Retinol-binding protein 4 in transthyretin-associated forms of cardiac amyloidosis: differences in the pathobiologies of mutant (ATTRm) and wild-type (ATTRwt) diseases

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Abstract

The role of retinol-binding protein 4 (RBP4), a natural binding partner of plasma circulating transthyretin (TTR), in TTR-associated cardiomyopathies is unknown. RBP4 is a small (21 kDa) protein that normally functions as a transporter for all-trans retinol (Vitamin A) and travels in the bloodstream as a ternary complex bound to TTR. The role of RBP4 in TTR amyloid fibril formation. Though increased serum concentrations of RBP4 have been reported in non-amyloidotic cardiomyopathies, there is little information about circulating levels in the inherited (ATTRm) and acquired (ATTRwt) forms of TTR-associated cardiac amyloid disease. Our study objectives were to investigate TTR amyloid-infiltrated cardiac tissue for the presence of RBP4 and to compare serum levels of the protein in ATTRm, ATTRwt and controls. We hypothesized that there would be histological and serological differences in RBP4 between ATTR patient and control samples. In the present study, we demonstrate that RBP4 is highly abundant in ATTRm cardiac tissue surrounding amyloid-infiltrated regions and, to a lesser extent, in ATTRwt specimens. Serum levels of RBP4 are significantly lower in ATTRm compared to ATTRwt (p < 0.0001) and healthy controls (p < 0.0011), with significant correlation between circulating RBP4 concentration and cardiac troponin-I (c-TnI) in our ATTRwt cohort. These data suggest, for the first time, that the pathobiologies of ATTRm and ATTRwt are dissimilar, and provide support for RBP4 as a serum biomarker of amyloid cardiomyopathy in ATTRm.

Introduction

Retinol-binding protein (RBP) is a protein that exists as one of five isoforms. The most abundant and only secreted form is RBP4, a 21 kDa protein almost exclusively produced by the liver, which transports all-trans retinol (Vitamin A) to target tissues in a ternary complex with transthyretin (TTR). RBP4-TTR complexes are formed in the ER [1-2] and their release from the liver into circulation is dependent on retinol binding to RBP4 [3]. Consequently, 95% of circulating RBP4 is bound to TTR [4,5]. The serum concentration is normally 42 µg/mL, but elevated levels have been described in type 2 diabetes [6], obesity [7], and inflammatory cardiomyopathy [8].

TTR, the natural binding partner of RBP4, is a homotetrameric protein predominantly expressed by the liver and present in plasma at physiologic concentrations of 200-400 µg/mL [9,10]. More than 100 point mutations in the TTR gene are associated with inherited forms of amyloidosis [11] known as mutant transthyretin-associated (ATTRm) amyloidosis [12]. While there is some variability in the disease phenotype, the majority of ATTRm patients feature amyloidotic cardiomyopathy.

Interestingly, wild-type TTR can also be amyloidogenic, forming amyloid fibrils that mainly infiltrate the myocardium [13]. The accumulation of variant and/or wild-type TTR cardiac amyloid deposits causes progressive worsening of organ function and leads to congestive heart failure [14-16].

It is widely held that ATTR amyloid diseases share a common pathobiology. Amyloid fibril formation is thought to result from a series of misfolding and aggregation steps, collectively described as the amyloid cascade [17]. The mechanism is initiated by TTR tetramer dissociation, believed to be the rate-limiting step, and followed by aggregation of non-native monomeric forms of the protein. A specific cause for destabilization and disassembly of TTR native quaternary structure is unknown; nonetheless, in vitro studies have shown that tetramer-released monomers can rapidly adopt misfolded conformations that self-associate, forming aggregates, and eventually insoluble amyloid fibrils [18].

Tissue deposited amyloid fibrils, the hallmark of all amyloid diseases, are highly organized, non-branching structures that bind Congo red dye. In TTR-associated amyloidosis, the heart is commonly a site for unremitting accumulation of extracellular amyloid fibril deposits. Diagnosis of ATTR amyloidosis frequently relies on biopsy proof of amyloid deposits, evidence of a pathologic TTR mutation in ATTRm disease, and more recently, the use of imaging techniques such as (99m)Tc-pyrophosphate scintigraphy [19,20]. Importantly, ATTR

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In the present study, we examined tissue and circulating forms of RBP4 in ATTRm and ATTRwt amyloidosis. Our interest was prompted by in vitro studies demonstrating the stabilizing effect of RBP4 on native TTR [21,23] and the limited information about RBP4 in ATTR cardiac diseases, despite extensive studies of the protein in other pathologies[6,24–26] particularly cardiomyopathy[8,27,28]. This report documents our 1) RBP4 histological and serological findings in ATTRm and ATTRwt; 2) assessment of baseline serological, clinical, and echocardiographic data of patients in each group; and 3) analysis of RBP4 as an indicator of ATTR amyloid disease type.

Materials and methods

Study Cohorts

Patient sera, tissue samples, and clinical data were obtained from the Boston University (BU) Amyloidosis Center repository with the approval of the BU Medical Center Institutional Review Board. Amyloid study samples were from 3 groups of patients presenting with amyloidotic cardiomyopathy: ATTRm, ATTRwt, and immunoglobulin light chain (AL) amyloidosis. Serum from patients seen at the BU Amyloidosis Center for a baseline evaluation between January 2009 and June 2013 were studied. A separate group of non-amyloid controls with cardiomyopathy were obtained from the BU Cardiovascular Center. Amyloidosis was diagnosed by Congo red staining of a tissue biopsy and identification of the amyloidogenic protein was accomplished by immunohistochemistry, immunogold electron microscopy, or mass spectrometry. ATTRm amyloidosis was established by the presence of a TTR variant protein on isoelectric focusing (IEF) and a pathologic gene mutation by DNA sequencing. For ATTRwt amyloidosis, TTR was identified in the amyloid deposit in the absence of a mutant TTR gene or variant protein sequence. Tissues from patients with immunoglobulin light chain (AL) amyloidosis were used as non-ATTR cases; AL amyloidosis was diagnosed by appropriate tissue amyloid typing, histopathologic identification of clonal plasma cells in the bone marrow, detection of a monoclonal immunoglobulin light chain (LC) by serum or urine immunofixation electrophoresis, and serum free LC testing. Clinical data was obtained from the electronic health record and included details of history, physical examination, and routine laboratory studies. Study design controlled for gender, age, and co-morbidities; patient study groups included males, 60 years of age or older, with triglyceride levels < 200 mg/dL. Triglyceride levels were ranging from 0 – 10 ng/mL, and serum dilutions of 1:100,000 were prepared in 0.1% tween-20. RBP4 protein (R&D) standard solutions, containing 0.1% tween-20, RBP4 protein (R&D) standard solutions, ranging from 0 – 10 ng/mL, and serum dilutions of 1:100,000 were prepared in 0.2% BSA, 0.025% tween. Standard, sample, or control antibody solutions. Tissue sections were counterstained with Harris’ hematoxylin, stained in alkaline Congo red, and viewed by bright field and polarized light microscopy.

Serum retinol-binding protein 4 measurement by sandwich enzyme-linked immunosorbent assay

To measure circulating levels of RBP4 in serum samples, a sandwich enzyme-linked immunosorbent assay (ELISA) was developed in our laboratory. Mouse monoclonal anti-human RBP4 and biotinylated polyclonal mouse anti-human RBP4 antibodies (R&D Systems, Minneapolis, MN) were used as the capture and detection antibodies, respectively. High-binding, flat bottom 96-well microplates (R&D Systems, Minneapolis, MN) were coated with 50 µL of 1 ng/mL mouse monoclonal anti-human RBP4 antibody and stored overnight at 4 °C. Plate wells were blocked with 1% BSA and washed extensively with PBS containing 0.1% tween-20. RBP4 protein (R&D) standard solutions, ranging from 0 – 10 ng/mL, and serum dilutions of 1:100,000 were prepared in 0.2% BSA, 0.025% tween. Standard, sample, or control solutions were added to microplate wells in triplicate. Solutions were shaken (300 rpm) for 2 hours at room temperature. Plates were washed and biotinylated polyclonal mouse anti-human RBP4 antibody (100 ng/mL) was applied. After 2 hours, streptavidin-HRP was added and the reaction proceeded for 30 minutes at room temperature with shaking (500 rpm). Plates were developed using 100 µL of TMB and the reaction was halted with 100µL of 1.5 N H₂SO₄. Chromogenic changes were detected with a Biotek ELx800 and data was analyzed by Gen5 software using the SPL curve fit. The intra-assay variability was less than 5% and the inter-assay variability was less than 15%.

Statistical analysis

Data from ELISA measurements of serum RBP4 were analyzed using Prism 5 software (Graphpad Software, La Jolla, CA). Gaussian distribution was assessed by running D’Agostino and Pearson omnibus normality testing. Continuous variables are described as mean ± standard error of the mean (SEM). Baseline demographic, serologic, and echocardiographic data were contrasted among the ATTRm, ATTRwt, and AL disease types using one-way ANOVA with multiple comparisons testing or Kruskal-Wallis tests. For baseline evaluation data in ATTRm and ATTRwt, associations of RBP4 with age, body...
mass index (BMI), triglycerides, BNP, c-TnI, interventricular septal thickness (IVST), LVEF, uric acid, or TTR were analyzed by Pearson correlation coefficients. All analyses were conducted with the use of SAS software, version 9.3 (SAS Institute). A p-value < 0.05 was used to determine statistical significance. No adjustments for multiplicity of testing were made.

Results

Serum retinol-binding protein 4 is present in ATTR cardiac tissue

Cardiac tissues from 4 separate cases of ATTRm amyloidosis were studied; these included ATTR-L58H (n = 2), ATTR-V30M (n = 1), and ATTR-V122I (n = 1). The presence of amyloid deposits in all cardiac tissue specimens was confirmed by Congo red staining. Representative results from a patient who was diagnosed with ATTR-L58H amyloidosis, an ATTRm disease featuring cardiac presentation,[31], are shown in Figure 1. In the Congo red treated section, stain-avid areas were observed by standard light microscopy (Figure 1a); a polarized view of the same field (Figure 1b) exhibited the characteristic ‘apple green’ birefringence, confirming the presence of amyloid. Immunohistochemical treatment of a serial section with polyclonal anti-human TTR antibody (Figure 1c) demonstrated strong reactivity in the area that stained with Congo red, providing evidence that the amyloid deposits were constituted from TTR protein. Testing for RBP4 showed strong cytoplasmic staining in cardiomyocytes adjacent to the amyloid deposits (Figure 1d).

RBP4 histological and immunohistochemical features in ATTRwt cardiac tissue samples (n=3) were also characterized. Congo red treatment provided evidence of extensive amyloid deposition in the myocardium and myocardial vasculature under standard (Figure 1e) and polarized light (Figure 1f). Immunohistochemical analyses of serial sections showed strong reactivity for TTR confirming the biochemical nature of the amyloid deposits (Figure 1g). Staining for RBP4 demonstrated the presence of RBP4 protein in the cardiomyocytes surrounding amyloid infiltrated vasculature (Figure 1h); however, the RBP4 staining in ATTRwt tissues was weaker and less uniform compared to the immunoreactivity observed in ATTRm heart samples (Supplemental Figure 1).

For comparison, autopsied heart tissue sections from a patient with cardiomyopathy in the absence of amyloid were analyzed. Congo red treatment of the tissue was negative when viewed by standard and polarized light microscopy, confirming that no amyloid fibrillar deposits were present (Supplemental Figure 2a, b). Moreover, there was no evidence of RBP4 in the tissue by immunohistochemical staining (Supplemental Figure 2c) indicating that the protein was absent in cardiac tissue lacking amyloid infiltration. Liver tissue sections from an AL (non-ATTR) amyloidosis case served as positive staining controls (Supplemental Figure 2d) since RBP4 is abundantly synthesized by hepatocytes; as expected, strong RBP4 staining was observed in the liver tissue and confirmed antibody specificity.

Serum concentrations of retinol-binding protein 4 in ATTR amyloidosis

To investigate the levels of RBP4 in amyloid patient sera and controls, we developed an ELISA method for reproducible quantification. We assessed serum levels of RBP4 in males over 60 years of age from the following groups: 1) healthy controls, 2) ATTRm amyloidosis featuring cardiomyopathy (CMP), 3) ATTRwt amyloidosis, 4) AL (non-ATTR) amyloidosis with CMP, and 5) non-amyloid with CMP. The ATTRm group consisted of patients heterozygous for the following TTR mutations: T60A (n = 11), V122I (n = 8), and individual cases of D18G, V20I, G47E, L68L, and S77Y. For our comparative analyses, we also chose to investigate RBP4 serum concentrations in patients with cardiomyopathy unrelated to ATTR amyloidosis, i.e. AL cardiac amyloidosis and non-amyloidotic cardiomyopathy. Control and patient serum RBP4 data are detailed in Figure 2. RBP4 was significantly decreased in ATTRm compared to healthy controls (29.11 µg/mL, p = 0.0011) and all patient groups, including ATTRwt (29.11 µg/mL, p < 0.0001). Conversely, RBP4 levels in patients with ATTRwt were comparable to controls and not significantly different from the AL (non-ATTR) or non-amyloid CMP groups.
Baseline demographic, serological and echocardiographic data in the cardiac amyloid patient groups (ATTRm, ATTRwt, and AL) are presented as mean values in Table 1A; analysis of variation among the groups is reflected in the p-values. Serum RBP4 levels (highlighted in gray) were significantly different in the group comparison with the lowest levels observed in ATTRm (29.11 ug/mL); ATTRwt concentrations were slightly lower than AL (42.83 vs. 47.84 ug/mL). Serum TTR levels were significantly different (p = 0.0020) in ATTRm, ATTRwt, and AL; consistent with the RBP4 data, the lowest levels of TTR were found in the ATTRm group (0.18 mg/mL). While elevated levels of cardiac biomarkers were present in ATTR and AL, there was significant variation in BNP (p < 0.0001) and cTn-I values (p < 0.0244) among the groups. Although the mean age was higher in the ATTRwt patient cohort, other measures including BMI, triglycerides, uric acid, and BMI were comparable among the cardiac amyloid groups.

To assess relationships between baseline RBP4 serum concentrations and other previously reported measures associated with survival in ATTR [30], correlation coefficients were calculated and adjusted for age and triglyceride-status prior to analysis. RBP4 and TTR were correlated in both ATTR amyloid diseases (Table 1B) as expected. More interestingly, a strong association between RBP4 and c-TnI was identified solely in the ATTRwt cohort (Pearson r = 0.3716, two-tailed p-value = 0.035). No other significant correlations with RBP4 were noted.

Discussion

This report documents the results of our histological and serological analyses of RBP4 in TTR-associated forms of cardiac amyloidosis. We provide evidence that RBP4 is present in cardiac tissues from patients with ATTRm and ATTRwt amyloidosis, although not associated with TTR in the amyloid fibril deposits. Our study of serum RBP4 shows that significantly lower levels are present in patients with ATTRm compared to ATTRwt (p < 0.0001) and healthy controls (p < 0.0011). Moreover, in a correlation analysis, we find that there is significant correlation between circulating RBP4 concentration and cardiac troponin-I (c-TnI) in patients with ATTRwt amyloidosis at baseline evaluation.

Previous in vitro studies[21,22] have provided evidence that RBP4 can extend the aggregation kinetics of thermally-denatured TTR, implying that RBP4 stabilizes the native conformation of TTR. Based on these findings, we theorized that TTR found in amyloid deposits would no longer be bound to its natural and stabilizing binding partner, RBP4. To investigate this hypothesis, we analyzed autopsied ATTR cardiac samples, heavily infiltrated with TTR amyloid, for the presence of RBP4. Congo red treatment of ATTRm and ATTRwt heart sections confirmed the presence of amyloid deposits in all tissues and immunohistochemical testing conclusively identified TTR as the amyloid fibril protein. Using antibodies specific for RBP4, the secreted form of the protein, immunoreactivity was observed in intracellular regions adjacent to amyloid deposits in ATTRm and ATTRwt cardiac tissue sections. The intracellular RBP4 appeared to be specifically concentrated in regions of the tissue surrounding amyloid laden blood vessels (Figure 1); there was no evidence of RBP4 in the extracellular amyloid deposits. In our study, the presence of RBP4 in cardiac tissue was a finding unique to the ATTR samples, as testing of heart specimens from non-amyloid cases of cardiomyopathy showed no evidence of the protein.

While RBP4 was observed in all ATTR cardiac sections, the staining profiles of ATTRm and ATTRwt samples were dissimilar. In the ATTRm samples, immunostaining was more abundant and of greater intensity compared to less pronounced results observed in the ATTRwt specimens. These differences may reflect weaker interactions between RBP4 and variant forms of TTR, i.e. decreased binding affinities specifically between RBP4 and L58H, V30M, or V122I. It has previously been reported that interactions in contact areas between RBP4 and TTR are highly sensitive to even subtle conformational changes in either protein[32,33]. Thus, it is possible that TTR mutant proteins in general, or those specifically associated with ATTR cardiomyopathy, feature structural configurations that decrease the binding affinity of the protein to RBP4 through steric hindrance, electrostatic or hydrophobic interference, or other factors that limit interactions. In this scenario, TTR free of RBP4 in the circulation of patients with ATTR amyloidosis would be available and could deposit in heart tissue more readily.

In the serological analysis (Figure 2), data from ELISA measurement of serum RBP4 showed significant differences between ATTRm and ATTRwt cases featuring cardiomyopathy (29.11 vs. 42.83 g/mL, p < 0.0001). In fact, circulating RBP4 levels were uniquely decreased in ATTRm compared to age- and gender-matched groups that included healthy controls, and cases of AL cardiac amyloidosis or non-amyloid cardiomyopathy. These data are consistent with a recently reported study by Arvanitis et al. showing that RBP4 levels in ATTR V122I amyloidosis were significantly lower compared with non-amyloid controls[34]. In ATTRm amyloidosis, it seems reasonable that lowered RBP4 levels correlate to increased amyloid fibril formation more likely to occur. Our immunohistochemical analysis demonstrating the absence of RBP4 in ATTRm cardiac amyloid deposits (Figure 1a-d) lends support to this supposition. It is important to note that serum concentrations of TTR were also lowest in the ATTRm group; whether this is due to increased amyloid deposition or an alternative cause is unclear. Conversely, RBP4 levels in the ATTRwt group were an unexpected finding. Concentrations were comparable to the healthy controls and not significantly different from the other patient groups (AL cardiac and non-amyloid cardiomyopathy). Wild-type TTR, unlike the mutant forms, contains a ‘normal’ amino acid sequence and higher
Our immune histochemical and serological results suggest that the role of RBP4 in the mechanism of TTR amyloidogenesis differs in ATTRm and ATTRwt cardiac amyloid diseases. In ATTRm amyloidosis, weakened TTR-RBP4 associations may result in lower levels of serum RBP4 and a higher degree of TTR amyloid deposition in cardiac tissue. In contrast, a tighter interaction between wild-type TTR and RBP4 would explain higher amounts of the circulating complexes in ATTRwt cardiac amyloidosis. Studies published over the last thirty years have demonstrated the existence of several RBP4-specific receptors in a variety of tissues including the choroid plexus, testes, kidney, and eye. One such receptor, STRA6, allows RBP4 to transfer vitamin A across cell membranes for binding to intracellular isoforms of RBP (CRBP) [36–40]. A similar receptor, with specificity for TTR-RBP4 complexes, may be present in cardiac tissue. In a cascade of events, conformational changes in RBP4 resulting from interactions with a cardiac cell surface receptor would lead to weakening of the TTR-RBP4 interactions, destabilization of tetrameric TTR, subunit dissociation, and initiation of the amyloid fibril formation pathway.

The present study included several factors influencing the results. Linear and nonlinear correlation analyses were performed to evaluate the association between biomarkers and clinical outcomes. A variety of statistical tools were used, including t-tests, ANOVA, and regression analysis. The results showed significant associations between serum RBP4 concentrations and clinical outcomes, such as survival and disease progression. The study also included a comparison between ATTRm and ATTRwt cardiac amyloidosis, revealing differences in the pathobiology of the two diseases. The data suggest that RBP4 may be a biomarker for ATTRm cardiac amyloid disease.

**Table 1A. Baseline demographic, serological, and echocardiographic data in cardiac amyloidosis**

| Characteristic | ATTRm | ATTRwt | AL | p-value |
|---------------|-------|--------|----|---------|
| N             | 25    | 34     | 27 |         |

Table 1B. Associations between baseline serum RBP4 concentrations and predictors of reduced survival in ATTR amyloidosis

| Predictor | ATTRm | ATTRwt |
|-----------|-------|--------|
| Age, y    | 0.17  | 0.27   |
| BMI, kg/m²| 0.14  | 0.20   |
| Triglycerides, mg/dL | 0.13  | 0.09   |
| Uric acid, mg/dL | -0.002| 0.01   |
| BNP, pg/mL  | 0.13  | 0.01   |
| c-TnI, ng/mL | -0.05 | 0.001  |
| TTR, mg/mL  | 0.53  | 0.01   |
| IVST, mm   | 0.29  | 0.12   |
| LVEF, %    | 0.26  | 0.10   |

Pearson correlation coefficients (p-values) for each analysis are shown; ATTRm, mutant TTR-associated cardiac amyloid disease; ATTRwt, wild-type TTR-associated cardiac amyloid disease; AL, immunoglobulin light chain-associated cardiac amyloid disease

* p < 0.05, ** p < 0.01

order structure that favors interactions with RBP4; hence, elimination of RBP4, a relatively small protein, through renal filtration should be prevented and result in higher circulating levels of RBP4. A stronger molecular association between RBP4 and wild-type TTR, compared to mutant forms of the protein, would be consistent with our observation of lower amounts of RBP4 in ATTRwt cardiac tissue specimens. While the analysis of serum RBP4 concentrations does not provide direct information about RBP4-mediated TTR stabilization, these data are in line with previous reports and our in vitro findings (data not shown) that RBP4 stabilizes TTR and slows the aggregation kinetics of the amyloidogenic protein.

Our immune histochemical and serological results suggest that the role of RBP4 in the mechanism of TTR amyloidogenesis differs in ATTRm and ATTRwt cardiac amyloid diseases. In ATTRm amyloidosis, weakened TTR-RBP4 associations may result in lower levels of serum RBP4 and a higher degree of TTR amyloid deposition in cardiac tissue. In contrast, a tighter interaction between wild-type TTR and RBP4 would explain higher amounts of the circulating complexes in ATTRwt amyloidosis, but suggests that factors other than TTR-RBP4 pairing must contribute to amyloid fibril formation in this disease. It seems plausible that the cardiac tissue environment may contain key features, yet to be defined, that contribute to the pathobiological dissimilarities we have observed in ATTRm vs. ATTRwt. In addition to differences in TTR-RBP4 pairing associations, variations in receptor-mediated targeting and/or cell signaling pathways may be important, especially in ATTRwt amyloidosis. Studies published over the last thirty years have demonstrated the existence of several RBP4-specific receptors in a variety of tissues including the choroid plexus, testes, kidney, and eye. One such receptor, STRA6, allows RBP4 to transfer vitamin A across cell membranes for binding to intracellular isoforms of RBP (CRBP) [36–40]. A similar receptor, with specificity for TTR-RBP4 complexes, may be present in cardiac tissue. In a cascade of events, conformational changes in RBP4 resulting from interactions with a cardiac cell surface receptor would lead to weakening of the TTR-RBP4 interactions, destabilization of tetrameric TTR, subunit dissociation, and initiation of the amyloid fibril formation pathway.

The present study was limited by small sample sizes in both the histological and serological analyses. Several constraining factors included the study selection criteria which were age > 60 years, male sex, and exclusion of all patients participating in ATTR clinical trials or taking investigational drugs at the time of blood draw. In addition, while sera are routinely collected at all baseline evaluations and readily available from our repository, cardiac tissues were restricted to a small set of autopsy specimens. Expansion of these studies to larger group numbers would confirm our findings and further demonstrate the utility of RBP4 as a biomarker of ATTRm cardiac amyloid disease.

Moreover, this study did not include examination of serum vitamin A levels, potentially important as retinol is a key player in the recognition and binding of RBP4 to TTR [4, 21, 41]. Unfortunately, accurate measurement of vitamin A is challenging as levels are homeostatically controlled [50–200 µg/dL] and serum retinol concentrations are only an indirect reflection of total vitamin A in the liver. Obtaining these data in future studies may be key as normal homeostasis of vitamin A is interrupted in several conditions where decreased levels (< 0.7 µg/dL) have been reported and retinoid insufficiency has been linked to alterations in the extracellular matrix of the heart which can affect adult cardiac function.

**Conclusion**

In summary, results of our study of cardiac tissues and sera from patients with ATTRm or ATTRwt amyloidosis suggest that the pathobiologies of the two diseases are dissimilar with respect to RBP4. We demonstrate, for the first time, that RBP4 is uniquely present in

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cardiac tissues from patients with TTR-associated cardiac amyloidosis. While not co-deposited with TTR fibrils, RBP4 appeared to be present in locations adjacent to the cardiac amyloid deposits more abundantly in ATTRm tissues compared to ATTRwt specimens. Moreover, circulating levels of RBP4 were significantly lower in ATTRm compared to ATTRwt (p < 0.0001) and healthy controls (p < 0.0011). Lastly, we found a correlation between circulating RBP4 levels and cardiac troponin-I (c-TnI) in ATTRwt, as well as expected strong correlations between RBP4 and TTR in both ATTRm and ATTRwt. These data suggest that RBP4 may have utility as a serum biomarker in ATTRm amyloidosis and provide evidence in support of a therapeutic strategy that takes advantage of the stabilizing effects of a natural binding partner of TTR.

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Conflict of interest

All authors declare no conflict of interest

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