

Clinical impact of circulating biomarkers in prediction of adverse cardiac events in patients with congenital heart disease. A systematic review

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ARTICLE INFO

Keywords:

Biomarkers

Prognosis

Congenital heart disease

Hs-TnT

CRP

ST-2

GDF-15

RDW

Albumin

Creatinine

Gal-3

Norepinephrine

BNP

NT-proBNP

MELD-XI

ABSTRACT

Introduction: Patients with congenital heart disease (ConHD) are at increased risk for adverse cardiac events. Predicting long-term outcomes and guidance of patient management might benefit from a range of (new) biomarkers. This is a rapidly evolving field with potentially large consequences for clinical decision making. With a systematic review of available biomarkers in ConHD we identified the clinical role of these markers, knowledge gaps and future research directions.

Methods: We systematically reviewed the literature on associations between blood biomarkers and outcome measures (mortality or composite adverse outcomes in patients with ConHD).

Results: The inclusion criteria were met by 102 articles. Biomarkers assessed in more than 3 studies are discussed in the main text, those studied in 3 or less studies are summarized in the supplement. Thus, we discuss 15 biomarkers from 92 studies. These biomarkers were studied in 32,399 / 10,735 patients for the association with mortality and composite adverse outcomes, respectively. Biomarkers that were studied most and had statistically significant associations with mortality or composite adverse outcomes were (NT-pro)BNP, MELD-XI score, Hs-CRP, creatinine, albumin and sodium. Most of these biomarkers are involved in intracardiac processes associated with inflammation or are markers of renal function.

Conclusion: For (NT-pro)BNP, clinical value for prediction of mortality and composite adverse outcomes in adult and paediatric ConHD has been shown. For MELD-XI, hs-CRP, albumin, creatinine, sodium, RDW, and GDF-15, correlations with mortality and composite adverse outcomes have been demonstrated in patient groups with mixed types of ConHD, but clinical utility needs additional exploration.

1. Introduction

Congenital heart disease (ConHD) is the most common birth defect, affecting around 1 % of newborns [1]. In recent decades the survival of

patients with ConHD has increased due to improvements in antenatal screening, diagnostics, catheter interventions, surgical techniques, and supportive care. For common types of ConHD over 60 % of patients have residual lesions after initial (surgical) correction [2–5]. The combination

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of residual abnormalities and increased prevalence of ConHD results in a growing population at risk for long-term adverse events [6,7]. As many as 1 in 13 patients with ConHD will develop HF during their lifetime, and the risk of death after the age of 18 years is 3.2 times higher among ACHD patients compared to controls [8,9]. A major challenge in the follow-up of paediatric and adult patients with ConHD is to stratify the risk of who will develop complications [10]. Furthermore, there is a need to find new targets for screening, monitoring, (guidance of) drug therapy and endpoints for (drug) trials in paediatric patients [11]. This is a rapidly evolving field in which several pathways and related blood biomarkers of neurohormonal activation, myocardial injury and stress, inflammation, fibrosis, remodeling, vascularization and non-cardiac organ dysfunction have been discovered in ConHD [12–16]. Of these markers, B-type natriuretic peptide (BNP) and N-terminal segment of pro-BNP (NT-proBNP) have been included [17] in current guidelines for patients with ConHD, contributing to assessment of prognosis and indications for interventions [11,18]. Recently several new markers have been reported, providing new and additional information on pathways related to outcome measures in ConHD [19]. In this paper, we examined the (potential) clinical role of these markers by systematic review, aiming to identify their current clinical role, knowledge gaps and potential future research directions.

2. Methods

2.1. Search strategy

A systematic search of the Embase, Medline, Web of Science Core Collection and Cochrane Central Register of Controlled Trials databases was performed on 14th of June 2023 to identify studies on the prognostic value of blood biomarkers in patients with ConHD. The exact search strategy used is given in *Supplementary Text 1*.

2.2. Eligibility of studies

All articles were screened by title and abstract and assessed independently by two reviewers (QD and HA). Discrepancies were resolved by discussion with a third reviewer (WG). Subsequently full-texts of included articles were read. Studies were included if they were written in English, studied patients with ConHD across the paediatric and adult age range, were cohort studies or clinical trials with a minimum follow-up of six months, evaluating the prognostic value of blood biomarkers on the prognostic endpoints mortality or composite adverse outcomes. We included studies if composite adverse outcomes was defined as the occurrence of all-cause mortality, heart failure, (re)hospitalization for cardiac reasons, arrhythmia, heart transplantation, pulmonary arterial hypertension (PAH), liver or kidney failure, impaired exercise tolerance (decreased peak oxygen uptake (VO₂) or cardiac re-interventions, or a combination of the above.

2.3. Inclusion

We list all studies that fulfilled the inclusion criteria and their data (*supplementary Table 1 & 18–37*) but, for practical reasons, we opted to discuss only those biomarkers that were studied in at least 4 different studies, irrespective of the type of patients included in the full text of this paper.

2.4. Data extraction

The following information was extracted from the included articles: type of ConHD, number of participants, male/female distribution, mean or median age, the biomarkers studied, prognostic endpoints, follow-up duration, and measures of the association between the biomarkers and the study endpoints.

2.5. Statistical analysis

Studies were aggregated that a) included a mixture of ConHD conditions, b) focussed on a specific ConHD condition and c) evaluated patients with ConHD complicated by PAH. Studies were classified as 'paediatric' if the mean or median age of study subjects was below 18 years, and as 'adult' otherwise. The Quality in Prognostic Studies tool was utilized to assess risks of bias of individual studies, screening was done by one of 3 authors (WG, QD, HA) [20].

In case 95 % confidence intervals (CIs) of effect sizes were not reported in the included studies, these were calculated using other data in the manuscript or extracted from graphs if possible from the available data. Results were considered significant if $p < 0.05$.

3. Results

3.1. Study characteristics

A total of 102 articles fulfilled the inclusion criteria (Fig. 1). A complete overview of the number of studies per biomarker is provided in Fig. 2. Two thirds of studies included have been published after 2017. Applying the criterion for discussion of at least 4 available studies per biomarker, we will report on 92 studies (44 in mixed types of ConHD, 9 on ConHD with PAH, 17 in specific ConHD types, of which 10 in Fontan patients). An overview of all included studies with their main characteristics is shown in *Supplementary Table 1 & 2* [12,21–115].

Detailed information on patient characteristics, type of biomarker, outcome measures and percentage of outcome reached for all the 92 included studies can be found in *Supplementary Table 3–37*. Results of the quality assessment can be found in *Supplementary Table 38–39*. We utilized the Quality in Prognostic Studies (QIPS) score to appreciate quality of the included studies [20]. On average studies done in adults were of higher quality (2.61 ± 0.44) compared to paediatric studies (2.35 ± 0.48) (Table 1). In general most studies scored well on most points except in correction for confounding.

From 92 articles, a total of 32,399 patients were included in mortality analysis and 10,735 patients were included for analysis of composite adverse outcomes. The number of patients per biomarker studied for the association with mortality and composite adverse outcomes is depicted in Fig. 3. On average 331 patients were included per study. The average number of patients per type of cohort was: 530 for mixed ConHD diagnostic types, 162 for studies on single ConHD types (165 for Fontan patients) and 102 for studies analyzing ConHD with PAH. Paediatric studies included a mean number 151 participants, in adults this was 371.

3.2. Biomarkers and mortality

The median follow-up between time of biomarker assessment and reaching an endpoint (mortality or composite adverse outcomes) was 4.25 years (IQR range 2.43–6.13 years). As shown in Fig. 3A, MELD-XI score, (NT-pro) BNP creatinine, albumin and sodium were all studied in large numbers of patients (> 3000 patients / marker included) and showed mostly significant associations with mortality. Hs-CRP, norepinephrine and RDW were studied in relatively large cohorts. These biomarkers also had statistically significant associations with mortality in almost all studies. Platelet count, and Gal-3 were studied in smaller cohorts, mostly with a significant association with mortality. Uric acid was studied in a relatively large cohorts (± 1000 participants in total) but showed no association with mortality in over half of the patients.

In mixed cohorts (Fig. 4A), results were very similar to the overall group. Fig. 5A shows that the outcomes in disease-specific cohorts differed from those of the overall population. In disease specific cohorts norepinephrine was frequently studied and uric acid was the marked that most often showed significant results. Biomarkers ST2, Hs-CRP, GDF-15, and Gal-3 were studied less frequently and their correlations

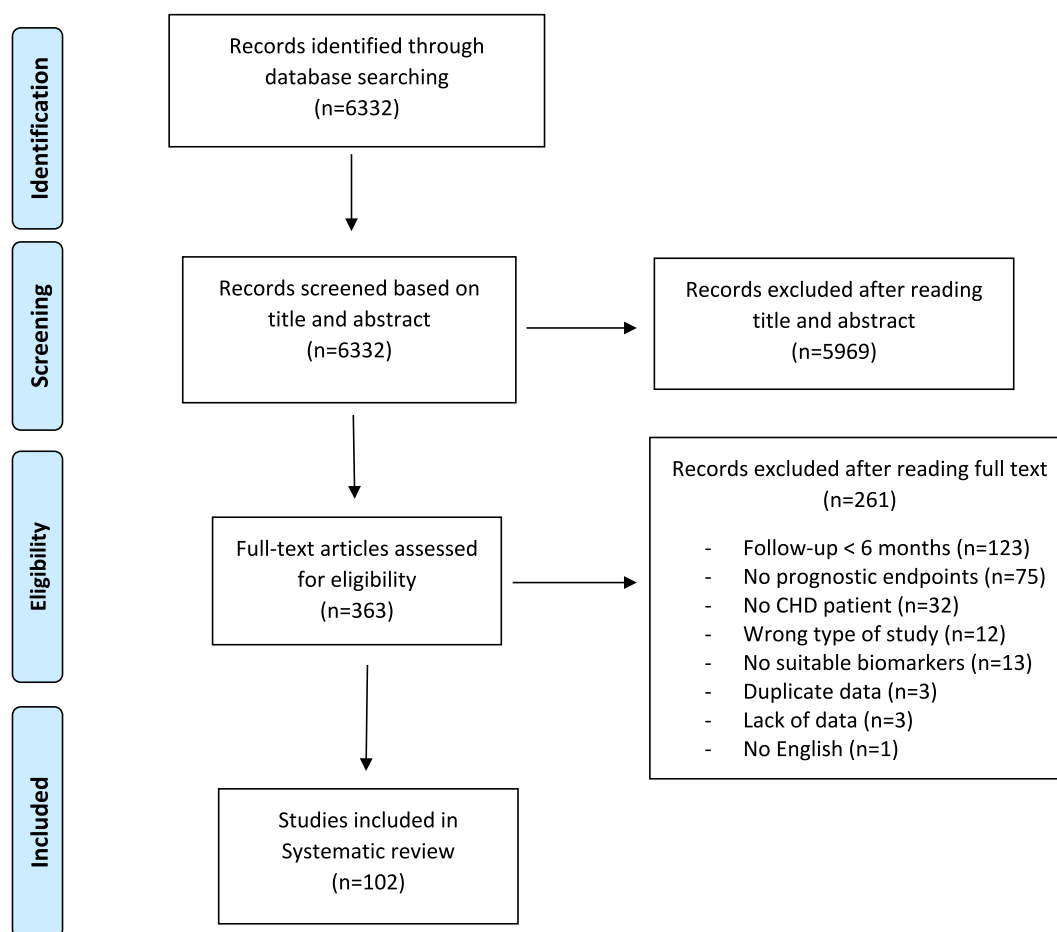


Fig. 1. Flowchart.

were not found to be significant in contrast to in overall populations.

In patients with PAH in ConHD, the biomarkers (NT-pro) BNP, hs-CRP, creatinine, hs-TnT, RDW, uric acid and Gal-3 were shown to have significant associations with mortality (Fig. 6A).

Fig. 7A and 8A show an overview of studies of biomarkers in different age groups. 29,676 adult patients were studied. As this is the most studied group these results are very comparable to the overall cohort. 2723 children were studied. NT-proBNP, norepinephrine and uric acid were studied in large patient numbers with significant association with mortality. For hemoglobin, Hs-CRP, GDF-15 and Hs-Tnt no association with mortality was noted. ST2, Gal-3 and platelet count showed positive association with mortality, in relatively small patient numbers.

3.3. Biomarkers and composite adverse outcomes

An overview of the number of patients studied with composite adverse outcomes as outcome marker is given in Fig. 3B. (NT-pro) BNP was by far the most studied biomarker with a significant association in most studies. In mixed cohorts (Fig. 4B) and disease specific cohorts (Fig. 5B) (NT-pro) BNP was most frequently studied, with relatively more non-significant studies in disease specific cohorts compared to mixed cohorts and patients with CHD-PAH (Fig. 6B). Hs-TnT, albumine and Hs-CRP were also studied frequently. For these markers a significant association with composite adverse outcomes was found in around 2/3 of patients. Albumine and Hs-CRP were not studied in disease specific cohorts (Fig. 5B). In mixed cohorts albumin and creatinine were extensively studied, however in around half of the patients no significant association was found. For Patients with PAH in ConHD (Fig. 6B

creatinine, Hs-TnT and (NT-pro) BNP showed a significant association with composite adverse outcomes in all.

4. Discussion

The study of biomarkers in ConHD is a rapidly evolving field with potentially large consequences for clinical decision making. This systematic review has provided an overview of available biomarkers in ConHD. The strongest evidence is for the predictive value of (NT-pro) BNP in both predicting mortality and composite adverse outcomes in adult and paediatric ConHD. For other markers reviewed, correlations with mortality and composite adverse outcomes have been demonstrated in patient groups with mixed types of ConHD. The evidence for these biomarkers to predict adverse outcomes varies by biomarker and by type of diagnosis.

A summary of the mechanisms of action of the biomarkers studied in this review is given in Table 2. In general, these mechanisms have been described extensively in acquired heart disease [116–141].

Blood biomarkers may play an important role in risk stratification for mortality and composite adverse outcomes and may improve monitoring, treatment guidance and the understanding of pathophysiology. The associations with mortality and composite adverse outcomes in ConHD will be discussed for the different biomarkers emerging from the systematic review in the following. Knowledge gaps exist for other roles of these biomarkers, as will be discussed subsequently.

4.1. Cardiac associated biomarkers

In ConHD (NT-pro) BNP has been studied extensively. This is the

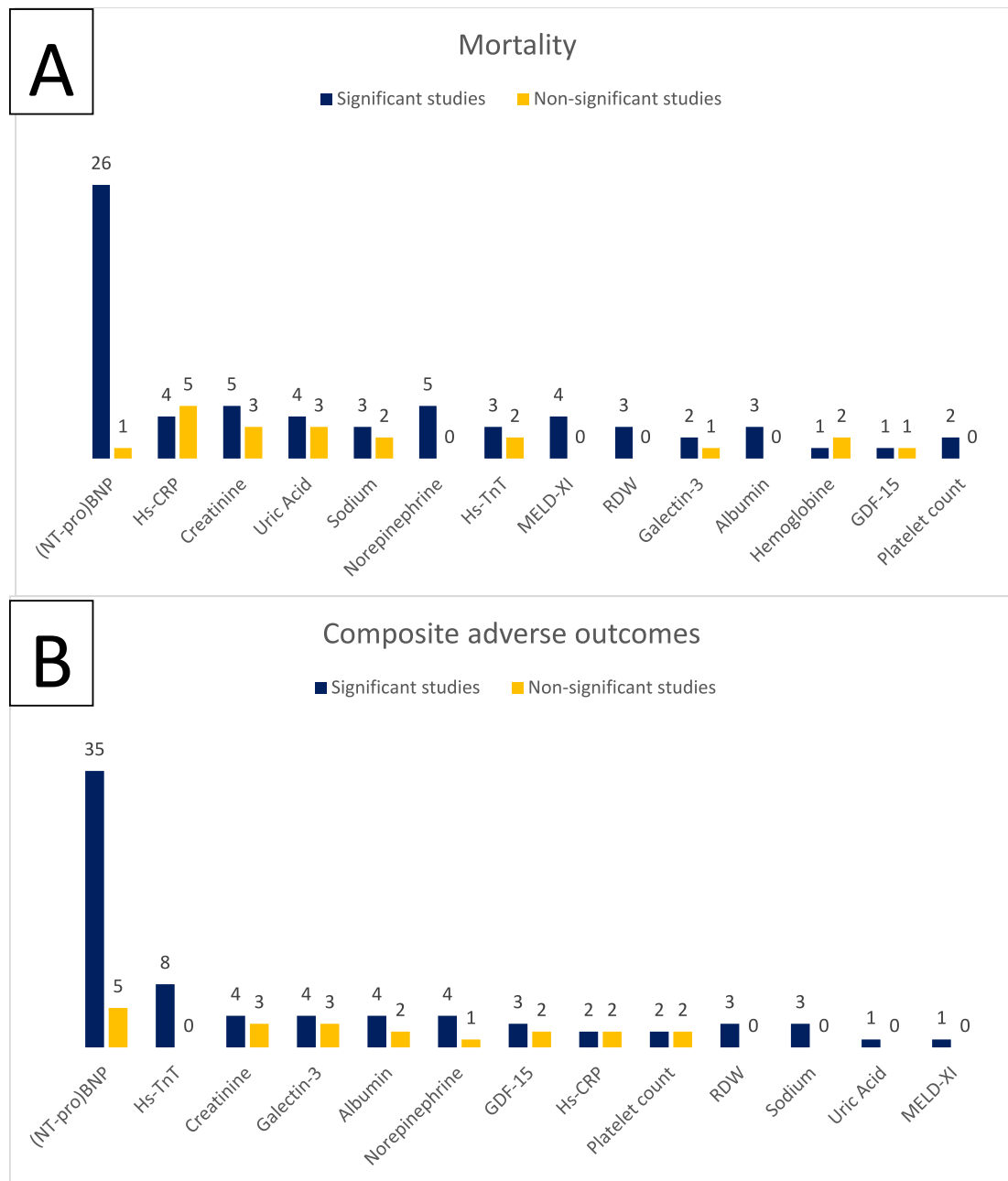


Fig. 2. Number and significance level for association with outcome measures mortality (panel A) and composite adverse outcomes (panel B) of studies per biomarker in ConHD.

Abbreviations: (NT-pro)BNP: N-terminal pro b-type natriuretic peptide; **Hs-TnT**: High-sensitivity troponin T; **Hs CRP**: High-sensitivity C-reactive protein; **RDW**: Red cell distribution width; **ST2**: suppression of tumourigenicity 2; **MELD score**: Model for End-Stage Liver Disease score; **GDF-15**: Growth differentiation factor 15; **ConHD**: congenital heart disease; **PAH**: pulmonary arterial hypertension. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

only biomarker currently included in adult ConHD guidelines [40,44,47,142–144]. It was shown to have strong associations with mortality and composite adverse outcomes in all different groups, both in adult and paediatric cohorts, making it the best available marker for adverse events in ConHD at present.

For hs-TnT there was a strong association between increased levels and both composite adverse outcomes and mortality in PAH-ConHD patients [65,77] and in patients with ccTGA, with univentricular hearts, and in cohorts with mixed types of ConHD disease. The added value of Hs-TnT in predicting composite adverse outcomes in patients with ConHD and PAH is less clear [21,31,39,65,77,87]. In paediatric cohorts the predictive value of Hs-TnT was low whereas in adults there

was predictive value. Hs-TnT is mainly released as a response to cell damage often as a result of inadequate oxygen supply to the myocardium [145]. We speculate that children might be better able to maintain adequate oxygen supply to the tissues where adults have developed more myocardial damage over time, e.g. resulting in more myocardial fibrosis, in combination with deterioration of coronary artery function [146–148].

Gal-3 was associated with mortality in relatively small cohorts of mixed ConHD. In relatively large mixed ConHD cohorts, Gal-3 was associated with composite adverse outcomes. No significant association with mortality was observed in disease-specific cohorts. See paragraph on limitations of the available studies.

Table 1
overview of the association of described biomarkers with different disease conditions.

	Myocardial stretch	Myocardial injury	Myocardial fibrosis	Chronic inflammation	Renal dysfunction	Hepatic dysfunction	Nutritional deficiencies	Endothelial dysfunction	Oxygen free radicals	Adverse remodeling	Coagulation
(NT-pro)BNP	+										
Hs-TnT		+	+	+							
Galactin 3				+							
Hs CRP				+							
RDW			+	+	+		+			+	
ST2	+			+	+			+			
Uric acid				+	+						
MELD score				+	+						
Albumin				+	+					+	
Norepinephrine				+	+						
Platelet count				+	+						
GDF-15		+		+	+	+					+

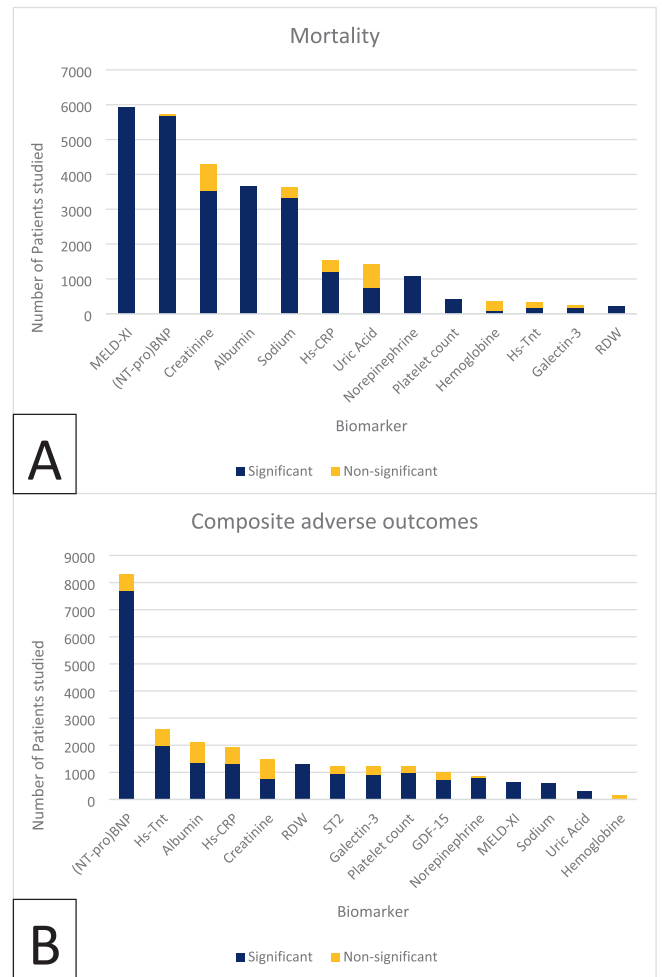


Fig. 3. Number of patients and significance level per biomarker for association with outcome measures mortality (panel A) and composite adverse outcomes (panel B), overall.

4.2. Markers of inflammation

For Hs-CRP associations with mortality were significant in a majority of patients for both mixed and PAH-ConHD cohorts. However, in the only disease specific cohort, patients with tetralogy of Fallot, no association with mortality was found, probably related to the small number of patients reaching the endpoint.

Significantly higher risk for mortality with increased RDW was found in mixed types ConHD and in patients with ConHD related PAH [45,109]. RDW was not studied in disease specific cohorts.

For ST2 60 % of the studies, including a large study in a mixed ConHD cohort, found an association with composite adverse outcomes. Two studies in disease specific cohorts did not find this association.

For hyperuricemia, 2 out of the 4 studies showed a significant association between mortality and higher uric acid levels for patients with a Fontan circulation and in ConHD patients with PAH [98,99]. The only mixed-ConHD cohort in which uric acid was studied did not show a significant association [81].

4.3. Renal function associated biomarkers

In general, creatinine showed strong associations with mortality in all types of cohorts. However, associations with composite adverse outcomes were less clear.

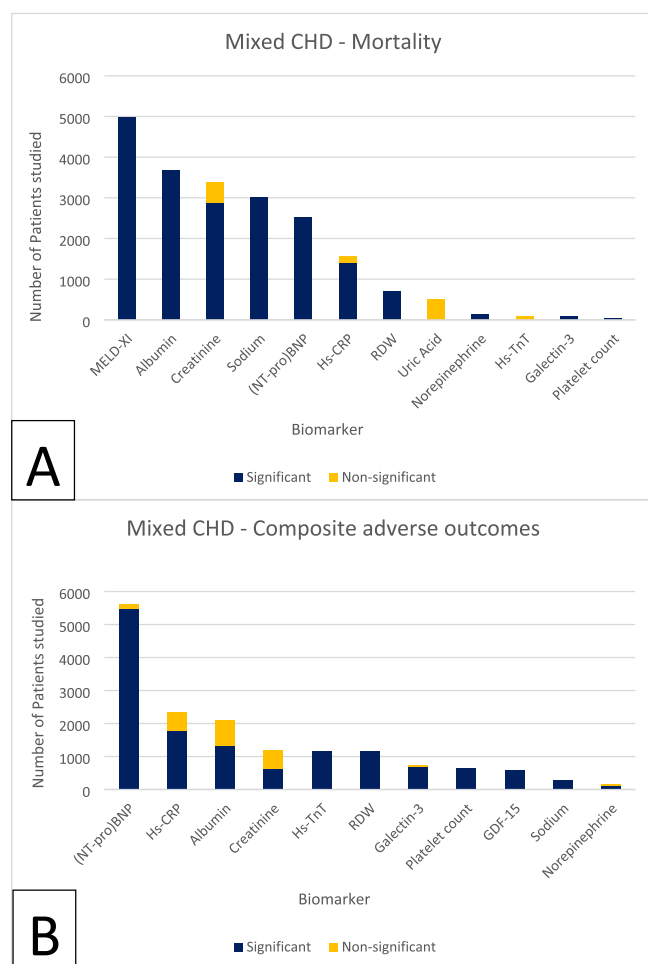


Fig. 4. Number of patients and significance level per biomarker for association with outcome measures mortality (panel A) and composite adverse outcomes (panel B) for the number of patients per biomarker, in studies of cohorts with mixed types of congenital heart disease.

4.4. Other biomarkers

Albumin has been associated with adverse events in ConHD, related to increased central venous pressure, a known complication of many different types of ConHD [149,150]. Increased central venous pressure may lead to a shift in albumin from intracellular to the extracellular space [84]. Hypoalbuminemia is probably also influenced by chronic inflammation and endothelial dysfunction [151], which have also been related to complications of different types of ConHD [152]. However, significant associations between albumin levels and mortality and composite adverse outcomes were only found in mixed cohorts of ConHD, probably related to the wide range of central venous pressures and inflammation levels in specific groups [152].

Our review showed an association between norepinephrine and mortality and composite adverse outcomes in almost 1000 patients with Fontan circulation [55,85,98]. However, norepinephrine was not studied in other ConHD populations. Because of the unique physiology of the Fontan circulation, we cannot extrapolate these results beyond Fontan patients.

Platelet count is associated with all-cause mortality in the general population as well as in patients with cardiovascular disease [153,154]. This is thought to be caused by changes in coagulation and inflammation [102]. An increased risk of composite adverse outcomes was found in Fontan patients [50]. However, composite adverse outcomes in this study was mainly driven by liver cancer, a complication of Fontan

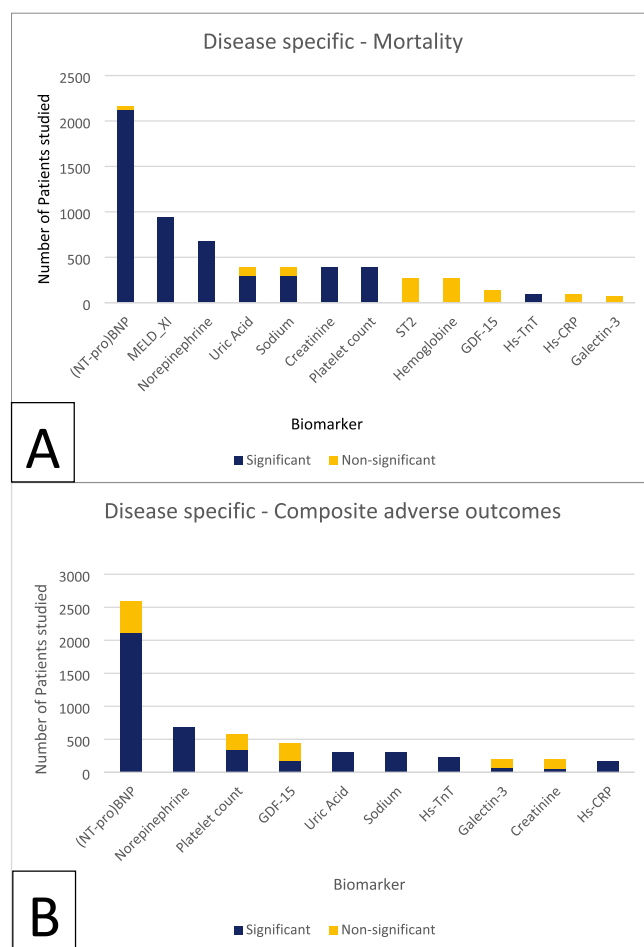


Fig. 5. Number of patients and significance level per biomarker for association with outcome measures mortality (panel A) and composite adverse outcomes (panel B) for the number of patients per biomarker, in studies of cohorts with specific ConHD.

associated liver disease [6]. Platelet count is decreased in patients with liver failure and portal hypertension [155]. Therefore, in Fontan patients, platelet count might reflect liver rather than cardiac.

The MELD-XI score is a combination of creatinine, bilirubin and INR, indicating hepatorenal function [3]. Whilst this biomarker score was studied in only 5 studies, the number of included patients was high (> 5000), mainly driven by large studies in mixed cohorts and in Ebstein's disease [113,156]. The predictive value of MELD-XI for mortality has also been shown in Fontan patients, known for liver problems [157]. Evidence for MELD-XI as risk predictor is lacking in patient groups without liver complications.

Considering its documented role in acquired heart disease GDF-15 is a promising biomarker with demonstrated association with all-cause mortality and composite adverse outcomes in mixed cohorts of ConHD patients [123–125]. However, in disease specific ConHD cohorts, mostly non-significant results have been shown. Therefore, it remains uncertain how useful GDF-15 is as a predictor for adverse events in ConHD.

4.5. Biomarkers studied in less than 4 studies

59 biomarkers were studied in less than 4 studies, 18 of these biomarkers were cardiac associated biomarkers, 9 were markers of inflammation, 8 were associated to renal function and 22 could be classified as other biomarkers (See Tables 19–37 in supplemental material). Often these biomarkers were closely related to biomarkers studied more frequently. For example bilirubin (studied 1 time) is part of

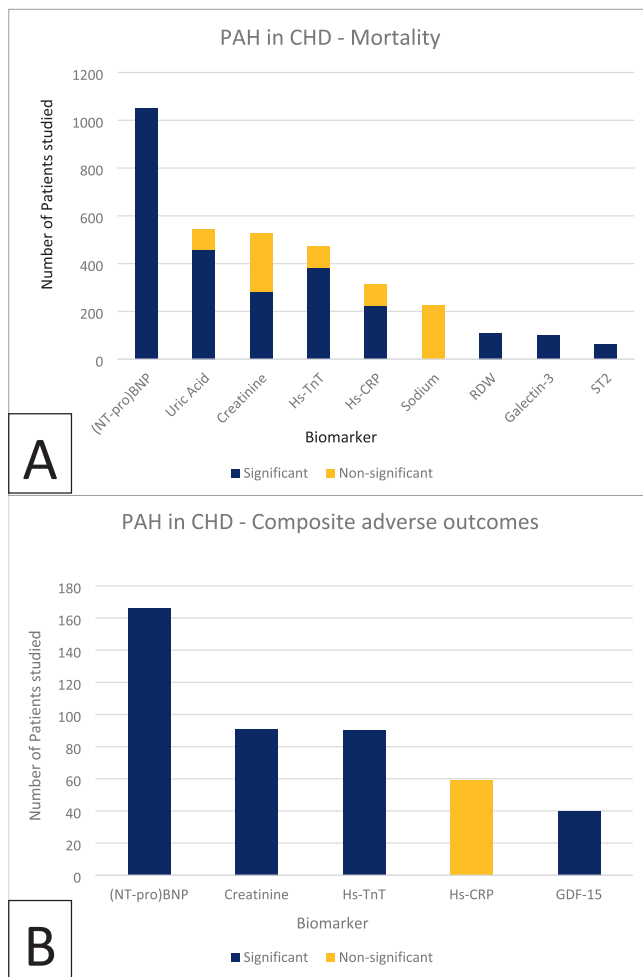


Fig. 6. Number of patients and significance level per biomarker for association with outcome measures mortality (panel A) and composite adverse outcomes (panel B) for the number of patients per biomarker, in studies of cohorts of ConHD with PAH.

the MELD-XI score and ANP (studied 3 times) is closely related to (NT-pro)BNP. The studies in this group of less frequently studied biomarkers in general were smaller (115 participants vs 330 participants for studies studying more frequently studied biomarkers). Often these less frequently studied biomarkers have only studied by one specific research group, which reduces generalizability and increases the risk for bias.

4.6. Knowledge gaps and directions for future research

Currently available studies in ConHD have not answered relevant questions like:

- i) What type of use of additional biomarkers in addition to (NT-pro)BNP is predictive of outcomes (e.g. baseline values, serial measurements, combinations of different biomarkers, changes with therapy)?

It has been suggested that combining several markers increases the predictive value for heart failure risk in adult ConHD [158,159]. However, most reviewed studies focused on a single biomarker and its association with mortality and/or composite adverse outcomes. Some studies (Supplementary Table 1 & 2) have tested and shown the additional predictive value beyond (NT-pro)BNP [12]. In a study by Baggen et al. the authors combined three different biomarkers (NT-pro)BNP, Hs-TnT and GDF-15) and showed that patients who had no biomarkers

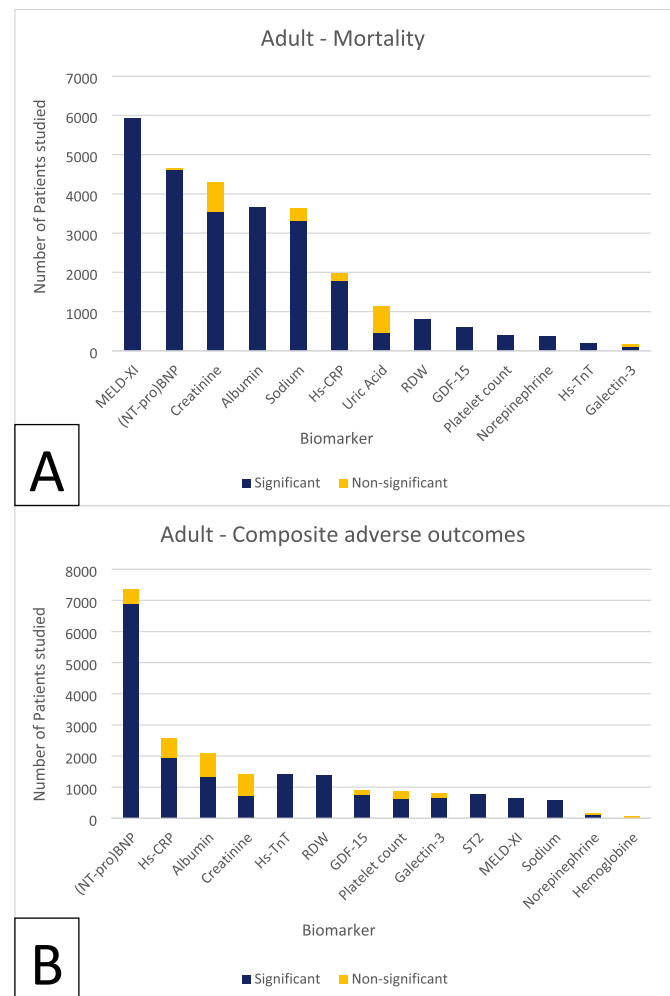


Fig. 7. Number of patients and significance level per biomarker for association with outcome measures mortality (panel A) and composite adverse outcomes (panel B) for the number of patients per biomarker, in studies of cohorts in adult ConHD patients.

elevated had an event free survival of 86 % over 4 years, whereas in patient where all 3 biomarkers were elevated only 27 % was without adverse events over 4 years. Often it has been difficult to show additive value of biomarkers over NT-proBNP [106]. Therefore we advise that all new biomarkers should at least be tested on their added value beyond NT-proBNP. Lastly we hypothesize that biomarkers based on different organ systems could provide added value when combined.

A versatile approach in managing ConHD in adults involves combining blood biomarkers with other clinical data to create a risk score, used for screening, monitoring, treatment guidance and risk prediction, as demonstrated in recent studies [160]. These aspects require further evaluation in adults and children with ConHD, shown by the lack of confounding correction seen in the studies included in this review.

- ii) what is the impact on clinical decision making of potential natural fluctuations in biomarker levels and of confounders in assessment (e.g. age, sex, obesity, renal function)?

In a recent review on the use of biomarkers in children with ConHD, McGinn et al. pointed towards the lack of age-related reference rates for many of the biomarkers discussed, except (NT-pro)BNP and troponin [161]. In acquired heart failure in adults, the importance of confounders of biomarker levels such as obesity has been stressed [162]. These

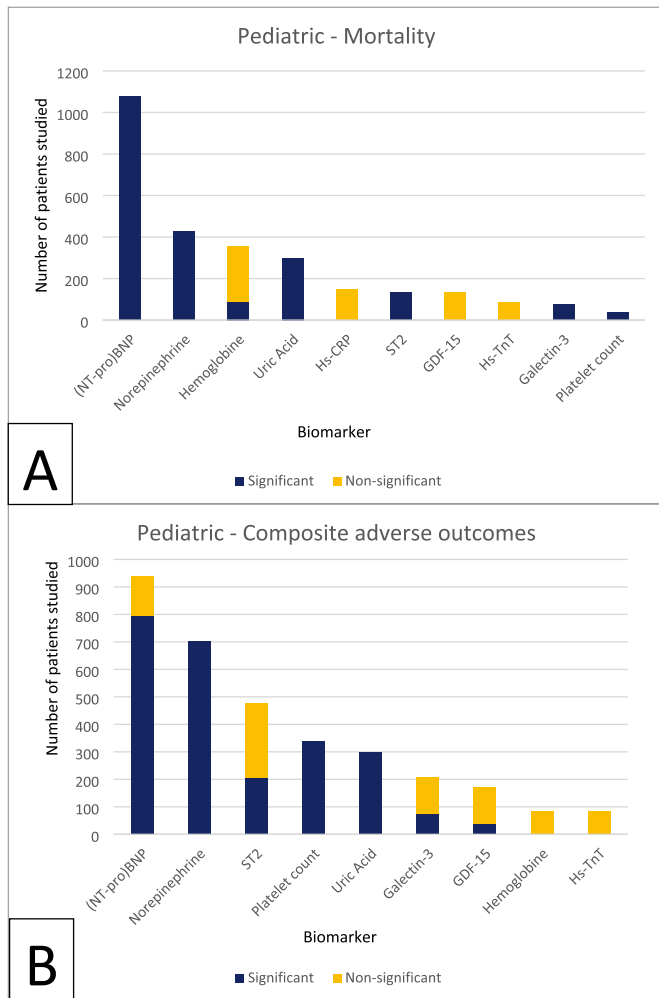


Fig. 8. Number of patients and significance level per biomarker for association with outcome measures mortality (panel A) and composite adverse outcomes (panel B) for the number of patients per biomarker, in studies of cohorts in paediatric ConHD patients.

factors have hardly been explored in patients with ConHD.

- iii) does clinical decision making based on serum biomarkers improve outcomes for patients?

This important step has not been studied yet in ConHD and therefore it is unknown if implementation of these biomarkers leads to better identification of at-risk patients.

- iv) which biomarkers may serve as endpoint or therapeutic targets in (drug) trials?

Biomarkers may have a role in developing optimal drug therapy with ConHD, e.g. by serving as surrogate endpoint [163]. Biomarkers, e.g. (NT-pro)BNP, have been established as adequate surrogate endpoints in adults with acquired with heart failure [164]. It is hard to reproduce similar studies in children with ConHD, mainly because hard endpoints are only obtained at long-term. In these cases biomarkers might be utilized as surrogate endpoints to bridge known effects in adults to children. In a few paediatrics drug trials, circulating biomarkers have been used as (surrogate) endpoints in cardiomyopathy [164]. In a randomized trial of ivabradine in children with heart failure from dilated cardiomyopathy (DCM) NT-proBNP was used a secondary outcome [165]. For the biomarkers in the population of ConHD, additional studies are required to determine their potential as bridging biomarker, since several of the required criteria cannot be derived from the reviewed studies [163]. NT-pro BNP in all groups, norepinephrine in Fontan patients, or Hs-TnT in patients with PAH in ConHD are promising in this respect, considering their strong patient-level association between the biomarker and FFS endpoints, particularly death and rein-terventions [163].

- v) what are the effects of interventions and changes in biomarkers on outcomes and what is the dynamic of these changes that are predictive of changed (improved) outcomes [166]?

This has not been studied yet.

- vi) which biomarkers can be implemented in clinical guidelines?

Of the biomarkers discussed in this study, clinical value for prediction of mortality and composite adverse outcomes in adult and paediatric ConHD has been shown for (NT-pro)BNP, which has translated into various clinical guidelines. For MELD-XI, hs-CRP, albumin, creatinine, sodium, RDW, and GDF-15, correlations with mortality and adverse outcome have been demonstrated in patient groups with mixed types of ConHD, making them potential candidates to be included in guidelines for adults with ConHD, but not (yet) in specific lesions or in children. Apart from the limitations discussed in the items i – v, heterogeneity of the test populations and the lack of validation cohorts independently confirming the relationship between specific biomarkers and outcomes are issues to be addressed before biomarkers can be included in clinical guidelines.

4.7. Limitations

Based on the available literature meta-analysis was not possible. The high variability in the reporting of outcome measures and biomarker levels, such as the use of quartiles versus continuous risk either in a linear or logarithmic scale, hampered meta-analysis of biomarkers. In NT-pro(BNP) sufficient studies reported a continuous risk to execute a meta-analysis. However hazard ratio appeared strongly dependent on baseline level of (NT-pro) BNP. Arranging results according to baseline levels of NT-proBNP would result in small number of studies per meta-analysis or in high levels of heterogeneity in a single meta-analysis. We advocate future studies to report on hazard ratios (HR) for

Table 2

Aggregated averages for quality scores of QUIPS (full scores in supplementary tables 37–38).

	Study participation	Study attrition	Prognostic factor measurement	Outcome measurement	Study confounding	Statistical analysis and reporting	Risk of bias
Paediatric studies							
Mean	2,53	2,65	2,76	2,65	2,00	2,65	2,35
SD	0,50	0,59	0,42	0,59	0,59	0,48	0,48
Adult Studies							
Mean	2,49	2,20	2,73	2,84	2,26	2,67	2,61
SD	0,57	0,48	0,44	0,37	0,69	0,62	0,44

biomarkers in a standardized way, preferably HR per increase of standard deviation on a logarithmic scale. Meta-analyses could also not be performed due to the heterogeneity in study populations, outcome definitions of the individual studies or lack of data.

Relatively few studies on blood biomarkers have been conducted in children. This hampers identification of early predictors of mortality and composite adverse outcomes in children with ConHD.

Most of the work we reviewed was based on single marker assessment. Potentially important markers may have been missed by this approach. Explorative tools, like e.g. targeted bioarray proteomics and metabolomics, may play an important role in further clarifying pathophysiology of heart failure in ConHD [67,167–172].

Most of the included studies in this systematic review are characterized by study populations heterogeneous for type of ConHD, age of the patients, follow-up duration and means of reporting of HR. Most of the studies involved relatively small study populations and may have been underpowered to establish significant associations bet.

een biomarker levels and the prognostic outcomes. Some studies included results on heart failure and mortality in the results for composite adverse outcomes.

5. Conclusion

Biomarkers in ConHD have been studied frequently and increasingly, mostly in mixed cohorts of adult patients with ConHD. For (NT-pro) BNP, clinical value for prediction of mortality and composite adverse outcomes in adult and paediatric ConHD has been shown, which has translated into various clinical guidelines. For MELD-XI, hs-CRP, albumin, creatinine, sodium, RDW, and GDF-15, correlations with mortality and composite adverse outcomes have been demonstrated in patient groups with mixed types of ConHD, not in specific ConHD types and paediatric cohorts.

Further studies are required to explore the role of additional blood biomarkers as a tool for screening, monitoring, treatment guidance and risk stratification in (paediatric) ConHD.

Disclaimer

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CRediT authorship contribution statement

W.J. van Genuchten: Writing – original draft, Methodology, Investigation, Formal analysis, Data curation, Conceptualization. **H. Aversch:** Writing – review & editing, Data curation. **Q.M. van Dieren:** Writing – review & editing, Formal analysis, Data curation. **D. Bonnet:** Writing – review & editing, Conceptualization. **M. Odermarsky:** Writing – review & editing, Investigation. **M. Beghetti:** Writing – review & editing, Investigation. **J.W. Roos-Hesselink:** Writing – review & editing, Investigation. **Z. Reinhardt:** Writing – review & editing, Investigation. **C. Male:** Writing – review & editing, Investigation. **E. Naumburg:** Writing – review & editing, Methodology. **D. De Wolf:** Writing – review & editing, Investigation, Conceptualization. **W.A. Helbing:** Writing – original draft, Resources, Formal analysis, Conceptualization.

Acknowledgement

The authors wish to thank M. Udo from the Erasmus MC Medical Library for developing and updating the search strategies.

Funding

This project has received funding from the Innovative Medicines Initiative 2 Joint Undertaking under grant agreement No 777389. The Joint Undertaking receives support from the European Union's Horizon 2020 research and innovation programme and EFPIA.

WJ van Genuchten was supported by the Netherlands Cardiovascular Research initiative: an initiative with support of the Dutch Heart Foundation and Hartekind (grant to WA Helbing as part of a consortium; CVON219-002 OUTREACH).

Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.ijcard.2024.132723>.

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