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**Platelet survival is associated with platelet surface
expression of CXCR4 and CXCR7 and left ventricular
functional recovery after ST-elevation myocardial
infarction**

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Meiner Familie

und

in Erinnerung an meinen Großvater

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Abbreviations

ACS	Acute coronary syndrome
ADP	Adenosine diphosphate
AMP	Adenosine monophosphate
AP	Angina pectoris
Apaf-1	Apoptotic protease activation factor-1
ATP	Adenosine triphosphate
Bcl-2	B-cell lymphoma 2
BH	Bcl-2 homology domain
BMS	Bare-metal stent
CAD	Coronary artery disease
CCS	Canadian Cardiovascular Society
DD	Death domain
DES	Drug-eluting stent
DTS	Dense tubular system
ECG	Electrocardiography
FACS	Fluorescence-activated cell sorter
FADD	Fas-associated protein with death domain
FITC	Fluorescein isothiocyanate
FSC	Forward scatter
GMP	Guanosine monophosphate
GP	Glycoprotein
IP ₃	Inositol triphosphate
LBBB	Left bundle branch block
LVEF	Left ventricular ejection fraction
MFI	Mean fluorescence intensity
MOMP	Mitochondrial outer membrane permeabilization
MRI	Magnetic resonance imaging
NO	Nitric oxide

Abbreviations

NSTEMI	Non-ST-segment elevation myocardial infarction
OCS	Open canalicular system
PAR	Protease activation factor
PBS	Phosphate-buffered saline
PBS-Ca ²⁺	Phosphate-buffered saline with calcium
PCI	Percutaneous coronary intervention
PE	Phycoerythrin
PFA	Paraformaldehyde
PGI ₂	Prostaglandin I ₂ / Prostacyclin
PLC β	Phospholipase C- β
PMT	Photomultiplier tube
PRP	Platelet-rich plasma
PS	Phosphatidylserine
PTCA	Percutaneous transluminal coronary angioplasty
SD	Standard deviation
SDF-1	Stromal-derived factor 1
SSC	Sideward scatter
STEMI	ST-segment elevation myocardial infarction
tBID	Truncated BID
TMRE	Tetramethylrhodamine ethyl ester
TNF	Tumor necrosis factor
Tpo	Thrombopoietin
TRAIL	TNF-related apoptosis-inducing ligand
vWF	Von Willebrand factor

1 Introduction

Responsible for 5.3% of all causes of death in Germany, acute myocardial infarction ranks in the second place of the ten most common causes of death, behind coronary heart disease [2]. Both of these diseases can be the result of coronary atherosclerosis and are only two different ways in which it manifests [3, 4]. Despite a long history of research and much effort to achieve knowledge, prevention, and therapy, further scientific efforts are still necessary. A closer look at the role of platelets in myocardial infarction has revealed that it can make a great contribution to cardiac repair mechanisms [5, 6].

1.1 Platelets

In recent years, knowledge about platelet function and fundamental mechanisms has increased rapidly [6]. Platelets are involved in many processes of tissue injury such as detection, inflammatory response, chemo attraction, angiogenesis, proliferation, remodelling, apoptosis and many more [6-9].

1.1.1 Morphology of platelets

Platelets play a crucial role in haemostasis [10]. They can be activated by various circumstances such as vessel injury, which leads to a change of their outer appearance and release of their content [10-12]. Having a diameter of 2.0 μm to 5.0 μm , the anuclear and discoid platelets are the smallest cellular content of circulating blood [13-15]. Under conditions of not being involved in endothelial lesion, thrombosis, haemostasis or another activating effect, their estimated lifetime is between 7 and 10 days [7, 15]. Current studies postulate that their lifespan might be mainly limited by the intrinsic pathway of apoptosis [7-9].

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Platelets are fragments of their progenitors, the megakaryocytes, which primarily reside in the bone marrow [16-18]. They are subdivided into four zones: the peripheral zone, the sol-gel zone, the organelle zone, and the platelet membrane system [15, 19].

The peripheral zone consists of the plasma membrane with a lipid bilayer and a glycocalyx with glycoprotein (GP) receptors, similar to other plasma membranes in the human body [15, 20]. However, unique for these blood cells is their rugose surface, similar to gyri and sulci in the brain, which is assumed to be an additional membrane, needed for platelet spreading after activation [15, 21]. The surface of these blood cells expresses a huge number of receptors, which are mostly involved in haemostasis and thrombosis, but also in other functions such as inflammation, tumor growth and angiogenesis [11, 22-27]. Important receptors for haemostasis are the GPIB-IX-V complex, the integrin $\alpha_{IIb}\beta_3$, the protease activation receptor (PAR) and the prostaglandin receptor [11, 15, 28-31]. One further receptor family of importance is the family of chemokine receptors, in particular its receptors CXCR4 and CXCR7 [11].

The sol-gel zone is responsible for platelets' shape, and contains the greatest amount of platelet volume, except for the organelles [15]. One component of the sol-gel zone is a circumferential band of microtubules [15, 32-37]. These microtubules stabilise the flat discoid shape of the platelets and act as a suspension for cell organelles [15, 32-37].

The organelle zone contains mitochondria and three types of secretory granules: α -granules, dense granules and lysosomes [12, 15]. The most numerous organelles in platelets are α -granules, which can release more than 300 types of proteins [12, 15, 38-41]. A selection of these proteins includes P-selectin, GPIb-IX-V, GPIIb/IIIa (also known as $\alpha_{IIb}\beta_3$) and GPVI [12, 40-44]. In addition, there are factors V, VII, IX, XIII, fibrinogen, von Willebrand factor (vWF), SDF-1, epidermal growth factor and many more [12, 40-44]. Dense bodies contain low molecular

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weight ingredients such as calcium, adenosine diphosphate (ADP), adenosine triphosphate (ATP), serotonin, histamine and pyrophosphate [12, 15, 44].

The platelet membrane system consists of an open canalicular system (OCS) and a dense tubular system (DTS) [12, 15, 45-47]. The OCS is responsible for cell spreading upon platelet activation, and the DTS stores calcium, thus playing an important role in platelet activation [12, 15, 45-47].

1.1.2 Platelet activation

The physiological function of platelets is to detect a disrupted endothelium and exposed endothelial matrix after vessel injuries [48]. To prevent platelets that are not activated from adhesion to the vessel's endothelium, the endothelium releases nitric oxide (NO), prostacyclin (PGI₂) and endothelial ecto-ADPase [48, 49]. Nitric oxide is released upon shear stress and is well documented as an inhibitor of platelet aggregation [48, 50]. Prostacyclin is also released upon shear stress after cyclooxygenase-1-dependent production [48, 51]. Both of these increase levels of cyclic guanosine monophosphate (GMP) and cyclic adenosine monophosphate (AMP), respectively, which leads to the blockage of phospholipase C [52, 53]. Blocked phospholipase C decreases the intracellular level of Ca²⁺ and subsequently inhibits platelet activation [52, 53]. Endothelial ecto-ADPase mediates the metabolism of ADP, which is the most potent stimulator of platelet aggregation [48, 49, 54]. Since ADP is the most potent stimulator of platelet activation, decreased levels of ADP preserve platelets from being activated in healthy vessels [48, 49, 54].

Clot formation is a sequence of overlapping processes with no clearly defined stages [55]. Nevertheless, a three-stage model comprising initiation, extension and stabilization is described with the purpose to form a stable platelet clot after vessel injury [55].

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The initiation step begins with a vessel injury. During this step, platelets form a plug to cover the vessel injury [55]. Detection of these injuries by circulating platelets leads to their adhesion. Participating agents are the GPIb-IX-V receptor complex and vWF, which can bind to collagen in the bare subendothelium even under high shear stress with a subsequent activation of GPVI and $\alpha 2\beta 1$ [29, 48, 56]. With the activation of GPVI and $\alpha 2\beta 1$, an intracellular cascade, including phospholipase C γ , IP₃ and Ca²⁺, is initiated, leading to platelet activation [55, 57-60]. This cascade is depicted in **Figure 1**. Upon platelet activation, the platelet's

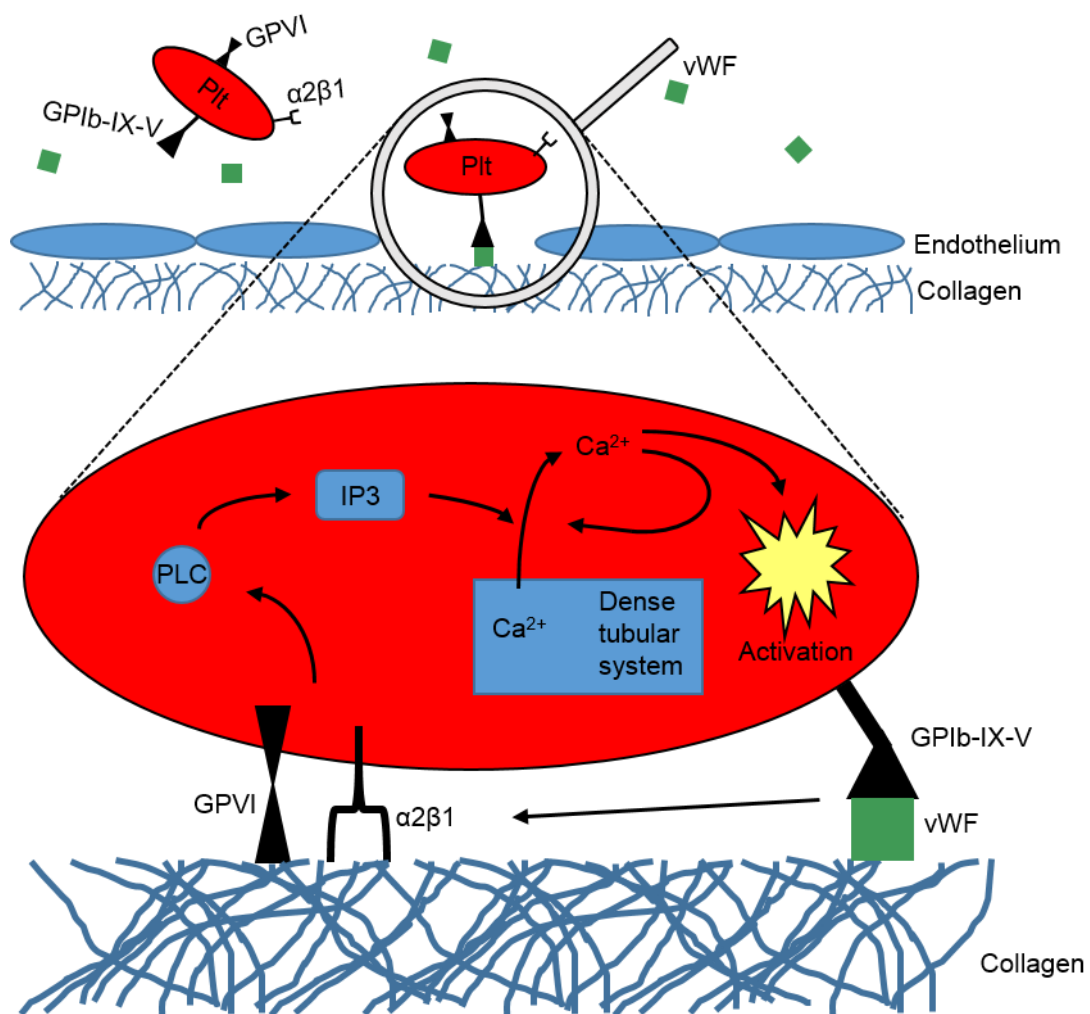


Figure 1: Initiation step after vessel injury in the endothelium. Plt: Platelet; vWF: von Willebrandt factor; PLC: Phospholipase C γ ; IP₃: Inositol triphosphate; GPIb-IX-V: Glycoprotein Ib-IX-V-platelet-surface receptor complex; GPVI: Glycoprotein VI-platelet surface receptor; $\alpha 2\beta 1$: Platelet-surface receptor.

A: Depiction of platelet adhesion to collagen, mediated by GPIb-IX-V and vWF after endothelial injury. Figure modified after Chen J et al., 'Interactions of Platelets with Subendothelium and Endothelium', *Microcirculation*, 2005.

B: Platelet receptor cascade after GPIb-IX-V and vWF have tethered to collagen. Figure modified after Sangkuhl K et al., 'Platelet aggregation pathway', *Pharmacogenetics and Genomics*, 2011.

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shape changes from a thin discoid cell into a rounded and vesicular cell with spines [57, 61]. Simultaneously, the granule's content is released into the extracellular space [59, 61]. These mechanisms are mediated by myosin phosphorylation, which leads to the reorganization of microtubules and actin filaments with a subsequent granule release [36, 45, 59, 61-63].

Release of the granule's content enables the second step of clot formation. The monolayer of platelets is enhanced by additional platelets and the procoagulant function of the platelets is elevated by auto-amplifying agents [55, 57]. These are phospholipase A₂, arachidonic acid, Cyclooxygenase-1, thromboxane A₂ and phospholipase C-β (PLCβ), all leading to a raise of intracellular Ca²⁺ [11, 55, 57, 64-66].

A further auto-amplifying agent is ADP, which is released from dense granules [12, 53, 55, 57]. Adenosine diphosphate auto-amplifies the activation signal by stimulating its G-protein-coupled receptors P2Y₁ and P2Y₁₂, thus activating PLCβ and increasing the plasma level of Ca²⁺ [11, 31, 55, 67].

The most potent activation amplifying agent is thrombin [55, 57]. As the inactive pre-stage prothrombin, it circulates in plasma [12, 68]. In a small amount, thrombin is already generated during the initiation step, due to prothrombin's exposure to the sub-endothelial expressed tissue factor [69, 70]. However, the vast majority of thrombin is produced during the step of clot formation as a second peak of thrombin generation [69, 71]: With the release of the granules' content, a large number of molecules is released, including factors V, VII, IX, and XIII [12, 41, 57]. In addition, after platelet activation phosphatidylserine (PS) is flipping from the inner leaflet of the platelet's membrane to the outer leaflet [8, 72, 73]. In the presence of PS and Ca²⁺, coagulation factors V and X form the prothrombinase complex, which cleaves prothrombin to thrombin [57, 69, 70, 74]. Thrombin associates with the G-protein-coupled receptor PAR1, activating PLCβ and increasing intracellular Ca²⁺ [11, 55].

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The third step of clot formation is called stabilization [55]. During this stage, a strong mesh of fibrin and vWF is formed within the platelets, which stabilizes the plug and forms a stable clot [55]. Besides fibrinogen and vWF, an essential role is played by the receptor $\alpha_{IIb}\beta_3$, which is released from the α -granules [12, 55, 75-77].

1.2 Coronary artery disease

1.2.1 Definition

There are a large number of known risk factors for developing coronary artery disease (CAD), including age, smoking, diabetes mellitus, hypertriglyceridemia, genetic predisposition and arterial hypertension [78, 79]. Atherosclerosis can affect the whole body, leading to strokes, peripheral artery disease or CAD [4, 80].

The main symptom of CAD is chest pain, called 'angina pectoris' (AP), which is caused by a mismatch of the myocardial supply and the demand for oxygen [78]. Coronary artery disease is categorized into stable AP and acute coronary syndrome (ACS). Stable AP appears in situations of physical or emotional stress and disappears at rest. The Canadian Cardiovascular Society (CCS) has published a classification to grade the severity of stable AP from grade 0 to grade IV [81]. If an AP appears *de novo*, has a CCS score of IV or increasing pain character, or does not disappear at rest, it is called an ACS.

Acute coronary syndrome describes a group of CAD manifestations which can be immediately life-threatening. These manifestations are not only distinguished by clinical signs but also by chemical parameters and the results of an electrocardiogram (ECG). Every ACS has to be treated as an emergency [82, 83].

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The clinical manifestation ranges from an unstable angina pectoris as the simplest type of ACS to a non-ST-segment elevation myocardial infarction (NSTEMI) to a fulminant necrosis through all the layers of the myocardium, called an ST-segment elevation myocardial infarction (STEMI) [78, 84-89]. A brief overview to distinguish between these manifestations is depicted in **Table 1**. Even if not mandatory fatal, an unstable AP can be a precursor to a myocardial infarction and has to be treated quickly. Both a STEMI and NSTEMI can lead to sudden cardiac death and are commonly treated with interventional heart catheterization or a bypass operation [82, 89].

Table 1: Manifestations of CAD. Based on 'ESC guidelines on the management of stable coronary artery disease', 2013 [78, 84-89].

	CCS	Troponin	ECG
Stable AP	Unaltered I–III	Negative	Normal
ACS Unstable AP	IV, raised or <i>de novo</i>	Negative	Normal, ST depression or T-inversion
NSTEMI	IV, raised or <i>de novo</i>	Positive	Normal, ST depression or T-inversion
STEMI	IV, raised or <i>de novo</i>	Positive	ST elevation or new LBBB > 20 min

1.2.2 History

Research into angina pectoris has a long history. The first clinical description was made by Heberden in 1772 [90, 91]. In 1812, Warren established the term 'angina pectoris', a combination of 'angina', meaning 'strangling', and 'pectus' meaning 'chest' [91, 92]. The first article reconciling the symptoms and pathology was published in 1879 by Hektoen, who concluded that the origin of angina pectoris is based on sclerotic changes in the coronary arteries, leading to a subsequent thrombosis [91, 93, 94]. The ECG was a further landmark in clinical cardiac research. The development of the ECG was based on findings of muscle electricity by Galvani in the 18th century, improved significantly by Einthoven, Wilson and Goldberger in the early 20th century [95-99]. Eventually, Pardee described the typical reactions recorded by an ECG during a myocardial infarction; in particular the elevation of the ST-segment, which became famous as 'Pardee's sign', and is known today as 'ST elevation' [100-102].

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The first approach in therapy was made by Herrick in 1912 [91]. Considering stress as the main factor triggering ACS, he recommended absolute bed rest in calm rooms for several days to prevent patients experiencing physical and emotional stress [91, 103, 104]. Many ACS patients died from arrhythmias and sudden cardiac death, which led to establishment of coronary care units in 1961 [103]. These units enabled constant monitoring and thus instant cardiac massage in case of ventricular fibrillation, which considerably reduced the death rate after myocardial infarction [91, 103, 105]. The next progress in treating myocardial infarction was the development of the heart catheter.

Research into vessel catheterization had already been conducted as early as 1844, and advanced in 1929 with Forssmann's first catheterization of a healthy human heart – conducted on Forssmann himself [91, 106-109]. Thereafter, progress in new approaches to treating coronary stenosis developed rapidly. The first 'percutaneous transluminal coronary angioplasty' (PTCA) was performed by Grüntzig in 1977 [103, 107, 110-118]. Ten years later Dotter implanted the first coronary stent [103, 107, 110-112, 114, 118, 119].

Besides the treatment of ACS, research into risk factors gained greater consideration when the Framingham study started in 1948 [91, 120]. It was designed as a long-term epidemiological study about the development of cardiovascular diseases [120]. In this study, elevated blood pressure, elevated serum cholesterol, smoking and being male were indicated as risk factors and led to the development of primary and secondary prevention mechanisms [120-122].

1.2.3 Therapy

A STEMI is treated according to the 2014 guidelines on myocardial revascularisation of the European Society of Cardiology and the European Association for Cardio-Thoracic Surgery [123]. To minimize the myocardial ischemia duration, the revascularisation should be performed as fast as possible

and preferably within the first two to three hours after becoming symptomatic [123]. Hospital structures are commonly organized with the purpose to reduce the time between a patient's arrival and the primary percutaneous coronary intervention (PCI) to less than 60 minutes [123]. The preferred reperfusion strategy is a percutaneous catheter intervention in which the stenotic passage is dilated with a balloon catheter, followed by the implantation of an endoluminal stent [123, 124]. Primary stent implantation can be conducted with bare-metal stents (BMS) or drug-eluting stents (DES) [123]. Drug-eluting stents have indicated a reduced risk for restenosis, but an elevated risk for in-stent thrombosis [125]. The long-term outcome of the new generation of DES is slightly better than the BMS [126, 127]. For stent implantation via the arteries, the radial approach is preferred and associated with a lower risk of acute bleeding events [123, 128].

Today, revascularisation via fibrinolysis has nearly been replaced by primary PCI as the preferred approach to restore the myocardial oxygen supply [103, 123]. Fibrinolysis should only be considered as a pre-hospital therapy if the expected period of time between the first medical contact and catheter intervention is longer than 120 minutes, and on condition that the onset of symptoms was within the last 120 minutes [123]. After arrival in hospital, a subsequent PCI is still recommended [123].

Besides PCI, for patients with a STEMI, dual antiplatelet therapy with acetylsalicylic acid and a P2Y₁₂ receptor blocker is recommended, if possible before intervention [123].

1.2.4 Myocardial regeneration

1.2.4.1 Estimating recovery of ventricular function

The extent of myocardial damage after an infarction influences cardiac performance, which results in a reduced left ventricular ejection fraction (LVEF) [129]. After a STEMI, a LVEF of <40% is a reliable predictor to estimate short- and long-term mortality in terms of the development of adverse events [129, 130].

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Due to heterogeneous healing and remodelling of the left ventricle, the early assessment of the LVEF has no reliable predictive value with regard to its long-term development [131, 132]. The best prognosis for long-term mortality can be assessed with the LVEF six months after a myocardial infarction, while studies are inconsistent concerning the benefit of early implantation of a cardioverter-defibrillator [133-136]. The current therapeutic regime to extend the long-term survival is the best medical treatment [137].

One approach to improving individual and early therapy is to search for early predictors of future LVEF outcomes [131]. Predictors of LVEF improvement are initial low levels of troponin and a low peak of creatine kinase [136, 138-141]. Another tool to estimate LVEF recovery is early evaluation of the acute infarct size via gadolinium-enhanced magnetic resonance imaging (MRI), which can detect a hibernating myocardium and is the best tool to predict long-term mortality, development of adverse events and recovery of LVEF [131, 142-150].

1.2.4.2 Platelet chemokines and the SDF-1-CXCR4/CXCR7 axis

In recent years, special attention has been given to chemokines and their influence on myocardial recovery. Chemokine proteins are a subgroup of cytokines, which can be secreted by many distinct cell types working as agents for cell communication [151]. They play a role as mediators in a wide range of processes such as inflammation, cardiovascular diseases, infections, survival, apoptosis, adhesion and many more [152].

Based on their molecular structure, chemokines can be divided into four groups: the CC, CXC, CX₃C and XC subfamilies [153]. Appending an 'L' for 'ligand' and a number, which represents the number of the corresponding chromosome nomenclature of its gene location, forms the official name of the chemokine [154]. The corresponding receptor is named with an 'R' as suffix [155]. Receptors are G-proteins coupled with seven transmembrane domains [155]. They have an affinity to more than one ligand and have another numeration system [154]. For example, suitable receptors for CXCL12 are CXCR4 and CXCR7 [154, 156]. Besides the official nomenclature, ligands also have commonly used names. For

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example, CXCL12 is commonly called 'SDF-1', since it is a product of the chromosome 'SCYB12' with the gene 'SDF-1' [154].

The chemokine SDF-1 has a high affinity to CXCR4 and CXCR7, and is expressed by a number of different tissues [156-159]. It is involved in the development, homing, proliferation and survival of hematopoietic progenitor cells or in neuromodulation [156, 158, 159]. Its receptors are expressed in many cell types, including platelets [160]. In mesenchymal stem cells, expression of this type of receptor is elevated under hypoxic conditions [161]. Expression on the platelet's surface remains on the same level without being affected by the platelet's state of being activated or resting [162].

The chemokine SDF-1 is a key player in the recruitment of CXCR4-expressing cells from bone marrow to damaged tissue for the purpose of better regeneration [163]. This process has been demonstrated in myocardial infarction and strokes [156, 164, 165]. In patients with CAD, high platelet expression of SDF-1 is concomitant with high levels of CXCR4 and CXCR7 [162]. Patients with ACS also exhibit higher levels of CXCR7 expression compared to patients with stable CAD [162].

The SDF-1-CXCR4-CXCR7 axis has an influence on myocardial regeneration. In ACS patients with a high expression of the platelet SDF-1, an amelioration of the LVEF could be detected after three months [166]. In patients with symptomatic CAD, low levels of CXCR4 correspond to a low overall mortality compared to patients with high levels of CXCR4 [167]. High levels of the platelet CXCR7 in ACS patients indicate a correlation with an improvement of the LVEF after three months [162].

1.3 Apoptosis

'Apoptosis' comes from the Greek word 'αποπτωση', which can be translated as 'pieces of leaves falling from the tree in autumn' [168]. This translation is commonly regarded as a metaphor for the removal of unwanted cells [169]. It is a complex process of programmed cell death and plays a crucial role in multicellular organisms [169]. From the beginning of embryonic development until our death, apoptosis has an influence on our growth, our homeostasis, and the development of diseases [170-179].

1.3.1 Morphological changes in apoptotic cells

The morphological changes of apoptosis are well known. In 1974, Kerr noticed two discrete stages [180]. The first stage comprises the formation of apoptotic bodies and during the second stage, cell fragments undergo phagocytosis [180].

More specifically, the first stage begins with the condensation of the nuclei, DNA fragmentation and shrinkage of the cell [169, 172]. This is followed by nuclear fragmentation and cell fragmentation in small apoptotic bodies, keeping the cell organelles intact [169, 172, 180].

During the second stage, apoptotic bodies become phagocytosed by the surrounding cells [169, 172]. These cells are macrophages, but also intact cells of the surrounding tissue [180]. During this process, none of the intracellular components are released into the extracellular space, and all organelles remain intact [169, 181].

Another morphological feature of apoptosis is the exposure of phosphatidylserine (PS) at the outer cell membrane [7]. The negatively charged PS is an ingredient of the plasma membrane, and is localized on its inner leaflet [182-185]. Upon activation, it flips to the outer leaflet of the cell membrane. The degree of PS exposure is assumed to be cell-type-specific, but nevertheless is a common

feature of nearly all apoptotic cells [186, 187]. Phosphatidylserine serves as a decoy for macrophages, leading to the recognition and removal of apoptotic cells [183, 184].

1.3.2 A molecular view of apoptosis

The process of apoptosis involves many molecular processes and can be caused by a great number of different stimuli. These stimuli can originate from the extracellular space or mitochondrion, categorized as extrinsic and intrinsic pathways, respectively [169]. Stimuli using the extrinsic pathway are mediated by different ligands, activating death receptors such as TNF1, Fas, and DR4 [188]. The intrinsic pathway involves stress factors such as heat, radiation, hypoxia and anticancer drugs [189]. Both extrinsic and intrinsic pathways result in the induction of a caspase cascade [181]. Caspases are mediators of the proteolytic cleavage of the cell's components, leading to morphological changes during apoptosis [190]. At the same time, a subgroup of caspases mediates an inflammatory cell response [191, 192].

1.3.2.1 The extrinsic pathway

The extrinsic pathway is initiated by the activation of cell-surface death receptors [169, 193]. These receptors are transmembrane proteins from the tumor necrosis factor receptor (TNF) superfamily [169, 188]. The triggers are extracellular death ligands such as the Fas receptor, TNF-R1, DR4 (also called TRAILR1) or DR5 [169, 194]. There is a great number of other death ligands [169].

The activation of death receptors activates an intracellular signal cascade. The intracellular ligand associates with a protein with death domain (DD), which subsequently recruits the Fas-associated protein with death domain (FADD) [195-202]. This facilitates binding of the procaspase 8 to the FADD, which results in a self-activation of procaspase 8 to caspase 8, thus inducing the cascade of caspases [200, 203, 204]. **Figure 2** depicts a simplified extrinsic pathway, including its cross talk to the intrinsic pathway.

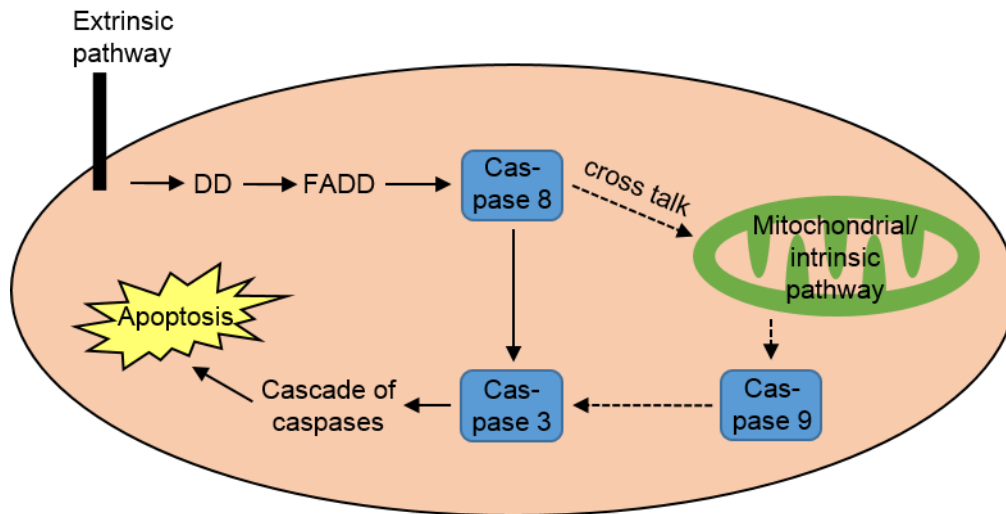


Figure 2: Extrinsic pathway of apoptosis. DD: Death domain; FADD: Fas-associated protein with death domain. The cross talk depicts the way from the extrinsic to the intrinsic pathway of apoptosis. Figure modified after Schultz D. R. et al., 'Apoptosis: Programmed Cell Death at a Molecular Level', Seminars in Arthritis and Rheumatism, 2003.

1.3.2.2 The intrinsic pathway

The intrinsic pathway is triggered by intracellular signals acting on intracellular targets [189]. Key players in this pathway are the mitochondria [189]. The mitochondria are organelles in eukaryotic cells, supplying them with chemical energy in the form of ATP [205, 206]. One crucial mitochondrial component of apoptosis is the protein cytochrome c [188]. Mitochondria consist of two membranes, an inner and an outer membrane [207]. The space between the outer and inner membranes is called the intermembrane space [188, 208]. In healthy cells, cytochrome c is stored in the intermembrane space [188, 208]. During apoptosis via the intrinsic pathway, cytochrome c is released into the cytosol [209].

Pro- and anti-apoptotic factors are located in the outer mitochondrial membrane, maintaining a state of equilibrium [169, 210]. With the ageing of the cell, anti-apoptotic factors withdraw, but the factors are also affected by stimuli such as toxins, radiation, free radicals, hypoxia, hyperthermia, or viral infections [189]. Disturbance of this state of equilibrium in favour of pro-apoptotic factors results in enhancement of mitochondrial outer membrane permeabilization (MOMP), which is attended by a reduction of the mitochondrial transmembrane potential

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($\Delta\Psi_m$) [169, 211, 212]. Potential dropping is an early event of apoptosis and marks a point of no return, whereafter transduction of the apoptotic signal passes off within five minutes, regardless of its initiating signal strength [212-214]. The dropping membrane potential leads to an increase of membrane permeability and pro-apoptotic proteins such as cytochrome c are released into the cytosol [169, 215]. The main actors in the balance of pro- and anti-apoptotic factors are proteins from the B-cell lymphoma 2 (Bcl-2) family, so called due to its first description in research about follicular lymphomas [169, 216]. B-cell lymphoma 2 proteins are a large and heterogeneous family, which, on the one hand, keeps cells from apoptosis, and on the other hand, triggers apoptosis [217, 218]. They are commonly located in the outer mitochondrial membrane [169, 219]. Based on their molecular domain motifs and their functional role in apoptosis, they are divided into inhibitors of apoptosis, promoters of apoptosis and BH3-only proteins [7, 217, 218, 220-223]. The best known inhibitors are Bcl-2 and Bcl-xL; the best known promoters are BAX and BAK [169, 209, 217, 218, 220, 224-228]. One BH3-only protein is BID, which is involved in extrinsic and intrinsic apoptosis [217].

With the initiation of apoptosis and the release of cytochrome c into the cytosol, the formation of the apoptosome is possible [169, 219, 229]. The apoptosome is a wheel-shaped enzyme complex, which serves as an activation tool for procaspase 9 [189, 230-233]. Upon release of cytochrome c, it binds to apoptotic protease activation factor 1 (Apaf-1), which allows an assembly of approximately seven Apaf-1 proteins to the apoptosome [189, 230-233]. The central structure of the apoptosome forms the procaspase 9 binding site, where initiator caspase 9 is activated, which initiates activation of downstream caspases [219, 231, 233-236]. **Figure 3** depicts an overview of intrinsic apoptosis.

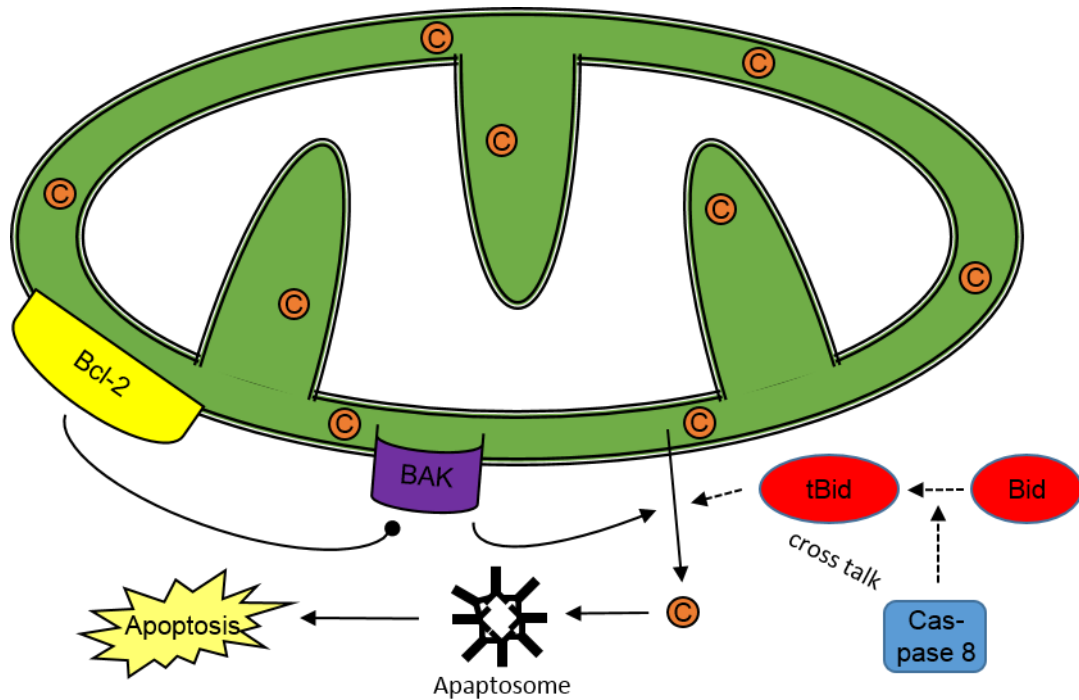


Figure 3: Intrinsic pathway of apoptosis. Bcl-2: B-cell lymphoma 2 are a group of anti-apoptotic proteins; BAK are a group of pro-apoptotic proteins; c: Cytochrome c release mediates formation of the apoptosome and thus apoptosis; The cross talk initiated by caspase 8 and by Bid and tBid links the extrinsic with the intrinsic pathway of apoptosis. Figure modified after Schultz D. R. et al., Apoptosis: 'Programmed Cell Death at a Molecular Level', Seminars in Arthritis and Rheumatism, 2003.

1.3.2.3 The caspases

Caspases are a group of proteolytic enzymes that play a key role in apoptosis [237]. The inactive pre-stage of the caspase is called the 'zymogen' or procaspase [192]. In nucleated cells they are expressed continuously, holding programmed cell death always ready to run [181, 238]. Activated caspases can cleave intracellular targets and help with the separation of the cell from its neighbours by cleaving the cell's skeleton [181, 239-244]. In addition, they activate other apoptotic enzymes with the purpose of supporting the caspase's depletion of the cell [243, 245-251]. All these caspase-triggered events are responsible for the morphological changes during apoptosis.

Depending on their role in the apoptotic cascade, caspases are divided into initiator and executioner (or downstream) caspases [252, 253]. Besides that, with caspases 1, 4, 5 and 11 a group of inflammation-regulating caspases is activated, leading to the release of inflammatory cytokines [191, 192]. The initiation of apoptosis is conducted by caspases 2, 8, 9 and 10 [192, 252, 254-256]. The

execution of apoptosis is conducted by caspases 3, 6 and 7 [192, 252, 254-256]. After activation, both the extrinsic and intrinsic pathways lead to activation of the caspase cascade by activating the executioner caspases, regardless of their trigger [169, 254]. Running the caspase cascade contains at least the activation of one initiator caspase, followed by the activation of one executioner caspase [255]. The change from procaspase to caspase is the result of the proximity-induced clustering to the dimeric active form [255].

Caspase 8 is capable of inducing a cross-talk between the intrinsic and extrinsic apoptotic pathways [189]. It cleaves the pro-apoptotic BH3-only protein BID, which is located in the cytosol [189, 257]. After cleavage, the henceforth called truncated BID (tBID) can translocate to the mitochondria, acting as a pro-apoptotic protein, thus releasing cytochrome c [254, 257]. With the release of cytochrome c, the mitochondrial pathway is activated, which amplifies the death receptor-induced signal [258].

1.3.3 Apoptosis in platelets

Apoptosis was formerly assumed to occur exclusively in nucleated cells, but platelets converting to apoptotic bodies was discovered as late as 1997, as a consequence of research into the short lifespan of platelets [180, 259-261]. Until the discovery that the intrinsic pathway plays an essential role in the platelet lifespan, several models about life restriction had been proposed and rejected [7-9, 262-269].

The first protein discovered to be involved in platelet lifespan was Bcl-x_L [7-9]. Further studies on platelets, taking a closer look at these particular apoptotic features, revealed that they contain caspase 3, caspase 9, Apaf-1 and cytochrome c [270]. Furthermore, morphological changes could be demonstrated in ageing platelets and upon activation of apoptosis as well [270-273]. This included an increasing exposure of PS and other phagocyte scavenger receptors

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in ageing platelets with the simultaneous condensation of cytoplasm and the ability of mitochondrial membrane potential dropping [271-273].

The current model explaining the platelet lifespan is called the 'molecular clock' [7-9]. It assumes that degradation of Bcl-x_L occurs more quickly than degradation of BAK [7-9, 274, 275]. This model is substantially based on findings that depression of single pro- and anti-apoptotic proteins leads to an increase or reduction of platelet lifespan [7-9, 276, 277]. According to this model, platelet survival is mainly regulated by a balance of Bcl-x_L and BAK, leading to apoptosis after disturbance of this state of equilibrium in favour of BAK [7]. There is no evidence that platelets undergo an extrinsic pathway of apoptosis [7, 278]. Indeed, mRNA for the TRAIL receptors DR4 and DR5 has been found in platelets, but on the cell surface no expression of any corresponding protein like Fas has been detected [279].

2 Aims

Despite progress in the management and therapy of myocardial infarction, mortality is still high [2]. Unfortunately, there is little knowledge about prognostic markers indicating a prognosis of cardiac function [280, 281]. Indeed, studies have been conducted on the functional cardiac recovery in a population of ACS patients in general, but no special consideration has been given to STEMI patients in particular [166]. Recent studies on ACS patients have indicated a correlation between elevated levels of CXCR7 platelet surface expression and an improvement of the left ventricular ejection fraction (LVEF) within three months [160, 162]. In addition, there is evidence that CXCR4 and CXCR7 are closely associated with platelet survival. For this reason, a more precise view on the prognostic markers of this particular disease is necessary.

For this study, a population of patients with a diagnosis of acute STEMI was recruited. The aim was to investigate whether there is a relation between the six-month development of the LVEF on the one hand, and platelet survival on the other hand.

The study sought to achieve the following:

- To measure the course of the LVEF using thoracic MRI as a parameter for the cardiac function in STEMI patients;
- To assess apoptosis in platelets, including measuring their survival with tetramethylrhodamine ethyl ester (TMRE) and their apoptosis with annexin V;
- To investigate the association between markers of platelet apoptosis and the platelet surface expression of CXCR4 and CXCR7; and
- To investigate the association between platelet apoptosis and the course of the LVEF after a STEMI.

3 Materials and methods

3.1 Patient characteristics

This pilot study was conducted with patients who had experienced ST-segment elevation myocardial infarctions (STEMIs). The diagnosis and treatment were according to the guidelines of the Task Force on the management of ST-segment elevation acute myocardial infarction of the European Society of Cardiology (ESC) [88]. All subjects were immediately treated with percutaneous transluminal coronary angioplasty (PTCA) in accordance with these guidelines [123]. This study was conducted in the Department of Cardiology of the University of Tübingen between August 2013 and June 2015 [1], and included 87 patients [1]. Blood was taken within 24 hours after intervention.

In each case 70 mL of blood was taken from a peripheral vein and immediately analysed for platelet survival and platelet apoptosis [1]. Contemporaneously, platelets were tested for the expression of the chemokine receptors CXCR4 and CXCR7 [1]. All samples were analysed with a fluorescence-activated cell sorter (FACS) [1]. Furthermore, the LVEF% was evaluated in each patient who had undergone a cardiac MRI during intrahospital stay, and six months later [1].

All subjects were informed in detail about the study and gave written informed consent. The study complies with the guidelines of the European Parliament and of the International Conference of Harmonization for good clinical practice, and with the Declaration of Helsinki [282-284]. The institutional ethics committee of the University of Tübingen gave its approval for this study (270/2011BO1).

The baseline characteristics are presented in **Table 2**. For 38 of the 87 subjects, no information about previous pharmacological therapy was available [1].

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Table 2: Baseline characteristics. n: number of subjects; \pm : standard deviation [1].

Basic data		
Number of subjects	87	
Number of males	63 (72.4 %)	
Age (mean \pm SD)	60.8 \pm 12.1	
Cardiovascular risk factors		
	Yes	No
Arterial hypertension	56 (64.4%)	31 (35.6%)
Hyperlipidaemia	23 (26.4%)	64 (73.6%)
Diabetes mellitus	20 (23.0%)	67 (77.0%)
Smoker	41 (47.1%)	39 (44.8%) ex: 7 (8.0%)
ACS in family	17 (19.5%)	70 (80.5%)
Prior CAD	18 (20.7%)	69 (79.3%)
Medication on admission		
	Yes	No
Acetylsalicylic acid	14 (16.1%)	73 (83.9%)
Clopidogrel	2 (2.3%)	85 (97.7%)
Prasugrel	1 (1.1%)	86 (98.9%)
Ticagrelor	3 (3.4%)	84 (96.6%)
Oral anticoagulant	2 (2.3%)	85 (97.7%)
ACE inhibitor	13 (14.9%)	74 (85.1%)
Angiotensin II receptor inhibitor	8 (9.2%)	79 (90.8%)
Calcium channel inhibitor	9 (10.3%)	78 (89.7%)
Beta blocker	20 (23.0%)	67 (77.0%)
Diuretic (thiazide, loop)	9 (10.3%)	78 (89.7%)
Potassium-sparing diuretic	4 (4.6%)	83 (95.4%)
Statin	12 (13.8%)	75 (86.2%)

3.2 Materials

Table 3 lists the used substances and agents. The consumable materials are listed in **Table 4**. The technical equipment is registered in **Table 5**.

Table 3: Used substances and agents.

Name	Description
PBS-Ca ²⁺	This study used Dulbecco's Phosphate-buffered Saline with MgCl ₂ and CaCl ₂ , liquid, sterile-filtered, suitable for cell culture from Sigma-Aldrich, St. Louis, USA; Cat-No D8862.
PBS	This study used Dulbecco's Phosphate-buffered Saline, modified, without calcium chloride and magnesium chloride, liquid, sterile-filtered, suitable for cell culture from Sigma-Aldrich, St. Louis, USA; Cat-No D8537.
PFA	This study used Paraformaldehyde (PFA) for synthesis, CAS-No 30525-89-4 from Merck-Schuchardt, 8011 Hohenbrunn, Germany; Cat-No 818715.
CXCR4	This study used anti-hCXCR4 PE conjugated from RD Systems Minneapolis, Minnesota 55413, USA; Cat-No FAB170P.
CXCR7	This study used anti-hCXCR7/RDC-1 PE conjugated from RD Systems, Minneapolis, Minnesota 55413, USA; Cat-No FAB42271P.
CD42b	This study used CD42b FITC conjugated from Beckman Coulter Company, 13276 Marseille, France; UNSPSC 41116015.
Thrombin	This study used Thrombin from human plasma, lyophilized from Roche, Basel, Switzerland; Cat-No 10602400001.
Annexin V	This study used Annexin V FITC conjugated from ImmunoTools, 26169 Friesoythe, Germany; Cat-No 314900013.
TMRE	In this study, TMRE was ordered from Invitrogen, Carlsbad, USA; Cat-No T669.

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Table 4: Consumable materials.

Name	Producer	Material
S-Monovette 7.5 ml Z-Gel	Sarstedt	Polypropylene
S-Monovette 8.5 ml CPDA	Sarstedt	Polypropylene
Safe-lock microcentrifuge tubes 1.5ml	Eppendorf	Polypropylene
Cellstar tubes 15 ml	Greiner Bio-One	Polypropylene
Falcon tube 5 ml	Corning	Polystyrene
Pipette Research 100 – 1000 µl	Eppendorf	-
Pipette Research 10 – 100 µl	Eppendorf	-
Pipette Research 0.5 – 10 µl	Eppendorf	-
Pipette tips 1000 µl	Ratiolab	Polypropylene
Pipette tips 200 µl	Sarstedt	Polypropylene
Pipette tips 10 µl	Biozym Scientific	Polypropylene
Pasteur-plast pipets, 2,5 ml, 150 mm, graduated	Ratiolab	Polyethylene

Table 5: Technical equipment.

Equipment	Description
Centrifuge	This study used a Heraeus® Multifuge® 1S from Thermo Fisher Scientific, Waltham, MA, USA.
Automated Hematology Analyzer	This study used the Automated Hematology Analyzer KX-21N™ from Sysmex, Kobe, Japan.
Shaker	This study used a Vortexer IKA Vibrofix VF2 from IKA-Werke, Staufen im Breisgau, Germany.
FACS	This study used the FACSCalibur™ Flow Cytometer 342975 from Becton Dickinson Biosciences, Heidelberg, Germany.
FACS-Program	This study used a BD FACStation™ Mac Pro computer with a 2.8-GHz Quad-Core Intel® Xeon® processor and 2 GB RAM was used as a working station. As an analysis software, this computer ran with BD CellQuest™ Pro, Version 5.d.7f6b.
MRI	This study used a Magnetom Avanto from Siemens Healthcare, Erlangen, Germany.
MRI-Program	Thus study used cvi42 from Circle Cardiovascular Imaging Inc, Calgary, AB, Canada.

3.3 Methods

3.3.1 Study design

This study was designed as a prospective study. It was based on the parallel measurement of apoptosis and the expression of distinct chemokine receptors in platelets. Measurements were taken within the first 24 hours after PTCA stent implantation in patients with a diagnosis of STEMI. Analysis and course of the LVEF was conducted twice with MRI. Examination took place during initial hospitalization after the event, and six months later.

Blood samples were taken from the peripheral veins of the subjects. Measurements of platelet apoptosis in platelet-rich plasma (PRP) were taken for each subject. In a parallel measurement in whole blood, the platelets' expression of chemokine receptors was analysed. Experiments were conducted in the laboratories of the University of Tübingen, Department of Cardiology (Department of Internal Medicine III), where a constant temperature of 22°C could be ensured. All experiments were conducted at this temperature.

3.3.2 Preparation of chemokine dye

3.3.2.1 Preparation of a dilution with 1% paraformaldehyde

To prepare 500 ml of a 1% paraformaldehyde (PFA) dilution, 5 g of PFA was taken and dissolved in 400 ml of phosphate-buffered saline (PBS); 40 ml of sodium hydroxide (NaOH) (with 1 N equivalent concentration) was added. Afterwards, the liquid was stored in a dark environment. All these steps were conducted under a fume hood. In an environment of 65°C, this liquid was stirred continuously, until the liquid was clear. After being cooled down to room temperature, the pH of the liquid was reduced to 1.4 with hydrogen chloride (HCl) (with 1 M molar mass) and was filtered through a 0.2 µm filter.

3.3.2.2 Analysis of chemokines CXCR4 and CXCR7

For the analysis of the platelets' CXCR4 and CXCR7 expression, whole blood was taken. The blood sample was decanted into a CPDA-coated S-Monovette. Afterwards, 20 μ l of the whole blood was put into a Cellstar tube and diluted with 980 μ l of PBS.

Amounts of 35 μ l of this dilution were transferred into each of two Falcon tubes. An amount of 5 μ l of PBS was added to each Falcon tube to increase the level of dilution. Into each dilution, 5 μ l of CD42b was added. An amount of 5 μ l of CXCR4 was added to one of the Falcon tubes, and 5 μ L of CXCR7 was added to the other Falcon tube. These two dilutions were mixed with the vortex mixer and incubated in a dark environment for 30 minutes.

The incubation of both samples was stopped with 300 μ l of the prepared PFA dilution and was immediately analysed with FACS. **Figure 4** depicts a schematic illustration of the experimental setup for the expression of chemokine receptors.

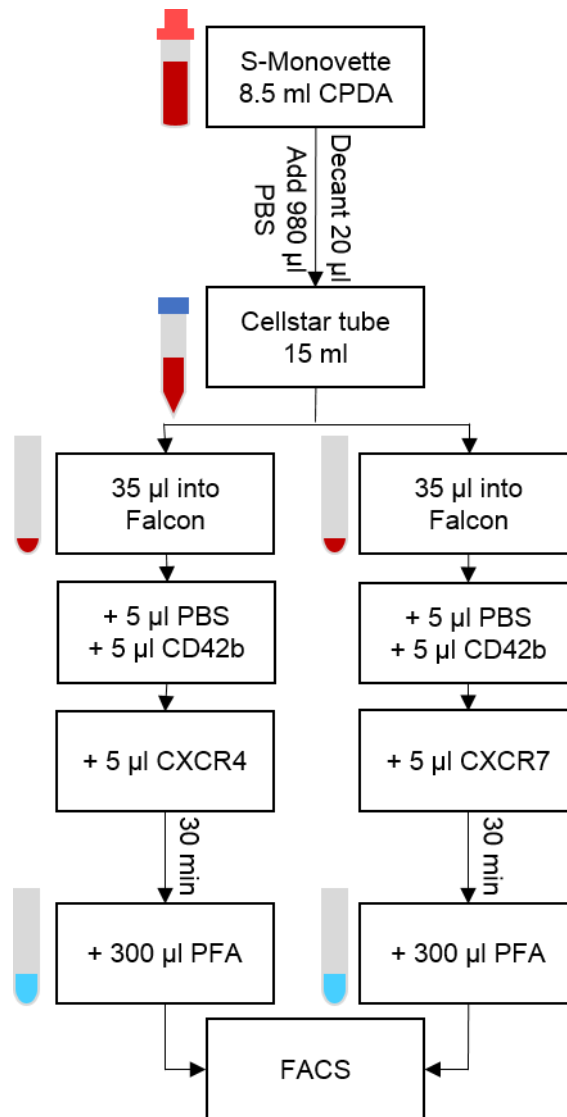


Figure 4: Preparation of chemokine dye.

3.3.3 Preparation of apoptosis dye

3.3.3.1 Preparation of platelet-rich plasma and of the dyes

The ordered thrombin was in a lyophilized state. In a preparatory step, it was dissolved in 2000 μL of PBS- Ca^{2+} . The ordered TMRE was delivered in a high concentration: 10 μL of the TMRE was diluted with 990 μL of PBS- Ca^{2+} .

After having taken a blood sample with a Z-Gel-coated S-Monovette, the sample was directly transferred into a Cellstar tube, which is appropriate for centrifugation. This sample was put into the centrifuge and was hydro-extracted with 1000 rounds per minute for 20 minutes. Subsequently, the supernatant PRP

was carefully transferred with a Pasteur pipette into another Cellstar tube. To ensure that only pure PRP would be transferred, 2 ml of PRP above the remaining effluent was not transferred.

In order to measure the platelet count in the PRP, 10 μL of the extracted PRP was transferred into a 1.5 ml Safe-lock microcentrifuge tube and diluted with 90 μL of PBS- Ca^{2+} . The platelet number in the resulting aliquot was measured with the Automated Hematology Analyzer. The platelet count was noted and the aliquot was rejected afterwards. With the measured platelet count, the final aliquot for the FACS analysis was formed. Via extrapolation, a PRP volume containing approximately 10^6 platelets was calculated. This volume was filled into each of four Falcon tubes. Afterwards, the prepared PRP was incubated with the other ingredients.

3.3.3.2 Induction of apoptosis and incubation with apoptosis-detecting agents

To approximate the whole potential of apoptosis in platelets, the samples were divided into two: One half to be incubated with thrombin and the other half to remain resting without thrombin.

Before the incubation of the platelets, an individually calculated volume of PBS- Ca^{2+} was added. The volume of PBS- Ca^{2+} depended on the added PRP volume in the step before, and also depended on the volume of the agents which had to be added afterwards. The purpose of this calculation was to configure a total volume of 50 μL for each tube. With this approach, it was guaranteed that the number and the concentration of platelets in each aliquot was the same. An amount of 5 μL of the thrombin dilution was added to two of the four Falcon tubes. To incubate this mixture, the samples were covered in a dark environment for 60 minutes. The two samples without added thrombin remained unaffected and were covered in a dark environment for 60 minutes as well. After incubation, 5 μL of TMRE was added to one of the thrombin-activated tubes and one of the resting tubes. Simultaneously, 4 μL of Annexin V was added to the remaining two tubes. These samples were incubated in a dark environment for 30 minutes. After

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incubation, 400 μL of PBS- Ca^{2+} was added to each tube. Analysis was conducted immediately with FACS. **Figure 5** depicts a schematic illustration of the experimental setup for measuring apoptosis.

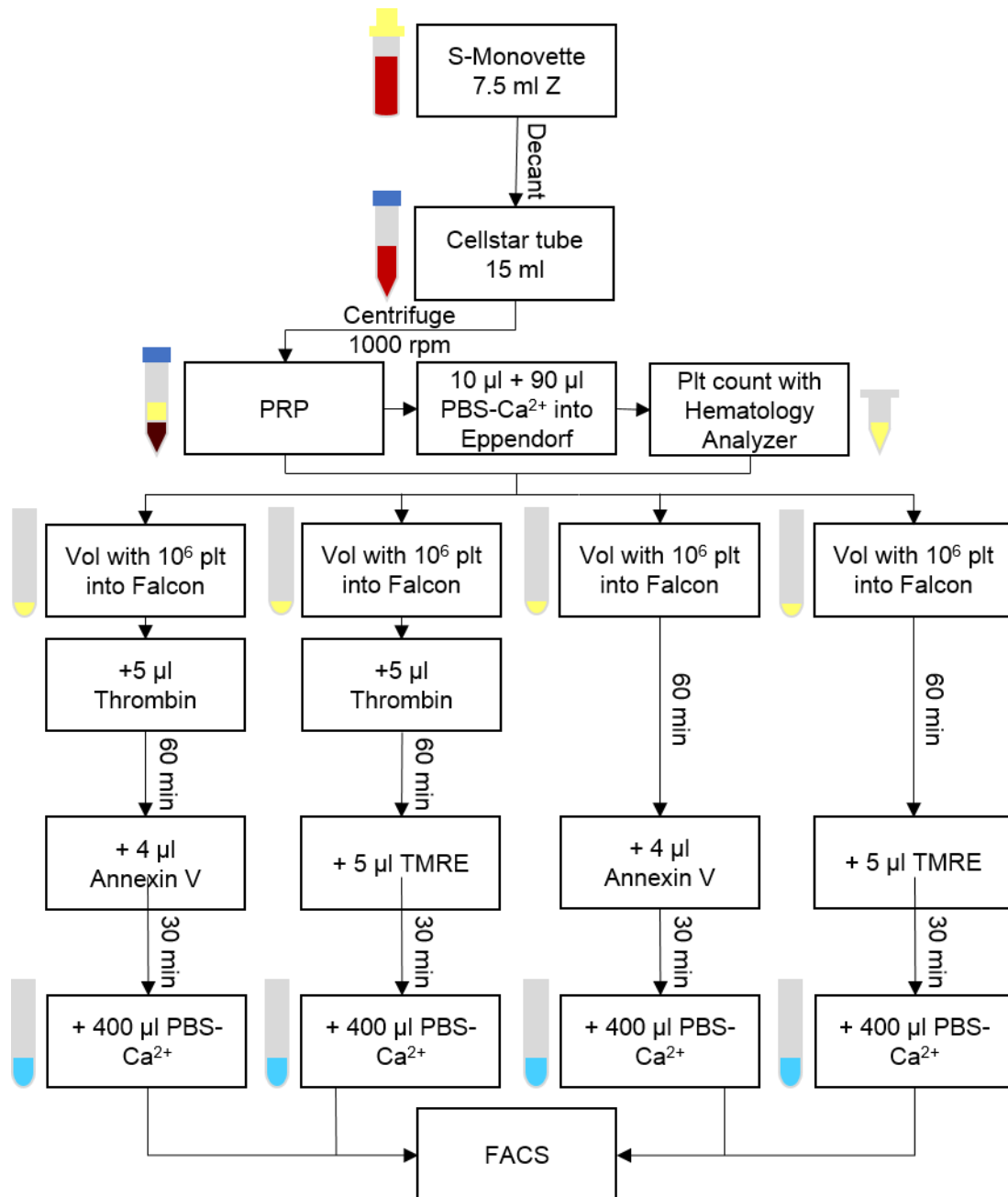


Figure 5: Preparation of apoptosis dye.

3.3.4 FACS analysis

The FACS that was used possesses two lasers with different wavelengths:

Blue laser: Air-cooled argon-ion laser, 15 mW with $\lambda = 488$ nm

Red laser: $\lambda = 635$ nm

This device has six detectors, which are calibrated for different wavelengths. The detectors are designed for forward scatter (FSC) and sideward scatter (SSC). Additional detectors for PE- and FITC-fluorescence, and for PerCP- and APC-fluorescence were installed. The fluorescence channels are called FL1 to FL4. Specifically, the used detectors and measured wavelengths are as follows:

FSC: Solid-state silicon detector

SSC: Photomultiplier tube (PMT)

FL1: Photomultiplier tube (PMT) for $\lambda = 530$ nm (FITC)

FL2: Photomultiplier tube (PMT) for $\lambda = 585$ nm (PE/PI)

FL3: Photomultiplier tube (PMT) for $\lambda > 670$ nm (PerCP)

FL4: Photomultiplier tube (PMT) for $\lambda = 661$ nm (APC)

3.3.5 MRI analysis and follow-up

To evaluate the recovery of myocardial function, subjects received two cardiac MRI examinations. The baseline evaluation was carried out during initial hospitalisation, at a median of three days after the acute event and under conditions of cardiorespiratory stability. The follow-up evaluation was carried out six months later. Both of these examinations were conducted using thoracic MRI.

For imaging, patients were placed in a supine position while breath was held in expiration. Imaging was ECG triggered. For acquiring images, the steady-state free precession technique was used. For evaluation of the LVEF, various axes were chosen. These axes were oblique images through the left ventricle, resulting in a four-chamber view, a two-chamber view, and a short axis scout. The LVEF was calculated with analysis software, in which the contours of the endocardium in end diastole and end systole, respectively, had to be drawn manually.

3.3.6 Statistical analysis

For the statistical analysis, SPSS statistics, version 24.0.0.1, from the IBM Corporation in Armonk, USA was used.

To determine the distribution of the samples, the data set was analysed with a graphical evaluation via histogram and box-plot.

The next step was to conduct a bivariate analysis: Since linear correlation requires linearity, a scatter plot was created for optical evaluation. Under conditions that the values were normally distributed, without any outliers, and that there was a linear correlation between the two samples, Pearson's correlation coefficient was calculated. Under conditions that the normal distribution had to be rejected or outliers were detected, Spearman's rank correlation coefficient was calculated.

The level of significance was determined as $p < 0.05$.

4 Results

A total of 87 patients were included in this study [1]. The mean fluorescence intensity (MFI) was measured in TMRE dyed platelets, Annexin V dyed platelets, CXCR4 dyed platelets, and CXCR7 dyed platelets. Platelets were sampled on the first day after the ST-segment elevation myocardial infarction (STEMI). In addition, the left ventricular ejection fraction (LVEF) was surveyed within a median of three days after the STEMI during hospital stay as a baseline, and six months later as a follow-up [1]. The LVEF was surveyed using thoracic MRI and under conditions of cardiorespiratory stability.

Since the values of the LVEF were evaluated twice and at two different dates in the same subjects, they were treated as related samples. The results were ratio scaled and thus different calculations, including division, were possible. The MFI results of the fluorescence-activated cell sorter (FACS) were ratio scaled as well.

The tetramethylrhodamine ethyl ester (TMRE) dyed platelets and Annexin V dyed platelets were both evaluated under resting conditions and under fully activated conditions simultaneously. **Table 6** presents a list and explanations for every calculation which was used for the final analysis. A similar approach was followed with the results of the LVEF evaluation. The precise approach of conducted calculations is also presented in **Table 6**.

Results

Table 6: Values calculated. Explanations of the henceforth used terms of the results.

Value	Explanation/Calculation
TMRE MFI resting	MFI of TMRE dyed platelets under resting conditions
TMRE MFI thrombin	MFI of TMRE dyed platelets thrombin stimulated
Ratio platelet TMRE MFI resting/thrombin stimulated	$\frac{[\text{TMRE MFI resting}]}{[\text{TMRE MFI thrombin}]}$
Annexin V MFI resting	MFI of Annexin V dyed platelets under resting conditions
Annexin V MFI thrombin	MFI of Annexin V dyed platelets thrombin stimulated
Ratio platelet Annexin V MFI resting/thrombin stimulated	$\frac{[\text{Annexin V MFI resting}]}{[\text{Annexin V MFI thrombin}]}$
LVEF baseline	LVEF (%) evaluated immediately after event
LVEF follow-up	LVEF (%) evaluated six months after event
Ratio LVEF follow-up/baseline	$\frac{[\text{LVEF follow-up}]}{[\text{LVEF baseline}]}$
Platelet CXCR4 MFI	MFI of CXCR4 dyed platelets
Platelet CXCR7 MFI	MFI of CXCR7 dyed platelets

4.1 Descriptive statistics

4.1.1 Dropouts

There were manifold reasons for dropouts in this study. The most limiting factor was attendance at evaluation of the LVEF with thoracic MRI. Initially, 87 subjects had been recruited [1]. During the course of this study, 35.6% of the subjects rejected the first thoracic MRI evaluation and thus 56 subjects remained in the sample [1]. The second evaluation six months later was rejected by 37.5% of the remaining subjects [1]. Eventually, 35 subjects attended evaluation of the LVEF twice [1]. Comparing the baseline characteristics of these three groups, no structural difference was found [1]. In most cases, the reason for rejection of MRI evaluation was lacking motivation to participate in this study. Thus, the missing data were treated as missing at random.

Experimental failure occurred completely without regularity and was independent of the subject population. Eventually, all dropouts were regarded as part of the

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overall collective and the reasons for dropout were coincidental [1]. Therefore, it did not harm the statistical analysis.

4.1.2 Evaluated results over time

4.1.2.1 Left ventricular ejection fraction

To quantify the alteration of cardiac function, MRI-based evaluation of the LVEF was conducted within a median of three days after the event as a baseline and six months later as a follow-up [1]. The images of one subject are depicted in **Figure 6**.

In 35 subjects, cardiac function could be evaluated twice [1]. The average LVEF at baseline was 53.09%, with a standard deviation of 9.12% and a maximum span between 38% and 71% [1]. The average LVEF at follow-up was 58.28%, with a standard deviation of 10.36% and a maximum span between 34% and 80% [1]. Both the baseline and follow-up LVEFs were assumed to be normally distributed.

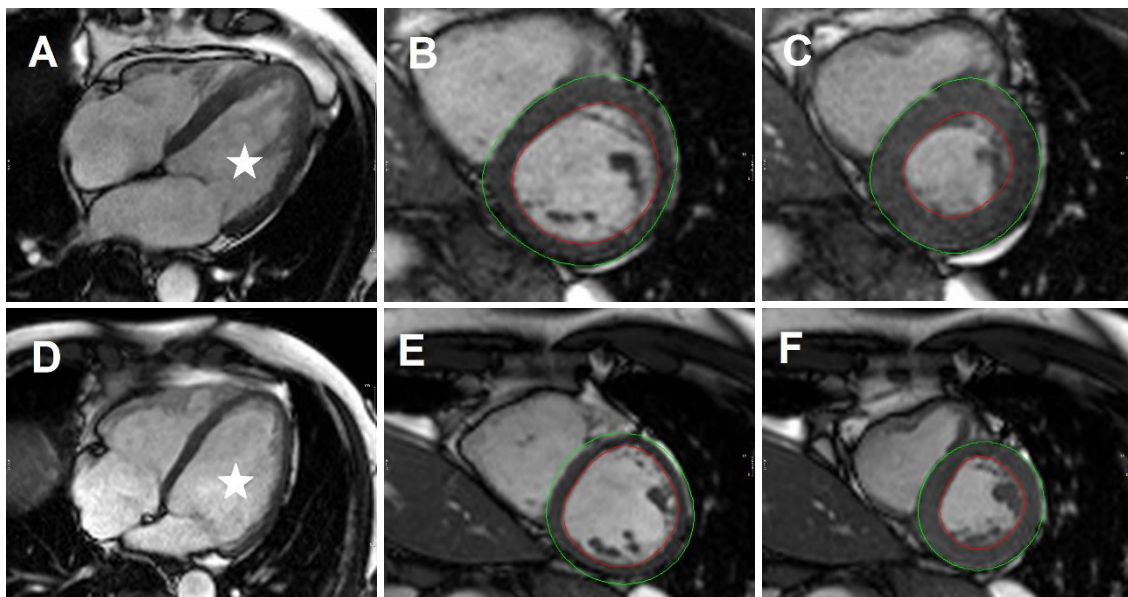


Figure 6: Measurement of LVEF baseline and follow-up. These images illustrate a typical result of the cardiac MRI. Here, imaging was conducted on a 58-year-old male patient with 52% LVEF at baseline and 58% LVEF at follow-up. **A – C:** Imaging at baseline; **D – F:** Imaging at follow-up; Star: left ventricular; green circle: pericardium of left ventricle; red circle: endocardium of left ventricle; **A & D:** four-chamber view; **B & E:** Short axis ventricle view, end diastolic in a middle layer of imaging; **C & F:** Short axis ventricle view, end systolic in a middle layer of imaging. These pictures were kindly provided by Patrick Krumm, M.D. from the Department of Radiology at the University Hospital Tübingen. White stars and letters were inserted for better explanation. For anonymization, the upper and lower edges of the picture were deleted [1].

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4.1.2.2 Tetramethylrhodamine ethyl ester

To illustrate a typical result of the FACS analysis, the results of resting platelets incubated with TMRE are presented in **Figure 7**. The fluorescence intensity of each cell in a sample was measured with FACS. The output value for each sample was the mean fluorescence intensity (MFI). **Table 7** presents the average results of platelet TMRE MFI.

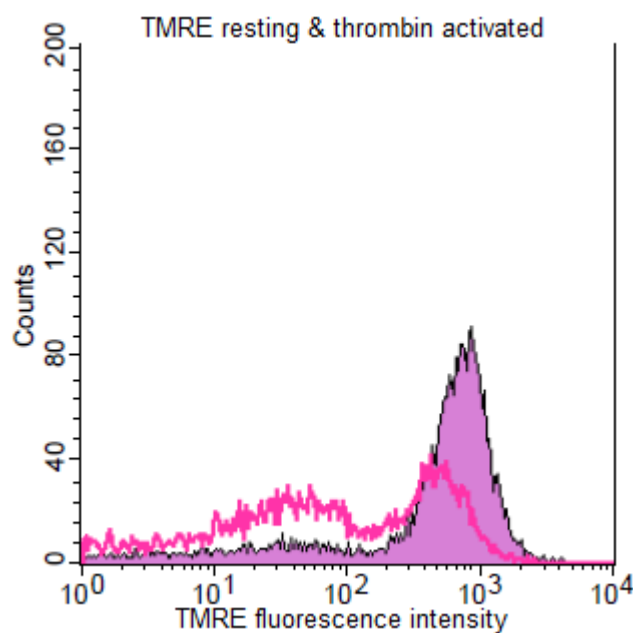


Figure 7: Fluorescence intensity of TMRE dyed platelets. The histogram indicates fluorescence intensity of TMRE dyed platelets. Graph with bright purple background illustrates fluorescence intensity of resting platelets and the dark purple line illustrates fluorescence intensity of thrombin-activated platelets. Both plots present the results sampled simultaneously in one patient.

In each plot, 10,000 events were counted. The intensity of fluorescence was detected with the second fluorescence channel of the photomultiplier tubes, calibrated for $\lambda = 585$ nm. The mean fluorescence intensity (MFI) in this exemplary measurement was, for **resting platelets**: 651.80 and for **thrombin activated platelets**: 218.78 [1].

Table 7: Results of TMRE dyed platelets. n: number of subjects; \pm : standard deviation; IQR: interquartile range; MFI: mean fluorescence intensity; In TMRE dyed resting platelets normal distribution was assumed, but it had to be rejected both in TMRE dyed thrombin-activated platelets, and in the calculated ratio of resting and thrombin-activated platelets. The results of TMRE MFI resting are depicted with mean and standard deviations. Both the results of the thrombin dyed platelets and the ratio between the resting and the thrombin-activated platelets are depicted with the median and interquartile range [1].

	n	Results
TMRE MFI resting	47	652.26 \pm 166.36
TMRE MFI thrombin stimulated	47	176.58 IQR 181.73
Ratio platelet TMRE MFI resting/thrombin stimulated	41	3.51 IQR 2.99

Results

4.1.2.3 Annexin V

To illustrate a typical result of the FACS analysis, the results of the resting platelets incubated with Annexin V are presented in **Figure 8**. The fluorescence intensity of each cell in the sample was measured with FACS. The output values for each sample was the MFI. **Table 8** presents the average results of platelet Annexin V MFI.

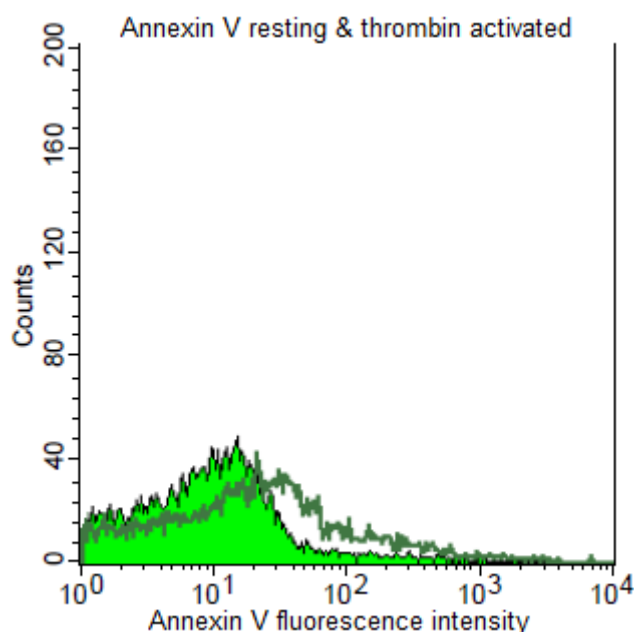


Figure 8: Fluorescence intensity of Annexin V dyed platelets. The histogram indicates the fluorescence intensity of Annexin V dyed platelets. The graph with bright green background indicates the fluorescence intensity of resting platelets and the dark green line indicates the fluorescence intensity of thrombin-activated platelets. Both plots illustrate the results sampled simultaneously in one patient.

In each plot, 10,000 events were counted. The intensity of fluorescence was detected with the first fluorescence channel of the photomultiplier tubes, calibrated for $\lambda = 530$ nm. The mean fluorescence intensity (MFI) in this exemplary measurement was, for **resting platelets**: 14.40 and for **thrombin-activated platelets**: 54.49 [1].

Table 8: Results of Annexin V dyed platelets. n: number of subjects; IQR: interquartile range; MFI: mean fluorescence intensity; none of the tested values of Annexin V dyed platelets was assumed as normally distributed. The results are depicted with the median and the interquartile range [1].

	n	Results
Annexin V resting (MFI)	59	12.69 IQR 7.09
Annexin V thrombin (MFI)	43	40.18 IQR 59.36
Ratio platelet Annexin V MFI resting/thrombin stimulated	43	0.35 IQR 0.45

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4.1.2.4 Chemokine receptors

Parallel with TMRE and Annexin V, the expression of CXCR4 and CXCR7 receptors was measured. The platelets were dyed with fluorescent agents, specifically for these receptors. To illustrate a typical result of the FACS analysis, the results are illustrated in **Figure 9**. The average results of detected MFI are depicted in **Table 9**.

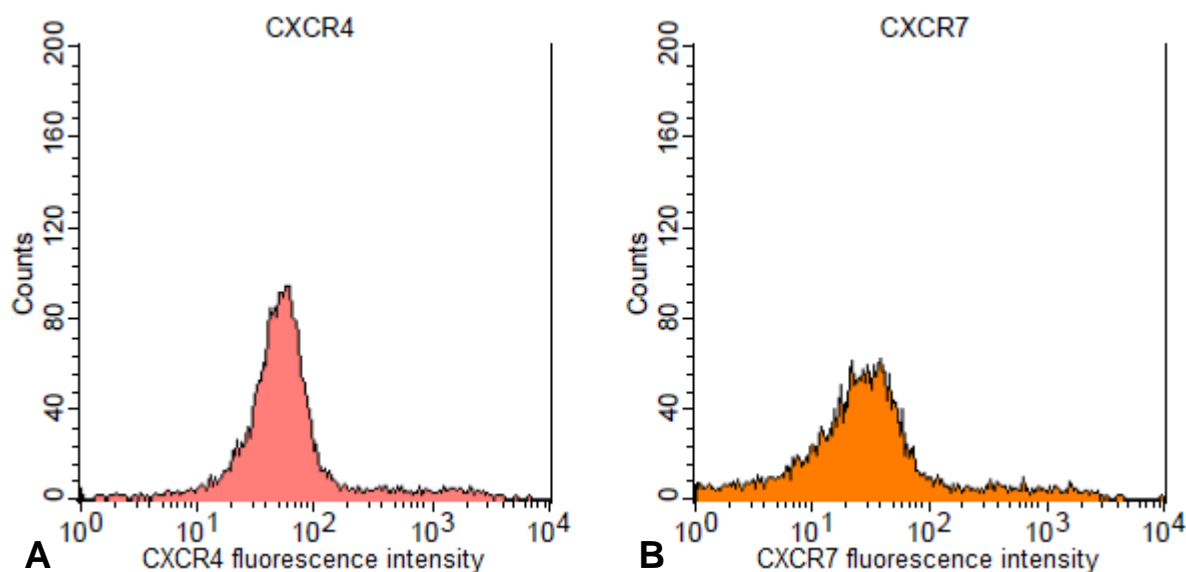


Figure 9: Fluorescence intensity of CXCR4 and CXCR7 dyed platelets. The histograms indicate the fluorescence intensity of CXCR4 and CXCR7 dyed platelets, respectively. In each plot, 10,000 events were counted. The intensity of fluorescence was detected with the second fluorescence channel of the photomultiplier tubes, calibrated for $\lambda = 585$ nm. The mean fluorescence intensity (MFI) in this exemplary measurement was, for **A**: 97.22 and for **B**: 75.14 [1].

Table 9: Chemokine receptor results of the FACS analysis. n: number of subjects; \pm : standard deviation; MFI: mean fluorescence intensity; all results were assumed to be normally distributed. The results are depicted with mean and standard deviations [1].

	n	Results
CXCR4 MFI	50	68.38 ± 21.56
CXCR7 MFI	53	56.08 ± 21.70

4.2 Exploratory data analysis

4.2.1 Association of chemokines and apoptosis

Bivariate analysis was conducted with the results of platelet chemokines with platelet apoptosis. The results are depicted in **Table 10**. The values of pure resting platelets dyed with apoptosis signalling agents revealed no significant correlation, but the ratio of resting and thrombin-stimulated platelets indicated a positive correlation for survival markers (TMRE), both with CXCR4 and CXCR7 [1]. For better presentation, the results are depicted in **Figure 10**. The ratio of resting and thrombin-stimulated platelets indicated a negative correlation for markers of apoptosis (Annexin V), both with CXCR4 and CXCR7 [1]. The results are depicted in **Figure 11**.

Table 10: Bivariate correlation of parameters of apoptosis with chemokines. MFI: mean fluorescence intensity; n: number of subjects; r: Pearson’s correlation coefficient; ρ : Spearman’s rank correlation coefficient; p: level of significance [1].

	Platelet CXCR4 MFI			Platelet CXCR7 MFI		
	Correlation	n	p	Correlation	n	p
Platelet TMRE MFI resting	r = -0.052	44	0.738	r = -0.216	45	0.154
Platelet Annexin V MFI resting	ρ = 0.121	60	0.121	ρ = 0.083	62	0.521
Ratio platelet TMRE MFI resting/thrombin stimulated	ρ = 0.365	41	0.019	ρ = 0.417	42	0.006
Ratio platelet Annexin V MFI resting/thrombin stimulated	ρ = -0.273	43	0.076	ρ = -0.401	45	0.006

Results

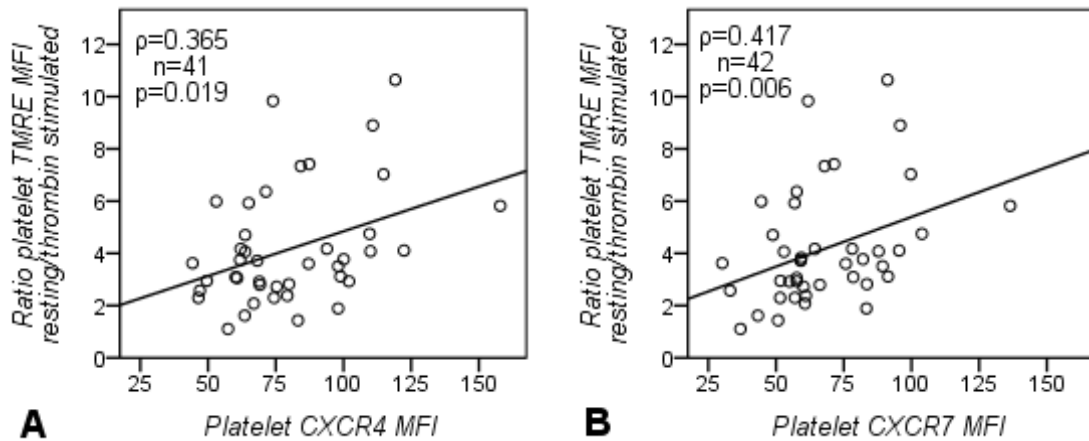


Figure 10: Association of the ratio of platelet TMRE MFI resting/thrombin stimulated with chemokine receptors. ρ : Spearman's rank correlation coefficient; n : number of subjects; p : level of significance; MFI: mean fluorescence intensity. **A**: with platelet CXCR4 MFI. **B**: with platelet CXCR7 MFI [1].

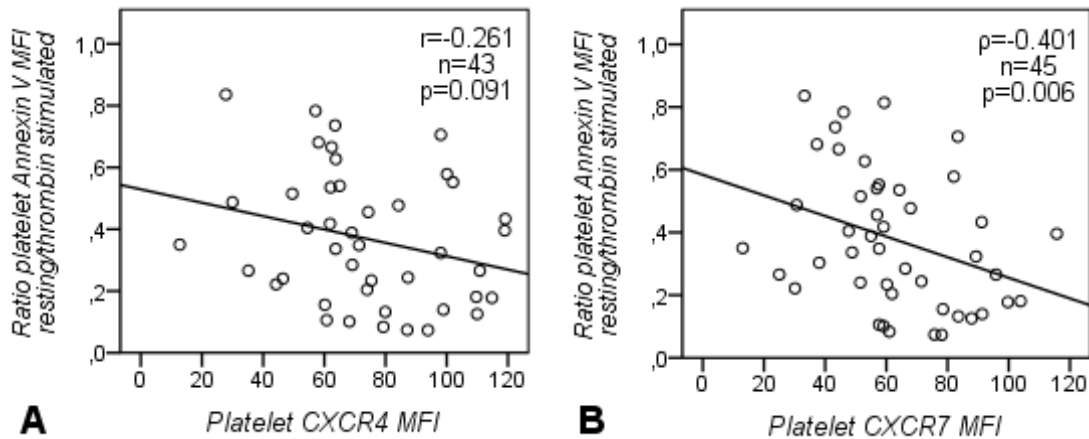


Figure 11: Association of the ratio of platelet Annexin V MFI resting/thrombin stimulated with chemokine receptors. r : Pearson's correlation coefficient; ρ : Spearman's rank correlation coefficient; n : number of subjects; p : level of significance; MFI: mean fluorescence intensity. **A**: with platelet CXCR4 MFI. **B**: with platelet CXCR7 MFI [1].

4.2.2 Association of LVEF and apoptosis

Bivariate analysis was conducted with the results of platelet apoptosis with the LVEF. It was tested whether there is an association between calculated apoptosis values on the one hand, and the LVEF at baseline or the ratio of the LVEF follow-up and the baseline on the other hand. The results are depicted in **Table 11**.

Results

Table 11: Bivariate correlation of parameters of apoptosis with the LVEF. MFI: mean fluorescence intensity; n: number of subjects; r: Pearson's correlation coefficient; ρ : Spearman's rank correlation coefficient; p: level of significance [1].

	LVEF baseline			Ratio LVEF follow-up/baseline		
	Correlation	n	p	Correlation	n	p
Platelet TMRE MFI resting	r = -0.228	40	0.158	r = 0.006	28	0.976
Platelet Annexin V MFI resting	ρ = -0.293	52	0.030	ρ = -0.367	32	0.039
Ratio platelet TMRE MFI resting/thrombin stimulated	ρ = -0.188	36	0.273	ρ = 0.490	25	0.013
Ratio platelet Annexin V MFI resting/thrombin stimulated	ρ = -0.243	39	0.137	ρ = -0.028	25	0.895

The ratio of platelet TMRE MFI between resting and thrombin-stimulated platelets indicated a positive correlation with the ratio between follow-up and baseline of the LVEF. The results are depicted in **Figure 12**. There was no correlation between the ratio of the LVEF and the ratio of platelet Annexin V MFI resting/thrombin [1].

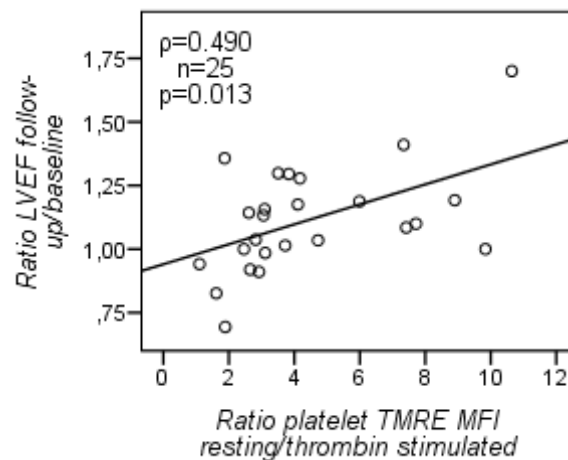


Figure 12: Association of the ratio of platelet TMRE MFI resting/thrombin stimulated with ratio of LVEF follow-up/baseline. ρ : Spearman's rank correlation coefficient; n: number of subjects; p: level of significance; MFI: mean fluorescence intensity [1].

Results

There was significant correlation between the Annexin V MFI of resting platelets, both with the LVEF (%) at baseline and with the ratio between the follow-up and baseline LVEF as well [1]. The results are depicted in **Figure 13**.

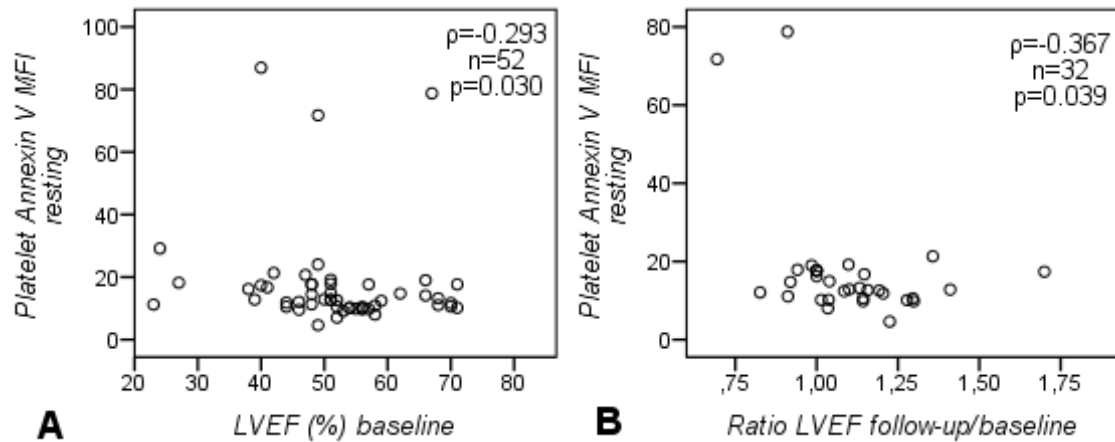


Figure 13: Association of platelet Annexin V MFI resting with LVEF. ρ : Spearman's rank correlation coefficient; n : number of subjects; p : level of significance; MFI: mean fluorescence intensity. **A:** Association with the LVEF at baseline. **B:** Association with the ratio between the follow-up and baseline of the LVEF [1].

5 Discussion

The major findings of this study are that platelets exhibit an expression of CXCR4 and CXCR7 after a STEMI. Chemokine receptors indicate no association with platelets, either dyed with TMRE or Annexin V [1]. However, the ratio between resting and thrombin-stimulated TMRE dyed platelets indicates a positive association with CXCR4 and CXCR7 [1]. Equally, the calculated ratio between resting and thrombin-stimulated Annexin V dyed platelets indicate a negative association with CXCR4 and CXCR7 [1].

Moreover, the ratio between resting and thrombin-stimulated TMRE dyed platelets exhibits a positive correlation with the six-month alteration of the LVEF, calculated as the ratio between the LVEF follow-up and baseline [1]. Pure resting platelets dyed with Annexin V or TMRE indicate no association to the LVEF or its recovery [1].

5.1 Methods

5.1.1 *Patient collective*

During the period of this study, nearly every patient with the diagnosis of a STEMI was interrogated to participate in this study and gave approval. Concerning baseline characteristics, we could not find any structural differences between dropouts and participants. Since dropouts only happened by coincidence, and abandonment or continuation revealed no regularity, this patient collective could be an approximate population of STEMI patients. Despite the small number of patients, the results of this study might have a high level of validity. A similar

approach concerning patient collective with a high quality was adopted elsewhere [285].

5.1.2 Evaluation of cardiac function using cardiac MRI

The LVEF is an established indicator of cardiac function after a STEMI and has a high predictive value concerning prognosis [129, 130]. Magnetic resonance imaging as a tool to evaluate cardiac function has one great disadvantage, which is poor approval by the subjects. Rejection during the course of this study caused a high percentage of dropouts. Conducting evaluation with transthoracic echocardiography is much easier, less expensive and has a higher rate of approval by the subjects. However, the quality of the results obtained in this manner is highly dependent on the expertise of the examiner. By comparison, imaging of the systolic function with cardiac MRI is of a very high quality and is accepted as the gold standard [286]. In the overall evaluation of the LVEF, cardiac MRI provides the best image quality and is the preferred technique [287]. In addition, it generates more supplementary information. For example, MRI gives exact information about the infarction size. As a consequence, a decision was made in favour of high-quality results instead of a larger sample size, and therefore cardiac MRI was chosen for the LVEF evaluation.

As an imaging technique, steady-state free precession imaging was used. This is commonly regarded as the superior technique for the generation of pictures with the highest quality [288, 289]. A similar approach has already been adopted successfully in other studies [166].

5.1.3 FACS analysis

With regard to flow cytometry, a well-established technique for the analysis of samples was used. In the laboratories of the University of Tübingen, Department of Cardiology (Department of Internal Medicine III), FACS is used frequently and there is a high level of expertise. For every single measurement, the same pattern

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of settings was used. The FACS machine was cleaned properly after every use. In cases of contamination or failed dye, the measurement was repeated.

To summarize, the experimental setup concerning the FACS analysis guarantees a very high quality of data with high explanatory power and a low susceptibility to errors.

5.1.4 Agents

5.1.4.1 CXCR4, CXCR7 and CD42b

At the University of Tübingen, Department of Cardiology (Department of Internal Medicine III), usage of these agents to detect chemokine receptors on the platelet surface is well-established [167, 290]. CD42b, which is also called GPIb, is part of the GPIb/IX complex and thus part of the receptor for von Willebrand factor [291]. Therefore, it might be an ideal target to detect platelets. The FACS calibration to obtain these results was similar to that of other studies [167].

5.1.4.2 Thrombin

The use of thrombin for the stimulation of platelets exhibits one disadvantage. Thrombin is commonly known as a potent platelet activator, called activation factor IIa [292, 293]. However, recent studies have revealed that thrombin is also capable of inducing platelet apoptosis [292, 294]. It can be detected both in the dropping of $\Delta\Psi_m$ and the exposure of PS [294]. Causing platelet activation or platelet apoptosis it is dependent on the concentration of thrombin. A low thrombin concentration leads to the activation of almost all platelets, whereas only a small percentage undergoes apoptosis [292, 293]. A high thrombin concentration leads to a high degree of apoptosis [292, 293]. To avoid this problem, in this study a high dose of thrombin was chosen. The purpose of the usage of thrombin was to calculate the ratio between unstimulated platelets and platelets under maximal stimulation.

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5.1.4.3 TMRE

The lipophilic cation tetramethylrhodamine ethyl ester (TMRE) exhibits the maximum fluorescence as long as it can accumulate in mitochondria with intact membrane potential ($\Delta\Psi_m$) [295]. During apoptosis and the subsequent dropping of $\Delta\Psi_m$, TMRE externalizes and disperses in cytosol, thus reducing its fluorescence [295]. With the extent of fluorescence measured in the FACS analysis, the extent of the survival of apoptotic cells can be approximated, but this needs careful interpretation [296].

The TMRE MFI is not a direct value for the number of intact mitochondria. The more negative the mitochondrial $\Delta\Psi_m$ is, the more TMRE will accumulate and will be detected [296]. This value is dependent of the individual conditions of the subject's platelets and mitochondria. For this reason, it is recommended to conduct a complementary approach, which can be set in relation to the TMRE dye [296]. For this purpose, thrombin as a platelet-stimulating agent was chosen. According to former studies, it can be assumed that with the used dose of thrombin nearly all platelets undergo apoptosis [292].

The results were related to each other. The TMRE MFI of resting platelets represents the actual survival of the dye. This was divided by the MFI of TMRE dyed and thrombin-stimulated platelets, which represents the remaining fluorescence after maximum apoptosis and thus minimum possible survival. Consequently, a high ratio could represent an elevated relative survival of platelets, whereas a low ratio could represent a reduced relative survival of platelets.

5.1.4.4 Annexin V

Annexin V is well established in measuring apoptotic cells. In the presence of a physiological calcium concentration, it binds to expressed phosphatidyl serine (PS) [297]. Phosphatidyl serine is naturally located at the inner leaflet of the cell membrane [182, 276]. The process of flipping from the inner cell membrane to the outer membrane is called externalization, which is dependent on the release

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of cytochrome c [182, 201, 276, 298, 299]. Cytochrome c release only occurs in apoptotic cells and is regarded as a general feature of this process [300]. After externalization, PS serves as a target for macrophages and the subsequent removal of apoptotic cells [183, 184, 298]. Consequently, a high MFI of Annexin V can be interpreted as a high level of platelets undergoing apoptosis.

This mechanism is similar in mammalian cell types [297]. Annexin V is a good and established tool to detect apoptotic cells in general. However, in the specific case of platelets, cytochrome c release is not the only mechanism leading to externalization of PS [8]. In platelets, PS is also exposed after activation, serving as a target for factors VIIIa, IXa and Ca^{2+} to form the tenase complex [8, 301]. The formation of the tenase complex leads to increased procoagulant activity of the platelets, which is an important step in platelet activation [298, 302]. This molecular overlap between activation and apoptosis is currently regarded as a process of activation-driven enhanced platelet senescence, with the purpose of the removal of thrombotic plaques after tissue healing [280].

One particular difference in the mechanisms of PS exposure is their dependence on Ca^{2+} [8, 276]. Apoptosis-driven PS exposure is dependent on the attendance of BAX or BAK and unaffected by the absence of Ca^{2+} , whereas activation-driven PS exposure needs extracellular Ca^{2+} and can be inhibited by Ca^{2+} -absorbing chelating agents [8, 259, 276]. In any case, with the used PBS- Ca^{2+} for dilution, the environment is already enriched with Ca^{2+} . Since Annexin V requires extracellular Ca^{2+} to detect PS, a calcium-rich environment is inevitable [297, 303]. In addition, Ca^{2+} is released by the platelets themselves [55]. For this reason, with PS exposure a definitive assertion whether the platelet is undergoing apoptosis or activation might be difficult [8]. Subsequently, a parallel detection of apoptosis and platelet activation is extremely likely, and an assessment of the measured MFI of Annexin V dyed platelets as apoptosis might not be possible. Rather, the results could represent a mixture of activation and apoptosis.

To adopt an approach comparable with TMRE evaluation, MFI of pure resting platelets dyed with Annexin V was set into relation with the Annexin V MFI of

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thrombin-stimulated platelets. Following from the premise of an appropriate dose of thrombin, which leads to maximal apoptosis, the result of this ratio can roughly be interpreted as a percentage of the maximum apoptosis. Of course, it might be difficult to distinguish between Annexin V detected platelet activation and apoptosis, but at least in the thrombin-stimulated samples it can be expected that nearly all platelets are undergoing apoptosis. Thus, the ratio might represent the proportion of activated and apoptotic platelets to the maximum of the apoptotic platelets.

5.2 Results

5.2.1 Platelet apoptosis with chemokines

In this study, it has been demonstrated that TMRE dyed resting platelets indicate no correlation with CXCR4 or CXCR7 [1]. This is in accordance with the knowledge that the pure value of TMRE gives no hint of the actual survival of platelets, and that a reference value is needed [296]. Annexin V, which was chosen to measure platelet apoptosis, and thus to have an opposite approach to platelet survival, did not reveal any correlation with CXCR4 or CXCR7 [1]. Due to the presentation of PS in platelet activation and apoptosis as well, this result is not surprising [8, 183, 184, 298, 300, 301].

After having set the values into relation, significant results can be detected. It could be illustrated that the calculated TMRE ratio has a positive correlation with the expression of CXCR4 ($\rho = 0.365$, $n = 41$, $p = 0.019$) and CXCR7 ($\rho = 0.417$, $n = 42$, $p = 0.006$), respectively [1]. Inversely, the Annexin V ratio exhibits a negative association with these chemokine receptors ($\rho = -0.273$, $n = 43$, $p = 0.076$ for CXCR4 and $\rho = -0.401$, $n = 45$, $p = 0.006$ for CXCR7, respectively) [1]. The TMRE ratio, which represents platelet survival, has a positive correlation with chemokine receptors. The Annexin V ratio partially represents platelet apoptosis and has a negative correlation with chemokine receptors. These contrary results

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are expected and can be interpreted as indicating the same result revealed with different tools [8, 292, 296, 297].

Here, platelet expression of CXCR4 and CXCR7 has a strong positive correlation with platelet survival (and, as a reverse argument, a strong negative correlation with platelet apoptosis) [1]. As a conclusion, platelet survival might be associated with platelet surface expression of CXCR4 and CXCR7 after a STEMI.

Both CXCR4 and CXCR7 are expressed permanently in every platelet, whilst the expression of CXCR4 is slightly higher [160]. After platelet activation and SDF-1 release, SDF-1-CXCR4 interaction leads to the homing of bone-marrow-derived cells to the place of the infarction, with the purpose of regeneration and further chemotactic recruitment [158, 163]. Upon SDF-1-CXCR4 interaction, expression of CXCR4 is decreased and expression of CXCR7 is increased [160]. The precise function of CXCR7 has not been revealed yet, but it could be demonstrated that the expression of CXCR7 is increased in mesenchymal stem cells under hypoxic conditions [160-162, 304]. It is also known that activated platelets have an influence on the regulation of apoptosis and the survival in surrounding cells via paracrine mechanisms [6, 305]. For example, SDF-1-driven CXCR4 suppresses apoptosis in bone marrow stromal cells [305, 306]. Subsequently, SDF-1 has been detected as a promoter of platelet survival [281, 290]. It could be demonstrated that the interaction of CXCR7 with the macrophage migration inhibitory factor, which is a chemokine-like ligand and released after platelet activation, increases the survival of platelets [280, 307, 308]. To sum up, with these results, it might be assumed that the influence on apoptotic processes is not only limited to surrounding cells, but acts on platelets too. At least the measured correlation provides evidence to suggest that there might be a crosstalk between chemokine expression and platelet survival. This assumption could indicate an approach for further studies.

The TMRE ratio has a slightly higher correlation coefficient with CXCR7 than with CXCR4 [1]. Similarly, the correlation coefficient of the Annexin V ratio with

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CXCR7 is considerable higher than with CXCR4 [1]. The difference in the correlation coefficients between CXCR4 and CXCR7 could be in accordance with a result from a former study conducted on renal progenitor cells [304]. This study revealed that not the expression of CXCR4, but rather the expression of CXCR7 is responsible for the survival of cells after SDF-1 induction [304]. In addition, it could be demonstrated that patients with ACS, compared with healthy subjects, exhibit increased platelet levels of CXCR7, but not of CXCR4 [160, 162]. These studies provide no evidence as to whether patients with ACS do exhibit higher levels of platelet survival as well [160, 162]. In this particular study, no assertion regarding this effect is possible, but it might be an interesting avenue for further studies.

It is also probable that the difference in correlation coefficients could have no meaning. The difference between the correlation coefficients of CXCR4 and CXCR7 each with the TMRE ratio is only slight [1]. Since both of these exhibit a strong positive correlation with either chemokine receptors, there is strong evidence that there is an association. To determine a possible difference, the sample size is too small. In further studies, a greater patient collective should be evaluated concerning this feature.

A considerable difference only occurs in the Annexin V ratio. This could be caused by the overlap of Annexin V as an indicator both for apoptosis and platelet activation. Perhaps in future studies, using a more appropriate test construction, this problem could be solved. In the meantime, the conclusion is that Annexin V is not an ideal marker to detect apoptosis in platelets.

5.2.2 Association of LVEF and apoptosis

The TMRE dyed platelets under resting conditions do not indicate any correlation either with the baseline LVEF or with the ratio between the follow-up and baseline LVEF [1]. These results are conclusive, according to the depicted limitations of TMRE under resting conditions [296]. Conversely, the Annexin V dyed platelets

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under resting conditions indicated a significant correlation coefficient, each with the baseline LVEF ($\rho = -0.293$, $n = 52$, $p = 0.030$), and the LVEF ratio between the follow-up and the baseline ($\rho = -0.367$, $n = 32$, $p = 0.039$) [1]. However, the graphic examination of these results in **Figure 13** reveals that there is actually only a very weak correlation [1]. Thus, there might not be any association between the Annexin V dyed platelets under resting conditions, neither with the baseline LVEF nor with the ratio of the LVEF. These results are in accordance with the limitations of Annexin V under resting conditions as well [8, 183, 184, 298, 300, 301]. With $\rho = 0.490$, $n = 25$, $p = 0.013$, the TMRE ratio indicates a strong and significant correlation with the LVEF ratio between the follow-up and the baseline, whereas the ratio of Annexin V exhibits no correlation [1]. Annexin V has the same limitations as depicted above [8, 183, 184, 298, 300, 301]. The findings on TMRE suggest that platelet survival might be associated with left ventricular functional recovery after a STEMI [1].

It has previously been demonstrated that interaction with CXCR4, driven by SDF-1, leads to the chemotactic recruitment of bone-marrow-derived cells and other cells with the purpose of regeneration [6, 158, 163]. A study in ACS subjects has revealed that elevated levels of platelet CXCR7 are associated with an amelioration of the LVEF [162]. A study in patients with CAD has revealed that patients with low platelet CXCR4 at baseline demonstrate poor overall mortality [167]. In addition, it has been demonstrated that the interaction of SDF-1 with CXCR4 and CXCR7 promotes platelet survival [281, 290]. Since the results from this current study suggest a crosstalk between platelet survival and the expression either of CXCR4 or CXCR7, the recognition that elevated survival levels are associated with a good prognosis might be a conclusive deduction.

6 Zusammenfassung

Hintergrund: Eine hohe Expression der thrombozytärer Chemokinrezeptoren CXCR4 und CXCR7 ist bei Patienten mit Koronarer Herzkrankheit mit gesteigerter myokardialer Reparatur und reduzierter Langzeitmortalität verbunden. Gleichzeitig verlängert sich das Überleben der normalerweise nach Aktivierung in Apoptose übergehenden Thrombozyten nach Stimulierung thrombozytärer CXCR7-Rezeptoren. Gegenstand dieser Studie ist die Frage, ob in einem Kollektiv aus STEMI-Patienten das thrombozytäre Überleben auch mit der Expression der thrombozytären Chemokin-Rezeptoren CXCR4 und CXCR7 assoziiert ist und ob eine Beziehung zwischen thrombozytärem Überleben mit der funktionalen Erholung der LV-Funktion besteht.

Methoden: Die Studie wurde an einem Kollektiv von 87 Patienten mit der akuten Diagnose eines ST-Hebungsinfarktes (STEMI) direkt nach erfolgter perkutaner Koronarintervention durchgeführt. Die Messung der linksventrikulären Ejektionsfraktion (LVEF) wurde mittels MRT während des Krankenhausaufenthalts und 6 Monate später durchgeführt. Mittels Durchflusszytometrie wurde die mittlere Fluoreszenz-Intensität (MFI) der thrombozytären Oberflächenrezeptoren CXCR4 und CXCR7 bestimmt. Außerdem erfolgte mit dieser Technik die Messung der thrombozytären Apoptose mittels Annexin V, sowie des thrombozytären Überlebens mittels TMRE zum einen nach erfolgter Stimulation mittels Thrombin zum anderen ohne Thrombin.

Ergebnisse: Ohne Stimulierung mittels Thrombin zeigt sich kein Zusammenhang zwischen der thrombozytären Apoptose mit der Expression der Chemokinrezeptoren oder der funktionalen Erholung der LV-Funktion.

Zusammenfassung

Das Verhältnis der MFI von Thrombin-stimulierten zu unstimulierten Annexin V gefärbten Thrombozyten korreliert negativ mit der Expression von CXCR4 ($\rho = -0.273$, $n = 43$, $p = 0.076$) und CXCR7 ($\rho = -0.401$, $n = 45$, $p = 0.006$). Analog reagiert das entsprechende Verhältnis TMRE gefärbter Thrombozyten positiv mit CXCR4 ($\rho = 0.365$, $n = 41$, $p = 0.019$) und CXCR7 ($\rho = 0.417$, $n = 42$, $p = 0.006$). Außerdem besteht eine positive Korrelation zwischen dem Verhältnis aus stimulierten und unstimulierten TMRE gefärbten Thrombozyten mit der Verbesserung der LVEF von der initialen Messung zur LVEF nach 6 Monaten ($\rho = 0.490$, $n = 25$, $p = 0.013$).

Schlussfolgerung: Mit dieser Studie konnte gezeigt werden, dass eine hohe thrombozytäre Expression sowohl von CXCR4 als auch CXCR7 mit einem gesteigerten Überleben der Thrombozyten nach STEMI einhergehen. Außerdem konnte gezeigt werden, dass ein gesteigertes thrombozytäres Überleben unmittelbar nach STEMI mit einer gesteigerten Erholung der linksventrikulären Funktion nach 6 Monaten einhergeht. Der Zusammenhang mit der Expression von CXCR4 und CXCR7 kann dabei einen Einfluss haben, eine andere Ursache ist mit dieser Studie jedoch nicht auszuschließen.

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8 Erklärung zum Eigenanteil

Diese Dissertationsschrift wurde in der Medizinischen Universitätsklinik und Poliklinik Abteilung Innere Medizin III unter der Betreuung von Herrn Prof. Dr. med. Tobias Geisler verfasst. Die Konzeption der Studie erfolgte in Zusammenarbeit mit Herrn Prof. Dr. med. Tobias Geisler (als Leiter der Forschungsgruppe), Herrn Dr. med. Dominik Rath, Frau Dr. rer. nat. Madhumita Chatterjee und Herrn Dr. med. Patrick Krumm.

Die Erhebung der Labordaten (exemplarisch dargestellt in **Figure 7 – Figure 9**) wurde nach Einarbeitung durch die Labormitarbeiterinnen Frau Dr. rer. nat. Madhumita Chatterjee und Frau Lydia Laptev in gemeinsamer Arbeit von Frau Nina Tekath und mir eigenständig durchgeführt.

Die Auswertung der MRT-Befunde (dargestellt in **Figure 6**) wurde in Kooperation und nach Einarbeitung durch Herrn Dr. med. Patrick Krumm aus der Abteilung für Radiologie am Universitätsklinikum Tübingen von mir eigenständig durchgeführt.

Die statistische Auswertung (inklusive der Darstellungen in **Figure 10 – Figure 13** und **Table 7 – Table 11**) wurde nach Einarbeitung von Herrn Dr. med. Dominik Rath und nach Beratung durch Frau Aline Naumann vom Institut für Biometrie durch mich eigenständig durchgeführt.

Die Ergebnisse dieser Dissertationsschrift waren Teil einer Publikation unter Leitung von Herrn Dr. med. Dominik Rath, welche in Zusammenarbeit mit Frau Dr. rer. nat. Madhumita Chatterjee, Frau Nina Tekath, Herrn Dr. med. Patrick Krumm, Herrn Dr. med. Constatnin Adams, Herrn Dr. med. Oliver Borst, Frau Dr. med. Karin Müller, Herrn Dr. med. Michael Droppa, Herrn Dr. med. Konstantin Nikolaou, Herrn Dr. med. Joachim Riethmüller, Herrn Prof. Dr. med. Meinrad Gawaz, Herrn Prof. Dr. med. Tobias Geisler und mir verfasst wurde. Die

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Die in dieser Publikation angefertigten Grafiken und Statistiken wurden von mir erstellt und mit dem Einverständnis des Erstautors und des korrespondierenden Autors für diese Dissertation verwendet.

Ich versichere hiermit, die Dissertationsschrift selbständig verfasst und keine weiteren als die von mir angegebenen Quellen verwendet zu haben.

Tübingen, den 25. März 2018

Christian Lennart Meyer

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