Imatinib, dasatinib, and nilotinib are the three FDA-approved TKIs for use as first-line chronic myeloid leukemia therapy in pediatrics. Also, sorafenib is used in young adults. The pathophysiological mechanism of TKI-induced cardiotoxicity is mitochondrial impairment and cardiomyocyte apoptosis. [10, 18, 20] Each of the above drugs is associated with a different type of cardiotoxicity. For example, dasatinib is more often associated with pleural effusion and less with hypertension, HF, pericardial effusion, and pulmonary hypertension. Nilotinib is associated with

peripheral artery disease, hypertension and prolonged QTc. In contrast, imatinib is related with less cardiotoxicity than the others TKIs.[7]

Anti- microtubule agents

Anti- microtubule agents, including docetaxel, paclitaxel and vinca alkaloids, prevent the polymerization or depolymerization of microtubules. [17] The clinical feature of the cardiotoxicity induced by anti-microtubule agents is mostly ischemia, and arrhythmia. Among all anti-microtubule agents in clinical use, paclitaxel induces the release of histamine, which in turn, activates specific cardiac receptors raising the myocardium's oxygen need, leading to coronary vasoconstriction. Also, Zhang et al reported that the frequency of spontaneous calcium concentration in cardiomyocytes was significantly increased after paclitaxel treatment. This finding could be of great significance as fluctuations of blood calcium levels are linked to arrhythmogenesis. [10,21]

Cisplatin

Cisplatin is an efficacious chemotherapeutic drug with a strong antitumor effect against a wide range of neoplasms. However, the drug's acute and cumulative cardiotoxicity, including electrocardiograph (ECG) abnormalities, angina and acute myocardial infarction, hypertension and hypotension, arrhythmias, myocarditis, cardiomyopathy, and congestive heart failure, is a significant factor that restricts cisplatin treatment. Cisplatin cardiotoxicity can be caused by reactive oxygen species generation, which leads to the creation of oxidative stress and the transition to a prothrombotic state, or by a direct toxic effect on cardiac myocytes.[10]

Monoclonal antibodies

Monoclonal antibodies, including bevacizumab (Avastin) and trastuzumab not used in children, inhibit angiogenesis. Bevacizumab blocks vascular endothelial growth factor (VEGF), while trastuzumab inhibits human epidermal growth factor receptor 2 (HER2) in cancer cells. Bevacizumab causes mostly hypertension, congestive heart failure, thromboembolic events of artery and vein through the mechanic of oxidate stress-induced by the cardiomyocyte apoptosis. Monoclonal antibodies are not widely used in children with malignancy, so we will not discuss them further. [10,18, 20]

Proteasome inhibitors

A new therapeutic option for the treatment of Acute Lymphoblastic Leukemia (ALL) includes proteasome inhibitors. [18, 20] Bortezomib and carfilzomib are two newly prescribed drugs with the potential to cause cardiac dysfunction. [22] Comparatively to carfilzomib (up to 25%), bortezomib has a lower incidence of heart failure (up to 4%). The pathogenesis of proteasome inhibitor cardiotoxicity is not currently well understood. Exposure to proteasome inhibitors in a prenatal mouse model has shown that they can induce oxidative stress leading to myocardial dysfunction. Carfilzomib is also known to induce renal toxicity and microangiopathy, as a consequence of endothelial dysfunction. Combining these studies reveals a complicated mechanism of cardiotoxicity linked to proteasome inhibitors, including alterations to the heart's muscle and vasculature that may be more severe with carfilzomib than bortezomib due to the irreversible nature of proteasome inhibition of carfilzomib. [23-24]

Risk factors

A significant number of factors contribute to the onset of cardiotoxicity. Several types of chemotherapeutic drugs may cause cardiotoxicity, as referred above. These drugs act on cancer cells through a variety of mechanisms and promote cardiotoxicity with distinctive clinical symptoms and underlying mechanisms. (Table 1) [7, 10, 16]

Table 1. Chemotherapeutic drugs and cardiovascular toxicity.(Adjusted by Rochette et al, Trends Pharmacol Sci. 2015 Jun;36(6):326-48 [112]).

| Medicine/Cardiotoxicity | Incidence (%) | Arrhythmia | Myocardial ischemia | Vascular toxicity | Heart failure | QT prolongation | Arterial Hypertension |
|-----------------------------------|---------------|------------|---------------------|-------------------|---------------|-----------------|-----------------------|
| Anthracyclines | | | | | | | |
| Doxorubicin | 3-26 | xxx | X | NE | Xxx | NE | X |
| Doxorubicin Liposomal | 2 | x | xx | NE | x | NE | X |
| Epirubicin | 0,9-3,3 | x | X | NE | X | NE | X |
| Daunorubicin | | xx | X | NE | X | NE | X |
| Idarubicin | 5-18 | xxx | X | NE | xx | NE | X |
| Antibiotics | | | | | | | |
| Mitoxantrone | 0,2-30 | xxx | xx | NE | Xx | NE | Xx |
| Mitomycin-c | 10 | Xx | xx | NE | Xx | | NE |
| Monoclonal antibody | | | | | | | |
| Trastuzumab | 1,7-8 | Xx | X | Xx | Xxx | NE | Xx |
| Bevacizumab | 1,6-4 | Xx | xx | xxx | xx | NE | Xx |
| Pertuzumab | 0,7-1,2 | | X | X | Xx | NE | x |
| dinutuximab beta | | NE | xx | NE | Xx | NE | Xx |
| Rituximab | | | X | xx | Xxx | X | NE |
| Tyrosine kinase inhibitors | | | | | | | |
| Dasatinib | 2-4 | xxx | xx | Xx | Xx | xx | Xx |
| Nilotinib | 1 | xx | NE | x | Xx | xx | Xxx |
| Vermurafenib | | xx | xx | Xx | x | NE | Xx |
| Sorafenib | 2-28 | | X | xx | Xx | NE | xx |
| Sunitinib | 2,7-15 | | X | xx | Xxx | x | Xxx |
| Erlotinib | 7-11 | NE | xx | Xx | NE | NE | NE |
| Lapatinib | 0,2-1,5 | NE | xx | X | NE | xxx | NE |
| Pazopanib | 7-11 | NE | xx | Xx | X | NE | Xxx |
| Imatinib | 0,2-2,7 | NE | xxx | Xx | Xx | NE | NE |
| Proteasome inhibitors | | | | | | | |
| Bortezomib | 2-5 | X | X | X | X | NE | X |
| Carfilzomib | 11-25 | Xx | xx | NE | X | NE | X |
| Antimetabolite | | | | | | | |
| 5-fluorouracil | 2-20 | xxx | xxx | NE | X | NE | NE |
| Capecitabine | | xxx | xxx | Xx | NE | NE | NE |
| Clofarabine | 27 | | | | | NE | |
| Alkylating agents | | | | | | | |
| Cyclophosphamide | 7-28 | NE | NE | X | NE | NE | NE |
| Ifosfamide | 0,5-17 | NE | NE | X | Xx | NE | NE |
| Cisplatin | rare | NE | NE | Xx | NE | NE | NE |
| Antimicrotubule agent | | | | | | | |
| Paclitaxel | <1 | xx | X | NE | X | NE | x |

| | | | | | | | |
|---------------------------|--------|----|----|----|----|----|----|
| nab-paclitaxel | | xx | NE | X | NE | NE | X |
| Docetaxel | 2,3-13 | xx | xx | NE | X | NE | Xx |
| Alkaloids of vinca | | | | | | | |
| Vincristine | 25 | xx | X | NE | NE | xx | X |
| Vinblastine | | NE | X | NE | NE | NE | X |
| Vindesin | | NE | NE | NE | NE | NE | NE |
| Vinorelbin | | NE | X | NE | NE | NE | NE |

The overall dose and mode of administration of each chemotherapeutic agent play an aggravating role in causing cardiac damage. For example, anthracycline induced cardiotoxicity is known to be both cumulative and dose-related, indicating that each administered dose induces sequential or additional damage. [18,25] The cumulative total anthracycline dose is the most important risk factor for cardiac dysfunction. [26]. In retrospective research, Von Hoff et al. [27] observed that when a patient receives a combined doxorubicin dose of 400, 550, and 700 mg/m², the incidence of cardiotoxicity is 3, 7, and 18%, respectively, with dose-limiting toxicity. Another study in adolescents found that even at dosages of 180–240 mg/m², 30% of participants experienced subclinical episodes 13 years after therapy [29]. These results imply that there is no anthracycline dose that is considered safe. Reduced cardiac function has been correlated with dosages as low as 100 mg/m². [28, 30-31].

In addition, female gender, age (under <5 years old), clinical condition of the patient (extent of disease, infection), genetic background, pre-existing cardiac disease, and combinations of cardiotoxic drugs, play an important role in causing cardiac damage [32] (Table 2). The pediatric population is more homogeneous, as a study population, since there are no confounding cardiovascular risk factors (diabetes, smoking, arterial hypertension). [7, 14, 33]

Table 2. Risk Factors.

| Risk factors related in child | Risk factors related in therapy |
|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <ul style="list-style-type: none"> • Female sex • Age < 5 years • genetic background • pre-existing cardiac disease • Cardiovascular risk factors (Diabetes, obesity, hyperlipidemia, Hypertension) | <ul style="list-style-type: none"> • Anthracycline > 250 mg/m² equivalent doxorubicin • Cumulative dose • Irradiation • combination of cardiotoxic drugs |

The risk of developing cardiotoxicity is also increased by concurrent radiation exposure to the chest. In addition to the myocardium, radiation therapy has the potential for damaging the pericardium, heart vessels, and conductive tissue. [18]

Carriers of certain genetic mutations are also more susceptible to cardiotoxicity. [9,18] Our understanding of genetic susceptibility to anthracyclines-related cardiotoxicity has been influenced by a sizable body of research, as we will describe below.

Imaging

Diagnostic approaches for chemotherapy-induced cardiotoxicity include electrocardiography and echocardiography, used as methods of monitoring cardiac function before, during and after treatment.

Electrocardiography (ECG) can be used to identify any early signs of cardiac toxicity, such as resting tachycardia, ST-T wave abnormalities, conduction disturbances, QT interval prolongation, or arrhythmias. However, these ECG findings could be induced by several factors unrelated to cardiotoxic treatment. These ECG abnormalities may be reversed and are not always related to the development of chronic cardiomyopathy. [7,26 ,33]

The two-dimensional Echocardiography (2D) is the most used imaging technique to monitor cardiac function. It is non-invasive, cheap, readily available and does not expose the patient to further radiation. However, standard echocardiographic parameters like LVEF may lack sensitivity for the detection of systolic dysfunction. [33]

Considering the poor sensitivity of 2D LVEF measurement, the use of global systolic longitudinal myocardial strain (GLS) analysis has become an area of interest. [33-34] A pathogenic percentage reduction of GLS greater than 15% from baseline is regarded as a sign of early LV dysfunction. When possible, it is preferable to use these sophisticated echocardiographic measures as the foundation for clinical decision.

Other methods in the monitoring of these patients are cardiac magnetic resonance imaging (CMR), nuclear cardiac imaging (MUGA) and myocardial perfusion imaging (MPI). There are 2 techniques for MPI: single photon emission computed tomography (SPECT) and position emission tomography (PET). All these methods take considerably longer than a follow-up echocardiography and might not be as accessible at all pediatric facilities. [33].

Biomarkers

Several biomarkers have been assessed for their efficacy in early prediction of patients' risk of cardiotoxicity and identification of cardiac dysfunction. The World Health Organization defines biomarkers as any element, structure, or process that can be detected in the body (or its byproducts) and affects or forecasts the development or course of a disease.

In accordance with the literature, troponin and natriuretic peptide are the most studied biomarkers for the detection of both early cardiotoxicity and its later follow-up. Lipshultz et al showed that the elevation of cardiac troponin T and N-terminal pro-brain natriuretic peptide (NT-pro-BNP) in children with acute lymphoblastic leukemia have been associated to notably reduced left ventricular (LV) mass, abnormal LV end-diastolic posterior wall thickness, and abnormal LV thickness-to-dimension ratios, all of which suggested LV remodeling, respectively, 4 years later. [35] However, further research hasn't pointed out an association between acute or chronic troponin release and left ventricular dysfunction, but in contrary found association with NT-pro-BNP in childhood cancer survivors. [36-39]

We should be especially careful in evaluating troponin and natriuretic peptide values in children < 1 year of age due to their normally elevated values at these ages. [40]

Other biomarkers investigated include inflammation markers, such as C-reactive protein (CRP) and growth/differentiation factor 15 (GDF-15) [37], oxidative stress markers such as myeloperoxidase, vascular remodeling markers such as placental growth factor and soluble Fms-like tyrosine kinase receptor 3, and fibrosis markers. (Galectin 3). [35,41-49] More Those conventional biomarkers usually show significant changes only after heart damage occurs.

To determine the proper use of these biomarkers in clinical practice, new prospective and multicenter studies with large populations, well-standardized dosing methodologies, well-defined time of sampling, and cardiologic end points are required.

Omics

In the last decades, new research and clinical studies have attempted to identify possible biomarkers of early cardiac damage by chemotherapeutic agents using omics technology. So, the omics science offers new opportunities for biomarker discovery in cardiotoxicity and may provide understanding of cardiotoxicity beyond traditional technologies. Omics technology includes genomics, transcriptomics, proteomics, and metabolomics.

Genomics

Cumulative anthracycline dose and other related risk factors seem not to be the exclusive risk factor responsible for the significant individual variation in the incidence and severity of heart failure in pediatric cancer survivors. Several studies have revealed how important host genetic

polymorphisms could lead to differential risk of cardiotoxicity among cancer survivors with otherwise identical clinical and treatment-related risk factors by using genome wide association or candidate gene approaches. [50-53] This explains why some patients experience cardiotoxicity while other patients can tolerate high doses of chemotherapy without heart damage.

The cumulative dose of anthracyclines and other risk variables do not appear to be the only ones contributing to the considerable individual variability in the occurrence and severity of heart failure in pediatric cancer survivors

Genomic polymorphisms are small changes in a specific part of the DNA chain. One or more polymorphisms can determine a range of patient characteristics such as their ability to metabolize and eliminate genotoxic substances. Cancer treatment-related cardiovascular toxicity risk may be influenced by genetic variation. Significant efforts using targeted and whole genome correlation studies have been made to reveal the pharmacogenomic causes of this predisposition. [50,68-62]

At least 45 SNPs located in 34 genes have been associated with anthracycline induced cardiotoxicity. Many of these associations require further investigation through replication and/or functional and mechanistic studies to make sure we confirm and better understand the roles of these associated variant in anthracycline-related cardiotoxicity (ACT).[9]

Polymorphisms in solute carrier transporter (SLC) genes are associated with ACT. One of the functions of the SLC family is acting as drug transporters for anthracyclines and thus provides biological support for these genetic associations. Research on childhood cancer survivors has discovered correlations between ACT risk and protective variants in SLC such as SLC28A3, SLC22A17, SCL22A7. These findings were successfully replicated [54-59]. Also, different studies reported protective variants in SLC10A2 and SLC22A1 [55]. SLC22A6 was first mentioned in the context of ACT by Sagi et al in patients treated for childhood ALL [57].

Retinoic acid receptor gamma (RARG) has been involved in cardiac development and remodeling through the repression of Top2b [60]. A recent genome-wide association study, Aminkeng [51] et al, uncovered a non-synonymous variant rs2229774 in RARG that was significant associated with ACT in survivors of childhood cancer. Specifically, rs2229774-carriers had a significantly increased risk of developing ACT as compared to non-carriers. [51]

Studies have also revealed an elevated risk brought on by a variation in the UGT1A6 gene, a member of the glucuronosyl transferase family. Through the glucuronidation path, the UGT1A6 play a significant role in the detoxification of drugs, including the metabolites of anthracyclines. [51,54,58]

Polymorphisms in adenosine triphosphate-binding cassette transporter (ABC) genes are related with cardiotoxicity in childhood patient cancers treated with anthracyclines. The ABC genes seems to play a role as efflux transporters of drugs including anthracyclines, so may have important effects in the myocardium. Eight variants in five genes (ABCB1, ABCB4, ABCC1, ABCC2, ABCC5) were associated with cardiotoxicity, especially with reduced ejection fraction. [57, 61-64]

Other studies investigated polymorphisms in carbonyl reductase genes, which have been associated with dose dependent increase in cardiomyopathy risk. Carbonyl reductase (CBR) will reduce anthracyclines to cardiotoxic alcohol metabolites. As Blanco et al has showed, among childhood cancer survivors, homozygosity for G allele in CBR3 leads to increased cardiomyopathy risk associated with low- to moderate-dose anthracyclines. Patients homozygous with the CBR3 V244M G allele have no safe cut off minimum dose. [65-66]

A recent study showed a gene environment interaction between single-nucleotide polymorphism on the CELF4 gene and higher dose of anthracyclines. [67] CELF4 (CUGBP Elav-Like Family Member 4) protein is responsible for pre-mRNA alternative splicing of TNNT2, the gene that encodes for cardiac troponin T.

Aminkeng [58] et al gathered the evidence-based clinical practice recommendations for pharmacogenomic testing and emphasized that the RARG gene rs2229774, UGT1A6 * 4 rs1786378, SLC28A3 rs7853758 have the potential to further discriminate patients at high and lower risk of ACT. A pharmacogenetic test for these genetic variations in RARG, SLC28A3, and UGT1A6 has been released at the British Columbia Children's Hospital since the publication of these guidelines. Based on genetic and clinical risk variables, tested patients were divided into several risk groups, and

therapy adjustments were chosen in accordance with this risk. Early evidence indicates that the British Columbia Children's Hospital's pharmacogenetic testing was effective in lowering the incidence of ACT in children, which should inspire additional clinics to utilize this pharmacogenetic test.

These findings might help develop prediction models that will spot patients who will be particularly susceptible to ACT and who need their therapy modified or who need closer monitoring. Further independent research may make it possible to identify people before treatment with a genetic predisposition to cardiovascular toxicity and for whom more thorough screening or perhaps preventive measures should be implemented. Replication analyses, however, occasionally failed to support the initial findings. Numerous factors, including the variability of cohorts, ambiguities in the definition of ACT, variations in procedures, and the type or dosage of the chemotherapeutic drugs used, may contribute to this. To increase the diagnostic and prognostic role in predicting ACT, more research is required.

Transcriptomics

Another interesting area is the integration of microRNAs in the early detection of cardiotoxicity. Recently, the potential use of circulating MicroRNAs (miRNAs) has been studied as possible specific biomarkers and therapeutic targets of cardiac disease. [73-79]

MicroRNAs are small endogenous non-coding RNAs of 21–24 nucleotides, acting as post-transcriptional gene regulators by inhibiting and/or degrading target messenger RNAs (mRNAs). Bioinformatics data suggest that each miRNA molecule can control hundreds of gene targets, thus indicating the potential effect of miRNAs on virtually any genetic pathway. MiRNAs play a significant role in different biological processes including proliferation, differentiation, development, and cell death. Furthermore, several miRNAs are involved in regulating heart development from embryonic to adult stage and their dysregulation leads to various heart diseases such as, arrhythmias, essential hypertension, heart failure, cardiomyopathy, cardiac hypertrophy, and atherosclerosis. [80-81]

Cardiotoxic effect of chemotherapeutic agents may lead to specific miRNAs with changed expression. These could be used to investigate the toxicity of potential drug candidates on cardiomyocytes and cell lines originating from the heart in a preclinical in vitro setting. The potential use of circulating miRNAs in plasma as indicators of drug-induced cardiotoxicity has undergone much research during the last several years.[80]

Nearly 30 circulating miRNAs have had their levels altered, both increasing and decreasing, and these changes have been linked to HF and associated pathologies. MiRNAs including miR-1, miR-133, miR-208a/b, miR-499, miR-29, and miR-34, which are substantially expressed in the myocardium compared to other tissues, are the ones that are primarily being researched [73]. In addition, a variety of harmful substances alter the miRNA profile in both plasma and cardiac tissue. Even at low toxin concentrations, where other tissue damage biomarkers are not discernible, alterations in miRNAs can be measured. [80] Most studies use data from experimental animals, while those utilizing clinical patient samples are limited.

MiR-1 is a skeletal muscle specific miRNA that has an important role in cardiac development, function, and disease. Abnormal miR-1 levels are associated with acute myocardial infarction, heart failure, arrhythmias, ventricular dysfunction, cardiac hypertrophy, and myocyte hyperplasia. [82]. MiR-499 and miR-208 are associated with acute myocardial infarction and HF. [82] Circulating levels of miR-133a have been associated with increased risk of cardiovascular diseases. Increased levels of miR-133a have been detected in patients with acute myocardial infarction earlier than cardiac troponin T increase. [83] MiR-133 are two miRNAs, named miR-133a and miR-133b, highly expressed in human heart and seems to be involved in heart development and myocyte differentiation.

The analysis of circulating miRNAs in breast cancer patients receiving doxorubicin (DOX) identified miR-1 as a potential candidate for the early detection of DOX-induced cardiotoxicity. [84] Leger et al investigated other possible markers of cardiotoxicity in children and young adults treated with anthracycline chemotherapy (AC). Candidate plasma profiling of 24 miRNAs was performed in

33 children before and after a cycle of AC or non-cardiotoxicity chemotherapy. MiR-1, miR-29b and miR-499 were reported to be upregulated in pediatric patients following acute initiation of AC.[85-86]. Monitoring the plasma levels of miR-208a and miR-208b showed an elevation in patients with myocardial damage and were even detected earlier than cardiac troponins [87]. This is concordance with the findings from other studies. [73-74,87-88] Table 3 provides a summary of the major miRNAs linked to drug-induced cardiotoxicity.

Table 3. Summary of major miRNAs link to drug-induced cardiotoxicity in people.

| MiRNA | Drug | Modulation | Species | System | References |
|-------------------------|----------------|------------|--------------------------|--------|-----------------------------------------------------|
| miR-1 | doxorubicin | increase | female patients | plasma | Rigaud et al, Oncotarget 2017 |
| miR-1, miR-29b, miR-499 | anthracyclines | increase | children and young adult | plasma | Leger et al, J Am Heart Assoc. 2017 |
| miR1254 | bevacizumab | increase | Humans | plasma | Zhao et al, Tumour Biol. 2014 |
| miR29 miR499 | doxorubicin | increase | Children | plasma | Oatmen et al, Am J Physiol Heart Circ Physiol, 2018 |
| miR208 | doxorubicin | nothing | female patients | plasma | Calvalho et al, J Appl Toxicol 2015 |

In addition to anthracyclines, other cytotoxic agents have shown cardiotoxic effects and biomarkers of their pathomechanism have been searched for, including miRNAs. Patients with bevacizumab-induced cardiotoxicity when compared with controls were found to have increased levels of five miRNAs. In the validation experiments, two of these (miR-1254 and miR-579) shown valuable specificity. MiR-1254 exhibited the strongest correlation with the clinical diagnosis of bevacizumab-induced cardiotoxicity [89].

With regards to a number of features of drug-induced cardiotoxicity, miRNAs appear to be a promising agent. A potentially successful method for preventing severe problems is the identification of patients with subclinical cardiotoxicity through the detection of cardio-specific miRNAs circulating in plasma that are not present under normal circumstances. A supposedly efficient method for identifying people with subclinical cardiotoxicity is the detection of cardio-specific miRNAs circulating in plasma that are not present under normal circumstances. [80]. Many other research studies should focus on how the miRNAs profile changes when interacting with drugs with proven cardiotoxicity.

Proteomics

The proteomic data available to date on chemotherapy-induced cardiac toxicity are limited, mainly involving anthracyclines, and related to experimental animal studies. [90]

Proteomics is the study of proteins, which are essential components of organisms and have a variety of functions. The proteome consists of all the proteins expressed by a cell, tissue, or organism. Proteomics could give us important information for a number of biological problems.

Ohyama et al identified cellular processes in mouse heart tissue from control rats and rats affected by different Adriamycin and docetaxel dosing protocols using a toxicoproteomic approach. They identified 9 different proteins that were expressed in the control and in the two treatment

groups, and were involved in energy production pathways, such as glycolysis, the Krebs cycle and the mitochondrial electron transport chain. [91]

Kumar et al in 2011 used a rat model of Doxorubicin-induced cardiotoxicity to show the differential regulation of several key proteins, including protein S that are stress responsive (ATP synthase, enolase alpha, alpha B-crystallin, translocation protein 1 and stress-induced phosphoprotein 1), and apoptotic/cell damage markers (p38 alpha, lipocortin, voltage-dependent anion-selective channel protein 2, creatine kinase and MTUS1). [86]

More recently, Desai et al pinpointed possible biomarkers of early cardiotoxicity in plasma from male B6C3F1 mice that have received weekly intravenous dose of 3 mg/kg doxorubicin (DOX) or saline (SAL) for 2, 3, 4, 6, or 8 weeks (corresponding to cumulative doses of 6, 9, 12, 18, or 24 mg/kg DOX). They suggested the neurogenic locus notch homolog protein 1 (NOTCH1) and von Willebrand factor (vWF) as early biomarkers of DOX cardiotoxicity, to address the clinically significant question of identifying cancer patients at risk for cardiotoxicity. [92]

Finally, Yarana et al using a mouse model of DOX-induced cardiac injury, quantified serum extracellular vehicles (EVs), assayed proteomes, counted oxidized protein levels in serum EVs generating following DOX treatment and examined the alteration of EV content. The release of EVs containing brain/heart glycogen phosphorylase (PYGB) before the increase in cardiac troponin in the blood following DOX therapy suggests that PYGB is an early indicator of cardiac damage, according to the proteomic profiling of DOX_EVs. [93]

To find out if these pathways could result in the discovery of early markers of cardiotoxicity, more research in that area is required.

Further studies are needed in that field to investigate if these pathways could result in the discovery of early markers of cardiotoxicity.

Metabolomics

Metabolomics is the upcoming new science with the potential to further increase our knowledge on cancer biology and the search for prognostic biomarkers. Up to now most studies use metabolomic data from experimental animals, while those utilizing clinical patient samples are extremely limited.

Metabolism is more directly related to the phenotype and physiology of a biological system. Metabolomics is the study of all cellular metabolites (hydrocarbons, amino acids, sugars, fatty acids, organic acids, steroids, peptides). It encompasses all levels of cellular regulation, that is, the regulation that occurs at the level of transcription, translation, and post-translational modifications, hence, they can closely reflect the phenotype of an organism at specific time. The human metabolome is thought to be composed of about 3,000 endogenous metabolites at current estimates (Human metabolome project). But the exact size of human metabolome is still debatable. It is also believed that nutritional compounds, xenobiotics, and microbial metabolites must be considered when defining the human metabolome [94]. Therefore, metabolome analysis can be a useful tool used to find diagnostic markers that will help us examine unknown pathological conditions, effectively.

Different analytical techniques can be used in measurement of the metabolites. Such methods are nuclear magnetic resonance (NMR) spectrometry, molecular mass spectrometry (MS), gas chromatography (GC), high performance liquid chromatography (LC) and tricarboxylic acid (TCA). The most common and more high throughput technologies are nuclear magnetic resonance (NMR) spectrometry and molecular mass spectrometry (MS).

Mass spectrometry is an analytical platform for metabolomic analysis. It is a highly sensitive, reproducible, and versatile method as it identifies molecules and their fragments by measuring their masses. This information is obtained by measuring the mass-to-charge ratio (m/z) of ions that are produced by inducing the loss or gain of a charge from a neutral species. The sample, which is comprised of up of a complicated mixture of metabolites, can be introduced into the mass spectrometer either directly or preceded by a separation approach (using liquid chromatography or gas chromatography). [95]

NMR spectroscopy utilizes magnetic properties of nuclei to determine the number and type of chemical entities in a molecule. Proton NMR spectroscopy can detect soluble proton-containing

molecules with a molecular weight of approximately 20 kD or less. The NMR spectra serve as the raw material for pattern recognition analyses, which simplifies the complex multivariate data into 2 or 3 dimensions that can be readily understood and evaluated. Both NMR and liquid chromatography-mass spectrometry (LC-MS) systems can be integrated to in vivo tissues or to biological fluids such as serum, plasma, urine, etc., obtained from humans. The advantages of NMR are that it requires relatively little sample preparation, it is non-destructive, and can provide useful information regarding the exact structure of metabolites. However, NMR sensitivity is related to the magnet's strength, while available instrumentation can unambiguously detect only the most abundant metabolites in plasma. On the other hand, the most important advantage of mass spectrometry coupled with upfront chromatography is of far greater sensitivity than NMR MS-based systems that have been used to resolve compounds in the nanomole to picomole and even femtomole range, whereas identification of compounds by ¹H-NMR requires concentrations of 1 nanomole or higher. [96-97]

The main methodologies that are used for metabolomic analysis are untargeted and targeted metabolomics. Untargeted metabolomics allow measuring a wider variety of metabolites present in an extracted sample without prior knowledge of the metabolome. The main advantage is that it provides with an unbiased way to examine the relationship between interconnected metabolites from multiple pathways. By contrast, targeted metabolomic analyses measure the concentrations of predefined set of metabolites and provides higher sensitivity and selectivity than untargeted metabolomics.

An overview of the main metabolomics associated with drug-induced cardiotoxicity detected in plasma/stem cell/heart in mice or people is given in Table 4. The role of carnitine in detection of cardiotoxicity was confirmed by a successive study in which Armenian et al compared a metabolomics analysis in 150 symptom-free childhood cancer survivors that received anthracycline treatment. Thirty-five participants were found to have cardiac dysfunction without symptoms. So, they compared the two groups (participants with cardiac dysfunction and with normal systolic function) and discovered 15 metabolites differentially expressed among patients. After adjusting for multiple comparisons, individuals with cardiotoxicity had significantly lower plasma carnitine levels in comparison with those with normal cardiac function. [98]

Table 4. Metabolomics associated with drugs-induced cardiotoxicity.

| Metabolite | Plasma | Stem cell | Heart | Mice | People | XRT | Medicine | Dose | Biomarker | References |
|--------------------------|--------|-----------|-------|------|--------|-----|-------------------------------------------------|-------------------------------------|-----------|--------------------------------|
| proline | ↓ // ↑ | | ↑ | Yes | no | | cyclophosphamide | 200mg/kg | | Li et al, J Proteome Res, 2015 |
| LPC 20:3 | ↓ | | | Yes | no | | cyclophosphamide | 200mg/kg | | Li et al, J Proteome Res, 2015 |
| linoleic acid | ↓ | | | Yes | no | | cyclophosphamide | 200mg/kg | | Li et al, J Proteome Res, 2015 |
| l-carnitine | ↑ // ↑ | | | Yes | no | | cyclophosphamide/doxo/isopterone/5-fluorouracil | 200mg/kg//20mg/kg//5mg/kg//125mg/kg | | Li et al, J Proteome Res, 2015 |
| 19-hydroxycorticosterone | ↑ // ↓ | | | Yes | no | | cyclophosphamide/doxo/isopterone/5-fluorouracil | 200mg/kg//20mg/kg//5mg/kg//125mg/kg | | Li et al, J Proteome Res, 2015 |

| | | | | | | | |
|-----------------|--------|-----|-----|---------------------------------------------|-------------------------------------|-----------------|---------------------------------------------------------------------|
| phytophingosine | ↓ | Yes | no | cyclophosphamide | 200mg/kg | | Li et al, J Proteome Res, 2015 |
| cholid acid | ↓ | Yes | no | cyclophosphamide | 200mg/kg | | Li et al, J Proteome Res, 2015 |
| LPC 14:0 | ↓ // ↓ | Yes | no | cyclophosphamide/doxorubicin/5-fluorouracil | 200mg/kg//20mg/kg//5mg/kg//125mg/kg | | Li et al, J Proteome Res, 2015 |
| LPC 18:3 | ↓ | Yes | no | cyclophosphamide | 200mg/kg | | Li et al, J Proteome Res, 2015 |
| LPC 16:1 | ↓ | Yes | no | cyclophosphamide | 200mg/kg | | Li et al, J Proteome Res, 2015 |
| LPE 18:2 | ↓ | Yes | no | cyclophosphamide | 200mg/kg | | Li et al, J Proteome Res, 2015 |
| LPC 22:5 | ↓ | Yes | no | cyclophosphamide | 200mg/kg | | Li et al, J Proteome Res, 2015 |
| LPC 22:6 | ↓ | Yes | no | cyclophosphamide | 200mg/kg | | Li et al, J Proteome Res, 2015 |
| LPC 22:4 | ↓ | Yes | no | cyclophosphamide | 200mg/kg | | Li et al, J Proteome Res, 2015 |
| LPC 20:2 | ↓ // ↓ | Yes | no | cyclophosphamide/doxorubicin/5-fluorouracil | 200mg/kg//20mg/kg//5mg/kg//125mg/kg | | Li et al, J Proteome Res, 2015 |
| PLE 20:3 | ↓ | Yes | no | cyclophosphamide | 200mg/kg | | Li et al, J Proteome Res, 2015 |
| pyruvate | ↑ | | | Doxorubicin | 20mg/kg | troponine T LDH | Andreadou et al, NMR Biomed, 2009 /Chauhari et al, Amino Acids 2017 |
| acetate | ↑ | ↑ | Yes | doxorubicin | 20mg/kg | troponine T LDH | Andreadou et al, NMR Biomed, 2009 /Chauhari et al, Amino Acids 2017 |
| formate | ↑ | | | Doxorubicin | 20mg/kg | troponine T LDH | Andreadou et al, NMR Biomed, 2009 |

| | | | | | | | | |
|------------|--------|--------|-----|------------|---------------------------------|---------|-----------------------|------------------------------------------------------------------------------------|
| | | | | | | | | /Chauhari et al, Amino Acids 2017 |
| succinate | ↑ | ↑ | Yes | | Doxorubicin | 20mg/kg | troponine T LDH | Andreadou et al, NMR Biomed, 2009 /Chauhari et al, Amino Acids 2017 |
| lactate | ↑ // ↑ | ↓ | Yes | | Doxorubicin | 20mg/kg | troponine T | Andreadou et al, NMR Biomed, 2009 |
| alanine | ↑ // ↑ | ↑ // ↑ | Yes | | Doxorubicin | 20mg/kg | troponine T | Andreadou et al, NMR Biomed, 2009 |
| glutamine | ↑ | ↓ | Yes | | Doxorubicin | 20mg/kg | troponine T | Andreadou et al, NMR Biomed, 2009 |
| glutamate | ↑ | no | Yes | ↑ | Doxorubicin | 20mg/kg | troponine T | Andreadou et al, NMR Biomed, 2009 |
| creatine | | no | Yes | | Doxorubicin | 20mg/kg | troponine T | Andreadou et al, NMR Biomed, 2009 |
| taurine | | no | Yes | ↓ | Doxorubicin | 20mg/kg | troponine T | Andreadou et al, NMR Biomed, 2009 |
| valine | ↑ | ↓ | Yes | ↑ | Doxorubicin | 20mg/kg | troponine T | Andreadou et al, NMR Biomed, 2009 |
| leuline | ↑ | ↓ | Yes | | Doxorubicin | 20mg/kg | troponine T | Andreadou et al, NMR Biomed, 2009 |
| isoleukine | ↑ | ↓ | Yes | ↑ | Doxorubicin | 20mg/kg | troponine T | Andreadou et al, NMR Biomed, 2009 |
| carnitine | ↓ // ↑ | ↓ | Yes | yes | anthracyclines//do xorubicin | | troponine T | Armenian et al, Cancer Epidemiol Biomarkers Prev. 2014 |
| threitol | ↓ | | | yes | anthracyclines | | | Armenian et al, Cancer Epidemiol Biomarkers Prev. 2014 |
| mannose | ↓ | | | yes | anthracyclines | | | Armenian et al, Cancer Epidemiol Biomarkers Prev. 2014 |

| | | | | |
|----------------------------------|---|-----|----------------|--------------------------------------------------------|
| pyroglutamine | ↓ | yes | anthracyclines | Armenian et al, Cancer Epidemiol Biomarkers Prev. 2014 |
| n-acetylalanine | ↓ | yes | anthracyclines | Armenian et al, Cancer Epidemiol Biomarkers Prev. 2014 |
| creatine | ↓ | yes | anthracyclines | Armenian et al, Cancer Epidemiol Biomarkers Prev. 2014 |
| eicosenoate | ↓ | yes | anthracyclines | Armenian et al, Cancer Epidemiol Biomarkers Prev. 2014 |
| stearidonate | ↓ | yes | anthracyclines | Armenian et al, Cancer Epidemiol Biomarkers Prev. 2014 |
| arachidonate | ↓ | yes | anthracyclines | Armenian et al, Cancer Epidemiol Biomarkers Prev. 2014 |
| dihomo-linoleate | ↓ | yes | anthracyclines | Armenian et al, Cancer Epidemiol Biomarkers Prev. 2014 |
| l-stearoylglycerophosphoinositol | ↓ | yes | anthracyclines | Armenian et al, Cancer Epidemiol Biomarkers Prev. 2014 |
| dehydroisoandrosterone sulfate | ↓ | yes | anthracyclines | Armenian et al, Cancer Epidemiol Biomarkers Prev. 2014 |
| pregnen-diol; disulfate | ↓ | yes | anthracyclines | Armenian et al, Cancer Epidemiol Biomarkers Prev. 2014 |
| pregn steroid monosulfate | ↓ | yes | anthracyclines | Armenian et al, Cancer |

| | | | | | | | |
|-----------------|----|----|-----|---|-------------|----------------|------------------------------------------------|
| | | | | | | | Epidemiol Biomarkers Prev. 2014 |
| arginine | ↑ | ↑ | Yes | | Doxorubicin | | Schnackenberg et al, Appl. Toxicol. 2016 |
| asparagine | ↑ | ↑ | Yes | | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| citrulline | ↑ | ↑ | Yes | | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| glycine | ↑ | ↑ | Yes | ↑ | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| histidine | ↑ | ↑ | Yes | | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| lysine | ↑ | ↑ | Yes | | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| methionine | ↑ | ↑ | Yes | | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| ornithine | ↑ | ↑ | Yes | | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| phenylalanine | ↑ | ↑ | Yes | | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| serine | ↑ | ↑ | Yes | | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| threonine | ↑ | ↑ | Yes | ↑ | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| trptophan | ↑ | ↑ | Yes | | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| tyrosine | ↑ | ↑ | Yes | | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| acetylornithine | ↑ | ↓ | Yes | | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| hydroxproline | ↑ | no | Yes | | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| citrate | no | no | Yes | | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |

| | | | | | | |
|----------------------------------------|----|----|-----|-------------|-------------|------------------------------------------|
| propionylcarnitine | ↑ | no | Yes | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| serotonine | no | ↑ | Yes | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| putrescine | no | ↑ | Yes | Doxorubicin | troponine T | Schnackenberg et al, Appl. Toxicol. 2016 |
| malate | ↑ | ↑ | Yes | Doxorubicin | | Tan et al, PLoS One 2011 |
| fructose | | ↑ | Yes | Doxorubicin | | Tan et al, PLoS One 2011 |
| glycose | | ↑ | Yes | Doxorubicin | | Tan et al, PLoS One 2011 |
| cholesterol | | ↑ | Yes | Doxorubicin | | Tan et al, PLoS One 2011 |
| alanine | | ↑ | Yes | Doxorubicin | | Tan et al, PLoS One 2011 |
| glutamine | | | Yes | ↓ | Doxorubicin | Tan et al, PLoS One 2011 |
| docosaehaenoic acid | | ↓ | Yes | Sunitinib | | Jencen et al, Metabolites. 2017 |
| arachidonic acid/eicosapentaenoic acid | | ↓ | Yes | Sunitinib | | Jencen et al, Metabolites. 2017 |
| 6-hydroxynicotinic acid | | ↓ | Yes | Sunitinib | | Jencen et al, Metabolites. 2017 |
| o-phosphocolamine | | ↓ | Yes | Sunitinib | | Jencen et al, Metabolites. 2017 |
| ethanolamine | ↑ | | Yes | Sunitinib | | Jencen et al, Metabolites. 2017 |
| xenobiotics | | | | | | |

More recently, Li et al [99] identified 39 biomarkers for detecting cardiotoxicity earlier than biochemical analysis and histopathological assessment. They used rats to create cardiotoxicity models in which the toxicity was caused by doxorubicin, isoproterenol, and 5-fluorouracil. The metabolomics analysis of plasma was performed by using ultraperformance liquid chromatography quadrupole time-of-flight mass spectrometry. They used a support vector machine (SVM) trying to deploy a predictive model to confirm more exclusive biomarkers with more important l-carnitine, 19-hydroxydeoxycorticosterone, lysophosphatidylcholine (LPC) (14:0) and LPC (20:2). [99]

Similarly, Schnackenberg et al attempted to discover molecular markers of early stage of cardiotoxicity induced by doxorubicin in mice before the onset of cardiac damage. They discovered 18 metabolites significantly altered in plasma, and another 22 metabolites were increased in cardiac tissue after a cumulative dose of 6mg/kg, while myocardial injury and cardiac pathology were not noticed until after 18 and 24 mg/kg cumulative doses, respectively.[100] Metabolomics analyses of plasma and heart tissue showed significant variations in the levels of many amino acids (among

arginine and citrulline), biogenic amines, acylcarnitine's (carnitine) and tricarboxylic acid cycle (TCA)-related metabolites (lactate, succinate e.g.) .

Tan et al conducted a study using gas chromatography-mass spectrometry to describe the metabolic profile of doxorubicin-induced cardiomyopathy in mice. They identified 24 metabolites, which were implicated in glycolysis, the citrate cycle and the metabolism of some amino acids and lipid and were selected as possible biomarkers for detection of cardiotoxicity. [101]

Andreadou et al used nuclear magnetic resonance (NMR) spectrometry to describe the metabolic profile of the acute doxorubicin cardiotoxicity in rats and to evaluate the metabolic alterations conferred by co- treatment with oleuropein.[90] The mice were divided into six groups: the first group included the control group, the second group received DOX, and the other four groups of mice received doxorubicin with oleuropein in a different dose and days, regarding oleuropein. Mice hearts were excised 72 hours after doxorubicin administration and H-NMR spectra of aqueous myocardium extracts were monitored. The results of analysis showed the increase of levels of acetate and succinate in DOX group compared to controls, while amino acids levels were lower. The conclusion of the article was that acetate and succinate constitute novel biomarkers for early detection cardiotoxicity. [102-103]

Geng et al in their study, used gas chromatography-mass spectrometry analysis of main targeted tissues (serum, heart, liver, brain, and kidney), to systemically evaluate the toxicity of DOX. Multivariate analyses revealed 21 metabolites in the serum, including cholesterol, D-glucose, D-lactic acid, glycine, L-alanine, L-glutamic acid, L-isoleucine, L-leucine, L-proline, L-serine, L-tryptophan, L-tyrosine, L-valine, N-methylphenylethanolamine, oleamide, palmitic acid, pyroglutamic acid, stearic acid, and urea, were changed in the serum in the DOX group. [104]

Tantawy et al identified lower plasma abundance of pyruvate and higher abundance of lactate in patients with carfilzomib-related cardiovascular adverse events. (CVAEs). They emphasized the significance of the pyruvate oxidation pathway associated with mitochondrial dysfunction. In order to better understand the mechanisms of Carfilzomib - CVAEs further investigation and validation is needed in a larger independent cohort. [105]

Yin et al proposed 15 different metabolites which play important role in cyclophosphamide induced cardiotoxicity. In this study, rat plasma samples were collected and analyzed one, three and five days after cyclophosphamide administration using ultra-performance liquid chromatography quadrupole time-of-flight mass spectrometry (UPLC-QqTOF HRMS). Of biomarkers studied, the proline, linoleic acid and glycerophospholipids changed significantly in three periods and the change associated with increasing time of occurrence of cardiotoxicity from cyclophosphamide. [106]

The study of Jensen et [107] al showed significant decreases in docosahexaenoic acid, arachidonic acid/ eicosatetraenoic acid, O-phosphocolamine and 6-hydroxynicotinic acid after sunitinib treatment with non-targeted metabolomics analysis of mice heart.[30] The same author, also, showed alterations in taurine/hypotaurine metabolism in the hearts and skeletal muscles of mice after sorafenib treatment. [108]

Except for the analysis of plasma and heart tissue, NMR spectroscopy-based metabolomics may detect low molecular weight metabolites in urine and cell culture media. For example, Chaudhari [109] et al showed reduction in the utilization of pyruvate and acetate, and accumulation of formate contrast to control culture medium of human induced pluripotent stem cell-derived cardiomyocytes exposed to doxorubicin. In contrast, Wang et al [110] showed in their study that tryptophan and phenylalanine metabolism in urine was also an important process in the systemic toxicity of doxorubicin. Also, Park et al identified 19 urinary metabolites in rats treated with doxorubicin. [111]

This technology is still under development, it seems obvious that metabolomics holds the potential to revolutionize our ability to profile samples in order to understand biological processes and find useful disease diagnostic biomarkers

Conclusions

Cardiovascular toxicity continues to be the major cause of drug failure during preclinical and clinical treatment models and contributes to drug withdrawal after approval. numerous medication

drugs that have been used frequently in adult clinical practice for a long time, have demonstrated potentially harmful effects on the heart in pediatric patients. The cardiotoxicity of these medications persists as a significant issue, having a negative impact on patients' quality of life as well as the overall survival. Several strategies for early detection of cardiotoxicity have been developed to reduce the numbers of patients with cardiac mortality and morbidity. Of importance, the biomarkers identified by the "omics" approach are considered new potential markers, especially in the scenario of diagnosis and risk stratification of acute coronary syndromes induced by chemotherapeutic drugs and may prove helpful in the early detection of anticancer cardiotoxicity.

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Abbreviations

| | |
|-----------|-----------------------------------------------------|
| ABC | Adenosine triphosphate-binding cassette transporter |
| ABCC2 | ATP Binding Cassette Subfamily C Member 2 |
| ACT | Anthracycline-related cardiotoxicity |
| ALL | Acute lymphoblastic leukemia |
| BNP | B-type natriuretic peptide |
| CBR | Carbonyl reductase |
| CELF4 | CUGBP Elav-Like Family Member 4 |
| CHF | Congestive heart failure |
| CP | Cyclophosphamide |
| CMR | Cardiac magnetic resonance imaging |
| CRP | C-reactive protein |
| CVAEs | Cardiovascular adverse events |
| DOX | Doxorubicin |
| ECG | Electrocardiography |
| EVs | Extracellular vesicles |
| GC | Gas chromatography |
| GDF-15 | Growth/differentiation factor 15 |
| GLS | Global systolic longitudinal myocardial strain |
| HER2 | Human epidermal growth factor receptor 2 |
| HF | Heart failure |
| IFO | Ifosfamide |
| LC | Liquid chromatography high performance |
| LC-MS | Liquid chromatography-mass spectrometry |
| LPC | Lysophosphatidylcholine |
| LV | Left ventricular |
| LVD | Left ventricular dysfunction |
| LVEF | Left ventricular ejection fraction |
| miRNAs | MicroRNAs |
| mRNAs | Messenger RNAs |
| MPI | Myocardial perfusion imaging |
| MS | Molecular mass spectrometry |
| MUGA | Nuclear cardiac imaging |
| NMR | Nuclear magnetic resonance spectrometry |
| NOTCH1 | Neurogenic locus notch homolog protein 1 |
| NT-proBNP | N-terminal pro b-natriuretic peptide |
| PET | Position emission tomography |
| PYGB | Glycogen phosphorylase |
| RARG | Retinoic acid receptor gamma |
| SAL | Saline |
| SLC | Solute carrier transporters |
| SNP | Single-nucleotide polymorphism |
| SPECT | Single photon emission computed tomography |
| SVM | Vector machine |
| TKI | Tyrosine kinase inhibitors |
| TCA | Tricarboxylic acid |

| | |
|-----------------|-------------------------------------------------------------------------------------|
| TnT | Troponin T |
| TOP2 | Topoisomerase II |
| Top2 β | Topoisomerase-II β |
| UGT1A6 | Glucuronosyltransferase family |
| UPLC-QqTOF HRMS | Ultra-performance liquid chromatography quadrupole time-of-flight mass spectrometry |
| VEGF | Vascular endothelial growth factor |
| vWF | Von Willebrand factor |
| 2D | Two-dimensional Echocardiography |
| 5-FU | Fluorouracil |

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