

STATE-OF-THE-ART REVIEW

Cardiac Troponin in Patients With Light Chain and Transthyretin Cardiac Amyloidosis



JACC: CardioOncology State-of-the-Art Review

Laura De Michieli, MD,^{a,b} Alberto Cipriani, MD,^{a,c} Sabino Iliceto, MD,^{a,c} Angela Dispenzieri, MD,^d Allan S. Jaffe, MD^{b,e}

ABSTRACT

Cardiac amyloidosis (CA) is an infiltrative disease caused by amyloid fibril deposition in the myocardium; the 2 forms that most frequently involve the heart are amyloid light chain (AL) and amyloid transthyretin (ATTR) amyloidosis. Cardiac troponin (cTn) is the biomarker of choice for the detection of myocardial injury and is frequently found to be elevated in patients with CA, particularly with high-sensitivity assays. Multiple mechanisms of myocardial injury in CA have been proposed, including cytotoxic effect of amyloid precursors, interstitial amyloid fibril infiltration, coronary microvascular dysfunction, amyloid- and non-amyloid-related coronary artery disease, diastolic dysfunction, and heart failure. Regardless of the mechanisms, cTn values have relevant prognostic (and potentially diagnostic) implications in both AL and ATTR amyloidosis. In this review, the authors discuss the significant aspects of cTn biology and measurement methods, potential mechanisms of myocardial injury in CA, and the clinical application of cTn in the management of both AL and ATTR amyloidosis. (J Am Coll Cardiol CardioOnc 2024;6:1-15) © 2024 The Authors. Published by Elsevier on behalf of the American College of Cardiology Foundation. This is an open access article under the CC BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).

Cardiac amyloidosis (CA) is an infiltrative disease caused by the deposition of amyloid fibrils in the extracellular matrix, progressively leading to organ damage and dysfunction.¹ CA now is an increasingly recognized cause of heart failure, particularly with preserved ejection fraction.² The 2 main types of amyloidosis that can involve the heart, amyloid light chain (AL) and amyloid transthyretin (ATTR) amyloidosis, are pathophysiologically different. In AL amyloidosis, the amyloid precursor proteins are immunologic light chains

(LCs) most commonly produced by a plasma cell clone. In ATTR amyloidosis, either variant (hereditary) ATTR (ATTRv) amyloidosis or wild-type ATTR (ATTRwt) amyloidosis,³ the precursor protein is ATTR, which is produced predominantly by the liver and serves as a carrier for thyroxine and retinol. Although the disease management, course, and prognosis of AL and ATTR CA are different, at present, effective treatment options exist for both.^{4,5} The efficacy of these treatment options is dependent on disease stage at the time of diagnosis, determined

From the ^aDepartment of Cardiac, Thoracic and Vascular Sciences and Public Health, University of Padua, Padua, Italy; ^bCardiovascular Department, Mayo Clinic and Medical School, Rochester, Minnesota, USA; ^cCardiology Unit, University Hospital of Padua, Padua, Italy; ^dDivision of Hematology, Mayo Clinic, Rochester, Minnesota, USA; and the ^eDepartment of Laboratory Medicine and Pathology, Mayo Clinic and Medical School, Rochester, Minnesota, USA.

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**ABBREVIATIONS
AND ACRONYMS**

AL	= amyloid light chain
ATTR	= amyloid transthyretin
ATTRv	= variant (hereditary) amyloid transthyretin
ATTRwt	= wild-type amyloid transthyretin
CA	= cardiac amyloidosis
CAD	= coronary artery disease
cTn	= cardiac troponin
hs-cTn	= high-sensitivity cardiac troponin
LC	= light chain
LV	= left ventricle/ventricular
NT-proBNP	= N-terminal pro-brain natriuretic peptide
sFLC	= serum free light chain

mainly by the severity of cardiac involvement,³ which can be assessed using cardiac biomarkers and imaging techniques. Cardiac troponin (cTn) is the biomarker of choice for the detection of myocardial injury,⁶ and it is frequently found to be elevated in patients with CA, in particular when high-sensitivity assays are used. The extent of abnormalities in cTn is of crucial prognostic significance in these patients.³ In this review, we elucidate the significant aspects of cTn biology, measurement, and interpretation when it is used adjunctively to assist in the management of patients with CA. The multiplicity of possible causes for myocardial injury in these patients are addressed, with particular focus on the relevant pathophysiological mechanisms and common clinical scenarios.

cTn: BASIC CONCEPTS

Since the development of cTn assays in the late 1980s and early 1990s, this marker has assumed a progressively more predominant role in cardiovascular care. It is currently the biomarker of choice for the detection of myocardial injury and for the diagnosis of myocardial infarction.⁶ It is significantly more specific than prior markers and much more sensitive, which allows the identification of many new processes that can damage cardiomyocytes.

The troponin complex consists of 3 different proteins, troponin C, troponin I, and troponin T, which are encoded by separate genes and are essential for proper excitation and contraction in cardiomyocytes (Figure 1).⁷ Because of their cardiospecific isoform expression, cTnI and cTnT can be used as highly specific markers of myocardial damage.⁸ Recent data suggest that the re-expression of cTnT (gene *TNNT2*) in patients with active myopathies and myositis can contribute to cTnT elevations in some patients.^{9,10} At present, there is no evidence of up-regulation or re-expression of cTnI in diseased skeletal muscle.¹⁰ cTn elevations occur not only in acute ischemic disease with overt necrotic cardiac damage but in many other situations (Table 1).⁶

Over time, assays for cTnT and cTnI measurements have improved in terms of analytical performance. The original assays were insensitive, but present-day high-sensitivity cTn (hs-cTn) assays provide increased sensitivity and greater precision at low concentrations, so that they are able to identify patients at higher risk for adverse events in multiple situations.¹¹ Analytical terminology and characteristics of hs-cTn assays are reported in Table 2.^{12,13}

HIGHLIGHTS

- Two forms of cardiac amyloidosis (CA), AL and ATTR affect the heart.
- Therapies are available for both forms of CA.
- High sensitivity cTn a marker of myocardial injury is often elevated with CA.
- Many mechanisms for myocardial injury in CA require recognition and consideration.
- cTn is diagnostic and prognostic aid in CA and in monitoring therapy.

The correlation between values measured with contemporary (non-high-sensitivity) cTn assays and hs-cTn assays is good at higher levels but poor at low concentrations, especially with hs-cTnT.¹⁴

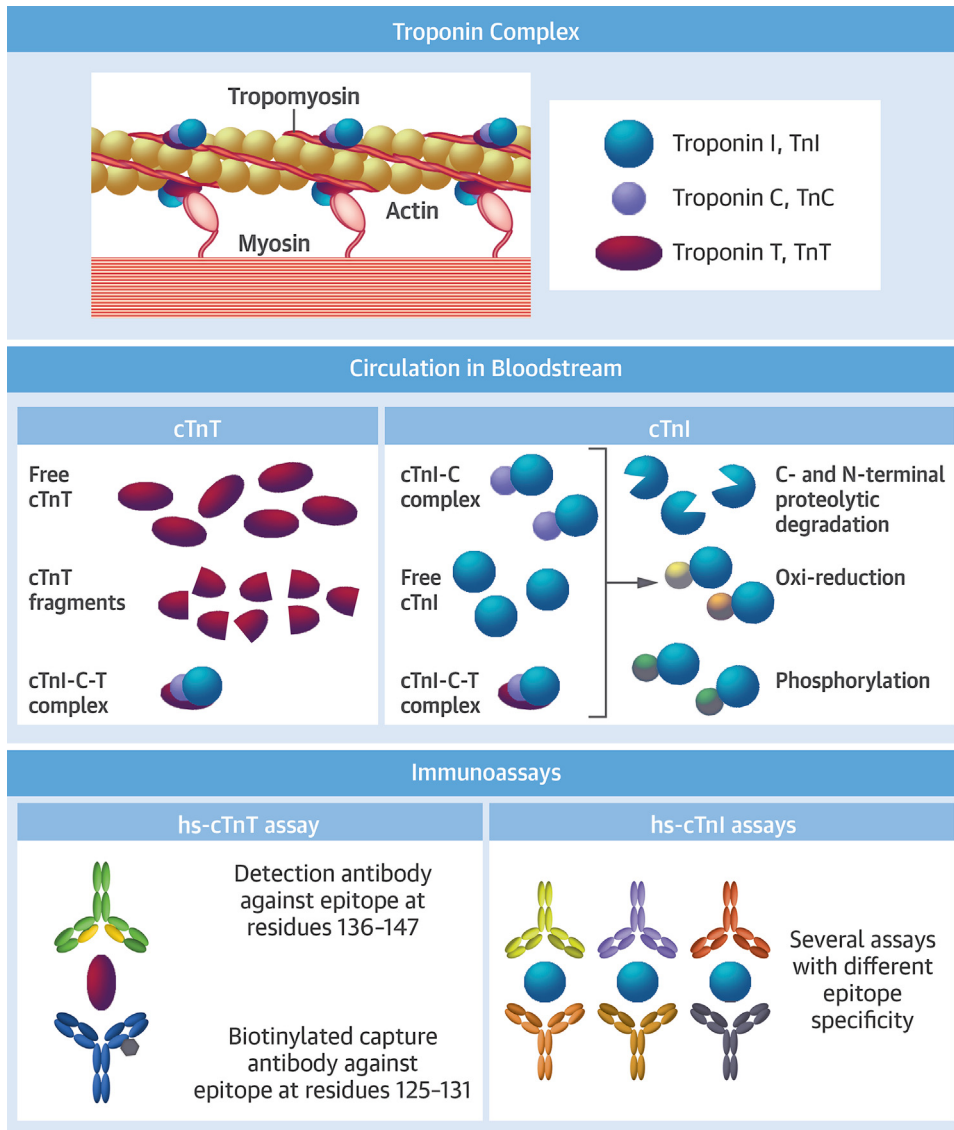
Although only 1 hs-cTnT assay is in use for clinical practice, many hs-cTnI assays have been developed; the same is true for older contemporary assays. Because of the different antibodies used in each assay and the unique mix of cTnI forms in each patient sample (Figure 1), there is no harmonization between the different hs-cTnI assays, and absolute values cannot be directly compared.¹⁵ Analytical characteristics of the different hs-cTn assays available on the market are reported in Table 3.

Myocardial injury is defined as a cTn value greater than the sex-specific 99th percentile upper reference limit,⁶ regardless of cause, and it is frequently present in patients with primary cardiovascular disease as well as those whose primary problems can be noncardiac conditions such as critical illness.¹⁶ Its prognostic relevance has been established in multiple clinical scenarios.^{5,16-20} With the transition from conventional to hs-cTn assays, the frequency of the detection of myocardial injury has increased,²¹ leading to a more precise stratification of patients at higher risk for adverse events but also escalating the challenge of identifying the reasons of cTn elevations in each patient. In some clinical situations, even values less than the 99th percentile upper reference limit have prognostic information, explaining why cTn helps in risk stratification when analyzed as a continuous variable and not only when relying on thresholds to define a normal range.^{22,23}

MYOCARDIAL INJURY IN CA

Since the 2000s, it has been appreciated that almost invariably, cTn levels are elevated in patients with CA.²⁴ Such elevations are common and manifest

FIGURE 1 Cardiac Troponin Structure, Release in Bloodstream, and Methodologies of Measurement



After a myocardial infarction, cardiac troponin I (cTnI) circulates in the blood mainly as a binary complex with cardiac troponin C, but also as free cTnI and a ternary complex, cTnI-C-T. Various molecular modifications and protein fragmentations have been described. cTnT circulates mainly in a free form, but cardiac troponin T (cTnT) fragments as well as the cTnI-C-T complex are also present. The high-sensitivity cTnT (hs-cTnT) assay uses fragment antigen binding portions of 2 cTnT-specific mouse monoclonal antibodies. For the fifth-generation assay, the original antibody has been re-engineered to produce a mouse-human chimeric detection antibody. Several high-sensitivity cTnI (hs-cTnI) assays are approved for clinical use, with a lack of harmonization related to different aspects, starting with the difference in epitope specificity of antibodies produced by different companies. Indeed, immunoassays that use anti-cTnI monoclonal antibodies are dependent on the epitope regions recognized by the antibodies incorporated into each assay. Moreover, cTnI-related characteristics such as biochemical modifications (eg, degradation, phosphorylation, oxide reduction) and different circulating complexes can lead to altered signal generation of sandwich-type cTnI immunoassays that use antibodies directed against these modified regions.¹⁵ Reproduced with permission from Brush et al.¹⁰⁰

TABLE 1 Possible Causes of Myocardial Injury According to the Fourth Universal Definition of Myocardial Infarction	
Myocardial injury related to acute myocardial ischemia	Atherosclerotic plaque disruption with thrombosis
Myocardial injury related to acute myocardial ischemia because of oxygen supply/demand imbalance	Reduced myocardial perfusion <ul style="list-style-type: none"> • Coronary artery spasm, microvascular dysfunction • Coronary embolism • Coronary artery dissection • Sustained bradyarrhythmia • Hypotension or shock • Respiratory failure • Severe anemia Increased myocardial oxygen demand <ul style="list-style-type: none"> • Sustained tachyarrhythmia • Severe hypertension with or without left ventricular hypertrophy
Other causes of myocardial injury	Cardiac conditions <ul style="list-style-type: none"> • Heart failure • Myocarditis • Cardiomyopathy (any type) • Takotsubo syndrome • Coronary revascularization procedure • Cardiac procedure other than revascularization • Catheter ablation • Defibrillator shocks • Cardiac contusion Systemic conditions <ul style="list-style-type: none"> • Sepsis, infectious disease • Chronic kidney disease • Stroke, subarachnoid hemorrhage • Pulmonary embolism, pulmonary hypertension • Infiltrative diseases (eg, amyloidosis, sarcoidosis) • Chemotherapeutic agents • Critically ill patients • Strenuous exercise
Ischemic and nonischemic causes of myocardial injury are shown. In each patient with cardiac amyloidosis, myocardial injury can be due to the amyloid infiltration process, but other concomitant factors (such as arrhythmias or hypotension) can also be present. Adapted with permission from Thygesen et al. ⁵	

important prognostic implications (discussed later). The exact frequency of myocardial injury is difficult to determine definitively and likely relates to differences in the populations studied, differences in the stages of the disease process, and the fact that so many different hs-cTn assays are available with unique cutoffs to define myocardial injury. Nevertheless, multiple pathophysiological mechanisms of

myocardial injury in patients with CA have been proposed (**Central Illustration**). It is important for clinicians to recognize that increases in hs-cTn values may have a multiplicity of causes, ischemic or non-ischemic (**Table 1**); in patients with amyloid infiltration, hs-cTn values reflect the extent of myocardial involvement while also being potentially influenced by all these other causes.

TABLE 2 Analytical Terminology to Describe hs-cTn Assay Performance	
Term	Description
Limit of blank	"Noise" signal inherent in the analytical system 95th percentile of analytical signal when no cTn is present, only matrix
Limit of detection	Lowest concentration detectable in 95% of measurements Imprecision at the limit of detection is often high High-sensitivity assays should measure cTn above the limit of detection in $\geq 50\%$ of healthy subjects
Limit of quantitation	Lowest cTn concentration that can be reported as a number with specified certainty (typically 20% imprecision)
Coefficient of variation	A measure of assay imprecision at any given concentration Coefficient of variation should be $\leq 10\%$ at the 99th percentile upper reference limit for hs-cTn assays
99th percentile upper reference limit	Universally endorsed as the reference cutoff to aid in the diagnosis of acute myocardial infarction. Key components: <ul style="list-style-type: none"> • The 99th percentile should be determined for each assay in a healthy population (≥ 300 male and 300 female subjects) • The 99th percentile for hs-cTn assays should be measured with an analytical imprecision of $\leq 10\%$ • The 99th percentile values should be reported as whole numbers only, in nanograms per liter
Analytical terminology to properly understand and describe the performance of hs-cTn assays is shown, as reported in Table 3 . cTn = cardiac troponin; hs-cTn = high-sensitivity cardiac troponin.	

TABLE 3 Analytical Characteristics and Reference Values of Available High-Sensitivity Assays

Company/Platform/Assay	LoB, ng/L	LoD, ng/L	99th Percentile, ng/L	CV at 99th Percentile
Abbott/Alinity i systems/Alinity i STAT High Sensitive Troponin-I; commercial (OUS)	1.0	1.6	Overall: 26.2 F: 15.6 M: 34.2	Overall: 4.0% F: 5.3% M: 3.5%
Abbott/ARCHITECT i systems/ARCHITECT STAT High Sensitive Troponin-I; commercial (U.S.)	0.9	1.7	Overall: 28 F: 17 M: 35	Overall: 4.3% F: 5.0% M: 4.1%
Abbott/ARCHITECT i systems/ARCHITECT STAT High Sensitive Troponin-I; commercial	0.7-1.3	1.1	Overall: 26.2 F: 15.6 M: 34.2	Overall: 4.0% F: 5.3% M: 3.5%
Beckman Coulter/Access 2, Dxl/Access hsTnI; commercial (OUS)	0.0-1.7	1.0-2.3	Overall: 17.5 F: 11.6 M: 19.8	Overall: 3.7% F: 4.2% M: 3.6%
Beckman Coulter/Access 2/Access hsTnI; commercial (U.S.): LiHep plasma	0.0-0.8	1.0-2.0	Overall: 17.5 F: 11.6 M: 19.8	Overall: 3.7% F: 4.2% M: 3.6%
Beckman Coulter/Access 2/Access hsTnI; commercial (U.S.): serum	0.0-0.8	1.0-2.0	Overall: 18.2 F: 11.8 M: 19.7	Overall: 6.0% F: 6.9% M: 5.8%
Beckman Coulter/Dxl, Access hsTnI; commercial (U.S.): LiHep plasma	0.0-1.7	1.5-2.3	Overall: 17.9 F: 14.9 M: 19.8	Overall: 5.2% F: 5.6% M: 5.0%
Beckman Coulter/Dxl, Access hsTnI; commercial (U.S.): serum	0.0-1.7	1.5-2.3	Overall: 18.1 F: 13.6 M: 19.8	Overall: 6.2% F: 6.5% M: 6.1%
BioMérieux VIDAS High Sensitive Troponin I; commercial	1.9	3.2	Overall: 19 F: 11 M: 25	7.0%
ET Healthcare Pylon hsTnI assay; China, FDA approved	0.8	1.2-1.4	Overall: 27 F: 21 M: 27	10%
ET Healthcare Pylon hsTnT; research	0.4	0.8	Overall: 13 F: 13 M: 14	4%
Fujirebio Lumipulse G G1200 and G600II hsTnI	1.2	2.1	Overall: 28.6 F: 22.4 M: 32.9 Serum overall: 26.9 F: 21.4 M: 29.4 LiHep plasma overall: 29.6 F: 27.8 M: 32.8	≤4.6%
LSI Medience PATHFAST cTnI; commercial	NP	1	Overall: 15.46 M: 16.91 F: 11.46	<6%
LSI Medience PATHFAST hs-cTnI/PATHFAST cTnI-II	1.23	2.33	Overall: 27.9 F: 20.3 M: 29.7	6.1%
Ortho/VITROS/hsTroponin I; commercial	0.14-0.51	0.39-0.86	Serum overall: 11 F: 9 M: 12 LiHep plasma overall: 11 F: 9 M: 13	<10%
Quidel/Alere TriageTrue hs-cTnI	0.4 (plasma) 0.5-0.8 (whole blood)	0.7-1.6 (plasma) 1.5-1.9 (whole blood)	Overall: 20.5 F: 14.4 M: 25.7	5.0%-5.9% at 21 ng/L (plasma) 5.9%-6.5% at 22 ng/L (whole blood)
Roche/cobas e601, e602, e411/cTnT-hs 18-min; commercial	2.53 (1.58 for e411)	3.16 (2.54 for e411)	Overall: 14 F: 9 M: 17	<10%
Roche/cobas e601, e602, e411/cTnT-hs STAT; commercial	2.36 (2.14 for e411)	2.85 (3.25 for e411)	Overall: 14 F: 9 M: 17	<10%
Roche/cobas e801/e402 cTnT-hs 18 min and 9 min; commercial	18 min: 2.21 9 min: 1.91	18 min: 2.97 9 min: 2.72	Overall: 14 F: 9 M: 17	<10%
Roche/cobas e601, e602, e411 cTnT Gen 5 STAT; commercial	2.5 (3 for e411)	3 (5 for e411)	Overall: 19 F: 14 M: 22	<10%

Continued on the next page

TABLE 3 Continued

Company/Platform/Assay	LoB, ng/L	LoD, ng/L	99th Percentile, ng/L	CV at 99th Percentile
Roche/cobas e801/cTnT Gen 5-9 min; commercial	2.5	3	Overall: 19 F: 14 M: 22	<10%
Siemens ATELLICA High-Sensitivity TnI (TnIH) (U.S. and OUS); commercial ^a	0.50	1.6	Overall: 45.4 F: 38.6 M: 53.5	<4.0%
Siemens ATELLICA VTLi hs-cTnI ^a	0.55	1.2 (plasma) 1.6 (whole blood)	Overall: 22.9 F: 18.5 M: 27.1	6.5% (plasma) 6.1% (whole blood)
Siemens ADVIA Centaur XP/XPT/CP High-Sensitivity TnI (TNIH) (U.S. and OUS); commercial ^{a,b}	0.50	1.6	Overall: 46.5 F: 39.6 M: 58.0	<4.9%
Siemens Dimension VISTA High Sensitivity TnI (TNIH) (OUS); commercial ^a	1.0	2.0	Overall: 58.9 F: 53.7 M: 78.5	<5.0%
Siemens Dimension ExL High Sensitivity TnI (TNIH) (OUS); commercial ^a	1.1	2.7	Overall: 60.4 F: 51.4 M: 76.2	<5.0%
Tosoh CL AIA-PACK cTnI; commercial	NP	1.61	Overall: ≤24 (Asian), ≤31 (Caucasian)	NP

Relevant analytical characteristics of high-sensitivity troponin assays available on the market inside and outside of the United States. Data were extracted from the International Federation of Clinical Chemistry and Laboratory Medicine Task Force on Clinical Applications of Cardiac Bio-Markers (version 052023). All data are listed as provided by the manufacturers. ^aManufacturers may have submitted assays they claim to be "high sensitivity" that do not meet the International Federation of Clinical Chemistry and Laboratory Medicine requirements of <10% CV at the 99th percentile and ≥50% measurable concentrations greater than or equal to the LoD for both men and women. ^bThe products and features mentioned herein are not commercially available in all countries.

CV = coefficient of variation; FDA = U.S. Food and Drug Administration; LiHep = lithium heparin; LoB = limit of blank; LoD = limit of detection; NP = not provided; OUS = outside the United States; F = Female; M = Male.

DIRECT CYTOTOXIC EFFECT OF AMYLOID PRECURSORS.

Amyloid precursors can have direct toxic effects on cardiomyocytes. In AL amyloidosis, the proteotoxicity of LCs has been extensively reported.²⁵ Interestingly, infusion of LCs from patients with severe AL amyloidosis with cardiac involvement causes marked impairment of ventricular relaxation on isolated mouse heart models.²⁶ Moreover, human amyloid LCs alter the cellular redox state in isolated cardiomyocytes.²⁷ This results in direct impairment of cardiomyocyte contractility and relaxation, independent of fibril deposition, associated with alterations in intracellular calcium handling.²⁷ Amyloidogenic LCs have been reported to provoke oxidative stress, cellular dysfunction, and apoptosis in isolated adult cardiomyocytes through activation of p38 mitogen-activated protein kinase,²⁸ which also mediates the transcription of brain natriuretic peptide.²⁵ A study performed on human cardiac fibroblasts and *Caenorhabditis elegans* reported a correlation between the overall conformational properties of native folded proteins (including flexibility, kinetic instability, and dynamic state) and the proteotoxicity of cardiotoxic LCs.²⁹ In vitro experiments suggest that whereas LC-derived amyloid fibrils exhibit inhibition of the cell growth and division, soluble LC proteins allow cell growth but cause cellular

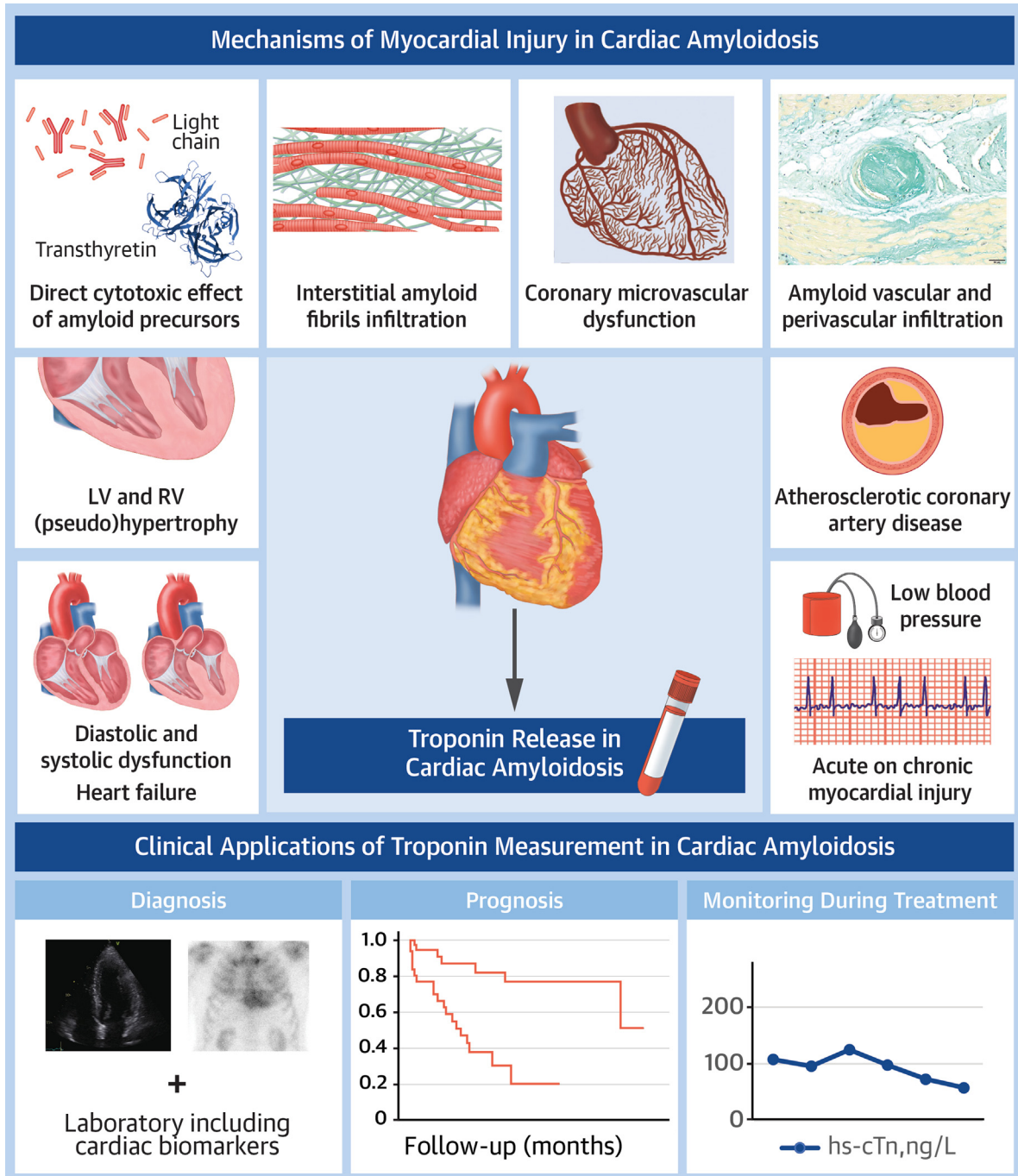
dysfunction and apoptosis in cardiomyocytes, suggesting that the mechanisms of cytotoxicity differ between soluble proteins and amyloid fibrils.^{25,30} Clinically, serum free LCs (sFLCs), particularly lambda, significantly correlate reasonably but not exactly with increases in cTnI and N-terminal-pro-brain natriuretic peptide (NT-proBNP) as well as echocardiographic parameters.^{31,32} Thus, the value of sFLCs provides additional prognostic information in conjunction with cTn and NT-proBNP, and therefore, they have been integrated into multi-parametric staging systems.^{33,34}

In ATTR CA, the literature is less extensive. Markers of tissue damage characteristic of inflammation, apoptosis, and the stimulation of reactive oxygen species have been found in tissues of human and transgenic mouse models carriers of mutant transthyretin variants, well before amyloid deposits are detected.³⁵ However, the direct toxicity of AL precursors is thought to be more marked than that of ATTR, possibly contributing to the different clinical profiles of the 2 conditions, such that AL amyloidosis has more rapid progression and worse prognosis if untreated than ATTR amyloidosis.³⁶

INTERSTITIAL AMYLOID FIBRIL INFILTRATION.

Amyloid infiltration in the heart results in disruption of tissue architecture and subsequent replacement fibrosis, with cardiomyocyte damage. The

CENTRAL ILLUSTRATION Myocardial Injury and Applications of Troponin Measurement in Cardiac Amyloidosis



De Michieli L, et al. *J Am Coll Cardiol CardioOnc.* 2024;6(1):1-15.

Several mechanisms are potentially involved in the development of myocardial injury (defined as an increase in cardiac troponin values greater than the 99th percentile sex-specific and assay-specific reference limits) in patients with cardiac amyloidosis. These mechanisms include amyloid-related causes, such as amyloid precursor toxicity, amyloid interstitial infiltration, and amyloid vascular involvement, but also non-amyloid-related causes such as diastolic and systolic dysfunction, heart failure, atherosclerotic coronary artery disease, and other causes of supply-demand imbalance such as tachyarrhythmias and hypotension. Regardless of the cause, measurement of cardiac troponin is important in the management of patients with cardiac amyloidosis for diagnostic, prognostic, and monitoring purposes. hs-cTn = high-sensitivity cardiac troponin; LV = left ventricular; RV = right ventricular.

extracellular fibrils have a significant impact on the mechanics and physiology of the target tissue.³⁷ In AL CA models, in vitro analyses have shown that amyloid fibrils rapidly surround cultured cardiomyocytes and recruit soluble LCs, triggering cytotoxicity.³⁷ Extracellular amyloid fibrils also appear to disrupt cardiac matrix homeostasis and alter extracellular matrix turnover, which is critical for the maintenance of myocyte-myocyte force coupling and proper myocardial function.³⁸ An overexpression of matrix metalloproteinases³⁹ also has been reported. Moreover, fibroblasts are able to internalize both aggregated transthyretin⁴⁰ and amyloidogenic LCs.⁴¹ These mechanisms contribute to the expansion of extracellular spaces and to the development of interstitial, reactive fibrosis in response to tissue damage, which may contribute further to myocardial disruption and damage.

A study by Pucci et al⁴² showed that extracellular volume evaluated on cardiac magnetic resonance correlates with the combination of amyloid deposition and interstitial fibrosis at endomyocardial biopsy of the left ventricle (LV). This combination of amyloid and fibrosis also correlates with hs-cTnT, in both AL CA ($r = 0.622$) and ATTR CA ($r = 0.533$), suggesting that infiltration, fibrosis, and extracellular disruption together contribute to myocardial injury. In a cohort of patients with ATTRwt CA, hs-cTnT correlated with native T1 and extracellular volume on cardiac magnetic resonance and modestly, if at all, with amyloid load ($r = 0.354$) on endomyocardial biopsy of the right ventricle⁴³; this poor correlation with amyloid load on endomyocardial biopsy is probably secondary to the selection bias inherent to biopsy itself, which provides focal information rather than an evaluation of total amyloid load.

CORONARY MICROVASCULAR DYSFUNCTION AND AMYLOID- AND NON-AMYLOID-RELATED CORONARY ARTERY DISEASE. Coronary microvascular dysfunction and amyloid- and non-amyloid-related coronary artery disease (CAD) also contribute to myocardial injury in patients with CA. Amyloid deposits can be found in the perivascular regions and in the media of intramyocardial coronary vessels,⁴⁴ with vascular and perivascular involvement being more frequent in AL CA.⁴⁵ Coronary microvascular dysfunction can be present, related to 3 possible mechanisms: structural (amyloid deposition in the vessels wall with thickening and stenosis, capillary rarefaction), extravascular (extrinsic compression of the microvasculature), and functional (autonomic and endothelial dysfunction).^{42,44,46} Dorbala et al⁴⁴ reported on 21 patients with CA without obstructive epicardial

CAD who underwent evaluation of coronary microvascular function with rest and vasodilator stress ¹³N ammonia positron emission tomography/computed tomography. Compared with 10 patients with LV hypertrophy, patients with CA had lower resting myocardial blood flow, lower stress myocardial blood flow, lower coronary flow reserve, and higher minimal coronary vascular resistance. Coronary microvascular dysfunction was associated with increased LV mass and myocardial relaxation abnormalities. Similarly, on echocardiography, patients with CA had significantly lower coronary flow reserve (together with lower rest and stress global longitudinal strain and lower myocardial work efficiency), and changes in coronary flow reserve and deformation capacity were strongly associated with exercise tolerance.⁴⁷ Also, in 20 patients with AL CA, stress-induced wall motion abnormalities on echocardiography were frequent (55%) despite the absence of significant epicardial CAD.⁴⁸ Other cardiovascular and non-cardiovascular comorbidities (such as diabetes) can contribute to endothelial abnormalities and coronary microvascular dysfunction, particularly in older and comorbid patients with ATTRwt CA. Therefore, coronary microvascular dysfunction is frequent in CA and likely is an important cause of myocardial injury.⁴⁹ Amyloid vascular and perivascular infiltration^{45,50} can also be a cause of chest pain, acute myocardial injury, and myocardial infarction in the absence of atherosclerotic CAD.⁵¹⁻⁵³ In addition, classic atherosclerotic epicardial CAD also can be present (particularly in older patients with ATTRwt CA²) and may contribute to a lower ischemic threshold. A recent study reported that chest pain is not an uncommon symptom in patients with CA (about 40%), and the etiology seemed to differ, with obstructive CAD more frequent in patients with ATTR CA, whereas amyloid vascular or perivascular involvement was more common in those with AL CA.⁵⁴ Management of acute coronary syndromes and the differential diagnosis for chest pain in these patients can be challenging because of the underlying chronic myocardial injury. Validated cTn-based algorithms for the rule-in and rule-out of myocardial infarction among patients presenting without ST-segment elevation⁵⁵ are difficult to apply in those who have baseline hs-cTn increases but are the only guidance available. However, the application of biologically significant relative changes between serial cTn values,⁶ which help in patients with acute coronary syndromes, have not been validated in those with CA.

DIASTOLIC DYSFUNCTION AND HEART FAILURE. Elevation of LV end-diastolic pressure can lead to

apoptosis and cTnI release, as shown in animal models,⁵⁶ and clinically, cTnT levels correlate with the LV end-diastolic pressure in patients with heart failure.⁴⁹ Because of the restrictive hemodynamic status of CA, increased LV filling pressure with elevated myocardial wall tension (an obstacle to subendocardial myocardial perfusion) can play an important role in the genesis of myocardial injury. This can result in a vicious circle characterized by elevated LV filling pressures causing subendocardial ischemia, reduced coronary perfusion pressure, and diastolic dysfunction, which in turn can lead to more elevations in LV end-diastolic pressure.⁵⁷ Moreover, both LV and right ventricular pseudohypertrophy can enhance these mechanisms. Group 2 pulmonary hypertension,⁵⁸ frequently detected in patients with CA,⁵⁹ can also contribute to cTn release through myocardial ischemia and cell death due to increased wall tension of the right ventricle with pressure and/or volume overload.⁶⁰

ACUTE ON CHRONIC MYOCARDIAL INJURY. Not only multiple mechanisms can contribute to the genesis and progression of myocardial injury in patients with CA, but these patients are also at risk for acute events related to CA or independent of CA as well. Some of these may be due to ischemia, although clear signs and symptoms of myocardial ischemia are often difficult to appreciate in this complex milieu. Acute exacerbations of heart failure or arrhythmias are frequently observed in CA, both of which are known causes of myocardial injury.⁶ Patients with CA also are particularly prone to developing atrial fibrillation or flutter and/or bradyarrhythmias.^{61,62} Nonsustained ventricular arrhythmias are also frequent, particularly in those with more advanced cardiac involvement.⁶³ Hypotension is also frequent in patients with CA, secondary to the primary cardiac disease but also to autonomic dysfunction, and similar to tachyarrhythmias and bradyarrhythmias, prolonged hypotension is counted among the possible causes of myocardial injury related to oxygen supply-demand imbalance.⁶ All of these are known causes of cTn elevations,⁶ particularly in individuals with underlying myocardial disease.⁶⁴

CLINICAL USE OF cTn IN CA

The use of cardiac biomarkers like cTn and natriuretic peptides to aid in the evaluation of patients with CA was developed over many years. Today, they have a fundamental role in the prognostic assessment of patients with CA, alone and integrated in multiparametric staging systems. Moreover, they are

helpful in monitoring the response to chemotherapy in AL amyloidosis. Emerging data suggest that they can also be of relevance when CA is suspected.⁶⁵ In the following paragraphs, the role of cTn in the different phases of CA management is discussed (see **Table 4** for AL CA and **Table 5** for ATTR CA).

DIAGNOSIS. Compared with its role in the prognostic assessment of patients with CA, the diagnostic value of cTn has been less extensively investigated, in part because the metrics necessary to distinguish CA from other disease entities that can cause myocardial dysfunction has not been probed adequately. In addition, there are many assays for cTnI, both conventional and high sensitivity, and finding exact thresholds for diagnosis requires large studies for each.

In patients at risk for developing AL amyloidosis, including those with monoclonal gammopathy of uncertain significance, periodical screening for potential cardiac involvement can be performed with NT-proBNP and cardiovascular imaging parameters.⁶⁶ Some studies have started investigating the role of hs-cTnT in the diagnostic algorithm of CA.⁶⁷⁻⁶⁹ Recently, Vergaro et al⁶⁵ reported that hs-cTnT, alone and in combination with NT-proBNP, was useful when CA was suspected to identify patients in whom the diagnosis is unlikely and those in whom it is much more likely. They suggested cutoffs of <180 ng/L for NT-proBNP and <14 ng/L for hs-TnT as optimal rule-out thresholds. hs-TnT \geq 86 ng/L was a good rule-in threshold, but the pretest probability of CA was high; about 60% of patients had confirmed CA in both derivation and validation cohorts. In the validation cohort, 74 patients had both biomarkers less than the cutoff values, with 4 false negatives. For the rule-in cutoff, about 143 patients were correctly classified, with 17 false positives. Further studies are needed to validate these findings in populations with lower prevalence of CA. Thus, for diagnosis, other features must be present at least adjunctively.

PROGNOSIS. AL amyloidosis. The prognostic role of cTn, particularly cTnT, in CA has been extensively validated.⁷⁰ Dispenzieri et al⁷¹ showed that survival of patients with AL amyloidosis was significantly worse among those with increased cTnT and cTnI values compared with those with undetectable cTn. On multivariable analysis, cTnT was a better predictor than cTnI. With the evidence that NT-proBNP also had a strong prognostic value in AL amyloidosis,⁷² a staging system was developed integrating cTnT and cTnI with NT-proBNP.⁷³ Patients were stratified in 3 stages (I, II, and III), with median survival times of

TABLE 4 Clinical Use of cTn in the Management of Patients With AL Amyloidosis

	Prognosis ^a		
	Prognostic Variables and Staging Systems	Estimated Survival	Response to Therapy
Diagnostic score to define cardiac involvement in AL amyloidosis ⁶⁷ <ul style="list-style-type: none"> • hs-cTnT >35 ng/L (1 point) • GLS \geq-17% (1 point) • RELAPS \geq0.9 (1 point) 	Mayo Clinic 2004 ⁷³ <ul style="list-style-type: none"> • cTnT \geq 0.035 μg/L or cTnI (Stratus CS) \geq0.1 μg/L or hs-cTnT \geq50 ng/L • NT-proBNP \geq332 ng/L 	Stage 1: 26.4 mo Stage 2: 10.5 mo Stage 3: 3.5 mo	Cardiac disease progression ⁸⁹ <ul style="list-style-type: none"> • cTn increase >33% or • NT-proBNP increase >30% and >300 ng/L or • LV ejection fraction reduction \geq10%
CA very likely in patients with suspected CA ⁶⁵ <ul style="list-style-type: none"> • hs-cTnT \geq86 ng/L 	Mayo Clinic 2012 ³³ <ul style="list-style-type: none"> • cTnT \geq0.025 μg/L or hs-cTnT \geq40 ng/L • NT-proBNP \geq1,800 ng/L • dFLC \geq180 mg/L 	Stage 1: 94.1 mo Stage 2: 40.3 mo Stage 3: 14 mo Stage 4: 5.8 mo	Cardiac disease response ⁸⁹ <ul style="list-style-type: none"> • NT-proBNP reduction >30% and >300 ng/L (if baseline \geq650 ng/L) or • \geq2-class reduction in NYHA functional class (if baseline III or IV)
CA unlikely in patients with suspected CA ⁶⁵ <ul style="list-style-type: none"> • hs-cTnT <14 ng/L and • NT-proBNP <180 ng/L 	European 2015 modification of Mayo 2004 ^{74,75} <ul style="list-style-type: none"> • cTnT \geq0.035 μg/L or cTnI (Stratus CS) \geq0.1 μg/L or hs-cTnT \geq50 ng/L • NT-proBNP \geq332 ng/L and >8500 ng/L 	Stage 1: NR Stage 2: 55% 3 y Stage 3a: 52% 3 y Stage 3b: 7 mo	Graded cardiac response ⁹¹ <ul style="list-style-type: none"> • CarCR: nadir NT-proBNP \leq350 pg/mL or BNP \leq80 pg/mL • CarVGPR: >60% reduction in NT-proBNP/BNP from baseline level not meeting CarCR • CarPR: 31%-60% reduction in NT-proBNP from baseline level not meeting CarCR • CarNR: \leq30% reduction in NT-proBNP from baseline level
	Boston University staging system ⁷⁶ <ul style="list-style-type: none"> • cTnI (assay not specified) >0.1 ng/mL • BNP >81 ng/L and >700 ng/L 	Stage 1: NR Stage 2: 9.4 y Stage 3: 4.3 y Stage 3b: 1 y	Restaging with Mayo 2004 and 2012 systems ^{94,95}
	Palladini et al ⁸⁰ <ul style="list-style-type: none"> • hs-cTnT >77 ng/L at baseline • cTnI (Advia Centaur CP, Siemens) \geq100 ng/L (integrated in staging system with NT-proBNP \geq332 ng/L) • Progression of NT-proBNP and increase >75% of hs-cTnT after therapy 	10.6 mo	
	ASCT candidates ^{70,82} <ul style="list-style-type: none"> • cTnT >0.06 μg/L or hs-cTnT >75 ng/L • NT-proBNP >5,000 ng/L 	Cutoffs for identification of patients at risk of early mortality after ASCT	

A summary of different thresholds for specific biomarkers (particularly cTn evaluated with different assays) for diagnosis, prognosis, and response to therapy in patients with AL amyloidosis is shown. For diagnosis, thresholds of specific markers, potentially useful to predict the likelihood of a final diagnosis of AL cardiac amyloidosis, are reported.^aIn prognostic staging systems, the stage for each patient is defined on the basis of the number of variables above the specified thresholds tabulated in the table. For some staging system, alternative cutoffs for BNP (instead of NT-proBNP) are available. Criteria for response to therapy in AL amyloidosis are reported for completion, even if they do not include cTn or high-sensitivity cTn. Criteria for hematological response are not reported.

AL = amyloid light chain; ASCT = autologous stem cell transplantation; BNP = brain natriuretic peptide; CA = cardiac amyloidosis; CarCR = complete cardiac response; CarNR = cardiac nonresponse; CarPR = cardiac partial response; CarVGPR = cardiac very good partial response; cTnI = cardiac troponin I; cTnT = cardiac troponin T; dFLC = difference between involved and uninvolved free light chains; GLS = global longitudinal strain; hs-cTnT = high sensitivity cardiac troponin T; LV = left ventricular; NR = not reached; NT-proBNP = N-terminal pro-brain natriuretic peptide; RELAPS = relative apical sparing; mo = months; y = years.

26.4, 10.5, and 3.5 months, respectively. The proposed cutoffs were \geq 332 ng/L for NT-proBNP, \geq 0.035 μ g/L for cTnT, and \geq 0.1 μ g/L for cTnI (using the Stratus CS assay). Subsequently, sFLC levels (particularly the difference between involved and uninvolved sFLCs) were integrated in the model (Mayo 2012 model), with stratification of patients in 4 groups. In this model, cTnT was used at a threshold cutoff of \geq 0.025 μ g/L.³³ A European modification of the first Mayo Clinic stage was also developed, subclassifying stage III patients in stages IIIa and IIIb according to NT-proBNP levels using a threshold of 8,500 ng/L.^{74,75} The Boston University staging system, instead, was based on brain natriuretic peptide (81 ng/L) and cTnI (0.1 ng/mL).⁷⁶ Comparing the available staging systems suggests that the European

2015 model had better prediction for 1-year mortality, but the Mayo 2012 model increased the ability to predict long-term survival.⁷⁷

When the hs-cTnT assay became available for clinical practice, a study from the Mayo Clinic⁷⁸ demonstrated that hs-cTnT numerical values could not merely be substituted for cTnT measurements in the original Mayo Clinic staging system. A threshold of 50 ng/L was derived with a quartic formula⁷⁹ and was integrated into the Mayo 2004 model. For the updated Mayo 2012 model, which is currently widely used, a threshold of hs-cTnT \geq 40 ng/L was validated.⁷⁹ Because of the improved analytical performance of the high-sensitivity assays,⁷⁸ the prognostic significance of this biomarker improved. Palladini et al⁸⁰ concurred and reported that a prognostic

TABLE 5 Clinical Use of cTn in the Management of Patients With ATTR CA

	Prognosis ^a		
	Prognostic Variables and Staging Systems	Estimated Survival	Response to Therapy
CA very likely in patients with suspected CA ⁶⁵ <ul style="list-style-type: none"> hs-cTnT \geq86 ng/L 	Grogan et al ⁸⁵ for ATTRwt CA <ul style="list-style-type: none"> cTnT >0.05 ng/mL NT-proBNP >3,000 pg/mL 	Stage 1: 66 mo Stage 2: 40 mo Stage 3: 20 mo	Disease progression in ATTR CA ⁹⁶ : at least 1 marker in each domain <ul style="list-style-type: none"> Clinical and functional domain Laboratory domain <ul style="list-style-type: none"> cTn increase >30% or NT-proBNP increase >30% or Advance in NAC stage ECG and imaging domain
CA unlikely in patients with suspected CA ⁶⁵ <ul style="list-style-type: none"> hs-cTnT <14 ng/L and NT-proBNP <180 ng/L 	Nakashima et al ⁸⁸ for ATTRwt CA <ul style="list-style-type: none"> hs-cTnT >50 ng/L BNP >250 ng/L eGFR <45 mL/min/1.73 m² NAC staging system for ATTRwt and ATTRv CA ⁸⁶ <ul style="list-style-type: none"> NT-proBNP >3,000 ng/L eGFR <45 mL/min² 	32 mo for high-risk group (score 2 or 3) Stage 1: 69.2 mo Stage 2: 46.7 mo Stage 3: 24.1 mo	

A summary of different thresholds for specific biomarkers (particularly cTn evaluated with different assays) for diagnosis, prognosis, and response to therapy in patients with ATTR CA is shown. For diagnosis, thresholds of specific biomarkers, potentially useful to predict the likelihood of a final diagnosis of ATTR CA, are reported. ^aIn prognostic staging systems, the stage for each patient is defined on the basis of the number of variables above the specified thresholds tabulated in the table. The NAC staging system has been reported for completeness, even if it does not include cTn or high-sensitivity cTn.
 ATTR = amyloid transthyretin; ATTRv = variant amyloid transthyretin; ATTRwt = wild-type amyloid transthyretin; eGFR = estimated glomerular filtration rate; NAC = National Amyloidosis Center; other abbreviations as in Table 4.

cutoff of 77 ng/L for hs-cTnT best predicted mortality at presentation. After chemotherapy, changes in NT-proBNP and a >75% increase in hs-cTnT were independent prognostic determinant. Thus, they suggested that hs-cTnT at baseline could be a simple and powerful tool for single subjects risk assessment and for patient stratification in clinical trials.⁸⁰

Dispenzieri et al⁸¹ reported also on the clinical use of soluble suppression of tumorigenicity 2 in the prognosis of patients with AL amyloidosis, together with cTnT, NT-proBNP, and differential sFLCs. It is unclear, however, how much these add to hs-cTnT alone.

Autologous stem cell transplantation has been shown to be an effective therapy for patients with AL amyloidosis, but only selected patients can undergo this procedure.⁵ Cardiac biomarkers at baseline are useful for risk stratification of early death following autologous stem cell transplantation. Specifically, cTnT >0.06 μ g/L (or hs-cTnT of 73/75 ng/L) and NT-proBNP >5,000 ng/L manifest optimal discrimination.^{70,82}

Even though in some of these approaches, thresholds for cTnI have been proposed,^{75,77,78,80} only 1 study has reported on a hs-cTnI assay (the Siemens Advia Centaur CP assay at a threshold of 100 ng/L). No other data exist for the other high-sensitivity assays, which are now the state of the art in clinical practice. It is likely that over time, with large studies, the optimal threshold values for each of the hs-cTnI assays will be defined.

ATTR CA. cTn, particularly cTnT and hs-cTnT, is now a recognized prognostic factor in ATTR CA, especially ATTRwt CA.^{83,84} Grogan et al⁸⁵ developed a prognostic staging system for ATTRwt CA on the basis of a cTnT threshold of 0.05 ng/mL and an NT-proBNP threshold of 3,000 pg/mL. The 4-year survival estimates were 57%, 42%, and 18% for stages I, II, and III, respectively, with stage III patients having an increased risk for mortality after adjustment for age and sex compared with stage I patients. A widely used staging system that avoids the issue of so many different cTn assays, was developed by Gillmore et al⁸⁶ for ATTRwt CA and ATTRv CA. It is based on NT-proBNP and estimated glomerular filtration rate. In 175 patients with ATTR CA (133 with ATTRwt CA and 42 with ATTRv CA), this staging system provided better prognostic accuracy compared with one using NT-proBNP and contemporary cTnI.⁸⁷ Recently, a staging system combining hs-cTnT (>50 ng/L), brain natriuretic peptide (>250 pg/mL), and estimated glomerular filtration rate (<45 mL/min/1.73 m²) was published, with good prediction of prognosis in 176 Japanese patients with ATTRwt CA.⁸⁸ As of now, consistent data on the prognostic value of hs-cTnI assays and of cTn when integrated with disease-specific biomarkers such as transthyretin and retinol-binding protein are not available.

RESPONSE TO TREATMENT. AL amyloidosis. Cardiac response to chemotherapy has been defined as a >30% decrease with a total decline of >300 ng/L in NT-proBNP or a decrease of at least two NYHA

functional classes (if baseline class III or IV).⁸⁹ Caution should be applied, however, in interpreting changes of NT-proBNP values <50% because of its high biological variability.⁹⁰ A recent paper reported that graded cardiac response (on the basis of the extent of NT-proBNP reduction) allowed a better assessment of cardiac improvement than the traditional binary response system.⁹¹ However, cTn was proposed to define cardiac disease progression at 6 months, on the basis of an increase of $\geq 33\%$.⁸⁹ However, the assays used were contemporary cTnT and cTnI assays, and the cTnI assay was not identified. Other criteria for disease progression were an NT-proBNP increase >30% and >300 ng/L or a reduction of $\geq 10\%$ in LV ejection fraction.⁸⁹

The current Mayo Clinic staging systems also are useful for restaging during treatment at 3 and 6 months after chemotherapy initiation. A worsening stage at 3 and 6 months is associated with worse survival than maintenance at the same stage.⁹² In patients with disease relapse after first-line therapy, both the Mayo 2004 and 2012 staging systems are useful for prognostic stratification with second-line therapy.⁹³

ATTR CA. A multiparametric approach is recommended to evaluate disease progression in ATTR CA; among the various criteria, an increase in cTn of >30% (which, however, is less than the reference change value, ie, the amount of change explained by conjoint analytical and biological variation) is considered indicative of disease progression.⁹⁴ Specific therapy for patients with ATTR CA has recently become available,³ but few studies have investigated the changes in cardiac biomarkers over time to verify the response to treatment. Most trials and clinical studies report the trend of NT-proBNP, not cTn, over time.^{4,95} Recently, a single-center French study⁹⁶ showed that tafamidis therapy stabilizes NT-proBNP and hs-cTnT levels over time, especially in patients with higher values at baseline. However, further studies on the role of hs-cTn for the monitoring of response to therapy in ATTR CA are needed.

FUTURE DIRECTIONS

The field of cardiac and noncardiac biomarkers in the clinical management of patients with CA continues to evolve. Many research gaps remain to be addressed. With regard to hs-cTn, hs-cTnI assays have not yet been validated in patients with CA. The specific metrics derived will vary depending on the assay being used. Moreover, new and improved assays will be available on the market in the near future, potentially further refining risk stratification in this

and other settings. In addition, novel markers are being elucidated as potential indicators of disease pathogenesis, course, and progression, for both AL CA (eg, LC glycosylation)^{97,98} and ATTR CA (including transthyretin, retinol-binding protein 4, non-native transthyretin, and neurofilament LC).⁹⁹ At present, their interactions with available prognostic biomarkers are yet to be determined. Finally, validated and new staging systems for prognostication will need to be explored in newer cohorts of patients with CA now that earlier diagnosis is more frequent and multiple effective therapies are available for all forms.

CONCLUSIONS

cTn has dramatically changed clinical practice in cardiology in the past 30 years. It is now the biomarker of choice for the detection of myocardial injury, and high-sensitivity assays can identify even modest increases in a multiplicity of clinical conditions. Myocardial injury is particularly frequent in patients with CA because of multiple synergistic mechanisms that contribute to cardiomyocytes damage. Evaluation of cTn, alone and integrated in staging systems, is helpful in the management of patients with CA, who are now being increasingly recognized and treated. There are still many potential areas of research in this field, including further addressing the diagnostic significance of cTn in suspected CA and its prognostic and monitoring role now that new effective therapies are available not only for AL CA but also for ATTR CA. Assay-specific thresholds (other than for cTnT and hs-cTnT) are also essential to allow a wider dissemination of troponin-based diagnostic and prognostic scores.

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Dr De Michieli has received honoraria from Pfizer, Alnylam Pharmaceuticals, and AstraZeneca. Dr Cipriani has received honoraria from Pfizer, Alnylam Pharmaceuticals, and AstraZeneca. Dr Dispenzieri has received research support from Alnylam Pharmaceuticals, Pfizer, Takeda, and Bristol Myers Squibb; participates on the data and safety monitoring board for Oncopeptides and Sorrento; and is on the advisory board and independent review committee for Janssen. Dr Jaffe has consulted or presently consults for most of the major diagnostics companies, including Beckman-Coulter, Abbott, Siemens, Ortho Diagnostics, ET Healthcare, Roche, Radiometer, SphingoTec, Amgen, and Novartis; and he has stock options in RCE Technologies.

ADDRESS FOR CORRESPONDENCE: Dr Allan S. Jaffe, Department of Cardiology, Gonda 5, Mayo Clinic, 200 First Street SW, Rochester, Minnesota 55905, USA. E-mail: jaffe.allan@mayo.edu. @LauraDemichieli, @albcipri6.

PERSPECTIVES

COMPETENCY IN MEDICAL KNOWLEDGE: Patients with CA, both AL and ATTR often manifest increases in cTn and natriuretic peptides. The causes for these increases include the cytotoxic effects of amyloid precursors, infiltration of interstitial amyloid fibrils, coronary microvascular dysfunction, amyloid/nonamyloid-related CAD, diastolic dysfunction, and heart failure. In addition, many clinical conditions, such as arrhythmias, hypotension, critical illness and decompensated heart failure, can add acute onto the chronic myocardial injury.

COMPETENCY IN PATIENT CARE AND

PROCEDURAL SKILLS: There are a variety of algorithms that can assist in diagnosing CA, helping to assess its prognosis and to allow one to follow patients' responses to a variety of treatments.

TRANSLATIONAL OUTLOOK: The metrics for the use of cTn and natriuretic peptides in this setting are distinct from the metrics applicable to patients with possible myocardial infarction. This needs to be kept in mind by clinicians in using the criteria proposed.

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