REVIEW



Inflammation and fibrosis in murine models of heart failure

Lucas Bacmeister^{1,2} • Michael Schwarzl^{1,2} • Svenja Warnke^{1,2} • Bastian Stoffers^{1,2} • Stefan Blankenberg^{1,2} • Dirk Westermann^{1,2} • Diana Lindner^{1,2}

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Abstract

Heart failure is a consequence of various cardiovascular diseases and associated with poor prognosis. Despite progress in the treatment of heart failure in the past decades, prevalence and hospitalisation rates are still increasing. Heart failure is typically associated with cardiac remodelling. Here, inflammation and fibrosis are thought to play crucial roles. During cardiac inflammation, immune cells invade the cardiac tissue and modulate tissue-damaging responses. Cardiac fibrosis, however, is characterised by an increased amount and a disrupted composition of extracellular matrix proteins. As evidence exists that cardiac inflammation and fibrosis are potentially reversible in experimental and clinical set ups, they are interesting targets for innovative heart failure treatments. In this context, animal models are important as they mimic clinical conditions of heart failure patients. The advantages of mice in this respect are short generation times and genetic modifications. As numerous murine models of heart failure exist, the selection of a proper disease model for a distinct research question is demanding. To facilitate this selection, this review aims to provide an overview about the current understanding of the pathogenesis of cardiac inflammation and fibrosis in six frequently used murine models of heart failure. Hence, it compares the models of myocardial infarction with or without reperfusion, transverse aortic constriction, chronic subjection to angiotensin II or deoxycorticosterone acetate, and coxsackievirus B3-induced viral myocarditis in this context. It furthermore provides information about the clinical relevance and the limitations of each model, and, if applicable, about the recent advancements in their methodological proceedings.

Keywords Myocardial infarction (MI) \cdot Ischemia/reperfusion \cdot Pressure overload \cdot Cardiac hypertrophy \cdot Neurohumoral activation \cdot Myocarditis

Cardiac inflammation and fibrosis in heart failure

Heart failure (HF) is a chronic disease that is associated with adverse outcome [290]. Due to demographic changes and the increasing prevalence of chronic HF, hospitalisation rates for acute HF are rising [54, 290]. Thus, HF is the leading cause for non-elective hospitalisations in developed countries of

Lucas Bacmeister l.bacmeister@uke.de

Diana Lindner d.lindner@uke.de

² DZHK (German Center for Cardiovascular Research), Partner site Hamburg/Kiel/Lübeck, Hamburg, Germany patients older than 65 [54]. HF results from injury to the myocardium due to a variety of causes, including, but not limited to, myocardial infarction (MI), hypertension, heart valve disease, and myocarditis [126]. On the one hand, a myocardial injury, regardless of its initiating cause, results in sterile myocardial inflammation [133], on the other hand, the infection with pathogens induces non-sterile myocardial inflammation of myocardial inflammation requires the activation of anti-inflammatory stimuli, which in turn initiate pro-fibrotic signalling [60]. Hence, most cardiac pathologies imply the development of myocardial fibrosis [21].

In the healthy heart, the extracellular matrix (ECM) provides a scaffold for cardiac cells and thereby ensures the structural integrity and function of the heart [52]. As shown in Fig. 1a, a decellularised murine heart maintains its morphology, which highlights the overwhelming significance of the ECM for the cardiac integrity. The magnification clearly

¹ Clinic for General and Interventional Cardiology, University Heart Center Hamburg, University Hospital Hamburg-Eppendorf, Martinistr. 52, 20246 Hamburg, Germany



Fig. 1 The cardiac extracellular matrix and its fibrotic remodelling upon injury. **a** Decellularised murine heart without atria that solely consists of extracellular matrix (ECM) from a macroscopic (left panel) and microscopic (right panel) perspective. The ECM covers every single cardiomyocyte and thus provides a scaffold to conduct the contractile force from each cardiomyocyte to the whole organ and maintains cardiac integrity. **b** Cross sections of left ventricular tissue (Masson-Trichrome stained) as representative examples of distinct categories of fibrotic extracellular remodelling. Left panel: reactive insterstitial fibrosis, mainly observed in the murine models

demonstrates that each cardiomyocyte (CM) is surrounded by ECM, which ensures the transmission of the contractile force from a single CM to the whole organ [133]. Moreover, the ECM provides a reservoir for a variety of latent growth factors [133]. On a pathophysiological level, fibrotic remodelling of the myocardium could disrupt appropriate myocardial excitation–contraction coupling and may promote arrhythmia, e.g. through the induction of re-entry circuits [280]. Furthermore, excessive fibrotic remodelling results in enhanced stiffness and is linked to systolic and diastolic dysfunction [116]. In contrast, dampened fibrotic remodelling in the context of MI may lead to insufficient scar formation, which can be associated with ventricular dilation or even rupture [211, 272].

of nonischemic HF such as transverse aortic constriction, subjection to angiotensin II or deoxycorticosterone acetate, or inoculation with CVB3. Interstitial cardiac fibrosis predominantly occurs as a result of repressing diffuse inflammation upon diffuse cardiac injury. Right panel: reparative fibrosis, mainly observed in the murine models of ischemic HF, such as permanent or transient LAD-ligation. The mechanistic rational of reparative fibrosis is to replace dead cardiac tissue by fibrotic scar tissue. Middle panel: healthy cardiac tissue. Blue: extracellular matrix, pink: cytoplasm

Since pathological fibrosis is characterised by excessive deposition of ECM proteins, several key pro-fibrotic stimuli, such as a variety of growth factors, most importantly transforming growth factor (TGF)- β , but also angiotensin II (Ang II) and aldosterone contribute to the development of cardiac fibrosis regardless of the underlying pathology [133]. However, the relative significance of a distinct molecular pathway depends on the type and the degree of the initial cardiac injury [133]. As shown in Fig. 1b, fibrotic remodelling may be divided into two paramount categories [213]: (1) 'reactive interstitial fibrosis', often seen in HF caused by prolonged ischemia to replace necrotic CMs by the formation of a myocardial scar.

Several murine models of HF exist, which either predominantly develop reparative or reactive interstitial fibrosis. Sterile inflammation and reparative fibrosis upon ischemia are observed in the murine models of (1) permanent or (2)transient ligation of the left anterior descending coronary artery (LAD). Sterile inflammation and reactive interstitial fibrosis upon pressure overload and neurohumoral activation are observed in the murine models of (1) transverse aortic constriction (TAC), (2) chronic infusion of angiotensin II (Ang II), and (3) chronic treatment with deoxycorticosterone acetate (DOCA) accompanied by unilateral nephrectomy and high salt diet. Non-sterile, excessive inflammation and subsequent interstitial fibrosis occurs in the murine model of viral myocarditis upon infection with coxsackievirus B3 (CVB3). Depending on the susceptibility of a certain mouse strain, viral myocarditis may further become chronic and thus result in dilated inflammatory cardiomyopathy.

This review aims to facilitate the challenging selection of a proper murine model of HF for a specific research question as it (1) provides an overview about the distinct molecular and cellular mechanisms that underlie cardiac inflammation and fibrosis in each of these six frequently used HF models. Further, it aims to provide information about (2) the methodological proceeding, (3) the clinical relevance, (4) the limitations and, if applicable, (5) the recent methodological advancements of each model.

Models of heart failure induced by ischemia

Myocardial infarction induced by permanent LAD-ligation

Murine model of heart failure and its clinical relevance

Permanent LAD-ligation in mice results in acute myocardial injury due to ischemia, which is defined as MI [163, 256, 289]. In humans, type 1 MI is caused by the disruption of an atherosclerotic plaque and results in myocyte necrosis, which is often accompanied by ST-elevations in the ECG (ST-elevated MI, STEMI). According to the ESC guidelines, prompt revascularisation for STEMI patients is recommended to reduce infarct size and mortality, but still a significant number of STEMI patients (approximately 15–25%) do not receive any form of reperfusion therapy [33, 76]. The current ESC guidelines for the treatment of STEMI additionally state that reperfusion is not recommended if hemodynamically stable patients present more than 12 h after the onset of symptoms [33, 110]. Thus, permanent LAD-ligation in mice particularly resembles the clinical situation of a significant fraction of patients that suffer from type 1 MI without timely revascularisation [256]. Even patients that undergo revascularisation can develop a form of ischemic cardiomyopathy resembling permanent LAD-ligation in animal models. In mice, the development of HF following permanent LAD-ligation is characterised by infarct wall thinning, left ventricular dilation and severe systolic and diastolic dysfunction. These hallmarks are frequently seen in post-MI patients as well. Since MI in mice is predominantly characterised by ventricular dilation, their HF resembles features of human HF with reduced ejection fraction (HFrEF) [94, 163, 273]. HFrEF in mice is observed within the first 24 h after MI and further aggravates over time [114, 187].

Methodological proceedings

Induction of HF utilising the model of permanent LADligation in small rodents was first described in 1979 in rats [207]. Initially, adult mice are anaesthetised and mechanically ventilated. Hereafter, surgical access to the heart is performed by an incision through the third left intercostal space [94]. Subsequently, the LAD is permanently ligated with a suture [94]. Figure 2a shows a schematic drawing and an intraoperative image of the procedure and the sufficient ligation, proven by blanching of the distal cardiac tissue [130]. Moreover, infarct-typical ECG-alterations, such as ST-elevation [75, 86], or elevated plasma levels of cardiac troponin in the cause of MI [1] can prove the ischemiainduced MI. The observation period following permanent LAD-ligation in mice typically ranges from a few hours up to several weeks [286].

To uncover the infarct zone of explanted hearts, the cardiac tissue can be incubated with the colourless, hydrophilic triphenyltetrazolium chloride immediately after harvesting. In viable cells, this dye is reduced to a deep red, hydrophobical formazan. Thus, only areas that contain viable CMs appear in deep red, whereas the infarct zone itself remains colourless [94, 155]. The mortality following permanent LAD-ligation in mice predominantly occurs from day 1 to day 7 as a result of left ventricular rupture, acute HF, or heart rhythm disturbances [39, 70, 86, 130, 156, 187].

Myocardial infarction induced by transient LAD-ligation (ischemia-reperfusion)

Murine model of heart failure and its clinical relevance

The murine model of transient LAD-ligation mimics type 1 MI followed by successful revascularisation [155], which reflects the clinical course of the majority of STEMI patients. According to the ESC guidelines, primary PCI, defined as percutaneous catheter intervention (PCI) without previous fibrinolysis, is recommended as the reperfusion strategy for STEMI patients within 90 min after diagnosis [110, 195]. Several clinical trials emphasise the Fig. 2 Myocardial infarction induced by ligation of the left anterior descending artery (LAD) in mice. a Left: schematic drawing that depicts the ligation of the LAD in mice and the resulting area at risk (white). Right: representative intraoperative image immediately taken after the LAD-ligation, in which the area at risk is demasked by the blanching of the tissue distal to the ligation. The size of the area at risk depends on the area of cardiac tissue that is supplied by the LAD. b Time course of the wave front of CM-necrosis (infarct zone, dark grey) within the area at risk (white) in mice. Ischemia of more than 20-30 min results in CM-necrosis, which is initially observed in the subendocardial layers and hereafter extends to the subepicardial layers; ultimately, after more than 60-90 min of ischemia, this results in the total infarction of the area at risk



strong association between delayed revascularisation and the degree of myocardial injury and its sequelae [109, 184, 256]. Mice that are subjected to transient LAD-ligation exhibit HF symptoms comparable to permanent LAD-ligation, such as dilated chamber dimensions, and thus resemble features of human HFrEF [257]. However, the severity of HF induced by transient LAD-ligation highly depends on the time of ischemia. As depicted in Table 1, the degree of cardiac dysfunction and the size of the infarct zone are typically less pronounced in ischemia-reperfusion (I/R) models than in models of permanent LAD-ligation [179]. Next to the ischemic damages in the I/R model, subsequent reperfusion exposes the cardiac tissue to oxidative stress, which exaggerates the tissue damage; this additional injury is entitled reperfusion injury [184, 256]. Thus, murine I/R models are likewise used for investigations that focus on the pathophysiology of this reperfusion injury.

Methodological proceedings

The surgical preparation for the transient LAD-ligation in mice is performed similar to the procedure of inducing permanent LAD-ligations. Commonly, one-step procedures are performed in which the surgical preparation is immediately followed by the transient LAD-occlusion [179]. A sufficient occlusion may be verified by blanching of the cardiac tissue distal to the ligation, as shown in Fig. 2a, or by an ST-elevation in the ECG [120]. In the literature, ligation is maintained for 20–60 min [120, 178, 202, 257, 287]. However, LAD-ligation of less than 30 min may not provoke ischemia-induced myocardial injury, whereas ligation of 60–90 min results in irreversible CM death and complete infarction of the area at risk as depicted in Fig. 2b [155]. Thus, 45–60 min of ischemia is recommended for the transient LAD-ligation. The observation period after subsequent reperfusion ranges from a few minutes up to several weeks [120, 178, 202, 257, 287], depending on the respective scope of the study.

The area at risk is defined as the area of non-perfused cardiac tissue, in which the wave front of the infarct zone subsequently extends (Fig. 2b). The size of the area at risk following transient LAD-ligation can be estimated by injecting Evans Blue into the circulation. This dye is subsequently distributed to the perfused tissue but leaves the non-perfused area at risk unstained. Perfusing the heart with Evans blue can either be performed in vivo or ex vivo [155, 273]. Afterwards, the size of the infarct zone can be estimated similar to permanent LAD-ligation using the triphenyltetrazolium chloride staining in the harvested cardiac tissue.

Table 1Comparison of MIinduced by permanent ortransient LAD ligation in mice

MI induced by						
Permanent i	schemia	Ischemia/reperfusion				
+++/++		++/+				
Transmural		Midmyocardial				
_		++				
+++		++				
+++		++				
+++	(† 24 h)	++/+	(† 6 h)			
+++	(† 7 d)	++/+	(† 3 d)			
+++	(† 7 d)	++/+	(† 3 d)			
++	(† 3 d)	++/+	(† 24 h)			
++	(† 7 d)	+/	(† 3 d)			
++	(† 7 d)	+/	(† 3 d)			
+	(† 7 d)	+	(† 6 h)			
++	(† 7 d)	+	(† 6 h to 3 d)			
+++	(† 4 d)	++	(† 3 d)			
28 d		14 d				
	MI induced Permanent is ++++/++ Transmural - ++++ +++ +++ +++ +++ +++ +++	MI induced by Permanent ischemia ++++/++ Transmural - +++ ++++ († 24 h) ++++ († 7 d) +++ († 4 d) 28 d	MI induced by Ischemia/reperfus Permanent ischemia Ischemia/reperfus ++++/++ Transmural Midmyocardial - ++ ++++ ++ +++ ++ +++ ++ +++ ++ +++ († 24 h) ++/+ +++ ++ +++ († 7 d) ++/+ +++ († 7 d) ++/+ +++ († 7 d) +/- ++ († 7 d) +/- ++ († 7 d) + +++ († 7 d) + +++ († 7 d) + 28 d 14 d 14 d			

Data from the following references were used: [42, 119, 262, 285] and [111]

 (\uparrow) peak of the event, + increased, - not changed

d days, h hours

Recent advancements LAD-ligation protocols are surgical procedures that, by themselves, may induce systemic inflammatory responses. To avoid that these systemic inflammatory responses interfere with local inflammatory processes during cardiac repair, the procedure may be split into a preparative surgery which is followed by the transient LAD-ligation several days after the initial surgery [120, 197]. Furthermore, the use of volatile anaesthetics during ischemia can precondition the murine heart to face reperfusion injury for up to 72 h [213]. Therefore, splitting the procedure avoids this preconditioning. However, systemic inflammatory responses after LAD-ligations may be limited to rather invasive procedures that include, e.g. thoracotomies and may not be seen in minimally invasive procedures. In line with this, a recent study concludes that minimally invasive sham LAD-ligations induce several pro-inflammatory genes but none of these genes passes the threshold required for increased protein expression [114].

Distinct characteristics of post-infarct remodelling in models of permanent and transient LAD-ligation

The degree and the temporal dynamics of post-infarct remodelling in mice depend on a proximal or distal ligation of the LAD and the use of reperfusion protocols [285]. Compared to permanent LAD-ligation, ischemia/reperfusion (I/R) subjects murine hearts to (1) a limited ischemic period and (2) a restored blood flow which allows effector cells to quickly take hold of the affected area and provides metabolites to the cells in the infarct wall. As a consequence, murine I/R models possess specific features that make them different from models of permanent LAD-ligation. These differences are summarised in Table 1. In general, murine I/R induces smaller infarct areas that are primarily located midmyocardial as compared to the transmural infarct areas observed after permanent LAD-ligation [93, 119, 285]. Further, reperfusion lowers the degree but accelerates the temporal dynamics of the postischemic remodelling [4, 285]. This is associated with less fibrotic remodelling and less impaired contractile function in the chronic phase after MI in I/R models [4, 285]. Beyond that, the time point of reperfusion may crucially affect the relative contributions of distinct cellular effectors to the postischemic pool of activated interstitial cells [4]. Due to their loss in the infarct zone, prolonged ischemia favours the recruitment of interstitial cells from the non-infarcted tissue or the blood stream, whereas quick reperfusion may favour the activation of viable resident myocardial cells [4].

Cardiac inflammation and fibrosis induced by ischemia

In mammals, adult CMs possess negligible self-regenerative capacity [53, 186]. Hence, cardiac remodelling after MI

ultimately results in the formation of a fibrotic scar. This process of reparative fibrosis consists of three sequential stages [190, 213, 245]: (1) during the initial inflammatory phase (acute phase), necrotic CMs release danger signals; this rapidly activates resident sentinel cells, which initiate the recruitment of pro-inflammatory cells to clear the infarct wall from the cellular debris and the defective ECM [61, 190]. The sufficient clearance of necrotic CMs, ECM fragments and apoptotic immune cells in the infarct wall promotes the resolution of inflammatory signals and thus the transition to the proliferative phase. (2) During the subsequent proliferative phase (subacute phase), cardiac fibroblasts (CFs) transdifferentiate into myofibroblasts (MFs) and secrete an abundant amount of collagen and non-structural matrix proteins to compensate the cellular loss [133, 190]. (3) During the maturation phase (chronic phase), the immature fibrillar collagens that have been secreted during the preceding proliferative phase are enzymatically crosslinked and thereby gain stability, whereas the cellular effectors mostly vanish [133, 190]. The degree of inflammation and fibrosis following MI in mice varies throughout distinct regions in the cardiac muscle, depending on their proximity to the infarct zone. Therefore, in models of murine MI the left ventricle is most often divided into three distinct zones: infarct zone, border zone and remote zone. In general, inflammation and fibrosis are most pronounced in the infarct zone, followed by the border zone, but can also spread into the remote zone in severe cases [94]. Table 1 provides an overview of the distinct temporal dynamics of cardiac inflammation and fibrosis in the murine models of permanent and transient LAD-ligation.

Acute phase after MI—the inflammatory phase

The inflammatory phase after MI plays an ambiguous role, as it is a pivotal basis of cardiac repair but also a crucial factor for a subsequent development of HF. Thus, the local and temporal restriction of inflammation is crucial for sufficient infarct healing. A schematic drawing of the tightly regulated interplay between pivotal cellular and molecular players of the inflammatory phase and its subsequent resolution is shown in Fig. 3.

Initiation of inflammatory cascades—danger signals, sentinel cells and pro-inflammatory mediators MI is pathologically defined as myocardial cell death due to prolonged ischemia [256]. Necrotic CMs release danger signals, the so-called damage-associated molecular patterns (DAMPs), such as mitochondrial DNA (mtDNA), the chromatin protein high mobility group box 1 (HMGB1), purine metabolites, sarcomeric protein fragments and S100 proteins [19, 57, 136, 158]. However, the cellular specificity of DAMPs is poorly understood [57]. They initially alarm a variety of sentinel cells, such as resident macrophages (MPs), endothelial cells (ECs), resident cardiac fibroblasts (CFs), surviving CMs in the border zone and subsequently also invading leukocytes [57]. The main pattern recognition receptors (PRRs) of the cardiac sentinel cells are (1) a variety of tolllike receptors (TLRs), (2) the receptor of advanced glycation end-products (RAGE) and (3) NOD-like receptors (NLRs) [211]. PRRs are either localised in intracellular compartments or on cellular surfaces. Therefore, specific PRRs play distinct roles in phagocytes and non-phagocytes. Recently, interactions of DAMPs and their corresponding PRRs in the context of MI have been comprehensively reviewed [211]. Downstream PRR-signalling activates the transcription factor nuclear factor kappa B (NF κ B), which subsequently elevates the expression of chemokines, cytokines and celladhesion molecules (CAMs) in the infarct wall [61, 80, 168, 211]. While secreted chemokines attract leukocytes, CAMs exposed on ECs facilitate their invasion [211]. Distinct chemokines determine the recruitment and composition of specific immune cells in the myocardium: while CC chemokines, such as CC-chemokine ligand (CCL)-2, -5, and -7, attract monocytes, lymphocytes and mast cells, CXC chemokines, such as CXCL-8, attract neutrophils [61, 133]. Pro-inflammatory cytokines, such as TNF- α and the interleukins (IL)-1β, -6, and -18, further amplify the inflammatory response by activating resident and invading effector cells [211]. In this context, interleukin (IL)-1 β may play a crucial role as it mediates myocardial chemokine synthesis and induces a pro-inflammatory M1 polarisation in MPs [22, 61, 133]. In line with this, IL-1 receptor type I deficient mice exhibit less inflammation and less fibrosis following MI [22]. Both, leukocytes and activated CFs, secrete the mature form of IL-1 β ; this requires the preceding caspase-1-mediated cleavage of its precursor pro-IL-1ß in a multiprotein complex entitled 'inflammasome'; the complex activation of inflammosomes is comprehensively reviewed elsewhere [233]. In the early phase after MI, IL-6 upregulates the expression of hyaluronan-synthases, which in turn provides a basis for MF transdifferentiation and thus cardiac repair; hence, IL-6 blockade may be detrimental after MI [188].

Oxidative stress and the reperfusion injury Next to the irreversible CM-necrosis in the infarct zone, the transient depletion of oxygen and other metabolites in the area at risk switches the metabolism of viable cells to anaerobic glycolysis and disrupts appropriate ion homeostasis [88]. This results in intracellular acidosis and accumulation of Ca^{2+} and Na^+ [88]. During acute ischemia, high intracellular Na^+ and H^+ protect the cell from mitochondrial damage as they close the mitochondrial permeability transition pore (MPTP) and prevent the deteriorating effects of myofibril hypercontracture [88]. Upon reperfusion, oxidative phos-

Mvocardial Fibrotic infarction remodellina day 0 day 7 Ischemia induces Alarmed by their PRRs, Thereby, they recruit invading Efferocytosis resolves inflammatory immune cells that efferocyte CM-necrosis and the resident sentinel cells are activated stimuli and induces a reprogramming in release of danger signals and initiate inflammatory cascades dead cells and fragment the ECM cardiac cells and thus a proliferative milieu



Fig. 3 The initiation of inflammation and its subsequent resolution in the context of myocardial infarction. Ischemia induces necrosis of cardiomyocytes (nCM) and thus the release of danger signals, such as mitochondrial (mt)DNA, the chromatin protein HMGB1, purine metabolites, sarcomeric fragments, S100 proteins and reactive oxygen species (ROS). Subsequently, resident sentinel cells, such as endothelial cells (ECs), cardiac fibroblasts (CFs) and macrophages (MPs) are activated (a) by their pattern recognition receptors (PRRs) and secrete pro-inflammatory mediators, e.g. interleukins (ILs), chemokines (CCLs), and tumor necrosis factor (TNF)- α . Further, ECs express cell-adhesion molecules (CAMs) upon activation and thereby facilitate leukocyte recruitment. Hereafter, leukocytes invade the injured tissue and are immediately confronted with the pro-inflammatory environment. Hence, they gain a pro-inflammatory signature (M1-polarised macrophages and N1-polarised neutrophils) and further amplify the inflammatory response by secreting pro-inflammatory cytokines, but also matrix metalloproteinases (MMPs). MMPs

phorylation and acid–base balance are rapidly restored within cells of the area at risk [88]. This leads to (1) opening of the mitochondrial permeability transition pore (MPTP) and thus mitochondrial damage, (2) myofibril hypercontracture and (3) excessive accumulation of reactive oxygen species (ROS) [88]. ROS further aggravate the reperfusion injury as they induce the dysfunction of the sarcoplasmic reticulum, oxidise membrane lipids and DNA and promote endothelial dysfunction [88, 170]. Moreover, ROS potently

cleave the matrix and thus generate matrix fragments (matrikines) which additionally serve as pro-inflammatory stimuli. Subsequently, M1-MPs and aCFs clear the wound from necrotic CMs but also from apoptotic neutrophils and matrikines and thereby resolve the proinflammatory environment. As they passage to the proliferative phase, the effector cells of the inflammatory phase undergo a second reprogramming and gain a reparative signature, by which they become myofibroblasts (MFs), N2-polarised neutrophils and M2-polarised MPs. Subsequently, these phagocytes, together with regulatory T-cells (Tregs), release anti-inflammatory stimuli such as transforming growth factor (TGF)-\u03b31, IL-10, lactoferrin, annexin A1, lipoxins and resolvins which further repress inflammation and provide a proliferative milieu. During the subsequent proliferative phase, predominantly MFs, but also M2-polarised MPs secrete abundant ECM proteins, e.g. collagens and matricellular proteins and thereby promote the formation of the myocardial scar

attract leukocytes [88, 170]. Accordingly, reperfusion of the area at risk following ischemia induces an additional reperfusion injury, which is of particular importance in the murine model of transient LAD-ligation. The current understanding of the pathophysiology of the reperfusion injury is reviewed in more detail elsewhere [88, 92].

Initial matrix remodelling and the first provisional matrix Activated resident cells, especially CFs and MPs, as

well as recruited leukocytes secrete and activate matrix metalloproteinases (MMPs) of both subgroups, the collagenases (e.g. MMP-8 and -13) and the gelatinases (e.g. MMP-2 and -9) [58]. Recently, the role of MMPs following MI has been comprehensively reviewed [154]. By degrading ECM proteins, MMPs generate matrix fragments (matrikines) which additionally serve as pro-inflammatory stimuli [133]. Well-characterised matrixines are (1) proline-glycine-proline (PGP), a collagen fragment which attracts neutrophils through binding to their CXC-receptor-1 and -2 and (2) a fragment of the glycosaminoglycane hyaluronan, whose delayed clearance is associated with excessive inflammation [58, 69, 264]. Beyond pro-inflammatory features, distinct matrikines may also orchestrate proliferative cellular responses, as, e.g. a fragment of collagen XVIII (endostatin) enhances fibroblast proliferation [58]. A plasma-derived, provisional matrix initially replaces the fragmented ECM; rapid upregulation of vascular endothelial growth factor (VEGF) in the infarct wall increases vascular permeability, which leads to the extravasation of the plasma proteins fibrin and fibronectin [5, 58]. This fibrin-based provisional matrix initially stabilises the infarct wall and provides a scaffold for leukocyte invasion [35, 44, 48]. Furthermore, through interactions with cellular integrins, it may alter the inflammatory profile of migrating leukocytes and CFs [58]. However, the specific regulatory effects of the early fibrin-based matrix remain poorly understood.

Endothelial cells Despite the common understanding of the cellular composition of the non-myocyte fraction in the mammalian heart during the last decades, recent findings revealed that ECs are the most abundant non-myocytes (>60%) in the healthy murine heart [209]. This changed the preceding doctrine that CFs are the most abundant ones. Upon stimulation with DAMPs and pro-inflammatory cytokines, ECs facilitate leukocyte extravasation into cardiac tissue [211]. Therefore, they (1) immediately mobilise preformed P-selectin from their weibel-palade bodies [211] and (2) subsequently upregulate the expression of CAMs, such as ICAM, VCAM and selectin on their cell surface. Further, ECs are a relevant source of cytokines and chemokines after MI [138, 292]. Recently published work shows that the proinflammatory activation of ECs after MI may be triggered by the CM-dependent release of natriuretic peptides, such as ANP and BNP [28]. This release is an immediate response after MI and correlates with the severity of the event [28]. Since ECs immobilise chemokines to glycosaminoglycanes on their surface, ECs also play a pivotal role for the local concentration of leukocyte attractors in the injured myocardium [61, 212].

Cardiac fibroblasts In the healthy murine heart, resident CFs represent less than 20% of non-myocytes [209]. Nonethe-

less, after MI their population undergoes marked expansion. In spite of the debate surrounding the origin of activated CFs in the infarcted murine heart, there is a growing body of evidence that the majority of activated CFs during MI originates from resident interstitial Tcf21⁺ fibroblasts [4]. In vitro experiments showed that CFs secrete pro-inflammatory mediators (IL-1 β , -1 α , -6, -8, and TNF- α) and matrixdegrading proteases (MMP-2, -3, -9, -12) upon stimulation with DAMPs or ROS [59, 154, 243]. However, their in vivo role in initiating cardiac inflammation after MI has just recently been shown. In isolated CFs, the loss of the G protein-coupled receptor kinase (GRK)-2 attenuates the expression of pro-inflammatory cytokines [278]. Thus, recent investigations in mice with a fibroblast-specific depletion of the G protein-coupled receptor kinase (GRK)-2 provide the first direct in vivo evidence for the pro-inflammatory role of CFs in this context, as these mice exhibit less neutrophils in their infarct walls, smaller infarct sizes and better contractile function 24 h after I/R [278]. The physiological signification of the pro-inflammatory feature of CFs may be that the secretion of inflammatory cytokines, especially IL-1β, prohibits premature MF transdifferentiation [229] and might therefore ensure the proper clearance of the infarct wound before scarring [59]. Beyond that, CFs mediate CM-apoptosis by secreting (1) pro-apoptotic inflammatory cytokines, (2) matrix-degrading MMPs, which deprive a sufficient ECM-environment, and (3) pro-apoptotic exosomes [59].

Neutrophils Neutrophils are the first cellular responders [61] and invade especially the border zone, mainly attracted by CXC chemokines [57, 211]. In comparison to other leukocytes, their number increases immediately after MI. While the highest neutrophil/leukocyte ratio (25%) in the infarct zone is seen at day 1, their total number increases until day 3 after MI (Table 1) [285]. Subsequently, the neutrophil/leukocyte ratio decreases to less than 5% at day 5 after MI [285]. Recent findings reveal that the degree of neutrophil recruitment into the infarcted cardiac tissue in mice shows a circadian kinetic, as it depends on the time-of-day onset of MI [232]. Confronted with the pro-inflammatory environment, infiltrating neutrophils exhibit an inflammatory N1-phenotype, which depends on the DAMP-mediated activation of TLR-4 [167]. Neutrophils play ambiguous roles during the post-MI remodelling: On the one hand, the number of inflammatory neutrophils positively correlates with infarct wall thinning in mice, which was related to their secretion of MMP-12 and -25 [167]. Furthermore, neutrophils secrete highly toxic mediators, such as myeloperoxidase, which worsen the postischemic remodelling and promote arrhythmia [182]. Beyond their direct proinflammatory actions, neutrophils mediate the recruitment of monocytes [57]. Recent findings reveal that this happens

in a synergistic cooperation with platelets, as human neutrophil peptide (HNP)-1 needs to form heteromers with the platelet-derived chemokine CCL-5 to fulfil its functions [3]. On the other hand, the recruitment of early inflammatory N1-neutrophils is essential for their subsequent N2-polarisation [100]. N2-polarised neutrophils were recently identified to contribute to the reparative M2-polarisation of MPs during the proliferative phase [100] and may thus be essential for a sufficient formation of the scar.

Monocytes/macrophages A relatively low number of nonmyocytes in the healthy cardiac tissue are assigned as leukocytes (approx. 7%); however, the vast majority of these cells are identified as resident MPs [49, 209, 291]. Thus, resident MPs play a crucial role as sentinel cells that initiate cardiac inflammation early after MI. Moreover, their absolute numbers increase following MI [285]. However, as neutrophils exhibit faster kinetics in invading the cardiac tissue during the acute phase, the MP/leukocyte ratio declines to approx. 60% until day 3 after MI [285]. Hereafter, further monocytes invade, whereas neutrophils undergo apoptosis; thus the MP/leukocyte ratio is reconstituted to more than 80% between day 7 and 14 after MI [285]. Hence, the cardiac recruitment of tissue-invading pro-inflammatory Ly6C^{hi}-monocytes, which later differentiate into MPs, is a key event in postischemic remodelling. The recruitment of monocytes is mediated in four sequential stages: (1) MI alters the hemodynamic conditions and thereby promotes the release of catecholamines from the adrenal glands into the blood stream. (2) high plasma levels of catecholamines repress stem-cell retention factors (e.g. CXCL-12) in bone marrow and spleen stem cell niches and thereby enable monocytes to leave their niches and enter the blood stream [48]. (3) the upregulated CCL-2 and CCL-7 expression in the injured myocardium attracts these pro-inflammatory monocytes and subsequently (4) mediates their invasion into the cardiac tissue [48]. The mechanistic significance of the initial catecholamine release from the adrenal glands is highlighted, as adrenergic β_2 -receptors are pivotal for the expression of monocytic CC-chemokine-receptor (CCR)-2 and thus for monocyte recruitment [81, 82].

Hereafter, MP colony-stimulating factor (M-CSF) promotes the differentiation of invading monocytes into MPs [63]. Once they are differentiated, MPs are confronted with the hypoxic environment in the infarct wall. This promotes the expression of hypoxia-inducible factor (HIF)-1, which alters MP metabolism towards glycolysis, a key feature of M1-polarisation [187]. Further, the high concentration of DAMPs, IL-1 β and TNF- α in the infarct wall contributes to their pro-inflammatory M1 polarisation [187]. Early M1 MPs secrete and activate MMPs and can thereby dig through the infarct zone to engulf and remove necrotic CMs [133, 154, 187]. However, to limit excessive

activation of inflammatory pathways in these early M1 MPs, several anti-inflammatory genes, such as schlafen (Slfn)-4 and CD-9, are concomitantly upregulated [187].

Mast cells The key feature of mast cells is their high cytoplasmic density of preformed granules. Thus, they are able to immediately respond to external stimuli. Despite that the number of mast cells may not be significantly increased after MI in mice [43], they are an important source of TNF- α and thus initiate the cytokine cascade and may orchestrate cardiac inflammation [62, 127]. However, mast cells play an ambiguous role in the postischemic remodelling: on the one hand, mast cell-deficient mice [145], as well as mice with a knockout of the mouse mast cell protease (mMCP)-4 [101], exhibit improved prognosis after MI. In the context of MI, the secretion of the chymase mMCP-4 promotes cardiac inflammation and subsequent fibrotic remodelling by activating IL-1β, angiotensin II, endothelin-1, MMP-9 and fibronectin [101]. On the other hand, mast cell deficiency has been shown to be beneficial in mice within the first hours after MI [18], but to induce larger infarct sizes and accentuated ventricular dilation during the chronic phase [30]. Further, a recent study concludes that mast cells preserve contractile function during MI by upregulating the Ca²⁺-sensitivity of myofilaments [196]. Thus, the role of mast cells in cardiac postischemic remodelling is still under debate.

Lymphocytes For decades, investigations focused on the role of the innate immunity in the postischemic remodelling. In contrast, the role of the adaptive immunity in postischemic remodelling following MI has been subject to less scrutiny [99]. Recently, the role of the adaptive immunity in the context of MI has been comprehensively reviewed [99]. Briefly, the invasion of adaptive immune cells into cardiac tissue reaches a peak at day 7 [99, 285]. In this context, the depletion of B-cells has been shown to be beneficial in murine models of MI [99]. However, knockout studies revealed that the depletion of CD4⁺ T-cells deteriorates cardiac function and worsens cardiac remodelling after MI in mice [99]. CD4⁺ T-cells in postischemic remodelling are predominantly Th1-polarised, whereas Th2- and Th17-polarised T-cells are not increased following MI [99, 285]. The role of regulatory T-cells (Tregs) in postischemic remodelling is highlighted as the depletion of Tregs deteriorates and the stimulation of Tregs vice versa ameliorates adverse cardiac remodelling [99, 253]. In general, Tregs downscale cardiac inflammation (1) by secreting anti-inflammatory mediators (e.g. IL-10 and TGF- β 1), (2) by cell-to-cell interactions that inhibit proinflammatory T-cells and (3) by cell-to-cell interactions that inhibit antigen presentation by, e.g. dendritic cells

[177]. The specific role of Tregs in cardiovascular disease has recently been reviewed comprehensively [177].

Resolution of cardiac inflammation and transition to the proliferative phase

The resolution of cardiac inflammation is mandatory for the transition to the proliferative phase and thus for a sufficient formation of the scar. The resolution of inflammation is associated with the clearance of dead tissue and the subsequent reprogramming of effector cells which in turn release anti-inflammatory molecules.

Removal of inflammatory stimuli and reprogramming of phagocytes Efferocytosis is the phagocytosis of dead cells and a key signal that dampens and finally resolves the inflammation in the injured myocardium after MI [211]. Efferocytosis implies the removal of (1) necrotic CMs in the infarct zone, (2) apoptotic CMs in the border zone, and (3) apoptotic leukocytes, especially neutrophils, which undergo apoptosis owing to their short life-span of 3–7 days [61, 194, 211]. Sufficient efferocytosis and thus sufficient resolution of cardiac inflammation depends on the myeloid-epithelial-reproductive protein tyrosine kinase (MERTK) and the milk fat globule-EGF factor 8 (MFG-E8) [102]. Moreover, MFs also contribute to the resolution of inflammation as they engulf dead cells similar to MPs in a MFG-E8-dependent manner [193]. The level of MFdependent efferocytosis is almost 40% of that of MPs [193]. Moreover, MPs internalise matrikines and thereby eliminate further pro-inflammatory stimuli [210]. However, efferocytosis does not only clear the wound from pro-inflammatory stimuli, but also promotes a reprogramming of MPs; initially, the early pro-inflammatory M1 MPs reprogramme and become intermediate MPs, which already perform oxidative phosphorylation, a key feature of M2-polarisation [187, 220]. Thus, the expression of AMP-activated kinase (AMPK), which mediates the metabolic conversion from glycolysis to oxidative phosphorylation [266], may display a crucial factor for MP reprogramming [187]. During further progression from the inflammatory to the proliferative phase, MPs undergo a second reprogramming by which they gain a reparative M2-signature. This second reprogramming depends on an IL-6 and IL-10 initiated pathway, the signal transducer and activator of transcription (STAT)-3 pathway. Additionally, TGF- β and IL-4 initiated signalling is required in this context [187, 263]. Similar to MPs, MFs acquire an anti-inflammatory signature during the progression from the inflammatory to the proliferative phase in an efferocytosisdependent manner [193].

Key mediators of anti-inflammatory, pro-fibrotic signalling Several anti-inflammatory and concomitantly profibrotic mediators orchestrate the transition from the inflammatory to the proliferative phase: (1) apoptotic neutrophils release lactoferrin and annexin A1, mediators that inhibit further leukocyte invasion [61]. (2) efferocytosis of dead cells stimulates MPs to secrete anti-inflammatory TGF-\u00b31, IL-10 and anti-inflammatory lipids like lipoxins and resolvins [61, 108, 246]. (3) Tregs additionally secrete TGF-β1 and IL-10 [57]. (4) MMPs limit leukocyte invasion by cleaving CCLs [274], which converts the pro-inflammatory leukocyte attractors to anti-inflammatory chemokine receptor antagonists and thus prohibits further leukocyte invasion [77, 275]. Furthermore, MMPs cleave CCL binding sites on ECs, the glycosaminoglycanes, which prohibits local chemokine accumulation [77, 274]. (5) heparin-binding domains on fibrin molecules of the provisional matrix provide a reservoir for anti-inflammatory, proliferative growth factors, such as TGFs, PDGFs, FGFs, and VEGFs [171].

Subacute phase after MI—the proliferative phase

The proliferative phase after MI is characterised by the proliferation of interstitial cells, a highly dynamic ECM and the vascularisation of the granulation tissue.

TGF-B1 and the transdifferentiation of cardiac fibro**blasts** The pivotal mediator of the proliferative phase is thought to be TGF- β 1 [56]. Though there are three TGF- β isoforms, TGF-β1 is the major and best characterised isoform in the cardiac tissue [56]. TGF- β 1 strongly represses inflammation and matrix degradation but enhances ECM enlargement and cell proliferation. It stimulates the antiinflammatory polarisation of leukocytes and the transdifferentiation of CFs [41]. Furthermore, it induces anti-inflammatory protease inhibitors, such as plasminogen activator inhibitor (PAI)-1 and tissue inhibitors of MMPs (TIMPs) [133, 231]. In the healthy heart, TGF- β 1 is bound to the ECM as an inactive, latent complex [6, 222]. The release of the bioactive form from latent stores is mediated by proteases like plasmin [165, 222] and MMPs, such as MMP-2 and MMP-9 [288]. Moreover, matricellular proteins, ROS and the acidic environment promote the release of mature TGF- β 1 from its latent form after MI [12, 165]. Beyond the mobilisation of TGF-β1 from latent stores, CFs, ECs, MPs, and platelets actively secrete mature TGF-\u00b31 [56]. TGF-\u00b31 binds to its type II receptor (T β RII) on the cell surface of various target cells; subsequently the receptor-ligand complex transphorylates the cytoplasmic domain of the type I receptor (TBRI) [242]. Hereafter, downstream intracellular signalling involves Smad2/3-dependent as well as Smadindependent pathways [133, 242]. TGF-β1 requires the concomitant presence of EDA-fibronectin to induce MF transdifferentiation [9, 238] by enhancing the Smad3-dependent α -smooth muscle actin (α -SMA) expression in CFs [45, 68, 97]. Beyond that, Smad3-signalling in MFs crucially regulates the integrin-mediated formation of organised MF arrays [134]. Furthermore, hyaluronan significantly contributes to MF transdifferentiation, as the loss of its main receptor CD44 results in lower collagen secretion after murine MI [106]. Though lineage tracing identified resident CFs as the major source of MFs [4], pericytes, ECs as well as circulating fibrocytes of bone marrow origin may also contribute to the population of MFs in cardiac repair [133, 183]. Key features of MFs are a highly active endoplasmic reticulum and α-SMA-containing stress fibres, which allow their contraction [95, 96]. Their stress fibres are connected to specific domains of extracellular fibronectin in an adhesion complex called 'fibronexus' [52]. Thereby, MFs are able to exert tensile force on the surrounding ECM [52]. This may explain the reduction of the scar size as compared to the initial infarct size by about 50% [262]. During the proliferative phase, MFs undergo a rapid expansion primarily in the border zone and secrete structural proteins, such as the fibrillar collagen-I and -III [31, 265]. Additionally, MFs secrete matricellular proteins upon stimulation with several growth factors or angiotensin II [58, 95]. Next to MFs, reparative M2 MPs were recently shown to deposit ECM proteins during the proliferative phase [187].

The second provisional matrix During the proliferative phase, the secretion of plasminogen and its subsequent activation disintegrate the provisional, fibrin-based matrix [36]. In line with this, plasminogen-deficient mice exhibit an impaired formation of the granulation tissue after MI (61). The first plasma-derived matrix is subsequently replaced by a second, cell-derived one: MFs and MPs mainly secrete fibronectin [20], but also hyaluronan, proteoglycans and matricellular proteins [58]. Since distinct growth factors, most importantly TGF- β 1, require interactions with matrix components for their signal transduction, the cell-derived matrix becomes a conductor of the remodelling orchestra during proliferative phase [58]. Further, the requirement for a cooperation between matrix components and growth factors restricts the proliferative signalling to the infarct wall [58].

Matricellular proteins Matricellular proteins display their highest expression in the border zone and serve as molecular bridges between ECM proteins and surface receptors [58]. Thereby, they limit inflammation and induce reparative reprogramming in interstitial cells [58]. Further, their requirement for growth factor and cytokine signalling prevents excessive fibrotic remodelling after MI since they localise proliferative stimuli to the infarct wall and temporarily restrict pro-fibrotic signals [58]. Well-characterised matricellular proteins are, e.g. thrombospondins (TSPs), tenascins, the CCN-family, periostin, osteopontin, and secreted

protein acidic and rich in cysteine (SPARC) [58, 133, 221]. TSP-1 and -2 are markedly upregulated in the border zone after MI [58]. They activate TGF-β1 and inhibit MMP-2 and -9, by which they build a barrier against the diffusion of pro-inflammatory molecules into the remote zone [64, 221]. CCN-2 has TGF-\u00b31 synergistic, proliferative effects on interstitial cells, but also promotes CM hypertrophy [213]. Osteopontin prevents excessive cardiac dilation by stimulating collagen deposition into the infarct zone [258]. The depletion of SPARC in the context of MI, however, results in higher rates of cardiac rupture and cardiac dysfunction due to immature collagen and disorganised granulation tissue [230]. Beyond that, the clearance of matricellular proteins supresses proliferative stimuli and thereby promotes the transition from the proliferative phase to the maturation phase [58]. In this context, the stabilin-1-mediated phagocytosis of SPARC is a well elaborated signal [140].

Vascular remodelling Vascular remodelling is a major process during the proliferative phase. Sprouting of former quiescent ECs upon hypoxia and high concentrations of VEGF induce the formation of neovessels [32, 61, 146]; myeloid cells secrete VEGF in an efferocytosis-dependent manner [102]. Neovessels function as distribution lanes that provide oxygen and nutrients to metabolically active cells in the wounded area. During the inflammatory phase, neovessels are hyperpermeable and lack pericytes and thus facilitate leukocyte extravasation [61]. However, during the proliferative phase, mature neovessels are covered by a pericyte coat, which prohibits leukocyte extravasation and the angiogenic potential of ECs [61, 215, 294]. The formation of a pericyte coat requires platelet-derived growth factor (PDGF)-signalling and is mandatory for the formation of a mature scar [294]. A lack of the pericyte coat owing to the disruption of PDGF-signalling prolongs immune cell extravasation and decreases the collagen content and thus the stability of the mature scar [61].

Chronic phase after MI—the maturation phase

The key feature of a mature murine scar is an ECM that consists of cross-linked collagens, whereas the cellular effectors, such as MFs and ECs, have vanished.

Formation of a mature scar The apoptosis of interstitial cells is induced by (1) the removal of matricellular proteins, (2) the deprivation of growth factors and (3) a reduced tensile force on MFs [215, 294]. The small proteoglycan biglycan orchestrates collagen organisation in the scar and an increased expression of lysil-oxydase catalyses the collagen cross-linking [55, 159, 273]. While collagen cross-linking increases the tensile strength of the myocardium and thereby may prevent chamber dilation, it can also induce

diastolic dysfunction owing to increased cardiac stiffness. The formation of a supportive scar depends on the processes that take place during inflammation and proliferation: initial overshooting of inflammatory signals favours matrix degradation and thereby facilitates cardiac rupture during the acute phase [61]. Prolonged inflammation, however, delays the proliferative phase and thus impairs the deposition of fibrillar collagen, which may lead to ventricular dilation during the chronic phase after MI, owing to impaired tensile strength of the scar [61]. There are several differences between murine postischemic scars and those of larger mammals: (1) mice display extraordinary thin post-MI scars, due to lesser collagen in their scars [44]. (2) MFs do not exhibit a complete evanescence after scar formation in human MI but rather align parallel to the endocardium and the epicardium and thereby support the extracellular scar tissue [276]. Both may explain, why mice are highly prone to cardiac rupture after MI, whereas the phenomenon of postischemic rupture is rare in humans [121]. (3) myocardial scar formation is fast in mice compared to larger mammals as a mature scar is present within 14 days in murine I/R models, whereas scar formation in canine I/R protocols requires 28 days [43].

Interstitial remodelling upon myocardial infarction

Sufficient scar formation resolves proliferative signalling in the infarct zone; however, hemodynamically relevant MIs induce cardiac pressure and volume overload. This in turn increases neurohumoral activation and ventricular wall stretch [58] and thus induces cardiac hypertrophy and interstitial fibrosis in the primarily unaffected remote zone of the myocardium. Moreover, inadequately restricted cardiac inflammation extends the infiltration of immune cells into the remote zone. Here, these immune cells initiate interstitial inflammation and subsequent fibrosis without a mechanistic signification, which worsens cardiac remodelling and thus impairs cardiac function [61]. However, interstitial remodelling in the remote zone after MI in mice is less pronounced than in the prototypical murine models of hypertrophy and interstitial fibrosis, such as transverse aortic constriction [216].

Models of heart failure induced by pressure overload and neurohumoral activation

Transverse aortic constriction

Murine heart failure and its clinical relevance

Transverse aortic constriction (TAC) mimics increased left ventricular afterload in patients, which in the clinical setting predominantly occurs owing to aortic stenosis or systemic hypertension. TAC, and its clinical correlates, subject the left ventricle to pressure overload and subsequently activate neurohumoral stimuli. As a reaction to pressure overload and neurohumoral activation, cardiac hypertrophy and interstitial fibrosis occur. However, the acute onset of pressure overload in the murine model of TAC does not perfectly resemble the patients' clinical course in which the pressure overload insidiously evolves over years or decades owing to aortic stenosis or hypertension. In mice, the model of TAC has been extensively used to examine signalling pathways that contribute to adverse cardiac remodelling and hypertrophy in the context of pressure overload and neurohumoral activation [112, 225, 252, 260]. As shown in Table 2, mice that are subjected to TAC exhibit cardiac remodelling in two sequential stages: (1) an initial stage of concentric hypertrophy and stable systolic but impaired diastolic function, thereby resembling features of human HF with preserved ejection fraction (HFpEF) [90, 252, 260]. (2) over time, the mice decompensate and exhibit severe systolic HF accompanied by eccentric hypertrophy, thereby resembling features of human HFrEF [90, 175, 252, 260]. However, the progression from HFpEF to HFrEF is not a mandatory sequela in the patients' course of HFpEF [47].

Methodological proceedings

As depicted in Fig. 4a, TAC is performed by constricting the transverse aortic arch between the brachiocephalic trunk and the left carotid artery [225]. First, mice are anaesthetised and artificially ventilated. Hereafter, the chest is entered by an upper partial sternotomy and a suture is tied around the aortic arch against a cannula [225, 252]. The use of a cannula ensures that the aortic lumen is constricted to a defined diameter. By removing the cannula, the inner aortic diameter is reduced to about 0.4 mm, which constricts the aortic arch by approximately 70% [225]. A sufficient constriction establishes a pressure gradient across the stenosis of more than 40 mmHg [147]. Non-invasive pulse-wave doppler analysis can be used to measure this gradient and to subsequently select only animals with comparable gradients for further evaluation, which should be done to ensure a sound study [147, 157, 199]. However, the analysis of pressure gradients using a doppler ultrasound is limited to a maximum gradient of 60 mmHg across the stenosis [157]. Data on periprocedural mortality of TAC-surgeries range from less than 10% [225, 252] to more than 50% [11, 181, 249]. To induce a less severe TAC, some groups tie the suture against a thicker 26-gauge needle, by which a lower pressure gradient is established [27, 89].

Limitations TAC-surgeries possess high procedural variability, depending on (1) the used mouse strain [13, 71], (2) the used sex [277], (3) the age at surgery and (4) the sever-

Table 2 Distinct characteristics of the murine models of heart failure induced by pressure overload and neurohumoral activation

	TAC late	e phase	TAC ea	rly phase	Ang II	high dose	Ang	II low dose	DOCA	1	DOCA	
	(>3 w)		(≤2 w)		(>1 mg	g/kg/d)	(0.2	mg/kg/d)	+ nepl + salt	hrectomy diet	alone	
Pressure overload	+++	1, 2	+++	1, 2	+	8, 9, 10	_	12	+	13, 14	+/-	16
Systemic hypertension	-		_		++	9, 11	_	12	+	13, 14, 15	_	16
Hypertrophy	+++	1, 2, 3, 4, 5	++/+	2, 3, 4, 5, 7	+++	9, 10, 11	+	12	++/+	13, 14, 15	+	16
Systolic dysfunction	+++	1, 3, 5, 6	+/-	1, 2, 3,4,5	++	8, 10, 11	-	12	_	13, 14, 15	_	16
Diastolic dysfunction	++	6	+	7	++	8, 10	+	12	+	13, 14, 15	_	16
Inflammation	+	5	++	6	++	8, 9, 10, 11	?	12	+	13, 14	_	16
Interstitial fibrosis	++/+	1,5	+++	6, 7	+++	8, 9, 10, 11	+	12	++	13, 14	-	16

Data were used from the following references: ¹[148], ²[166], ³[27], ⁴[131], ⁵[280], ⁶[216], ⁷[261], ⁸[17], ⁹[189], ¹⁰[271], ¹¹[240], ¹²[214], ¹³[162], ¹⁴[244], ¹⁵[118], ¹⁶[180]

+ increased, - not changed, ? not determined

d days, w weeks

ity of the constriction [175, 260]. Thus, the increase in left ventricular mass varies highly even within the same strain; in one study the increase in left ventricular mass ranges from + 42% to + 376% after 8–9 weeks [175]. Figure 4b shows a hypertrophic murine heart, whose ventricular mass is enlarged by approx. 200% 6 weeks after TAC. In addition, the temporal dynamics of the transition from HFpEF to HFrEF vary even within the same experimental group [90]. Further, in a recent study only a subset of mice (28%) that were subjected to TAC develop signs of HFrEF [181], thus indicating that some mice will not proceed from HFpEF to HFrEF after TAC at all. Therefore, the authors suggest to examine cardiac function by means of echocardiography prior to further investigations and select a subgroup of animals, whose cardiac function indicates that they refer to the scope of investigation [90].

Recent advancements In an attempt to reduce procedural variability, Melleby et al. recently introduced o-ring aor-



tic banding with fixed diameter constrictors [175]. In their study, utilising o-rings with an inner diameter of 0.66 mm induces progressive concentric hypertrophy and interstitial fibrosis accompanied by left ventricular stiffness but stable contractile function (HFpEF). However, when subjected to TAC utilising o-rings with an inner diameter of 0.61 mm the mice showed early progression from concentric hypertrophy to dilated congestive HF (HFrEF) [175].

Chronic subjection to angiotensin II

Murine heart failure and its clinical relevance

As depicted in Fig. 5, chronic subjection to angiotensin II (Ang II) in mice mimics chronic hypertension due to neurohumoral activation as seen in HF patients. Neurohumoral activation includes the activation of the renin–angiotensin–aldosterone-system (RAAS), which results in elevated plasma levels of its main effector hormones Ang II and



Fig.4 Transverse aortic constriction (TAC) in mice. **a** Schematic drawing depicting the constriction of the transverse aortic arch between the brachiocephal trunk (1) and the other two branches, the left common carotid artery (2) and the subclavian artery (3). After tightening the suture against a 26- or 27-gauge cannula, the latter is

removed, which leaves a defined aortic lumen behind. **b** Representative images of biventricular hearts (without atria) of mice that were either assigned to sham-surgery or to transverse aortic constriction (TAC) for 6 weeks demonstrate the enlarged ventricular mass observed after TAC in mice

Fig. 5 Reciprocal interactions between pressure overload, RAAS activation, and heart failure. Simplified schematic drawing that visualises the reciprocal interactions between pressure overload, activation of the reninangiotensin-aldosterone-system (RAAS), and heart failure (HF) in men and mice. The murine model of TAC mimics aortic stenosis in patients, in which cardiac pressure overload precedes the activation of the RAAS. The models of DOCA- or Ang II infusion, however, mimic the clinical course of patients, in which the development of HF, independent from its etiology, precedes the activation of the RAAS. However, sustained activation of the RAAS induces hypertension, thus RAAS activation also induces cardiac pressure overload. This initiates a selfperpetuating vicious circle of HF, RAAS activation and pressure overload. Further, since all here described murine models induce RAAS activation and pressure overload, the revelation of a distinct injurious stimuli in the development of HF and cardiac inflammation and fibrosis in these models is hampered



aldosterone. Thus, in spite of that it may often be seen as a hypertension model, chronic Ang II infusion resembles the myocardial injury and the subsequent cardiac remodelling owing to both, (1) cardiac pressure overload and (2) direct cardiac effects of Ang II and aldosterone. However, high Ang II levels that are observed in failing hearts do not only result from the activation of the systemic RAAS, but also from the activation of an intracardiac RAAS [205]. It has been postulated that the intracardiac cleavage of Ang I to Ang II in a local chymase-dependent manner is responsible for more than 75% of Ang II in failing hearts [133, 205]. Thus, it remains questionable, whether the infusion of supraphysiological doses of Ang II truly reflects the patients' clinical setting, in which an already failing heart is confronted with Ang II from both, local and systemic, origin [260, 272].

As shown in Table 2, mice subjected to Ang II infusion display cardiac remodelling in the presence [16, 271] as well as in the absence [214] of concomitant hypertension. This highlights that remodelling pathways exist that are independent from pressure overload. Depending on the dose and the time frame of subjection to Ang II, mice either exhibit concentric hypertrophy accompanied by diastolic dysfunction and thus resemble features of human HFpEF [17, 214], or they exhibit dilative, congestive HF accompanied by reduced ejection fraction and thus resemble features of human HFrEF [17, 271]. Likewise, the degree of hypertension depends on the dose and the time frame that mice are subjected to Ang II. Gomolak et al. analysed this dose-dependence in a study in which low Ang II-doses (approx. 0.1 mg/kg per day) do not induce hypertension, while intermediate doses (approx. 0.5 mg/kg per day) increase blood pressure over time and high doses (approx. 1.4 mg/kg per day) induce immediate and severe hypertension [78].

Methodological proceeding

Methodologically, osmotic minipumps, which subsequently release Ang II at a defined rate, are implanted subcutaneously [240]. Anaesthesia is maintained during surgery. However, no artificial ventilation is needed during the procedure, thereby making it less difficult than, e.g. the induction of MI or TAC. Typically, mice are subjected to Ang II for a time frame of 2–8 weeks [16, 17, 206, 240]. Sham controls are treated with volume-matched saline-vehicles. Though it would enhance reproducibility, blood levels of circulating Ang II are most often not measured [260].

Limitations Even at a given infusion rate (1.4 mg/kg per day) and at a given time of subjection to Ang II (8 weeks), the degree of cardiac remodelling and dysfunction strongly depends on the specific mouse strain: while C57BL/6 mice in this context display concentric hypertrophy and preserved contractile function, Balb/c mice display congestive HF [206]. Strikingly, this is independent from the degree of hypertension, as both strains exhibit comparable blood pressures upon Ang II infusion [206]. This may in part be attributable to the predominantly Th1-mediated immune responses that are observed in C57BL/6 mice, whereas Balb/c mice predominantly exhibit Th2-mediated immune responses [206].

Chronic subjection to deoxycorticosterone acetate

Murine heart failure and its clinical relevance

As shown in Fig. 5, chronic subjection to the aldosterone analogue deoxycorticosterone acetate accompanied by unilateral nephrectomy and high salt diet (DOCA) promotes hypertension and the subsequent development of HF in mice [162, 244]. In the clinical setting, the combination of hypertension and elevated aldosterone levels promotes hypertensive heart disease and its transition to HFpEF, which is present in up to 50% of HF patients [98, 180, 201]. Strikingly, mice subjected to DOCA display mild hypertension and impairment of diastolic function accompanied by exercise intolerance but do not show signs of systolic dysfunction [118, 162, 260]. Thus, they are seen as one of the few murine models that truly resemble the features of human HFpEF [118, 162, 260]. In DOCA models, hypertension is known to appear in two phases, an initial peak in blood pressure, which is followed by sustained hypertension [14].

Methodological proceedings

Animals are anaesthetised and hereafter a unilateral nephrectomy is performed [162, 244, 260]. After the procedure, drinking water that contains 1% sodium chloride is applied. Few days after nephrectomy, pellets that release the aldosterone analogue DOCA at a constant rate are implanted subcutaneously [162, 244]. The development of diastolic dysfunction depends on the onset of hypertension, as mice that are solely subjected to DOCA without unilateral nephrectomy and high salt diet do not exhibit hypertension or impaired diastolic function [244].

Cardiac inflammation and fibrosis induced by pressure overload and neurohumoral activation

The common feature of the murine models of TAC or the subjection to Ang II or DOCA (in the following summarised as models of nonischemic HF) is the induction of HF and reactive interstitial and perivascular fibrosis upon pressure overload and neurohumoral activation. Neurohumoral systems, such as the sympathetic nervous system (SNS) and the renin-angiotensin-aldosterone-system (RAAS) are activated by low cardiac output to maintain hemodynamic homeostasis. However, low cardiac output is a feature of HF, thus HF itself leads to chronic and sustained activation of the SNS and the RAAS (Fig. 5). In a vicious circle, this initially adaptive response induces self-perpetuating cardiac remodelling and subsequent HF [65]. Table 2 provides an overview of the distinct characteristics of the murine models of nonischemic HF, such as (1) TAC in the late remodelling phase, (2) TAC in the early remodelling phase, (3) Ang II administered at high dose, (4) Ang II administered at low dose, (5) DOCA accompanied by concomitant unilateral nephrectomy and high salt diet, as well as (6) DOCA alone.

The initiation of inflammation and fibrosis in nonischemic as compared to ischemic models of heart failure

The development of interstitial fibrosis in the models of nonischemic HF is in contrast to the induction of reparative fibrosis in the models of ischemic HF, such as permanent MI or I/R: here, reparative fibrosis is the mechanistically rational process of wound healing; the circumscribed area of the unrecoverable infarct wall is replaced by fibrotic scar tissue to maintain cardiac structure. Thus, there is an immense disparity in the pathogenesis of cardiac inflammation and fibrosis in models of nonischemic and ischemic HF: (1) in ischemic HF, inflammation typically occurs prior to fibrosis in a sequential manner. This is in contrast to models of nonischemic HF, in which inflammation and fibrosis typically coexist. (2) in models of ischemic HF, the initial pro-inflammatory stimulus is precisely elaborated: necrotic cell death upon ischemia triggers the subsequent release of danger-signals, which in turn immediately alert the fire brigade of resident sentinel cells. Opposed to this, the significance of distinct pro-inflammatory stimuli in models of nonischemic HF is less elaborated: (i) pressure overload increases myocardial stretch and thereby induces pro-inflammatory cascades [153], (ii) pressure overload may induce a limited amount of cell death and may therefore induce the release of danger-signals, and (iii) the concomitant activation (and/or the preceding external infusion) of neurohumoral stimuli directly promotes fibrotic reprogramming in several cardiac cells and recruits monocytes from stem cell niches in spleen and bone marrow [26]. (3) the resolution of inflammation is precisely elaborated in the murine MI model: immune cells and CFs efferocyte dead cells and phagocyte danger signals; subsequently, they undergo a reparative reprogramming and thus provide a proliferative milieu. In models of nonischemic HF, however, interstitial fibrosis is a chronic and progressive epiphenomenon of the sustained repression of non-circumscribed, self-perpetuating inflammation and the concomitant chronic activation of pro-fibrotic stimuli.

Reciprocal interactions between pressure overload, RAAS activation and heart failure

As depicted in Fig. 5, the sequential occurrence of pressure overload and neurohumoral activation differs in the models of nonischemic HF: since pressure overload markedly activates the RAAS, pressure overload precedes RAAS activation in the TAC model [23]. In contrast, in the murine models of chronic administration of either Ang II or DOCA, RAAS-activation is mimicked, which precedes the induction of hypertension and thus pressure overload. However, as illustrated in Fig. 5, reciprocal interactions between pressure overload, RAAS activation, and heart failure exist; this hampers the revelation of the specific contribution of a distinct stimulus for the induction of inflammation and fibrosis in the models of TAC, Ang II and DOCA. Therefore, the pathogenesis of inflammation and fibrosis in these models is less understood [133].

The renin-angiotensin-aldosterone-system

As depicted in Fig. 6, the renin–angiotensin–aldosteronesystem (RAAS) modulates vasotonus, renal water elimination, salt appetite, and the sympathetic activity of the central nervous system to ensure adequate organ perfusion. The RAAS consists of the interplay between distinct endocrine peptides. The liver secretes angiotensinogen, which is subsequently cleaved by the kidney-derived renin. Upon cleavage, angiotensinogen becomes the decapeptide angiotensin I (Ang I). Hereafter, Ang I is shortened to the octapeptide Ang II by the angiotensin-converting-enzyme (ACE), a membrane-bound metalloproteinase that is predominantly expressed on ECs of pulmonary vessels [205]. However, Ang I within the cardiac tissue is not only cleaved by ACE, but also by local, intracardiac chymases; a significant one is mast cell chymase [205]. In fact, more than 75% of the Ang II concentration in failing hearts may originate from a local RAAS [133].

As depicted in Fig. 6, Ang II activates the angiotensin receptor type 1 (AT1R) pathway in the kidneys, the adrenal glands, the brain, the vessels, and directly in cells of the cardiac tissue and thereby mediates its adverse cardiovascular

effects. (1) AT1R-signalling contracts vascular smooth muscle cells, which increases the peripheral resistance. (2) Ang II induces vascular remodelling, which results in endothelial dysfunction and further increased peripheral resistance. (3) Through actions in the circumventricular organs of the brain, such as the area postrema and the anteroventral third ventricle, Ang II alters the drinking behaviour, promotes sympathetic activation, and induces vasopressin release from the pituitary gland [84, 172]. Vasopressin is a potent vasoconstrictor and decreases the renal water clearance. (4) Ang II induces the release of the mineralocorticoid aldosterone from the adrenal gland. Aldosterone, and its analogue DOCA, are lipophilic molecules, which bind to their intracellular mineralocorticoid receptor (MR). In the kidneys, activation of the MR leads to increased retention of water and sodium and thereby induces hypervolemia. In the brain, activation of the MR promotes salt appetite and thus increases plasma osmolality [74, 84]. In a vicious circle, the circumventricular organs in turn detect the increased plasma osmolality and subsequently accentuate their sympathoexcitation and the release of vasopressin [84]. This further aggravates hypertension. The fact that lesions of the anteroventral third ventricle almost abolish a hypertensive phenotype in mice highlights the significance of the brain for the development of hypertension upon Ang II treatment [172]. Moreover, a recent study concludes that pro-inflammatory T-cell populations are essential for the development of hypertension upon neurohumoral activation [85]. As illustrated in Fig. 6, Ang II and aldosterone/DOCA directly mediate cardiac inflammation and fibrosis via actions on cells of the myocardium.

The direct impact of angiotensin II on cardiac cells

Pro-inflammatory Ang II-signalling Ang II is a potent proinflammatory and pro-fibrotic hormonal mediator [272]. The downstream cascade of the G protein-coupled AT1R includes the activation of janus-kinase and signal transducers and activators of transcription (JAK-STAT) pathways, which ultimately result in the activation of transcription factors from the AP1-family and NFκB [40]; NFκB activation in this context further requires the presence of ROS [248]. However, Ang II itself induces this generation of ROS [125, 133]. In line with this, mice that overexpress a mitochondria-targeted catalase are protected from cardiac fibrosis after Ang II infusion [37]. If stimulated by Ang II, CMs and CFs release TNF- α and IL-6 [143, 283]. IL-6 may be a specific player of the induction of cardiac fibrosis in the context of Ang II infusion in mice, as the disruption of IL-6 prevents cardiac fibrosis in spite of not affecting hypertension or cardiac hypertrophy in this model [79]. IL-6 further orchestrates the expression of cytokines, chemokines and CAMs and thus promotes the activation of resident inter-



Fig. 6 The initiation of cardiac inflammation and fibrosis upon activation of the RAAS. The precursor of the renin–angiotensin–aldosterone-system (RAAS), angiotensinogen, is released by the liver; subsequently it is cleaved by the kidney-derived renin to angiotensin I, which is hereafter cleaved by the angiotensin-converting enzyme (ACE) or by intracardiac chymases, such as mast cell chymase, to form angiotensin II (Ang II). The latter evokes the release of aldosterone from the adrenal glands. Ang II and aldosterone (or its analogue DOCA) promote cardiac inflammation and fibrosis through the activation of systemic and intracardiac pathways. They provoke

stitial cells and the recruitment of immune cells. One example is the CCL-2 dependent recruitment of CD34⁺ CD45⁺ monocytic fibroblast precursors [46, 87]. These precursors mature into collagen-secreting fibroblasts in a TNF-receptor (TNFR)-1-dependent manner [46, 87]. In CFs, TNFR1-signalling upregulates AT1R expression and thereby sensitises these cells to Ang II stimulation [83]. Interestingly, Ang II or TNF- α alone do not induce the maturation of fibroblast precursors [46]. This indicates that the Ang II-mediated induction of fibroblast reprogramming under nonischemic conditions depends on initial inflammatory co-stimuli. Additionally, enhancer regions of MMP and TIMP genes possess AP-1 and NF κ B binding sites [40], which promotes

(1) salt appetite, sympathoexcitation, and the release of vasopressin in the brain, (2) increased vascular resistance, and (3) hypervolemia and hypernatremia through actions in the kidney and thereby induce hypertension. Subsequently, hypertension results in increased mechanical stretch on cardiac cells, which initiates cardiac remodelling through stretch-dependent signalling, especially in cardiac fibroblasts (CFs) and cardiomyocytes (CMs). Further, Ang II and aldosterone (or DOCA) directly initiate cardiac inflammation and fibrosis through actions in CFs, CMs, immune cells (ICs) and endothelial cells (ECs)

matrix remodelling upon Ang II stimulation. Further, circulating Ang II mediates the recruitment of the later-invading monocytes from their splenic reservoir [144].

Pro-fibrotic Ang II-signalling The development of fibrosis upon Ang II infusion has two origins: (1) it is a sequela of the suppression of diffuse inflammation on the cost of activating TGF- β 1-dependent, pro-fibrotic cascades. (2) Ang II directly promotes proliferative signalling especially in ECs, CFs and CMs. In these cells, it induces the release of growth factors such as PDGF, and TGF- β 1, and the secretion of matricellular proteins, such as CCN-2 [143]. Like in other cardiac pathologies, TGF- β 1 is one of the main conductors of fibrotic remodelling in the context of Ang II infusion. It was shown, that the upregulation of TGF- β 1 through AT1R-signalling is mandatory for the development of cardiac fibrosis upon Ang II stimulation in mice [237]. Furthermore, the endothelial release of endothelin (ET)-1 upon Ang II stimulation is necessary for the development of HF in this model [2]. ET-1 enhances proliferation and attenuates apoptosis in CFs [133]. Moreover, ET-1 promotes matrix expansion as it stimulates collagen deposition and downregulates collagenase activity [133]. However, to selectively target pro-fibrotic pathways to prevent cardiac fibrosis is detrimental in the model of Ang II infusion, as it increases the susceptibility to cardiac rupture due to unrestricted inflammation [234]. However, the pro-fibrotic effects of cardiac AT1R-signalling may be fine-tuned by the anti-fibrotic effects of cardiac AT2R-signalling. In line with this, the CM-specific overexpression of AT2R decreases the degree of cardiac fibrosis but does not affect hypertension or cardiac hypertrophy in the context of chronic Ang II infusion in mice [139].

The direct impact of aldosterone/DOCA on cardiac cells

Aldosterone, and its analogue DOCA, directly mediate cardiac inflammation and fibrosis as they bind to their MRs in CMs, CFs, MPs, ECs and smooth muscle cells [133]. After nuclear translocation, MRs enhance the expression of proteins that are associated with cardiac inflammation and fibrosis, such as MMPs, TIMPs, TGF- β 1, plasminogen activator inhibitor (PAI)-1, CCN-2, fibronectin and the collagens I, III, and IV [40, 255]. Furthermore, there is a growing body of evidence that various non-canonical pathways contribute to the transdifferentiation of MFs upon MR-signalling: both, MR-dependent ROS-generation and MR-dependent transactivation of several growth factor receptors, such as epidermal growth factor receptor (EGFR) and PDGF receptor (PDGFR), were shown to directly promote MF transdifferentiation [255].

MR-signalling in macrophages On a cellular level, MPs seem to be crucially affected by MR-signalling in the murine models of nonischemic HF as MR-signalling critically polarises them towards a pro-inflammatory M1-phenotype [219, 259]. In line with this, MR-deficient MPs rather differentiate into M2-polarised MPs in a murine model of nonischemic HF [239]. Further, pro-inflammatory c-Jun NH₂-terminal kinase/activating protein-1 (JNK/AP-1) and NF κ B transactivation pathways depend on MR-signalling [239, 255]. However, it remains unclear, whether MRs in MPs are also mandatory for their recruitment: in a DOCA model, MP-specific MR-deficient mice are not protected from cardiac

MP recruitment, but from hypertension [219]. In contrast, in a murine model that combines the infusion of the nitric oxide synthase inhibitor L-NAME with Ang II infusion, MP-specific MR-deficient mice are protected from cardiac MP recruitment, but not from hypertension [259]. However, MP-specific MR-deficient mice show less inflammation and less fibrosis in both models [219, 259].

MR-signalling in cardiomyocytes CM-dependent MR-signalling additionally contributes to cardiac inflammation and fibrosis: depletion of CM-specific MR-signalling does not prevent immune cell infiltration but results in lesser collagen deposition in a murine DOCA model [217]. Interestingly, CM-specific MR-deficient mice show increased expression of decorin, an inhibitor of the pro-fibrotic mediators TGF- β 1 and CCN-2; they further show increased MMP-2 and MMP-9 activity [217]. Taken together, the loss of CM-specific MR-signalling may not prevent immune cell recruitment, but may disrupt a pro-fibrotic vicious circle as it modulates TGF- β /CCN-2- and MMP-2/MMP-9-dependent pathways. In contrast, in models of TAC, CM-specific MR-depletion does not prevent cardiac inflammation and fibrosis [160].

MR-signalling in cardiac fibroblasts CFs may also contribute to cardiac remodelling in a MR-dependent manner, as aldosterone in vitro stimulates the proliferation of CFs and increases their collagen synthesis [133]. Moreover, ECs express cytokines and CAMs, such as VCAM-1, in a MRdependent manner [161]. In line with this, cardiac inflammation and fibrosis are less pronounced in EC-specific MRdeficient mice that are challenged with DOCA [161, 218]. However, as with the CM-specific MR-depletion, the ECspecific MR-depletion does not affect cardiac remodelling in murine TAC models [228].

Adrenergic signalling

Increased contractile demands of the myocardium and the concomitant upregulation of RAAS hormones in the models of TAC, Ang II, and DOCA activate the sympathetic nervous system (SNS) and thus promote the release of catecholamines. Moreover, sustained activity of the SNS is a key feature of human HF, hence β -blockade is a pivotal basis of its therapy [72]. The two major sources of catecholamines are (1) the adrenal glands, which predominantly release epinephrin upon sympathetic stimulation and (2) cardiac adrenergic nerve endings, which release norepinephrin [164]. On the one hand, activation of the SNS promotes fibrotic cardiac remodelling as it evokes the release of renin from the kidneys [24] and thus contributes to further RAAS-activation and hypertension. On the other hand, catecholamines directly promote adverse cardiac remodelling, as they bind to the G protein-coupled

 β -adrenergic receptors (β -AR)-1 and -2 on cardiac cells [164]. Recently, the complex downstream signalling of these receptors in the context of HF has been comprehensively reviewed [38].

There are three types of β -ARs, of which the β_1 -AR is predominantly expressed in healthy cardiac tissue (75-80%), followed by the β_2 - (15–20%) and the β_3 -AR (2-5%) [164]. However, chronic activation of the SNS upon HF in mice and men downregulates the β_1 - and desensitises the β_1 - and the β_2 -AR [164]. In turn, adrenergic signalling upon HF upregulates the G protein-coupled receptor kinase (GRK)-2, an enzyme that is known to promote fibrotic remodelling in experimentally induced HF [56]. Further, chronic activation of the β -ARpathway in CMs directly induces hypertrophy and apoptosis and activates the transcription of a fetal gene programme [169]; further, it upregulates the myocardial expression of the pro-inflammatory cytokines IL-1 β , -6, -18 and TNF- α in a NFkB-dependent manner [25, 191]. Upon chronic adrenergic stimulation, TNFR-1-signalling worsens cardiac remodelling, whereas TNFR-2-signalling may be cardioprotective [72]. Moreover, a recent study concludes that excessive CMspecific β_1 -AR-signalling generates ROS and thereby activates the inflammosome, which is required for the release of IL-18 [281]. However, incubating CFs with the supernatant of hypertrophic CMs, previously stressed by adrenergic stimulation, induces MF transdifferentiation and the subsequent secretion of collagens [91]. This paracrine effect is attributed to the ROS-dependent secretion of growth factors by CMs, such as CCN-2 [91]. Thus, a cross talk between CMs and CFs may be of overwhelming importance in the context of β -AR-mediated MF transdifferentiation. In contrast to the adverse effects of β_1 - and β_2 -AR-signalling, $\alpha 1$ - (subtype A) [104, 198] and β_3 -AR-signalling [91] may have antipathetic effects as they ameliorate adverse remodelling and the transition to HFrEF upon pressure overload.

Mechanical stretch upon pressure overload and hypertension

Increased hemodynamic load after TAC, or to a minor degree upon Ang II- or DOCA-induced hypertension, increases the mechanical stretch on myocardial cells.

Cardiac fibroblasts—the sentinel cells that initiate inflammation upon mechanical stretch Resident CFs are connected to the ECM through their transmembrane-spanning integrins; thus, they may be the sentinel cells that detect increased mechanical stretch [40, 153, 192, 280]. Mechanical stretch activates the downstream extracellular signalregulated kinase (ERK), p38 and NF κ B; subsequently, CFs secrete pro-inflammatory cytokines and chemokines of the CC-family, such as CCL-2, CCL-7, and CCL-8 [153]. Strikingly, chemokine secretion is not observed in CMs or ECs upon mechanical stretch [153]. Beyond that, mechanical stretch leads to sustained induction of the pro-inflammatory proteases MMP-2, MMP-9 and the CAM Cx43 [254]. In reaction to pro-inflammatory signals, CFs and immune cells further increase their MMP activity, which enables leukocytes to dig through the basal membrane and infiltrate the ECM [275]. This aggravates cardiac inflammation, as ECM fragments (matrikines) amplify the vicious circle of inflammation and leukocyte invasion [153, 221, 247, 280]. Interruption of this vicious circle of inflammation requires the activation of pro-fibrotic stimuli. Thus, inflammatory proteases, such as MMPs, activate TGF-B1 from latent stores; TGF-β1 is the key stimulus by which inflammatory CFs reprogramme to a matrix synthesising phenotype [143,153]. In contrast to models of ischemic HF, in models of pressure overload only few of the activated, collagen-secreting CFs are α -SMA-positive MFs [185]. This highlights that CFs may not only follow one line of action in the murine models of HF, but possess a variety of reprogramming phenotypes that depend on the underlying pathophysiology. Though bone marrow-derived fibrocytes and endothelialto-mesenchymal transition have long been under debate to significantly contribute to the population of activated CFs in pressure overloaded hearts, recent evidence suggests that activated CFs in this setting arise from a lineage of resident endothelial and epicardial fibroblasts [185]. Additionally, mechanical stretch directly upregulates TGF-\u00b31 [153] and the matricellular protein CCN-2 [254]. Strikingly, CCN-2 inhibition does not affect cardiac remodelling in Ang IIinfused mice, but prevents adverse left ventricular remodelling and cardiac dysfunction in mice subjected to TAC [143, 2501.

Cardiomyocytes Mechanical stretch induces CM hypertrophy and activates an intracardiac RAAS in CMs, which results in the myocardial accumulation of Ang II [56, 205]. Moreover, mechanical stretch in CMs promotes AT1Rsignalling in a ligand-independent manner [293]. Beyond that, generation of oxidative stress in CMs upon pressure overload induces MR translocation in a ligand-independent manner [255]. Further, a limited amount of necrotic cell death may occur upon TAC [26]; the subsequent release of DAMPs induces an immediate inflammatory response through TLR-dependent pathways [133].

The transition from HFpEF to HFrEF in models of pressure overload The development of compensatory concentric hypertrophy upon pressure overload initially is a response to the increased contractile demands of the myocardium. As a consequence, the increased stiffness of the hypertrophied myocardium leads to diastolic dysfunction and exercise intolerance (HFpEF). However, the subsequent transition to eccentric hypertrophy accompanied by severe systolic dysfunction (HFrEF) upon pressure overload is a pure pathologic sequela of self-perpetuating inflammation and fibrosis. Thus, the players that promote this transition are of great interest as they may depict key targets for clinical translation. Recent findings in a murine TAC model reveal that the early recruitment of monocytes is a key event in this context: pressure overload increases the number of CCR-2⁺ monocytes and their subsequent cardiac recruitment in the stage of HFpEF [204]. Hereafter, these cells differentiate into MPs and mediate the cardiac invasion of T-cells by presenting cardiac antigens on their surfaces [204]. This is of eminent importance, as the activation of CD4⁺ T-cells was shown to (1) depend on antigen presentation [122] and (2) to be pivotal for the transition from HFpEF to HFrEF in the context of pressure overload [142]; thus, the CCR-2-dependent recruitment of monocytes may be seen as an upstream mechanism of chronic inflammation that crosslinks innate and adaptive immunity, which ultimately results in HFrEF. On a molecular level, a tightly regulated, reciprocal interplay between matrix-degrading MMPs and matrix-preserving TIMPs may be responsible for the transition from HFpEF to HFrEF [93, 133]. In HFpEF, MMPs remain rather quiescent, whilst they are active in HFrEF [113]. In contrast, pro-fibrotic stimuli such as TGF-B1 remain rather quiescent in HFrEF [133].

Vascular remodelling upon pressure overload Pressure overload in mice induces cardiac angiogenesis and extensive vascular remodelling [174, 280]. As pressure overload increases the blood pressure in coronary arteries, perivascular fibrosis may depict a beneficial response to resist these higher perfusion pressures [132]. Furthermore, in pressure overload models coronary arterioles promote myocardial fibrosis as they develop adventitial inflammation in an Ang II-dependent manner; inflammation subsequently extends and promotes perivascular fibrosis [173, 280]. However, vascular cells additionally possess anti-fibrotic features: ECs secrete interferon- γ -inducible protein (IP)-10/CXCL10 and hypoxia-inducible factor (HIF)-1; both of them suppress TGF- β -signalling [133].

Model of heart failure induced by viral myocarditis

Murine heart failure model and its clinical relevance

Myocarditis is an inflammatory disease of the myocardium that is predominantly caused by infectious agents [275]. The clinical course of myocarditis has a broad spectrum of outcomes, ranging from complete recovery to dilated cardiomyopathy (DCM) in the context of viral elimination or chronic inflammatory cardiomyopathy in the context of viral persistence. About 10% of patients with biopsy-proven viral myocarditis develop inflammatory cardiomyopathy [7]. However, myocarditis may remain asymptomatic and thus displays a significant cause of sudden death that occurs especially in young adults [34, 50, 236].

Depending on their susceptibility to the virus, mice, likewise humans, exhibit a variety of outcomes after infection with Coxsackievirus B3 (CVB3) [129, 151]. All strains, susceptible (e.g. BALB/c) or not (e.g. C57BL/6), initially suffer from acute myocarditis as characterised by systolic and diastolic dysfunction alongside interstitial remodelling [115, 129, 267]. Opposed to this, solely susceptible strains develop chronic myocarditis, also described as inflammatory cardiomyopathy, as a result of virus persistence or nonresolved cardiac inflammation. This is characterised by cardiac inflammation, reactive fibrosis alongside deteriorated cardiac function. Inoculating mice from susceptible strains with CVB3 therefore resembles features of inflammatory cardiomyopathy in humans.

Methodological proceedings

Coxsackievirus B3 is a non-enveloped, single-stranded, positive-sense RNA virus within the family of the picornaviredae. Commonly, $10^4 - 10^7$ plaque-forming units of the heart passaged enterovirus CVB3 are injected intraperitoneally [15, 16, 129, 203]. Several CVB3 viral strains exist; a frequently used one is the Nancy-strain [123, 176]. The potency and the efficacy of a certain viral strain may vary, which should be considered in the experimental setup [8,107]. As depicted in Fig. 7, the genome of CVB3 encodes 11 proteins, 4 structural proteins and 7 non-structural proteins. Since CVB3 is a positive-sense RNA virus, its RNA genome is directly translated into proteins upon host cell infection and hereafter functions as the template for viral replication. Appropriate viral infection can be proven by a loss of bodyweight and the detection of viremia in small serum samples, which typically occurs around day 2-4 post injection (p.i.). Tissue sampling and hemodynamic assessments should be performed either from day 6 to 8 during the acute phase of myocarditis or around day 28 during the chronic phase, depending on the curiosity for acute or chronic cardiac remodelling, respectively [15, 16, 150, 151, 274].

Cardiac inflammation and fibrosis induced by viral myocarditis

Upon inoculation with CVB3 in mice, myocarditis occurs in three sequential stages [66, 151]: (1) virus-mediated myocarditis from day 2 to 4, (2) immune-mediated myocarditis from day 4 to 14, and (3) the subsequent chronic stage of cardiac remodelling or chronic inflammation, which typically occurs around day 28. Virus-mediated myocarditis is characterised



Fig. 7 The course of viral myocarditis induced by CVB3 in mice. Inoculating mice with CVB3 results in viral myocarditis, which, likewise in humans, occurs in three sequential stages. The first stage, virus-mediated myocarditis, is observed from day 2 to 3 p.i. in mice. During this stage, the non-structural viral proteases 2A and 3C directly damage cells of the cardiac tissue and thereby initiate inflammation. Hereafter, sentinel cells are alarmed and recruit a first wave of immune cells into cardiac tissue, as shown in the representative hematoxylin and eosin (H&E) staining [15]. This results in the second stage, immune-mediated myocarditis, from day 4 to 14. During this phase, pro-inflammatory leukocytes either eliminate the virus

by viral entry and the translation of viral proteins in cardiac host cells. During this phase, the translated viral proteins themselves exert direct cytotoxic damage; however, infiltrating immune cells are absent [151]. The subsequent stage

(non-susceptible strains) or solely cease viral replication (susceptible strains); however, the leukocytes severely harm the cardiac tissue in both cases. In the case of viral elimination, cardiac remodelling is reversible and may either result in complete recovery or in dilated cardiomyopathy, depending on the degree of cardiac damage during acute myocarditis. In the case of viral persistence, a second wave of immune cells is recruited and mediates a chronic cardiac inflammation that is accompanied by dilated cardiomyopathy, known as inflammatory cardiomyopathy. Additionally, molecular mimicry may induce the development of a self-reactive autoimmunity in susceptible strains, which equally results in inflammatory cardiomyopathy

of immune-mediated myocarditis is characterised by a first wave of immune cell infiltration and the active replication of the virus in its host cells [151]. Activation of the host immunity eliminates the virus in non-susceptible strains or ceases viral replication in susceptible strains; however, this happens on the cost of initiating a thunderous storm of cardiac inflammation. The stages (1) and (2) are frequently summarised as acute viral myocarditis. The chronic stage in susceptible strains is characterised by a second wave of immune cell infiltration, which ultimately results in inflammatory cardiomyopathy. At this stage, viral replication has ceased, but the viral genome persists in cardiac tissue [151]. Beyond the genetic determination of the mouse strain, several other preconditions, such as age, nutrition status, gestation, and sex hormones were found to further determine the susceptibility to CVB3 infections in mice [284].

Virus-mediated acute myocarditis

CVB3 typically enters its host via the gastrointestinal or the respiratory tract [235], whereas it is inoculated intraperitoneally in the experimental setup. CVB3 enters CMs upon viremia 2-3 days after inoculation and subsequently exerts cell-to-cell spread [129, 235, 284]. Beyond that, in vitro evidence suggests a role for infected CFs [141, 151]. CVB3 enters CMs through interactions with the coxsackie-adenovirus receptor (CAR) [66, 129] and uses the decay acceleration factor (DAF) as a coreceptor [284]. In line with this, depleting the CAR in CMs [241] or treating CVB3-inoculated mice with soluble CARs [208] blunts acute myocarditis in this model. Higher CAR-expression in younger individuals may further determine their higher susceptibility to viral myocarditis [284]. After entering its host cell, the progeny viral RNA is translated and replicated and subsequently harms its host cells predominantly via two viral proteins, the proteases 2A and 3C. Both proteases are encoded within the viral genome and are immediately translated upon host cell infection. They cleave a variety of intracellular target proteins [66, 73]: (1) 2A and 3C break down protein biosynthesis by cleaving an initiation factor of translation (eIF-4G); (2) 2A and 3C activate caspase-8, by which apoptosis is induced; (3) 2A cleaves the transcription factor serum response factor (SRF) and thereby decreases the expression of contractile proteins; (4) 2A cleaves structural proteins such as dystrophin and dysferlin and thereby destabilises cellular and organellar membranes. This implies severe consequences for cardiac contraction as the dystrophin-cleavage affects the integrity of the sarcolemma membrane [10]. (5) 2A and 3C cleave autophagy-associated molecules. Finally, all these damages result in the lysis of host cells [73]. Strikingly, mice develop dilated cardiomyopathy after induced CM-restricted expression of the viral protease 2A in the absence of concomitant CVB3 infection [282]; further, they develop CVB3-induced myocarditis after the depletion of T- and B-cells [29]. Thus, host immunity is not required for the development of an initial virus-mediated cardiac injury. However, several resident cardiac cells, such as ECs, CMs, CFs and MPs are activated upon (1) viral entry into host cells and (2) virus-mediated cardiac injury and subsequently release pro-inflammatory mediators [275]. Thereby, they promote leukocyte invasion, fuel inflammation, and thus initiate the immune-mediated stage of viral myocarditis.

Immune-mediated acute myocarditis (antiviral immunity)

The fire brigade of innate immune cells invades the cardiac tissue 4–5 days p.i. and draws the virus into a fight for survival [103]. At this point, the initial virus-mediated myocardial inflammation proceeds to an immune-mediated one. Subsequently, cells of the adaptive immunity arrive 6–7 days p.i. to support the innate immune cells [284]. However, their invasion is associated with overshooting inflammation and subsequent adverse remodelling [284]. CM death and the infiltration of immune cells peak from day 9 to day 12 p.i., respectively [129]. The stage of immune-mediated acute myocarditis is critical for the long-term outcome, as viral elimination may permit reverse cardiac remodelling, whereas viral persistence results in sustained inflammation and non-reversible cardiac remodelling [284].

Activation of the innate immune system Pattern recognition receptors (PRRs) of the innate immune system, such as TLR-3 [67] and TLR-7/8 [284] detect viral single-stranded RNA and intermediate viral double-stranded RNA in infected cells [73]. Sex differences in TLR-expression may explain the high susceptibility of male mice as compared to females [223, 224]. After CVB3 infection, TLR-3, -7, -8, and -9, all predominantly localised in endosomes, are upregulated in mice [16, 251]. Hence, they are of predominant importance after viral internalisation in phagocytes. However, upon viral replication in host cells, the doublestranded intermediate RNA predominantly accumulates in the cytosol [284]. Therefore, RNA helicases, such as retinoic acid-induced protein I (RIG-I) and melanoma differentiation-associated gene 5 (MDA-5), detect cytosolic dsRNA in host cells [124]. Once alarmed by pro-inflammatory cascades downstream of their TLRs, innate immune cells secrete antiviral type I interferones (IFNs), such as IFN- α and IFN- β [66]. Type I IFNs possess antiviral features and inhibit viral replication. Murine knock out studies revealed that in the context of CVB3-induced myocarditis the depletion of the type I IFN-receptor worsens mortality [269]; however, as viral replication in cardiac tissue remains unaffected, this may rather be an epiphenomenon of higher hepatic viral replication. Pivotal regulators of IFN- β are protease-activated receptor (PAR)-1 and PAR-2: while the activation of PAR-1 enhances IFN- β secretion, the activation of PAR-2 downregulates IFN-β expression [268]. Moreover, innate immune cells, but also CFs [151], secrete pro-inflammatory cytokines such as IL-1 and TNF-

 α ; this amplifies cardiac inflammation and the recruitment of further immune cells [73]. IL-1 and TNF- α may display pro-inflammatory key stimuli, as prolonged and sustained secretion of IL-1 [149] and TNF- α [105] has both been shown to exaggerate viral myocarditis in mice. Recent evidence suggests that specifically neutralising IL-1 β reduces cardiac inflammation and fibrosis and prevents chronic viral myocarditis in mice [135]. Further, the treatment with a soluble TNF-receptor [137] or a neutralising chemokineantibody [274] lowers cardiac inflammation and protects the heart from myocardial injury. Moreover, the expression of a variety of MMPs is upregulated upon stimulation with pro-inflammatory cytokines [275]. The secretion of MMP-9 by invading immune cells is one example of how they break through the basal membrane [275]. Moreover, the collagenturnover shows both qualitative and quantitative changes during acute myocarditis: beyond a tiny reduction in absolute cardiac collagen content, the fraction of soluble collagen is increased during acute myocarditis in mice [275]. A high fraction of soluble collagen is associated with impaired collagen cross-linking, which is known to promote adverse remodelling [279]. Further, soluble collagen itself may promote vascularisation of the injured tissue and may permit myocyte-slippage, a process thought to play a crucial role in the development of DCM even after complete viral elimination [275]. In line with this, pro-inflammatory chemokines (e.g. CCL-2, -5, and -7) and cytokines (e.g. IL-1, IL-6, and TNF- α) are upregulated alongside pro-fibrotic players (e.g. TGF-β1, TIMP-1, collagen-1 and -3, as well as CCN-2) during acute myocarditis [15]. Strikingly, the expression levels of the aforementioned proteins return to baseline after acute myocarditis in non-susceptible strains, whereas they remain elevated in susceptible strains.

Activation of the adaptive immune system Once they engulfed viral particles and dead cells, MPs and dendritic cells present viral fragments on their surface and thereby recruit players of the adaptive immune system. However, T-cell recruitment is an ambiguous event in the context of viral myocarditis: on the one hand, T-cells are essential for the sufficient elimination of the virus, on the other hand, viral elimination includes the clearance of infected CMs. Since CMs possess negligible regenerative capacity in mammals, the T-cell mediated clearance of infected CMs (1) reduces the contractility of the myocardium and (2) induces cardiac remodelling to replace CMs with ECM. In line with this, T-cell infiltration is temporarily accompanied by pathological damage in CVB3-infected cardiac tissue [128]. Accordingly, the depletion of $CD4^+$ and $CD8^+$ T-cells has been shown to dampen mortality and inflammation in CVB3-induced myocarditis [200]. Strikingly, complete T-cell depletion does not affect cardiac viral titres [284]. This highlights the role of the innate immunity, which precedes the time-consuming activation of antigen-specific immunity, in limiting early replication and propagation of CVB3 [284]. However, excessive innate-mediated inflammation may favour the early recruitment of circulating fibroblast precursors; in the inflamed myocardium, these precursors transdifferentiate and, as contrarily described in ischemic HF, may exhibit a major source of MFs in the context of myocarditis-related cardiac remodelling [56]. This is in line with a recent study which shows that the depletion of MPs is associated with higher viral titres during acute myocarditis, but with fewer MFs and thus less fibrotic long-term remodelling in the chronic phase [117, 152].

The chronic stage of myocarditis (viral elimination or viral persistence and anti-self immunity)

Viral elimination As depicted in Fig. 7, the chronic stage after CVB3-induced myocarditis is highly variable in mice. In non-susceptible strains, sufficient viral elimination may either result in complete recovery or lead to sterile DCM, depending on the severity of cardiac injury during acute myocarditis. Thus, after the first wave of immune cells has vanished, reversible cardiac remodelling is frequently observed in non-susceptible strains [15]. However, cardiac contractility may remain impaired despite reversed morphologic remodelling in non-susceptible strains [15]. One may speculate that this is observed, at least in part, due to the loss of CMs during acute myocarditis owing to (1) myocyte-slippage and (2) to the lytic cycle of the virus. Since myocardial fibrosis is reversible in this context, it is likely that collagen cross-linking does not appear during the acute stages of myocarditis in non-susceptible strains, as it would impair the collagenase-mediated collagen cleavage that is required for reverse cardiac remodelling [15, 227]. In line with this, an increased fraction of soluble collagen is observed during acute myocarditis [275].

Viral persistence In stark contrast, in susceptible strains, a second wave of immune cells enters the cardiac tissue after the first wave has vanished [51] due to at least two different reasons, virus persistence or autoimmunity. Susceptible mouse strains can only hinder viral replication but do not completely eliminate the viral genome from their cardiac tissue. As these strains develop inflammatory dilated cardiomyopathy following acute viral myocarditis, it is likely that the persistence of the virus is, at least in part, responsible for such sequelae. In line with this, the CM-specific expression of a replication-incompetent CVB3 cDNA in mice leads to (non-replicating) viral persistence in CMs, which in turn results in inflammatory dilated cardiomyopathy [270, 284].

Due to the lysis of cells as the final step in the lytic cycle of viral replication, intracellular host proteins may induce the development of a self-reactive autoimmunity, which additionally contributes to the development of inflammatory dilated cardiomyopathy [66]. In this context, molecular mimicry initiates the autoimmune responses [226]; the exposure of sarcomeric fragments (e.g. myosin) that cross-react with CVB3-antigens on MHC molecules recruits self-reactive immune cells, which attack CMs through direct cytotoxicity or through the secretion of auto-antibodies [66]. As with viral persistence, the initiation of autoimmunity upon CVB3-infection is exclusively seen in susceptible mouse strains [51, 226].

Conclusion

Several murine models that develop HF accompanied by cardiac inflammation and fibrosis are frequently used in experimental setups. As summarised in Table 3, each model has features that makes it unique when compared to the other models. Thus, it may remain challenging to select an appropriate model that fits both, (1) the exploration of the scientific hypothesis in a setting with clinical relevance and (2) the technical capacity of the laboratory or its cooperatives.

While the ischemic models of HF and TAC require elaborate surgical skills owing to the need for artificial ventilation and access to the thorax, the subjection of mice to Ang II or

Table 3 Distinct characteristics of the murine models of heart failure

CVB3 does not require this and is therefore less complex. As the DOCA regimen includes a unilateral nephrectomy but no access to the thorax, it may stand in between the aforementioned models regarding its feasibility. Further, the initial injurious stimuli and thus the subsequent development of HF, inflammation and fibrosis may be divided into two paramount categories in the here-described murine models of HF: On the one hand, ischemic models exhibit a local injury which predominantly initiates reparative fibrosis and results in HFrEF. This injury is transmural in the model of permanent ischemia but rather midmyocardial in the model of ischemia-reperfusion. On the other hand, nonischemic models exhibit a diffuse injury of the myocardium that subsequently initiates interstitial fibrosis and either results in HFpEF or HFrEF, depending on the severity and the duration of the injurious stimulus. Hypertrophy, however, is most pronounced in the models of TAC and Ang II, but may also be seen in the remote zone after MI. However, TAC is the model, which is preferentially used for the analysis of hypertrophy in mice. Concerning the initiation of inflammation, pattern recognition receptors (PRRs) play a pivotal role in the models of ischemic HF due to the release of danger-associated molecular patterns (DAMPs) upon CM-necrosis. However, PRRs additionally play a crucial role in the nonischemic model of viral myocarditis due to

	Ischemic HF		Nonischemic HF						
	Permanent LAD-ligation	Transient LAD-ligation	TAC	Ang II	DOCA	CVB3			
General considerations									
Clinical relevance	MI, no reperfusion	MI, reperfusion	Aortic stenosis/ NH activation	Hypertension/ NH activation	Hypertension/ NH activation	Viral myocarditis			
Heart failure	HFrEF	HFrEF	HFpEF/HFrEF	HFpEF/HFrEF	HFpEF	HFrEF			
Complexity of the procedure	+++	+++	+++	+	++	+			
Localisation of the injury	Local, transmural	Local, midmyocardial	Diffuse, LV > RV	Diffuse	Diffuse	Diffuse			
Cardiac remodelling									
Reparative fibrosis	+++	++	+/-	_	-	+/			
Interstitial fibrosis	++/+	+/	+++/++	+++/++	++/+	+++			
Hypertrophy	++/+	+/	+++	++	+	-			
Initiation of inflam- mation									
DAMPs	+++	++	+	+/-	+/-	+			
PAMPs	-	-	-	-	-	+++			
Neurohumoral activa- tion	++/+	+/-	++	+++	++	+/-			
Pressure overload	++/+	+/	+++	++	+	+/			

NH neurohumoral, *LV* left ventricle, *RV* right ventricle, *DAMP* danger-associated molecular pattern, *PAMP* pathogen-associated molecular pattern, *HFrEF* heart failure with reduced ejection fraction, *HFpEF* heart failure with preserved ejection fraction

the pathogen-associated molecular patterns (PAMPs) that alarm sentinel cells and invading leukocytes. However, a limited amount of CM-necrosis and thus PRR-signalling may also occur in the murine models of TAC, Ang II and DOCA owing to cardiac pressure overload, which induces the release of DAMPs upon CM-necrosis or apoptosis. The prototypical model that mimics neurohumoral activation is Ang II; however, concomitant neurohumoral activation may additionally occur in all here-described models due to cardiac pressure overload and the development of HF (as shown in Fig. 5). Likewise, cardiac pressure overload is a prototypical characteristic of the TAC model, but may concomitantly occur in all other here-described models, either due to neurohumoral activation and subsequent hypertension, or due to impaired cardiac function.

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Compliance with ethical standards

Conflict of interest The authors declare no conflict of interest.

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