



**Impact of polyunsaturated omega-6 and omega-3  
fatty acids on cardiomyogenesis and vasculogenesis  
of mouse embryonic stem cells**

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in partial fulfillment of the doctoral degree in human biology at  
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## Dedication

I am writing my thesis and would like to dedicate it to my parents. However, they passed away while I was studying. Also, I would like to dedicate it to:

my sons Ali, Majd, and Yaman.

my lovely wife Rasha.

*Amer*

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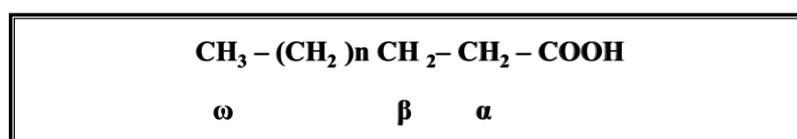
### 1. Introduction

#### 1.1 Essential fatty acids

The term essential fatty acids (EFAs) refers to those polyunsaturated fatty acids (PUFAs) that have the necessity to be provided by food because they cannot be synthesized in the human body. PUFAs are important for health and may contribute significant advantages in addition to fundamental nutrition. There are two major families of EFAs, omega-3 ( $\omega$ -3) (i.e.  $\alpha$ -linolenic acid (ALA)) and omega-6 ( $\omega$ -6) (i.e. linoleic acid (LA)). These EFAs produce very long-chain PUFAs (more than 22 C-atoms) during cascade of elongation reaction (Das, 2006; Kaur et al., 2014). In the late 1920s, they were first reported as essential in the diet (Shireman, 2003).  $\Omega$ -3 and  $\omega$ -6 PUFAs are beneficial elements of oils with specific dietary and functional characteristics. Their therapeutic and health-promoting impacts have already been discovered in several chronic inflammatory and autoimmune diseases through various mechanisms, including modifications in cell membrane lipid composition, gene expression, cellular metabolism, and signal transduction (Balić et al., 2020).

#### 1.2 Fatty acid structure and classification

Fatty acids consist basically of chainlike structure with a methyl group at one end of the molecule (designated omega,  $\omega$ ) and a carboxyl group at the other. The second carbon atom located after the carboxyl group is called  $\alpha$ -carbon atom, and the next one called  $\beta$ -carbon atom. According to the  $\omega$ -x notation, the letter  $n$  indicates the position of the double bond nearest to the methyl end ( $\omega$ ) in carbon chains (figure 1.1). The nomenclature of fatty acids may also depend on the location of double bonds with a citation to the carboxyl group ( $\Delta$ ) (figure 1.2) (Rustan & Drevon, 2001).



**Figure 1.1: The basic structure of fatty acids.** The fatty acid chain has two different ends, (COOH) refers to the carboxyl end, and the other end is a methyl group (CH<sub>3</sub>) named by the last letter of the Greek alphabet ( $\omega$ ). The letter ( $n$ ) indicates the location of the

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double bond nearest to the  $\omega$  methyl group. Figure adapted with permission from (Rustan & Drevon, 2001); copyright 2001 John Wiley and Sons.

Fatty acids present in two main groups: Saturated fatty acids, in which the carbon atoms are bound by hydrogen atoms. Most saturated fatty acids are straight hydrocarbon chain molecules, i.e. Palmitic acid (PA;16:0) and Stearic acid (18:0). PA is the most well-known saturated fatty acid estimating for 20–30% of total fatty acids in the human body, and can be obtained in the diet or synthesized endogenously through de novo lipogenesis. (Carta et al., 2017; Rustan & Drevon, 2001). The other group comprises of unsaturated fatty acids such as monounsaturated fatty acids (MUFAs), which have one carbon-carbon double bond. Their carbon-carbon double bond can occur in different positions. The most common monoenes have a double bond in cis configuration, which means that the hydrogen atoms on the same side of the double bond are oriented in the same direction. Cis configuration in the structure gives it a “kink” to form a different secondary structure, which may change their physical properties (Jump et al., 2012). PUFAs have more than one carbon-carbon double bond. There are two families of PUFAs,  $\omega$ -3 and  $\omega$ -6.  $\Omega$ -3 fatty acids have in common a closing carbon-carbon double bond in the  $\omega$ -3 position, the third bond from the methyl end of the fatty acid, while  $\omega$ -6 fatty acids have it in the  $\omega$ -6 position i.e. the sixth bond from the methyl end of the fatty acid (figure 1.2) (Kaur et al., 2014).

$\omega$ -characteristics	Methyl end	Carboxyl end	Saturation	$\Delta$ -characteristics
PA 16:0	CH <sub>3</sub>	COOH	saturated	16:0
Stearic 18:0	CH <sub>3</sub>	COOH	saturated	18:0
Oleic 18:1, $\omega$ -9	CH <sub>3</sub>	COOH	monoene	18:1 $\Delta$ 9
LA 18:2, $\omega$ -6	CH <sub>3</sub>	COOH	polyene	18:2 $\Delta$ 9, 12
ALA 18:3, $\omega$ -3	CH <sub>3</sub>	COOH	polyene	18:3 $\Delta$ 9, 12, 15

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Eicosapentaenoic acid (EPA) 20:5, $\omega$ -3		polyene	20:5 $\Delta$ 5, 8, 11, 14, 17
Docosahexaenoic acid (DHA) 22:6, $\omega$ -3		polyene	22:6 $\Delta$ 4, 7, 10, 13, 16, 19

**Figure 1.2: Schematic overview of the structures of different unbranched fatty acids with a methyl and a carboxyl (acidic) end.** Figure adapted with permission from (Rustan & Drevon, 2001); copyright 2001 John Wiley and Sons; Panel PA adapted from (Shah & Cox, 2009) licensed under Creative Commons Attribution 2.0 Generic (CC BY 2.0).

### 1.3 Metabolism of PUFAs

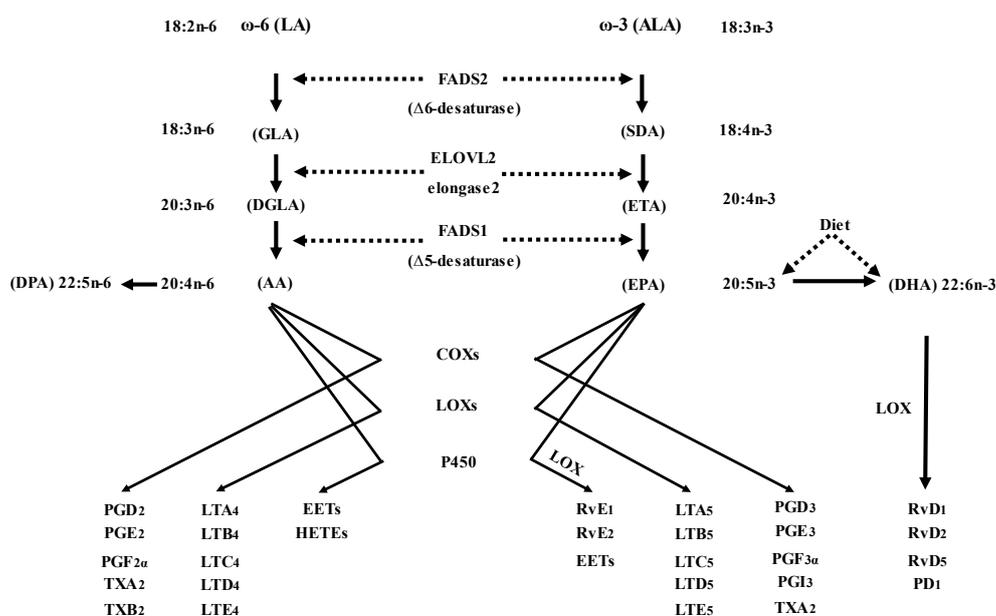
$\Omega$ -3 and  $\omega$ -6 PUFAs are provided from different food sources in triglyceride form. When consumed in the diet, triglycerides undergo digestion in the small intestine, which allows absorption, transport via blood and assimilation in the different organs of the body like the brain, heart, and other tissue. These fatty acids can be metabolized in different metabolic pathways:

- (1)  $\beta$ -oxidation to give energy in adenosine triphosphate (ATP) form.
- (2) Esterification into various cellular lipids as triglycerides, cholesterol ester and phospholipids.
- (3) Transformation into longer chain and more unsaturated products, which are formed by a series of desaturation and elongation reactions. This process takes place mainly in the liver and to a minor degree in other tissues (Kaur et al., 2014).

LA and ALA can be only obtained from dietary sources because the human body lacks the enzymes  $\Delta$ -12 and  $\Delta$ -15-desaturase, which are present in plants and are necessary for *de novo* synthesis of PUFAs (Kaur et al., 2014; Wallis et al., 2002). Hence, the LA and ALA undergo a series of desaturation and elongation reactions (Jump et al., 2012; Nakamura & Nara, 2004). Desaturation is the addition of a double bond to the fatty acid structure by the presence of fatty acid desaturase enzymes (FADS), i.e.,  $\Delta$ -5 desaturase (D5D) FADS1 ( $\Delta^5$ -desaturase) and  $\Delta$ -6 desaturase (D6D) FADS2 ( $\Delta^6$ -desaturase) which perform acyl chain desaturation of PUFAs. Elongation is a condensation reaction, which is catalyzed by elongase enzymes referred to as elongation of very long-chain fatty

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acids (ELOVLs) which add two carbon atoms. Elongation is required for the generation of very long-chain fatty acids (VLCFAs) (Denic & Weissman, 2007; Lee et al., 2019; Moon et al., 2001; Oh et al., 1997). There are three types of (ELOVLs): ELOVL2, ELOVL4, and ELOVL5 being selective for PUFAs (Agbaga et al., 2008; Kitazawa et al., 2009; Leonard et al., 2000; Moon et al., 2001; Tamura et al., 2009; Tvrđik et al., 2000; Wang et al., 2005). LA and ALA are digested by desaturase and elongase enzymes during metabolic reactions, and both serve as precursors to produce essential longer PUFAs such as  $\omega$ -6 gamma-linolenic acid (GLA), dihomo-gamma-linolenic acid (DGLA), arachidonic acid (AA) and docosapentaenoic acid (DPA) from LA, while  $\omega$ -3 stearidonic acid (SDA), eicosatetraenoic acid (ETA), and EPA from ALA, which play an essential roles in the physiological metabolism (figure 1.3) (Jump, 2002; Jump et al., 2012; Nakamura & Nara, 2004).



**Figure 1.3: Long chain unsaturated fatty acid synthesis from the EFAs such as LA and ALA in mammals.** Eicosanoids such as prostaglandins (PGs), leukotrienes (LTs), thromboxanes (TXs), resolvins (Rvs), epoxyeicosatrienoic acids (EETs) and 5-hydroxyeicosatetraenoic acids (HETEs) are synthesised from AA and EPA by lipoxygenases (LOXs), cyclooxygenases (COXs) and cytochrome P450 (CYP) enzymes. Furthermore, Rv, as well as protectins D1 (PD1), are synthesised from DHA. Figure adapted with permission from (Kang et al., 2014); copyright John Wiley and Sons.

### 1.4 Sources of PUFAs

#### 1.4.1 Sources of $\omega$ -3 as EPA

ALA is plentiful in flaxseed as well as in hemp, walnut, soybean, and canola oil (Hunter, 1990). It is mainly present in the chloroplast of green leafy vegetables. Furthermore, ALA can be converted during metabolism into other important PUFAs such as EPA. Fish and fish oil are major sources for EPA, the content of EPA in fresh and saltwater fish can be estimated to 39%-50%. Additionally, EPA is considered as a parent of series 3-eicosanoid hormones. DHA is also present in red-brown algae. The human brain consists of 65% fat, half of this is DHA (Kinsella, 1990).

#### 1.4.2 Sources of $\omega$ -6 as LA

LA is plentiful in safflower, sunflower, and corn, while medium amounts are present in soybean, sesame and almonds. Canola, peanut and olive oils contain LA in small amounts, and coconut and palm kernel contain only very low amounts. LA can be converted to other  $\omega$ -6 fatty acids depending on the body's requirements (Kaur et al., 2014). Evening primrose oil contains GLA, which is one of the products of LA metabolism, to about 7-10%, (Høy et al., 1983). Black currant and borage contain (15–20g/100g) and (18–26g/100g), respectively (Lawson & Hughes, 1988). Organ meat and breast milk contain a small quantity of it. AA is produced in more LA's metabolism. It is present in meat, eggs and dairy products. (Horrobin, 1990).

### 1.5 Stem cells

Stem cells are undifferentiated cells that can renew and give daughter cells by keeping their properties as stem cells or use the process of differentiation dependent on signals taken from their neighborhood. They have two main properties, first: they have the potential to retain their self-renewal infinitely, that means, keeping their undifferentiated state through infinite cycles of cell division, and second: differentiation potency, their potential to convert into different organo-typic cells (Friel et al., 2005).

### 1.5.1 Classification of stem cells based on their differentiation potency

Dependent on the differentiation potential of stem cells, they can be classified into different types. Therefore, the classes of stem cells regarding their plasticity are: Totipotent, pluripotent, multipotent, oligopotent and unipotent (figure 1.4) (Majo et al., 2008; Niwa, 2007; Rajabzadeh et al., 2019).

**Totipotent stem cells:** these stem cells are capable of developing into a complete organism or in other words, these cells can differentiate into any cell or tissue (Condic, 2013). Totipotent are cells that arise from zygotes (a one-cell embryo) and exist until the morula stage of embryogenesis (early stage of the blastocyst). A totipotent cell is also a one-cell embryo; that is, a cell that is capable of generating a globally coordinated developmental sequence (Condic, 2013; Niwa, 2007). Somatic stem/progenitor cells and primitive germ-line stem cells originate from totipotent stem cells (Weissman, 2000; Zakrzewski et al., 2019).

**Pluripotent stem cells:** Pluripotent stem cells have the potential to generate many but not all types of cells, because some of their differentiation potency has been lost. They arise from totipotent stem cells and exist in the inner cells mass (ICM) (Niwa, 2007). Pluripotent stem cells can differentiate into approximately all cells, i.e., cells derived from any of the three germ layers (ectoderm, mesoderm and endoderm), and these pluripotent cells are considered true stem cells because they own self-renewal and differentiation potential for all cell types of the adult organism (Baker & Pera, 2018; Horie et al., 2011). For example, embryonic stem cells (ES cells) have been listed within this classification. Therefore, human pluripotent stem cells are an invaluable tool for *in vitro* studies of human embryogenesis (Shamblott et al., 1998).

**Multipotent stem cells:** Multipotent stem cells are characterized by the capability of self-renewal, and differentiate into various types of cells. They can only differentiate into closely related cells in a specific tissue, organ or system (Niwa, 2007). Furthermore, multipotent stem cells can only differentiate into a limited number of cell types, for example, hematopoietic stem cells. Moreover, multipotent stem cell produce multiple

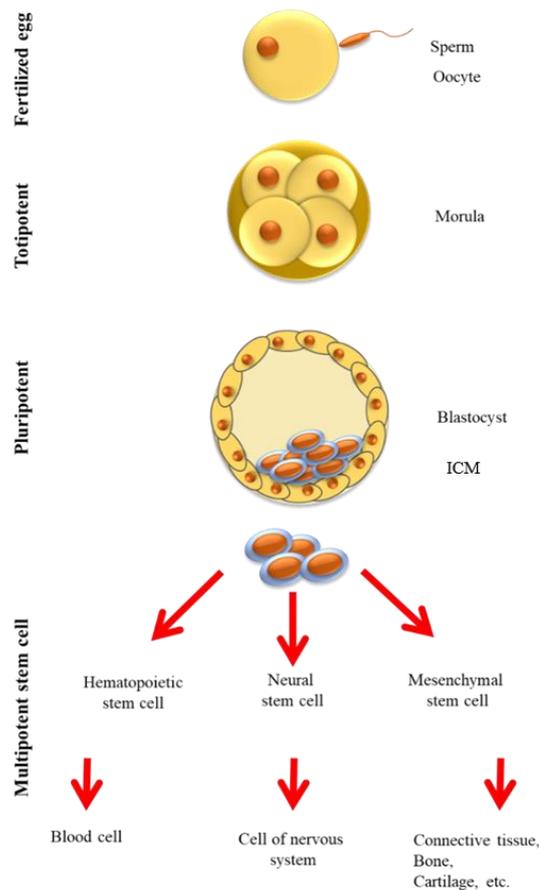
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blood cell progenitors (Quintana & Grosveld, 2011). Additionally, adipose tissue is considered as one of the sources for multipotent stem cells (Zuk et al., 2002).

**Oligopotent stem cells:** Oligopotent stem cells can differentiate into only very few cell types, including stem cells residing on the mammalian ocular surface (Majo et al., 2008).

**Unipotent stem cells:** Unipotent stem cells are capable to generate cells with only one lineage differentiation, example is: muscle stem cells (Kalra & Tomar, 2014). They have self-renewal property as well. Most epithelial tissues self-renew throughout adult life due to the presence of unipotent progenitor cells (Blanpain et al., 2007).

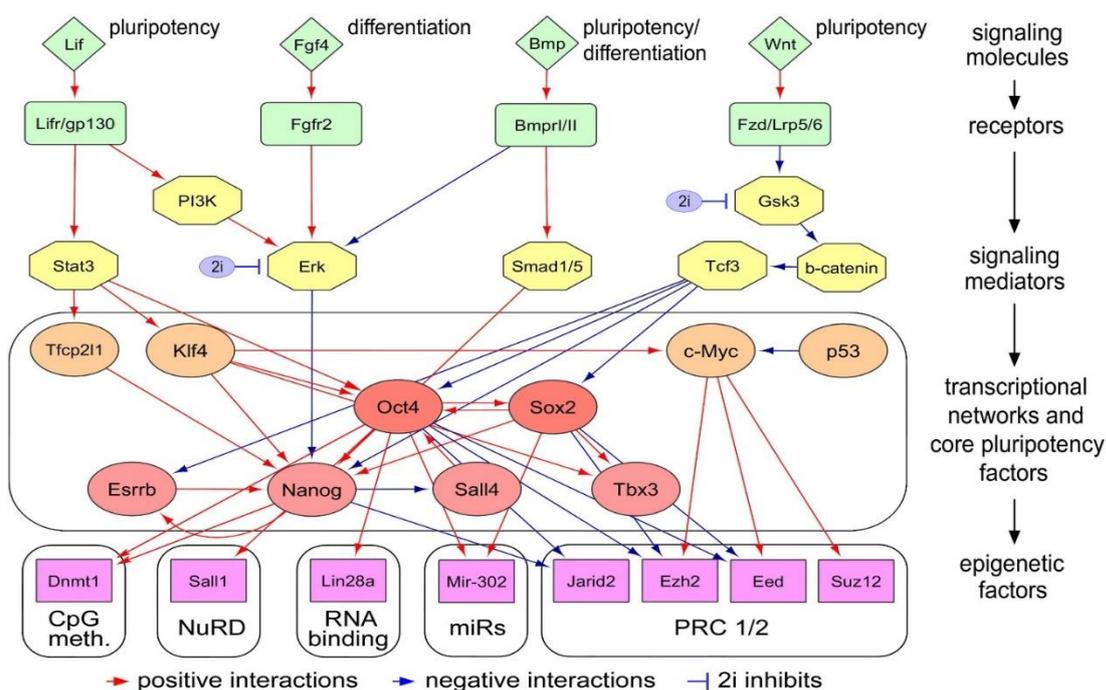


**Figure 1.4: Diagram overview of stem cells types regarding their potency.** This figure was modified from (Rajabzadeh et al., 2019).

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### 1.5.2 ES cells

ES cells are pluripotent stem cells, which have a high capability to undergo unlimited proliferation and self-renewal *in vitro* for an extended time period, and differentiate into all cell types in the body. Thus, they have potential for the therapy of various degenerative diseases (Khan et al., 2018; Papatsenko et al., 2018). Their pluripotent states can be maintained *in vitro* by a network of signaling, transcriptional and epigenetic regulatory synergies known as the pluripotency gene regulatory network (PGRN) (Davidson et al., 2015; Papatsenko et al., 2018). In cell culture conditions, cytokines and other factors have been demonstrated to preserve stem cell pluripotency by activation of specific transcriptional and epigenetic regulators (Morey et al., 2015). PGRN has been described as a series of hierarchical levels, determined according to the propagation of biological signals in ES cells, which are initiated with cell signaling to transcription and epigenetic factors (figure 1.5) (Papatsenko et al., 2018).



**Figure 1.5: Hierarchical organization of the PGRN.** Signal pathways represent the first hierarchical level (in green). Signal pathway mediators (in yellow) represent the second hierarchical level. Components of the transcriptional pluripotency network (red) represent the following levels. Lowest levels of the pluripotency network (in violet) include components of epigenetic complexes involved in chromatin modification, DNA

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methylation and regulation at posttranscriptional levels, as indicated. Red and blue arrows show positive and negative interactions correspondingly; blue T-arrows show targets of inhibitors for “2i” conditions. Figure adapted from (Papatsenko et al., 2018) licensed under Creative Commons Attribution Non-Commercial No Derivatives (CC-BY-NC-ND).

ES cells are able to remain in pluripotency state by providing them with nutrients and cytokines, such as leukemia inhibitory factor (LIF) and respond to WNT, bone morphogenetic protein (BMP) and fibroblast growth factors (FGF). (Itoh et al., 2014; Lakatos et al., 2014; Malaguti et al., 2013). The first responses to extracellular signals are mediated by cellular receptors such as LIF, WNT, BMP, and FGF. Therefore, these molecules may be supposed as the top hierarchical level, which downstream transfers the input information within the PGRN (Papatsenko et al., 2018). The second hierarchical level is initiated by factors such as signal transducer and activator of transcription 3 (Stat3), Smads,  $\beta$ -catenin, and extracellular signal-regulated kinase (Erk) which are effectors of LIF, BMP, WNT and FGF signaling pathways, respectively (Itoh et al., 2014; Lakatos et al., 2014; Tai & Ying, 2013; Van Oosten et al., 2012). Signals from the effectors are forwarded to many downstream targets, including the core pluripotency network (CPN), which includes transcription factors such as octamer-binding transcription factor 4 (Oct4), Sex-determining region Y/Box 2 (Sox2), nanog homeobox protein (Nanog), estrogen-related receptor beta (Esrrb), T-box transcription factor 3 (Tbx3), c-Myc and others, which control inputs from signaling systems and perform decisions associated with self-renewal or differentiation (Boyer et al., 2005; Dunn et al., 2014; Xu et al., 2014). The lower position in the PGRN network is occupied by the epigenetic regulation. Moreover, the transcription events of pluripotency include changes in the epigenetic landscape, including chromatin remodeling and deoxyribonucleic acid (DNA) methylation (figure 1.5) (Kobayashi & Kikyo, 2015). Recently, evidence has been provided, that in addition to growth factors and extracellular matrix cues, many metabolic pathways definitively provide essential signals for the self-renewal and differentiation potential of stem cells (Hu et al., 2016). Historically, murine ES cells were isolated and established in 1981 (Evans & Kaufman, 1981). Later, in 1998, Thomson and his team were the first to derive human embryonic stem cells (hESCs) from human blastocysts (Thomson et al., 1998). After a few years, in 2006, Takahashi and Yamanaka demonstrated

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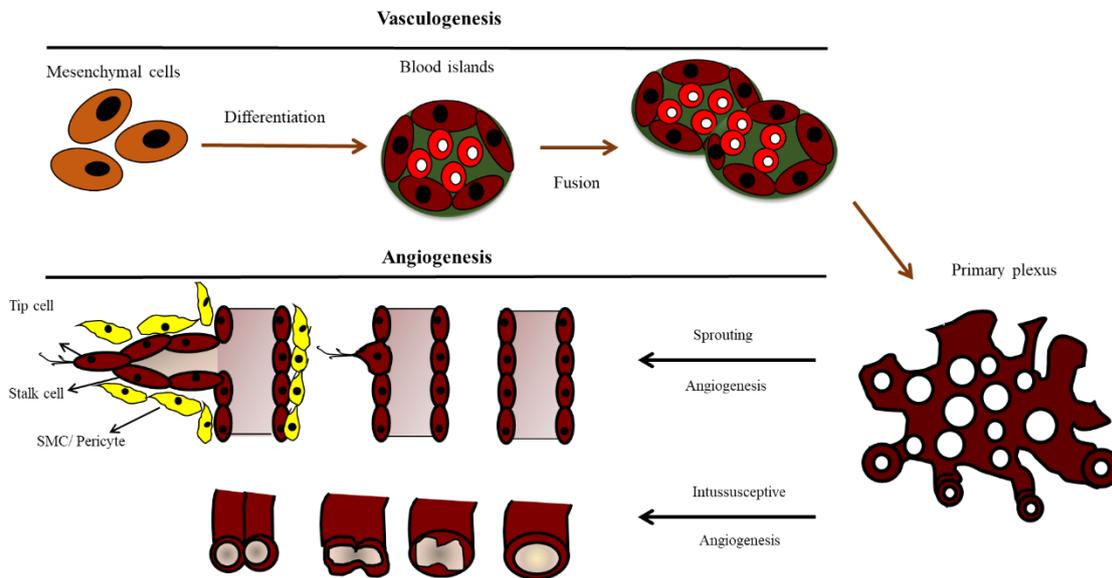
transdifferentiation of pluripotent stem cells from mouse embryonic or adult fibroblasts by introducing four factors, Oct3/4, Sox2, c-Myc, and kruppel-like factor 4 (Klf4), under ES cell culture states. These cells were called induced pluripotent stem cells (iPSCs) (Takahashi & Yamanaka, 2006).

### 1.6 Vasculogenesis

The creation of new blood vessels is a necessary step in the development of tissues. During embryogenesis and in *in vitro* models, the blood vessels provide the growing tissue with nutrients and oxygen. They remove waste products, transport macromolecules (e.g. hormones, growth factors and cytokines), exchange liquids between tissues and organs, and transport immune cells to sites of inflammation or infection. Other functions of blood vessels are the maintenance of steady body temperature and pH to keep homeostasis. (Adams & Alitalo, 2007; Aird, 2006; Jakobsson et al., 2007; Sweeney & Foldes, 2018). Vasculogenesis refers to *de novo* blood vessel formation from progenitor cells, while angiogenesis denotes migration, branching, and remodelling of pre-existing blood vessels to form complex vascular networks and capillary beds (figure 1.6) (Risau & Flamme, 1995). Endothelial cells are the basic unit of blood vessels, and are capable of creating a tube-like structure (Stratman et al., 2009; Sweeney & Foldes, 2018). Doetschman and his colleagues demonstrated in 1985, that endothelial cell development and subsequent vascular morphogenesis in differentiating ES cells cultures occurs in an *in vivo*-like fashion (Doetschman et al., 1985). During embryogenesis, blood vessel formation takes place in the yolk sac originating from groups of cells, which are called primitive endothelial cells and express endothelial markers such as vascular endothelial growth factor receptors (VEGFRs), vascular endothelial cadherin (VE-Cadherin), and cluster of differentiation 31 (CD31) (Platelet endothelial cell adhesion molecule (PECAM-1)) (Breier et al., 1996; Risau & Flamme, 1995). During the first steps of vasculogenesis primitive endothelial cells derive from the mesodermal layer of the embryo, migrate to form blood islands and differentiate into either hematopoietic cells in the central core or angioblastic cells (the precursors of endothelial cells) at the periphery of the blood islands (Gonzalez-Crussi, 1971). Angioblasts flatten and create intercellular connections to form a circumferential

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layer of primitive endothelial cells (figure 1.6) (Heinke et al., 2012; Risau & Flamme, 1995). Vascular development in differentiated ES cell cultures progresses through the same steps of vasculogenesis and angiogenesis as in the *in vivo* situation (Risau, 1997).



**Figure 1.6: Illustration of vasculogenesis and angiogenesis during blood vessel formation.** Vasculogenesis requires the differentiation of endothelial cells from angioblast cells to form a primitive vascular plexus. Mesodermal cells differentiate toward haemangioblasts, which more progress into angioblast. Developing processes of these pre-existing vessels is termed angiogenesis, which occurs in two manners: Intussusceptive angiogenesis and/or sprouting angiogenesis. Figure adapted with permission from (Heinke et al., 2012); copyright 2012 Frontiers in Bioscience.

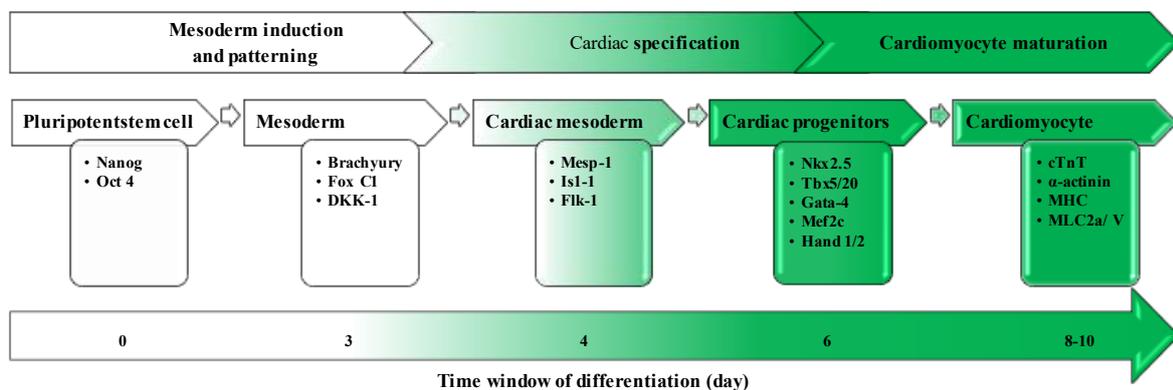
Angiogenesis occurs by two fundamental mechanisms: i) non-sprouting (intussusceptive) angiogenesis, which involves splitting of existing blood vessels by the creation of translumen pillars and growing of vessels. ii) Sprouting angiogenesis, where in growing branches of blood vessels, endothelial cells proliferate behind tip cells in response to cytokines, and lumens are formed by vacuole fusion. Angiogenesis requires the recruitment of smooth muscle cells (SMCs) or pericytes to fix the nascent vessels (Heinke et al., 2012; Risau, 1997).

### 1.7 Cardiomyogenesis

Cardiomyocytes arise from differentiation of pluripotent ES cells. These cells include all specialized cell types of the heart, such as atrial-like, ventricular-like and sinus nodal-like

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cells. The process of cardiomyogenic differentiation occurs in early stages of embryo development and is called cardiomyogenesis (Boheler et al., 2002). During the early stage of differentiation within a three-dimensional tissue of ES cells, named embryoid body (EB), the cardiomyocyte is characterized by single nucleus with a small and round shape which expresses on day 3 of differentiation the cardiac transcription factors Nkx2.5, myocyte enhancer factor 2C (MEF2C) and GATA4 (Wei et al., 2005). Growing myofibrils are sparse and irregularly organized. During further differentiation parallel bundles of myofibrils showing A and I bands are occurring (Westfall et al., 1997). During maturation cardiomyocytes become elongated with well-developed myofibrils and sarcomeres. Spontaneously contracting cardiomyocytes contain cell-cell junctions and are primarily mononucleated and rod-shaped (Westfall et al., 1997). In the final stage of differentiation, the bundles of myofibrils are densely packed and well-organized, and A, I bands and Z disks are organized into sarcomeres (Fassler et al., 1996; Westfall et al., 1997). In addition, cardiac-specific genes and proteins are expressed in parallel with ion channels (Boheler et al., 2002). Transcription factors such as zinc-finger GATA (GATA-4) protein and Nkx-2.5 regulate heart development together with several factors and signaling molecules (Sachinidis et al., 2003). Cardiac cell formation is organized by the sequential expression of multiple signal transduction proteins and transcription factors operating in a combinatorial manner. Numerous studies have reported that Wnts/Nodal, BMPs, and FGFs are implicated in the development of specialized cardiac subtypes (figure 1.7) (Marvin et al., 2001; Zhang & Bradley, 1996).



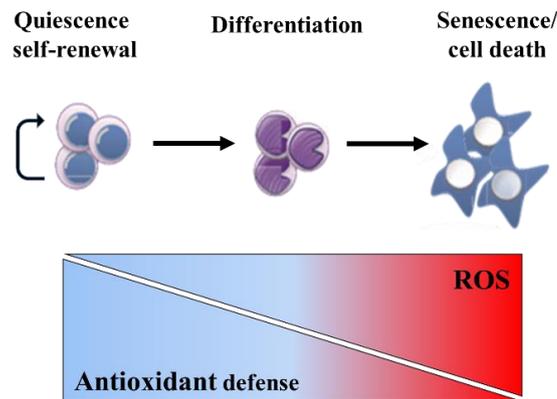
**Figure 1.7: Sequential steps within the differentiation time window of pluripotent ES cells to obtain matured cardiomyocytes.** The figure was modified from (Rajala et al., 2011; Willems et al., 2009).

### 1.8 Reactive oxygen species

Stem cell function during development and tissue homeostasis throughout life is regulated by fitting balance between self-renewal and differentiation. Recently, stem cell studies suggested that this balance is partly regulated by reactive oxygen species (ROS), which are reactive molecules that play significant roles in the regulation of several cell functions and biological processes (Bigarella et al., 2014; Chen et al., 2018). On the other hand, ROS play an important role in stem cell fate. Low concentration of ROS is required for stem cells to undergo self-renewal and proliferation. Increasing ROS generation reduces self-renewal of stem cells by enhancing stem cell proliferation and differentiation, and a further increase in ROS generation can cause stem cell senescence and apoptosis. Furthermore, excess ROS production is involved in development of pathological states and diseases by causing stem cell damage. Hence, stem cell self-renewal, proliferation, differentiation, senescence, and apoptosis are regulated dose-dependent by ROS (figure 1.8) (Chen et al., 2018; Ezashi et al., 2005; Juntilla et al., 2010; Kinder et al., 2010; Lewandowski et al., 2010; Owusu-Ansah & Banerjee, 2009; Sauer & Wartenberg, 2005; Wong et al., 2004; Zhou et al., 2014). Moreover, increased generation of ROS by various NOXs plays a critical role in cardiovascular differentiation of ES cells (Li et al., 2006; Sauer & Wartenberg, 2005; Schmelter et al., 2006). Peroxisome proliferator-activated receptor alpha (PPAR- $\alpha$ ) is a ligand for many fatty acids and plays a vital role in several physiological functions such as regulation of vascular tone, inflammation, and glucose homeostasis (Krey et al., 1997). Activation of PPAR- $\alpha$  is necessary for cardiomyogenesis of ES cells, which is associated with ROS-generation (Sharifpanah et al., 2008). Also, AA-induced vasculogenesis of ES cells is dependent on ROS generation (Huang et al., 2016). In summary, ROS play a very significant role in the fate decision of various stem cells in a concentration-dependent manner.

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**Figure 1.8: Effects of the intracellular ROS level on stem cell fate.** Figure adapted with permission from (Bigarella et al., 2014); copyright 2014 The Company of Biologists.

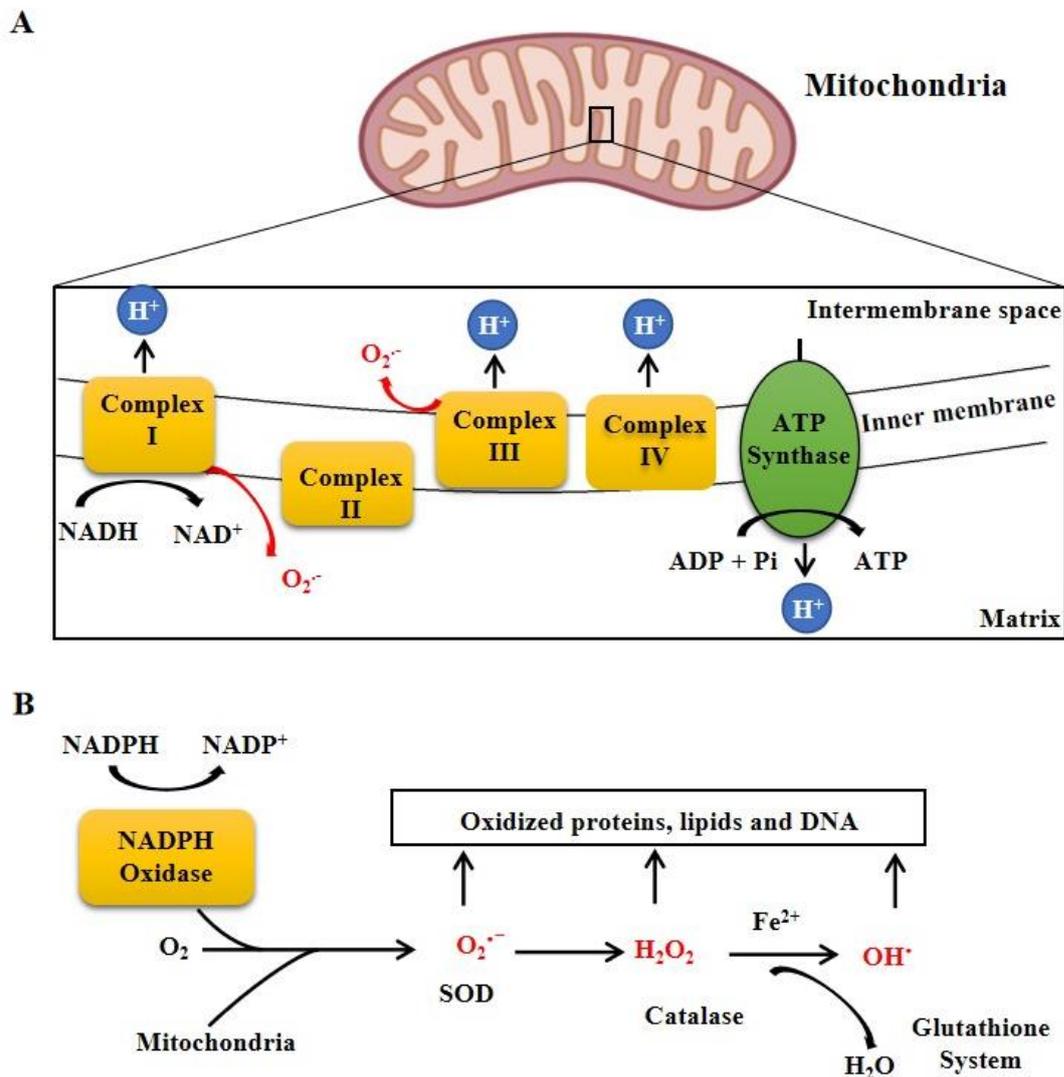
Chemically, ROS are small molecules, which refer to a number of reactive molecules and free radicals derived from molecular oxygen. Intracellular ROS comprise of three different forms: superoxide anions ( $O_2^{\cdot-}$ ), hydrogen peroxide ( $H_2O_2$ ) and hydroxyl radicals ( $OH\cdot$ ).  $H_2O_2$  is involved in many intracellular signal pathways and acts as a second messenger, integrating and delivering exogenous inducers to the downstream signal transduction cascade (Holmström & Finkel, 2014).  $H_2O_2$  forms through the rapid reduction of superoxide anion by the antioxidant enzyme superoxide dismutase (SOD).  $H_2O_2$  is further reduced to  $H_2O$  and  $O_2$  by catalase and various cellular antioxidants (figure 1.9 B).  $H_2O_2$  has a longer half-life and is able to produce other forms of ROS via diffusion through membranes (Holmström & Finkel, 2014).

### 1.9 Sources of ROS

Mitochondria have been considered as the main source of ROS in the cell. About ~0.1-0.2% of  $O_2$  consumed during mitochondrial respiration by electron transport chain (ETC) activity forms ROS through leak electron flow to  $O_2$ . The main role of the ETC is to produce the proton motive force for ATP production. This occurs through ATP synthase in a process known as oxidative phosphorylation. ROS are produced in the ETC complexes I and III (figure 1.9 A) (Tahara et al., 2009). The amount of ROS generation may differ greatly depending on the cell type, environment and the activity of mitochondria (Murphy, 2009). Another major source of ROS formation are membrane-bound nicotinamide

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adenine dinucleotide phosphate (NADPH) oxidases (NOXs). They are enzymes responsible for ROS generation and regulate many redox-dependent signaling pathways. Furthermore, they contribute to a variety of cell processes such as cell differentiation, growth, proliferation, and embryonic development (figure 1.9 B) (Forrester et al., 2018; Nathan & Cunningham-Bussel, 2013; Skonieczna et al., 2017).



**Figure 1.9: Intracellular ROS generation.** The schemes display the formation of three different isoforms of intracellular ROS: superoxide ( $\text{O}_2^-$ ), hydrogen peroxide ( $\text{H}_2\text{O}_2$ ) and

## Introduction

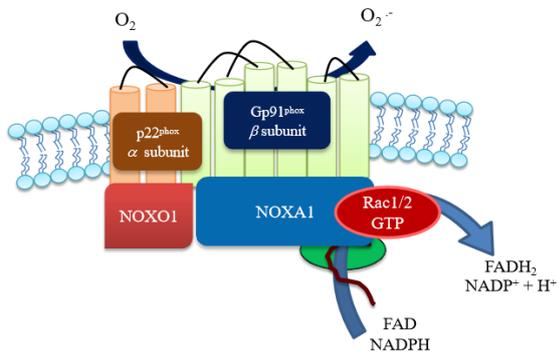
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the highly reactive hydroxyl radical (OH $\cdot$ ) from various sources including: (A), mitochondria and (B), NADPH oxidase. Figure adapted with permission from (Bigarella et al., 2014) copyright 2014 The Company of Biologists.

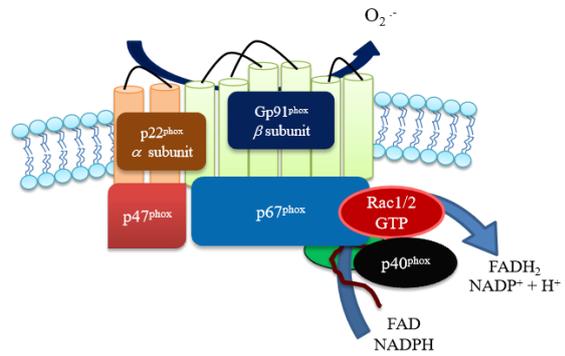
NOX enzymes have the ability to modulate the proliferation and differentiation of stem cells. Therefore, they play an essential role in stem cell therapies, tissue engineering and regenerative medicine. Several studies have been reported that NOX-mediated redox signaling played an important role in neovascularization *in vivo*, which is required for the generation of functional tissues (Elsa C Chan et al., 2009; Jiang et al., 2008; Lee et al., 2014; Skonieczna et al., 2017). The NOX family is a transmembrane protein group, participating in ROS generation (i.e. superoxide anion (O $_2^{\cdot-}$ ) and hydrogen peroxide (H $_2$ O $_2$ )) via reducing oxygen by transport electrons from NADPH (Bedard & Krause, 2007). There are seven NOX isoforms in mammals NOX1-NOX5 and the others two are Duox1 (dual oxidase 1) and Duox2. Rat and mice lack the NOX5 (figure 1.10) (Holterman et al., 2014). NOX isoforms distribute in cell membranes such as the plasma membrane, endoplasmic reticulum, mitochondrial membrane, and nuclear membrane, as well as in specialized membrane microdomains like caveolae and lipid rafts, focal adhesions, and invadopodia (BelAiba et al., 2007; Block & Gorin, 2012). Although mitochondria and NADPH oxidase are considered major sources of ROS, there are other potential sources such as (CYP), xanthine oxidase (XO), or nitric oxide synthase (NOS) which also contribute to the redox potential (Roy et al., 2015). Moreover, intracellular ROS can be generated by other organelles within the cell such as peroxisomes, which are very dynamic and metabolically active organelles. Peroxisomes have the potential to rapidly generate and scavenge H $_2$ O $_2$  and O $_2^{\cdot-}$ , which enables to regulate dynamic shifts in ROS levels (Sandalio et al., 2013).

## Introduction

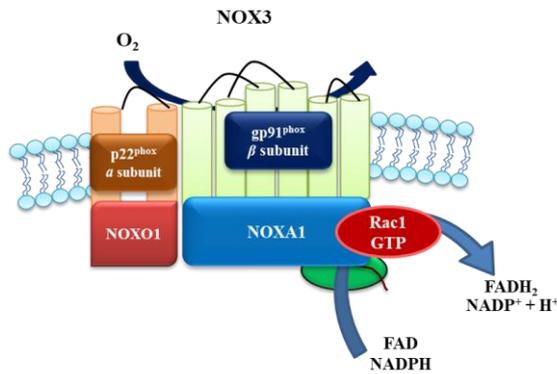
NOX1



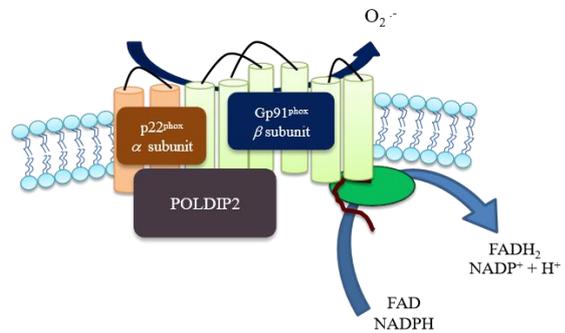
NOX2



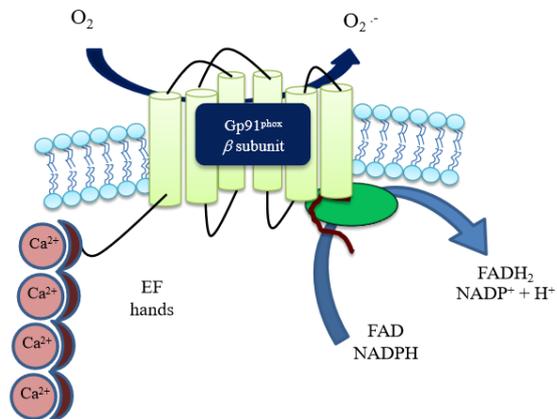
NOX3



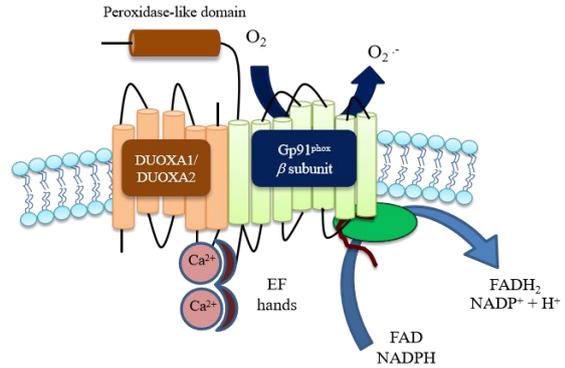
NOX4



NOX5



DUOX1/DUOX2



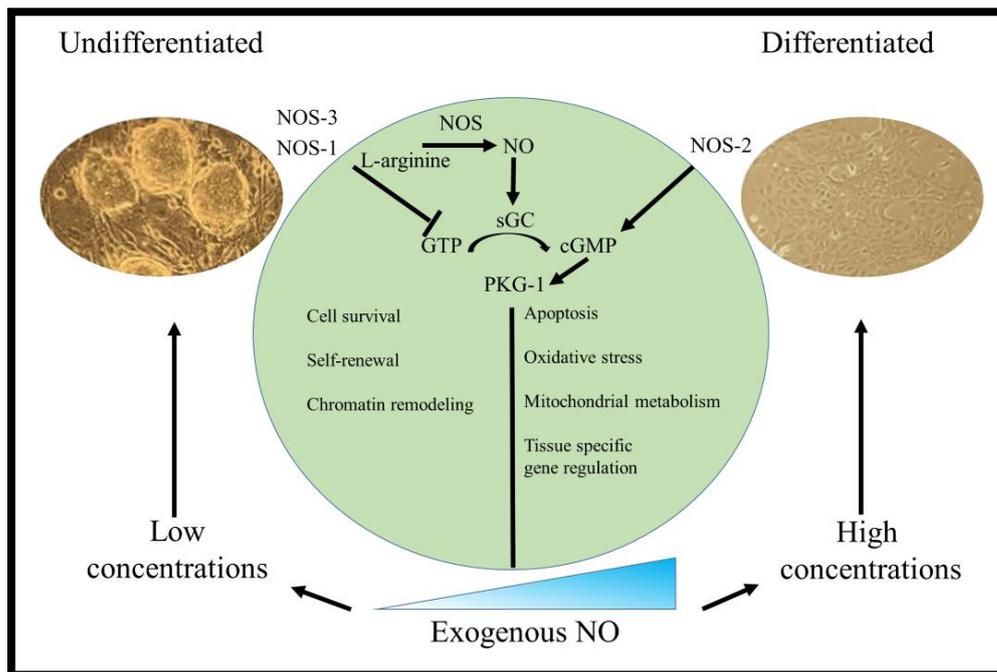
**Figure 1.10: Structure of NOX isoforms (NOX1 to NOX5) and (DUOX1 and DUOX2).** Figure adapted with permission from (Skonieczna et al., 2017); copyright 2017 Hindawi.

### 1.10 Nitric oxide

Nitric oxide (NO) is a short-lived free radical, synthesized enzymatically from the amino acid L-arginine. There are three NOS isoforms: neuronal NOS-1 (nNOS), inducible NOS-2 (iNOS), and endothelial NOS-3 (eNOS). (Marletta, 1994; Rochette et al., 2013; Tejedo et al., 2010). NO interacts with molecules such as oxygen, superoxide or metals, nucleic acids, and proteins (Cosby et al., 2003). NO serves as a second messenger and has numerous biological effects related to a variety of physiological functions in mammals, such as the regulation of blood pressure via smooth muscle relaxation and restraint of platelet aggregation (Enikolopov et al., 1999; Y Kots et al., 2011). It has been determined that it changes gene expression at the level of transcription and translation and regulates cell survival and proliferation in different cell types. In addition, NO is involved in growth, survival, proliferation, differentiation. Moreover, NO plays a significant role in the maintenance of fundamental features required for embryonic development and expansion of ES cells. In ES cells, NO has a dual function: it controls differentiation, survival, and pluripotency (Beltran-Povea et al., 2015; Caballano-Infantes et al., 2017; Tejedo et al., 2012), and promotes ES cell differentiation. Notably, it increases cardiac differentiation of mouse embryonic stem cell (mES cell) by positive and negative selection (Kanno et al., 2004; Padmasekar et al., 2011). The effects of NO on cellular processes are dose-dependent. Lower concentration of NO has been proposed to stimulate cell proliferation and survival processes, while a higher concentration of NO exerts oxidative and nitrosative stress (Alderton et al., 2001; Lundberg & Weitzberg, 2009; Thomas et al., 2008). In addition, it has been shown in ES cells, that lower concentrations of NO maintains pluripotency and enhances mitochondrial biogenesis, whereas at high concentrations, NO is a potential differentiation agent in several cell lines (Caballano-Infantes et al., 2017). Nitrate and nitrite form quickly by oxidation of NO and are now considered as non-inert end products because they are a source of NO through their reduction by a reverse pathway that describes an alternative source of NO while the endogenous L-arginine/NOS pathway is dysfunctional (Cosby et al., 2003). NOS-1 and NOS-3 are expressed in ES cells, and they produce a low level of NO. Cyclic guanosine monophosphate (cGMP) synthesis and protein kinase G (PKG-1) are inhibited in case of the low expression of the NO receptor

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gene. These changes in gene expression promote pluripotency, self-renewal and cell survival. Low doses of exogenous NO can stimulate differentiation by activation of soluble guanylate cyclase (sGC) and subsequent synthesis of cGMP. When ES cells are treated with high doses of NO apoptosis and changes in mitochondrial metabolism and tissue-specific gene regulation occur which depend on culture conditions and differentiation protocols (figure 1.11) (Mora-Castilla et al., 2010; Tejedo et al., 2010; Tejedo et al., 2012).



**Figure 1.11: NO signaling in ES cells.** Exogenous and endogenous NO effect on the ES cell status depend on the concentration. Low concentration of NO promotes pluripotency, self-renewal, and cell survival, while high concentration of NO leads to activation of apoptosis, mitochondrial metabolism and tissue-specific gene regulation. Figure adapted with permission from (Tejedo et al., 2012); copyright 2012 Springer Nature.

### 1.11 Intracellular calcium

The Intracellular calcium ( $\text{Ca}^{2+}$ ) ion is the crucial second messenger in extra/intracellular signaling cascades. Transient changes in intracellular  $\text{Ca}^{2+}$  concentration  $[\text{Ca}^{2+}]_i$ , occur in microseconds ( $\mu\text{s}$ ) or s, and  $\text{Ca}^{2+}$  can diffuse through cells and tissues. Spontaneous events

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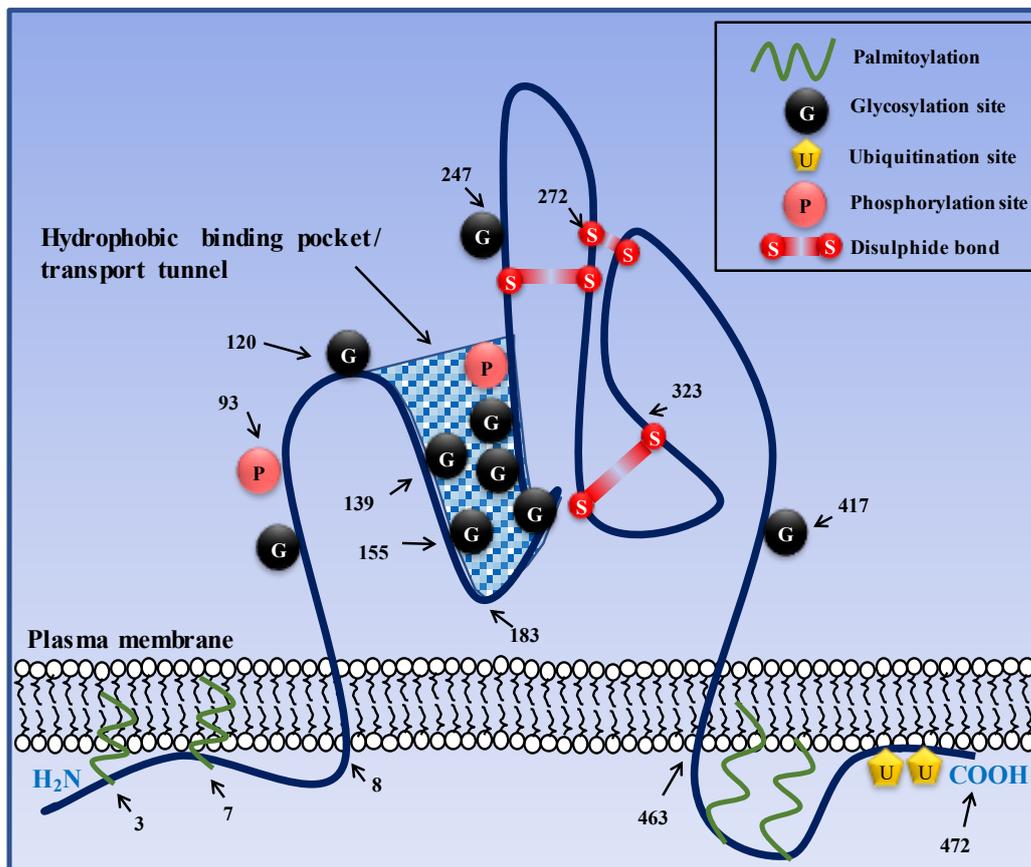
or regulated signal transduction cascades control free intracellular  $\text{Ca}^{2+}$  levels, and these signalling networks operate in different ways to contribute to various differentiation stages and stem cell fates. Moreover,  $\text{Ca}^{2+}$  plays an essential role in cellular functions such as muscle contraction, secretion, metabolism, gene expression, cell survival, as well as in cell life and death decisions. (Bootman et al., 2001; Görlach et al., 2015; Sun et al., 2007; Tonelli et al., 2012). There are two ways to increase intracellular  $\text{Ca}^{2+}$ : either by the flow of  $\text{Ca}^{2+}$  across the plasma membrane or release from intracellular stores of both the endoplasmic reticulum (ER) and sarcoplasmic reticulum (SR) in muscle.  $\text{Ca}^{2+}$  release takes place via activation of inositol trisphosphate (IP3) receptors or ryanodine receptors (RyRs) either by chemical or mechanical interaction between receptors/ion channels in the plasma membrane and  $\text{Ca}^{2+}$  stores (Berridge et al., 1999; Tumelty et al., 2007).

### **1.12 Scavenger receptor class B member 3 or CD36 receptor**

Scavenger receptor class B member 3 (SCARB3) or CD36 is a widely expressed transmembrane glycoprotein. It belongs to the class B of the scavenger receptor family (i.e., scavenger receptors with two transmembrane domains), which includes scavenger receptor-B1 (SR-B1) that is involved in binding of low-density lipoprotein (LDL) and high-density lipoprotein (HDL) particles, and lysosomal integral membrane protein II (LIMP2) (SR-B3). CD36 is now officially designated as SR-B2. CD36 consists of two major transmembrane domains, the large extracellular loop and the small cytoplasmic tails. The extracellular part has 9 *N*-linked glycosylation sites (positions 79, 102, 134, 205, 220, 231, 235, 247 and 417) and 2 phosphorylation sites (positions 92 and 237). Disulfide bonds linked between extracellular cysteines, which are present between amino acid residues 243–311, 272–333, and 313–323 (figure 1.12). Basically, in heart and muscle, CD36 is participating in long-chain fatty acid uptake across the plasma membrane. This occurs by vesicle-mediated reversible trafficking between intracellular membrane compartments and the cell surface (Luiken et al., 2016; PrabhuDas et al., 2014). Furthermore, CD36 has been involved in dysregulated fatty acid and lipid metabolism in pathophysiological conditions, especially in high-fat diet-induced insulin resistance and diabetic cardiomyopathy. (Glatz & Luiken, 2018). There is some evidence that CD36 facilitated transport of EPA, and DHA. CD36 also mediates fatty acid-induced signaling to increase intracellular  $\text{Ca}^{2+}$  in

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various cell types. The membrane-impermeable sulfo-N-succinimidyl oleate (SSO) irreversibly binds CD36 and has been extensively utilized to inhibit CD36-dependent fatty acid uptake and its effect on  $\text{Ca}^{2+}$  signaling (Franekova et al., 2014; Glatz & Luiken, 2015; Kuda et al., 2013).



**Figure 1.12: Two-dimensional structure and membrane topology of glycoprotein CD36.** Figure adapted from (Glatz & Luiken, 2018) licensed under Creative Commons Attribution 4.0 International (CC BY 4.0) and with permission from (Luiken et al., 2016); copyright 2016 Elsevier.

### 1.13 5' adenosine monophosphate-activated protein kinase

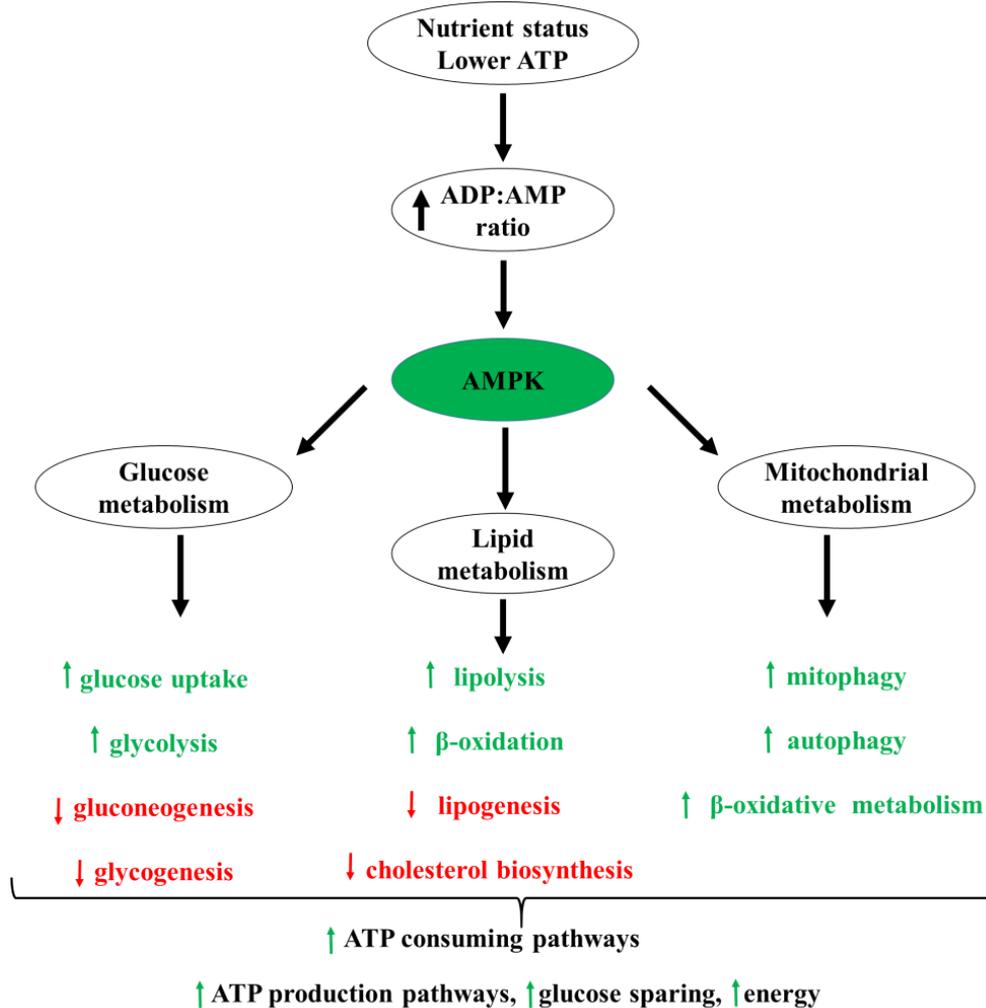
5' adenosine monophosphate-activated protein kinase (AMPK) is a serine/threonine-protein kinase. It is a master energy sensor, that regulates cellular metabolism in response

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to nutritional and environmental variations, and plays an important role in energy homeostasis. It activates and induces energy production via stimulation of glucose uptake and lipid oxidation during a deficit in the nutrient condition. It reestablishes energy balance by turning off energy-consuming processes including glucose and lipid production while activating energy-producing processes. Moreover, AMPK regulates glucose homeostasis in multiple tissues in the body such as skeletal muscle, liver, adipose tissue, and pancreatic  $\beta$  cells and assists as an inter-tissue signal integrator by responding to various hormonal signals including leptin and adiponectin (Long & Zierath, 2006; Lyons & Roche, 2018). AMPK is a heterotrimeric protein complex which consists of a catalytic  $\alpha$  subunit, a scaffolding  $\beta$  subunit, and a regulatory  $\gamma$  subunit. The  $\alpha$  and  $\beta$  subunit is encoded by two genes ( $\alpha 1$  and  $\alpha 2$  or  $\beta 1$  and  $\beta 2$ ), while three genes encode the  $\gamma$  subunit ( $\gamma 1$ ,  $\gamma 2$ , and  $\gamma 3$ ) (Cheung et al., 2000; Stapleton et al., 1996; Thornton et al., 1998). AMPK acts as a nutrient sensor, which is activated by a low energy state. ATP is an energy source, which is broken down to produce adenosine diphosphate (ADP) and then adenosine monophosphate (AMP), respectively. When the ADP:AMP ratio is higher and ATP level is lower, AMPK activates pathways that produce ATP through glucose, lipid and mitochondrial metabolism pathways, consequently increasing ATP levels. Conversely, pathways that deplete ATP are inhibited by AMPK (figure 1.13) (Garcia & Shaw, 2017; Hardie, 2015; Lyons & Roche, 2018; Ma et al., 2012). AMPK can be activated in two ways: i) directly, by binding with its activators (e.g. 5-aminoimidazole-4-carboxamide riboside (AICAR)), which stimulate structural changes in the AMPK complex through direct interaction with a specific subunit of AMPK. This takes place without any significant change in cellular ATP, ADP, or AMP levels, ii) indirectly (e.g. metformin or ALA) by accumulation of AMP and calcium (Kim et al., 2016). AMPK activity is principally determined by intracellular energy levels and is very sensible to the cellular ROS level (Choi et al., 2001). ROS can induce AMPK activity even without a reduction in cellular ATP (Quintero et al., 2006; Wu et al., 2012). Oxidative modification of the AMPK- $\alpha$  subunit seems to be an essential mechanism by which AMPK is stimulated under states of oxidative stress. Hence, every modulator which is able to cause intracellular ROS generation can activate AMPK without an associated decrease in ATP levels (Zmijewski et al., 2010).

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**Figure 1.13: Overview of AMPK in cellular metabolism.** Green arrows represent upregulated processes, red arrows represent downregulated processes. Figure adapted from (Lyons & Roche, 2018) licensed under (CC BY 4.0).

### 1.14 PPAR- $\alpha$

PPAR- $\alpha$  is a ligand-activated transcription factor which belongs to the nuclear receptor family of PPARs. This nuclear receptor family consists of the subfamily receptors PPAR- $\beta$  and PPAR- $\gamma$  (Kersten, 2014; Lee et al., 2009). Several PPAR- $\alpha$  target genes are associated with fatty acid metabolism. PPAR- $\alpha$  is highly expressed in a variety of tissues with high energy demand such as the liver, heart, skeletal muscle, brown adipose tissue, and kidney. It is also expressed in other tissues like intestine, vascular endothelium, smooth muscle and

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immune cells like monocytes, macrophages, and lymphocytes. It is significantly increased during cardiomyogenesis in ES cells (Ding et al., 2007; Pawlak et al., 2015). PPAR- $\alpha$  regulates the genes encoding mitochondrial fatty acid  $\beta$ -oxidation enzymes during cardiac developmental processes and under physiological/pathophysiological conditions. Additionally, inhibition of PPAR- $\alpha$  diminishes expression of cardiac sarcomeric proteins and specific genes during cardiac differentiation (Ding et al., 2007). PPAR- $\alpha$  is a nutritional sensor; thus PPAR- $\alpha$  plays a significant role in metabolic regulation and modifies the rates of fatty acid catabolism and lipogenesis (Hashimoto et al., 2000; Kersten, 2014; Lee et al., 2009). PPAR- $\alpha$  is a transcriptional regulator of genes implicated in peroxisomal and mitochondrial  $\beta$ -oxidation as well as fatty acid transport (J. Xu et al., 2002). The development and progression of atherosclerosis, plaque rupture, and thrombus formation could be abolished via activation of PPAR- $\alpha$  by its ligands in the cardiovascular system (Marx et al., 2001; Zandbergen & Plutzky, 2007). PPARs are required for multiple physiological functions (Kota et al., 2005), that are controlled by a lot of endogenous and exogenous compounds, including fatty acids (e.g. n-3 and n-6 fatty acids) and their metabolites. Eicosanoids and some endocannabinoids and phospholipids have been classified as endogenous ligands of PPARs. These PPAR- $\alpha$  activators include 8-epoxyeicosatrienoic acids (8-EETs), the AA lipoxygenase metabolite leukotriene B<sub>4</sub> (LTB<sub>4</sub>), and the arachidonate monooxygenase metabolites EETs (Cowart et al., 2002; Guan, 2002). Exogenous PPAR ligands e.g. clofibrate, gemfibrozil, fenofibrate, bezafibrate, and ciprofibrate have been developed and applied in experimental studies and clinical applications. For example, fibrates, which are PPAR- $\alpha$  ligands, are utilised to ameliorate microvascular risks associated with metabolic syndrome (Fidaleo & Sartori, 2008; Liss & Finck, 2017; Wang et al., 2013).

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### **Aims of the study**

PUFAs regulate many cellular functions, such as proliferation, cell growth, and differentiation via a complicated network of signaling pathways. Nevertheless, the role of PUFAs in the differentiation of ES cells into the cardiovascular lineage, e.g. cardiomyocytes and endothelial cells remains unexplored.

In detail, the goals of this study are:

- To unravel the impact of PUFAs, i.e. LA ( $\omega$ -6) and EPA ( $\omega$ -3) on cardiovascular lineage differentiation of ES cells.
- To identify intracellular messengers in the PUFA-induced signal transduction cascade such as ROS, NO and intracellular calcium ( $\text{Ca}^{2+}$ ), which have been identified in previous studies to play an important role in the regulation of ES cell differentiation into the cardiovascular lineage.
- To investigate, how ROS, NO, and  $\text{Ca}^{2+}$  raised by PUFAs interfere and interact with energy sensing molecules such as PPAR- $\alpha$ , and AMPK- $\alpha$ .

### 2. Materials and methods

#### 2.1 Materials

##### 2.1.1 Companies

**Table 1: Company list**

Company	Address
Abcam	Cambridge, UK
Abnova	Heidelberg, Germany
Air Liquide	Vitry-sur-Seine, France
Amsbio	Abingdon, UK
Ansell	Brussels, Belgium
AppliChem	Darmstadt, Germany
Aqua B. Braun Melsungen AG	Melsungen, Germany
BD Biosciences	Heidelberg, Germany
BD Falcon	New York, USA
Bemis Flexible Packaging	Neenah, USA
Biochrom	Berlin, Germany
BioLegend	Koblenz, Germany
Biometra	Göttingen, Germany
Biorbyt-Biozol	Eching, Germany
Biotec-Fischer	Reiskirchen, Germany
BioVision	Milpitas, USA
Biozym Scientific	Hessisch Oldendorf, Germany
Boehringer, Ingelheim,	Ingelheim am Rhein, Germany
Brand	Wertheim, Germany
Calbiochem	San Diego, USA
Carl Roth	Karlsruhe, Germany

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Carl Zeiss	Göttingen, Germany
Castel MAC SpA	Castelfranco Veneto, Italy
Cayman	Ann Arbor MI, USA
Cell Signaling Technology	Danvers, USA
Chemicon	Hampshire, UK
Corning	New York, USA
Edmund Bühler	Hechingen, Germany
Enzo	Farmingdale NY, USA
Eppendorf	Germany
Falcon	Franklin Lakes, USA
Fisher Scientific	Schwerte, Germany
Focus Biomolecules LLC	Plymouth Meeting, USA
Gentaur	Aachen, Germany
Gilson	Middleton, USA
Greiner bio-one	Frickenhausen , Germany
Hanna Instruments	Kehl am Rhein, Germany
Heirler-Cenovis	Radolfzell, Germany
Heraeus	Hanau, Germany
HLC BioTech	Bovenden Germany
Holzner	Nussloch, Germany
IKA	Wilmington, USA
Integra Biosciences	Fernwald, Germany
Integra Biosciences	Fernwald, Germany
Invitrogen by Life Technologies	Darmstadt, Germany
Invitrogen by Thermo Fisher Scientific	Dreieich, Germany
InvivoGen	San Diego, USA
ISO Lab	Wertheim, Germany
Jackson ImmunoResearch,	Cambridgeshire, UK

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Labexchange	Burladingen-Hausen, Germany
Lauda	Lauda-Königshofen, Germany
LeicaMicrosystems	Wetzlar, Germany
Life Technologies	Darmstadt, Germany
Lonza	Rockland, USA
Merck-Millipore	Darmstadt, Germany
Mettler Toledo	Greifensee, Switzerland
Millipore	Darmstadt, Germany
PAA	Cölbe, Germany
Peqlab Biotechnologie	Erlangen, Germany
R. Langenbrinck	Emmendingen, Germany
Sarstedt	Nümbrecht, Germany
Sartorius	Göttingen, Germany
Schleicher & Schuell	Dassel, Germany
Siemens	Munich, Germany
Sigma-Aldrich	Steinheim, Germany
Southern Biotech	Birmingham, USA
Stuart	London, UK
Tecan	Männedorf, Switzerland
Thermo Fisher Scientific	Dreieich, Germany
Tocris	Nordenstadt, Germany
Vasopharm	Würzburg, Germany
VWR International	Darmstadt, Germany

## Materials and Methods

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### 2.1.2 Materials

**Table 2: Materials list**

Materials	Supporting Company
24-well cell culture plate	Sarstedt
96-well ELISA plate	Falcon
Bacterial culture dish 100, suspension	Greiner bio-one
Blotting chamber Xcell II Blot module	Invitrogen by Life Technologies
Cellspin spinner flask stirrer	Integra Biosciences
Conical flasks	Fisher Scientific
Conical tubes 15ml & 50ml	BD Falcon
CryoPure Tube 1.8ml	Sarstedt
Filter paper circles	Schleicher & Schuell
Filter paper for blotting	Biotec-Fischer
Filtropur S 0.2	Sarstedt
Gel electrophoresis chamber	Invitrogen by Life Technologies
Gel electrophoresis power supply	Invitrogen by Life Technologies
Glass Pasteur pipettes	VWR International
Magnetic stirrer	IKA
Magnetic stirrer for spinner flask	Integra Biosciences
Micropipettes	Gilson and Eppendorf
Microscope Cover Slips	Fisher Scientific
Microscope slides, ground edges	R. Langenbrinck
Micro-tubes 0.5ml & 1.5ml	Sarstedt
Nalgene Mr. Frosty	Thermo Fisher Scientific
Nitrile gloves	Ansell
PAGEr EX Gels	Lonza, Rockland
Parafilm	Bemis Flexible Packaging
Pipette filter tips	Biozym Scientific

## Materials and Methods

Pipette tips	Sarstedt
PVDF Millipore transfer membrane	Merck-Millipore
Serological glass pipettes	ISO Lab
Serological pipettes wide tip	Corning
Serological plastic pipettes	Greiner Bio-One
Spinner flask	Integra Biosciences
Sponge Pad for blotting	Invitrogen <sup>®</sup> by Life Technologies
Tissue culture treated dish 100, standard	Sarstedt
Tissue culture treated dish 60, standard	Sarstedt
Tissue culture treated dish 60, suspension	Sarstedt

### 2.1.3 Instruments

**Table 3: Instruments list**

Instruments	Supporting Company
+4 °C refrigerator	Siemens
-20 °C freezer	Thermo Fisher Scientific
-80 °C freezer	Heraeus
Analogue tube rollers	Stuart
Autoclave	Holzner
Cell culture bench	Heraeus
Cell culture incubators	Heraeus
Centrifuge 1S-R	Heraeus
Centrifuge 5417C	Eppendorf
Chemiluminescence imaging system	Peqlab Biotechnologie
Confocal laser scanning microscope (CLSM) TCS-SP2, AOBS	LeicaMicrosystems
ELISA reader	Tecan
Heating block	Biometra
Homogenizer	Sigma-Aldrich

## Materials and Methods

Ice machine	Castel MAC SpA
Inverted light microscope (Zeiss Axiovert 40 C)	Carl Zeiss
Liquid nitrogen tank	Air Liquide
Microscope heating stage TRZ 3700	Labexchange <sup>®</sup>
Milli-Q Advantage A10 System	Millipore
Pellet pestle 1.5 ml	Fisher Scientific
pH-meter	Hanna Instruments
Safety Bunsen burner	Integra Biosciences
Shaker	Stuart and Edmund Bühler, Hechingen
Vacuum pump	HLC BioTech
Vortex-Genie <sup>®</sup>	VWR International
Water bath	Lauda
Weighing machine-AB265S	Mettler Toledo
Weighing machine-TE153S	Sartorius

### 2.1.4 Chemicals

**Table 4: Chemicals list**

Materials	Supporting Company
1,5-bis (2-(di-methylamino) ethylamino)-4,8-dihydroxyanthracene-9,10-dione (DRAQ5)	Cell Signaling Technology
2'7'-dichlorodihydrofluorescein diacetate (H <sub>2</sub> DCF-DA)	Thermo Fisher Scientific
4-(6-Acetoxymethoxy-2,7-difluoro-3-oxo-9-xanthenyl)-4'-methyl-2,2'-(ethylenedioxy) dianiline-N,N,N',N'-tetraacetic acid tetrakis (acetoxymethyl) ester (Fluo-4-AM)	Thermo Fisher Scientific
4-amino-5-methylamino-2',7' dichlorofluorescein diacetate (DAF-FM) diacetate	Thermo Fisher Scientific
Acetic acid	Sigma-Aldrich
Bovine serum albumins (BSA) (powder)	Sigma-Aldrich

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Collagenase B	Boehringer, Ingelheim
Copper sulfate	Sigma-Aldrich
Dimethyl sulfoxide (DMSO)	Millipore
Di-Sodium hydrogen phosphate dihydrate	Carl Roth
Dulbecco's modified Eagle medium (DMEM) high glucose medium	Biochrom
Dulbecco's 1x phosphate buffered saline (PBS) without Ca <sup>2+</sup> & Mg <sup>2+</sup> ions	PAA
EmbryoMax <sup>®</sup> 0.1% gelatin solution	Merck-Millipore
EPA	Focus Biomolecules LLC and Tocris
ESGRO <sup>®</sup> LIF (107 units)	Chemicon
Ethanol	Carl Roth
Ethylenediaminetetraacetic acid (EDTA)	Carl Roth
Fetal calf serum (FCS)	Sigma-Aldrich
Fluoromount-G <sup>®</sup>	Southern Biotech
Folin-ciocalteau reagent	Sigma-Aldrich
Hydrochloric acid (HCl)	Sigma-Aldrich
Hydrogen peroxide solution (30%)	Sigma-Aldrich
Iscove's basal medium	Biochrom
LA	Enzo
L-glutamine (200mM)	Biochrom
Luminol	Sigma-Aldrich
Methanol	Carl Roth
Mitomycin-C	Calbiochem
Non-essential amino acids (NEAA) (100X)	Biochrom
Non-fat dried milk powder	Heirler-Cenovis
Nonidet P-40	AppliChem
Novex <sup>®</sup> Sharp Pre-stained protein standard	Life Technologies
NuPAGE Antioxidant	Invitrogen by Thermo Fisher Scientific

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NuPAGE <sup>®</sup> LDS sample buffer (4x)	Invitrogen by Thermo Fisher Scientific
NuPAGE <sup>®</sup> Sample reducing agent (10x)	Invitrogen by Thermo Fisher Scientific
P-Coumaric acid	Sigma-Aldrich
Penicillin-Streptomycin (100X)	Biochrom
Phosphatase inhibitor cocktail 3	Sigma-Aldrich
Plasmocin <sup>™</sup> Prophylactic 25mg/ml	InvivoGen
Ponceau-S	AppliChem
Potassium chloride	Carl Roth
Potassium dihydrogen phosphate	Carl Roth
Potassium sodium tartrate tetrahydrate	Merck-Millipore
ProSieve EX- running buffer	Lonza
ProSieve EX- transfer buffer	Lonza
Protease inhibitor cocktail	BioVision
Restore Western blot stripping buffer	Thermo Fisher Scientific
Sigmacote <sup>®</sup>	Sigma-Aldrich
Sodium carbonate	Carl Roth
Sodium chloride	Carl Roth
Sodium deoxycholate	Sigma-Aldrich
Sodium dodecyl sulfate (SDS)	Sigma-Aldrich
Sodium hydroxide	Carl Roth
Sodium pyruvate (100mM)	Biochrom
Sterile distilled water (DW)	Aqua B. Braun Melsungen AG
Tris	Sigma-Aldrich
Triton <sup>®</sup> X-100	Sigma-Aldrich
Trypsin/EDTA (1x)	Life Technologies
Tween-20	Sigma-Aldrich
β-Mercaptoethanol	Sigma-Aldrich

## Materials and Methods

### 2.1.5 Buffer compositions

**Table 5: Buffers and solutions compositions**

<b>Stock Buffers</b>			
<b>Buffers</b>	<b>Volum</b>	<b>The final molar concentrations of the 1x solution are:</b>	<b>Materials</b>
10x PBS	1 Liter	2.7mM 2.0mM 137mM 10.0mM	KCl KH <sub>2</sub> PO <sub>4</sub> NaCl Na <sub>2</sub> HPO <sub>4</sub> · 2H <sub>2</sub> O
10xTBS	1 Liter	25mM 137mM	Tris base NaCl
Radioimmunoprecipitation assay (RIPA)-Buffer	1 Liter	50mM 150mM 1% 0.5% 0.5%	Tris. pH 7.4 NaCl Nonidet P-40 SDS Sodium deoxycholate (excluded in the Phosphoprotein assay)
1xTris-HCl	1 Liter	100mM	Adjust pH at 7.4 using concentrated HCl
<b>Mix-ups</b> Dissolve the materials in 800ml distilled H <sub>2</sub> O, adjust pH to 7.4, complete volume to 1L with additional distilled H <sub>2</sub> O, and then sterilize by autoclaving.			
<b>Washing Buffers</b>			
1xPBS		100ml of 10x PBS diluted in 900ml distilled H <sub>2</sub> O	
0.01% PBS-Triton X-100		100µl of Triton X-100 added to 1000ml 1x PBS	
0.1% TBS-Tween20		(100ml of 10xTBS) + (1ml of Tween20) diluted in 900ml distilled H <sub>2</sub> O	
<b>Permeabilization buffer</b>			
1% PBS-Triton X-100		1ml of Triton X-100 added to 100ml 1x PBS	
<b>Blocking buffers</b>			
10% FCS		1ml of FCS added to 10ml 0.01% PBST	
10% Non-fat dry milk		1g of milk dissolved in 10ml 0.01% PBST	
5% Non-fat dry milk		5g of milk dissolved in 100ml 1xTBS-Tween20	
<b>Protein extraction buffers</b>			

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Protein extraction buffer	RIPA buffer with sodium deoxycholate	1x
	Protease inhibitor cocktail	1x
Phospho-protein extraction buffer	RIPA buffer without sodium deoxycholate	1x
	Protease inhibitor cocktail (500x)	1x
	Phosphatase inhibitor cocktail (100x)	1x
	EDTA (pH 8)	1mM
	Glycerophosphate	1mM
<b>Ponceau-S staining solution</b>		
Ponceau-S staining (w/v)		0.1%
Acetic acid (v/v)		5%
<b>Enhanced chemiluminescence (ECL) detection solution</b>		
Luminol		1.25mM
p-Coumaric acid		225 $\mu$ M
Hydrogen peroxide solution (30%) (v/v)		3mM
Tris-HCl (pH 8.5)		100mM
<b>Lowry solutions</b>		
Solution A: 2% Sodium carbonate (w/v) in 0.1N sodium hydroxide solution		
Solution B: 1.25% Copper sulphate (w/v)		
Solution C: 3.37% Potassium sodium tartrate tetrahydrate (w/v)		
Solution D: Solution A, solution B, and solution C mixed-up in the ratio 100:1:1		
Solution E: 1N Sodium hydroxide solution		
Solution F: 2N Folin & ciocalteu's phenol reagent		

### 2.1.6 Cell culture media components

**Table 6: Media composition**

MEF medium	Final concentration
DMEM high glucose medium	1x
Heat-inactivated (FCS)	10%
$\beta$ - mercaptoethanol	0.1mM
Sodium pyruvate (100mM)	1.0mM

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L-glutamine (200mM)	2.0mM
NEAA (100mM)	0.1mM
Penicillin-Streptomycin (10000U/ml)	50U/ml
<b>LIF-Plasmocin-Prophylactic (LIF-pLpro) medium</b>	<b>Final concentration</b>
Iscove's basal medium	1x
Heat-inactivated (FCS)	16%
$\beta$ - mercaptoethanol	0.1mM
Sodium pyruvate (100mM)	1.0mM
L-glutamine (200mM)	2.0mM
NEAA (100mM)	0.1mM
Plasmocin Prophylactic	2.5 $\mu$ g/ml
LIF	1000U/ml
<b>CCE differentiation medium</b>	<b>Final concentration</b>
Iscove's basal medium	1x
Heat-inactivated (FCS)	16%
$\beta$ -mercaptoethanol	0.1mM
Sodium pyruvate (100mM)	1.0mM
L-glutamine (200mM)	2.0mM
NEAA (100mM)	0.1mM
Penicillin-Streptomycin (10000U/ml)	50U/ml
<b>Freezing Medium</b>	<b>Final concentration</b>
MEF medium	50%
Heat-inactivated (FCS)	40%
DMSO	10%

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### 2.1.7 Cell lines

Mouse embryonic fibroblasts (MEFs) (Amsbio).

CCE S103 cell line: Mouse ES cell line isolated from mouse embryos strain129/sv (Robertson et al., 1986).

### 2.1.8 Immunocytochemistry antibodies

**Table 7: Immunocytochemistry antibodies**

Primary antibody	Animal	Concentration	Supporting Company/ Catalog Number (Cat. No.)
Anti- $\alpha$ -actinin	Mouse	1:100	Abcam/ ab9465
Anti-CD31	Rat	1:100	Merck-Millipore/ CBL1337
Anti-phospho PPAR- $\alpha$ (S12)	Rabbit	1:200	Abcam/ ab3484
Secondary antibody	Animal	Concentration	Supporting Company/ Cat. No.
Alexa Fluor 488 anti-rat IgG	Donkey	1:100	Life Technologies/ A-21208
Alexa Fluor 647 anti-mouse IgG	Sheep	1:100	Jackson ImmunoResearch/ 515-605-072
Alexa Fluor 488 anti-rabbit IgG	Donkey	1:100	Life Technologies/ A-21206
Alexa Fluor 488 anti-rat IgG	Donkey	1:100	Jackson ImmunoResearch/ 712-546-153
DyLight 650 anti-rat IgG	Goat	1:100	Abcam/ ab98408

### 2.1.9 Western blot antibodies

**Table 8: Western blot antibodies**

Primary antibody	Animal	Concentration	Supporting Company/ Cat. No	KDa
<b>Endothelial cell markers</b>				
Anti-VE-Cadherin (CD144)	Rat	1:1000	BD Biosciences/ 555289	115
Anti-PECAM1 (CD31)	Goat	1:1000	BioLegend/ 910003	130
<b>Progenitor marker</b>				
Anti-fetal liver kinase-1(FLK-1)	Rabbit	1:1000	Cell Signaling Technology/ 9698	230
<b>Cardiac protein markers</b>				
Anti- $\alpha$ -actinin	Mouse	1:1000	Sigma-Aldrich /A 5044	107
Anti-Myosin regulatory	Mouse	1:1000	Thermo Fisher	19

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light chain 2, atrial isoform (MLC2a)			Scientific/PA5-30789	
<b>Phosphoproteins</b>				
Anti-p-AMPK- $\alpha$ (Thr172)	Rabbit	1:1000	Cell Signaling Technology/2535S	62
Anti-p-eNOS (Ser1177)	Rabbit	1:1000	Cell Signaling Technology/9571S	140
Anti-p-PPAR- $\alpha$ (pSer12)	Rabbit	1:1000	Abcam/ab3484	52
<b>Housekeeping proteins</b>				
Anti-GAPDH	Rabbit	1:1000	Abcam/ab9485	36
Anti-Vinculin	Mouse	1:1000	Sigma-Aldrich/V9131	116
Anti- $\beta$ -actin	Rabbit	1:2000	BioLegend/622102	45
<b>Secondary antibody</b>	<b>Animal</b>	<b>Concentration</b>	<b>Supporting Company/ Cat. No</b>	
HRP anti-Rat	Goat	1:1000	Cell Signaling Technology/7077	
HRP anti-Rabbit	Goat	1:1000	Abcam/ab205722	
HRP anti-Mouse	Horse	1:1000	Cell Signaling Technology/7076	
HRP anti-Goat	Donkey	1:1000	Abnova/PAB10570	

### 2.1.10 Substances

#### 2.1.10.1 Preparing PUFAs and mES cell treatment

The PUFAs were supplied as minimum purity >99% and were prepared by dissolving in DMSO (final concentration of DMSO  $\leq$  0.1% in the culture medium). Once prepared, PUFAs were stored at -20 °C as recommended in datasheets. To obtain dose-response relations, physiological concentrations of 1, 5, 10, 50, and 100 $\mu$ M PUFAs were applied. In vehicle (DMSO) groups, the identical concentration of DMSO as included in PUFA treated samples was added to the culture medium.

#### 2.1.10.2 Fluorescence substances

##### 2.1.10.2.1 H<sub>2</sub>DCF-DA

H<sub>2</sub>DCF-DA is an apolar, nonfluorescent, cell-permeable substance and serves as an indicator of various ROS. After diffusion into the cell, the diacetate moiety is cleaved by intracellular esterases. The resulting polar DCFH is nonfluorescent and cell-impermeable and is rapidly oxidized to the highly fluorescent 2',7'-dichlorofluorescein (DCF) by

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intracellular ROS. The fluorescence intensity of DCF, which is excited at 488 nanometer (nm) wavelength, serves as an indicator for the intracellular amount of ROS, the fluorescence emission is detected at a wavelength of  $\geq 540$  nm (Bass et al., 1983).

### 2.1.10.2.2 DAF-FM DA

The fluorescent marker DAF-FM DA is an indicator for the measurement of intracellular NO. The cell-permeable diacetate, DAF-FM can passively diffuse across the cell membrane into the cell. Once in the cell, the diacetate group is cleaved by cell-specific esterases. The resulting product DAF-FM is weakly fluorescent. It cannot leave the cell and reacts in the presence of NO to a highly fluorescent benzotriazole derivative, which can fluoresce up to 160-fold more intense than DAF-FM (Kojima et al., 2001). Maximum excitation wavelength of the oxidized form is at 495nm, and fluorescence emission is detected at wavelengths  $\geq 515$ nm. DAF-FM is dissolved in DMSO.

### 2.1.10.2.3 Fluo-4-AM

For the visualization (imaging) of intracellular  $\text{Ca}^{2+}$  transients in living cells, the cell-permeable dye Fluo-4- acetoxymethyl ester (Fluo-4-AM) was used. Fluo-4-AM diffuses into the cell, in which cell-derived esterases split off the AM group to trap the compound in the cells. On the other hand, the hydrolysis is necessary to enable calcium-binding and fluorescence emission of the dye. Fluorescence intensity is dependent on the amount of bound calcium and therefore suitable for quantitative measurements. Fluo-4-AM is used to quantify calcium concentrations ranging from 100nM to 1 $\mu$ M. The absorption maximum of the green fluorescent Fluo-4 is 494 nm; the emission maximum at 516 nm (Gee et al., 2000).

### 2.1.10.2.4 DRAQ5

DRAQ5 is a cell-permeable far-red fluorescent DNA dye that can be used in fixed or non-fixed live cells. DRAQ5 staining can be applied in flow cytometry, live-cell imaging, and cell-based assays. DRAQ5 is excited at 633nm, and emission is recorded at  $> 655$ nm (Cheedipudi et al., 2019).

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### 2.1.10.3 Inhibitors

To determine the impact of PUFAs and their metabolism on vasculogenesis and cardiomyogenesis, differentiating ES cells were pre-incubated from day 3 to 10 with specific inhibitors at the indicated concentrations (Table 9) either in presence or absence of PUFAs. Subsequently, generation of signaling molecules, protein expression and activation of enzymes, transcription factors, and cardiovascular proteins were evaluated by microfluorometry, immunocytochemistry and western blot.

**Table 9: Inhibitors used in detection of signaling pathways**

Inhibitor	Target	Final concentration	Soluble in	MW	Supporting Company/ Cat. No.
BAPTA-AM	Calcium chelator	10 $\mu$ M	DMSO	764.7	Abcam/ ab120503
GW6471	PPAR- $\alpha$ inhibitor	5 $\mu$ M	DMSO	619.67	Tocris/ 4618
N $\omega$ -Nitro-L-arginine methyl ester (L-NAME)	NO-Synthase	100 $\mu$ M	DW	269.7	Sigma-Aldrich/ N5751
N-(2-mercaptopropionyl)-glycine (NMPG)	Radical scavenger	100 $\mu$ M	DMSO	163.2	Sigma-Aldrich/ M6635
Trolox	Radical scavenger	100 $\mu$ M	DMSO	264.3	Sigma-Aldrich/ 93510
VAS2870	NOX inhibitor	50 $\mu$ M	DMSO	360.3	Vasopharm/ F3395
SSO	SCARB3	200 $\mu$ M	DMSO	482.6	Cayman/ 11211
CD36 antibody	SCARB3	2-3 $\mu$ g/ml	-	-	Antibodies-Online.com/abin343926

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### 2.1.11 Softwares

- 1- Adobe Acrobat Reader DC, version 2020.013.20064 (San Jose, USA)
- 2- ChemiCapt 500-Western blot imaging software version 15.01a, Vilber Smart Imaging, (Peqlab Biotechnologie, Eberhardzell, Germany).
- 3- Cisco AnyConnect Cisco Secure Mobility Client 4.1.0401 (Cisco Systems, San Jose, USA).
- 4- Endnote X9 [Bld. 12062], (Clarivate Analytics, Boston, USA).
- 5- GraphPad Prism 8-Statistical analysis software (GraphPad Software, San Diego, USA).
- 6- Image J-Western blot analysis software (National Institutes of Health (NIH), Bethesda, USA).
- 7- Leica Application Suite Advanced Fluorescence (Leica Microsystems, Wetzlar, Germany).
- 8- Metamorph Microscopy Automation & Image Analysis Software (Molecular Devices, San José, USA).
- 9- Microsoft Office 2013-2016 - Data documentation (Microsoft, Seattle, USA).
- 10- Tecan Magellan - ELISA Software (Tecan, Männedorf, Switzerland).

### 2.2 Methods

#### 2.2.1 Cell culture methods

##### 2.2.1.1 Preparing MEFs

MEFs have been used in this study as feeder layer cells for ES cells culture. Cultivation on feeder layers is important for maintenance of ES cells in a stemness state (Park et al., 2015), and promotes ES cell proliferation by providing the cell culture medium with extracellular secreted growth factors and LIF (Llames et al., 2015). The MEFs are stored in liquid nitrogen (-196 °C). MEFs were thawed in a water bath at 37 °C with gently shaking, till a small piece of ice in the tube was left. Immediately, they were transferred to a falcon tube containing 20ml pre-warmed MEF medium at 37 °C. Subsequently, the tube was centrifuged for 5 min at 209g. After centrifugation, the washing medium was aspirated, and cells were cultivated at a density of  $5 \times 10^5$  cells in 5 ml MEF medium per Petri dish (diameter 60mm). Feeder cells were cultivated at a temperature of 37 °C, 5% CO<sub>2</sub>, and relative humidity of 95%. Upon 80-90% confluency the feeder cell layer was mitotically inactivated by incubation with fresh MEF medium containing 10µg/ml Mitomycin-C for 2-3h in a dark incubator at 37 °C, 5% CO<sub>2</sub>. This step was important to avoid overgrowth of cells in the culture. Subsequently, the inactivation medium was removed, cells were washed three times with MEF medium and further incubated with pre-warmed fresh MEF medium. The washing step is critical to remove the rest of Mitomycin-C, which is a bioreductive alkylating agent, derived from *Streptomyces lavendulae* bacteria and has been clinically used as an antibiotic drug and in antitumor therapy (Mao et al., 1999). Mitomycin-C has the ability to crosslink with double strand-DNA and blocks DNA synthesis and nucleus division (Verweij & Pinedo, 1990). The MEF medium was changed every two days. Inactivated feeder layers can be stored under cell culture conditions for a maximum of one week.

##### 2.2.1.2 Cultivation and maintenance of ES cells.

In this study the CCE-S103 mES cells line derived directly from mouse embryos was used (Robertson et al., 1986). ES cells were cultivated on mitotically inactivated MEFs to keep their stemness properties in presence of LIF. LIF is a member of the interleukin-6 (IL-6)

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family of cytokines, and keeps mES cells in pluripotency and undifferentiated state, thus enabling long-term culture (Nicola & Babon, 2015). mES cells stored in liquid nitrogen tank (-196 °C) were thawed in water bath 37 °C for few seconds. Then the cell suspension was transferred to a conical tube containing 15ml CCE medium and centrifuged for 5 min at 209g to remove freezing medium. The freezing medium was aspirated and 200µl of LIF-pLpro medium was added to the pellet and resuspended by gently pipetting up and down. mES cells were then seeded on an inactivated MEF feeder cell layer, which was previously prepared. The mES cells were cultivated at a temperature of 37 °C, 5% CO<sub>2</sub>, and relative humidity of 95%. The mES cells were passaged regularly every 2-3 days to avoid any extra-confluence, stress and unfavorable differentiation processes. For each passage, the mES cells were enzymatically dissociated, and transferred to a new plate on a layer of MEF feeder cells. For differentiation experiments, mES cells were seeded into spinner flasks at a density of 3 x 10<sup>6</sup> cells/ml, which contained CCE differentiation medium. This system allows the mES cells to form a small, spherical three-dimensional structure called the EBs (Ali et al., 2018)

### 2.2.1.3 Surface coating of tissue culture plates and spinner flasks

Gelatin was used to enhance mES cell and EB attachment on the surface of tissue culture plates or coverslips. Incubation with gelatin 0.1% in PBS was performed for 1h at 37 °C. Subsequently, gelatin was removed and coated tissue culture plates or coverslips were dried in the clean bench. Spinner flasks and magnetic stirring bars were coated with 10ml of Sigmacote, which is a silicone solution, coating glass surfaces with microscopically thin silicone film to prevent attachment of mES cells to spinner flasks and stirring bars. After coating, the spinner flasks and magnetic stirring bars were dried in an oven at 60 °C for 1h, washed for three times with DW, and autoclaved.

### 2.2.1.4 Trypsinization of mES cells, spinner flask preparation, and EB formation

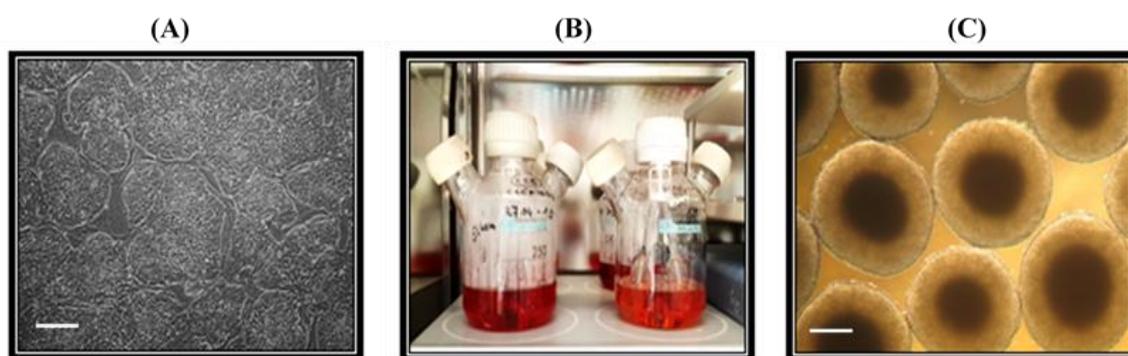
The mES cells were passaged every 2-3 days, when confluence reached about 80-90% (figure 2.1A). Briefly, mES cells in 60mm tissue culture plates were washed for one time with PBS and incubated with 2ml pre-warmed trypsin/EDTA solution for 2 min at 37 °C. Subsequently, mES cell colonies were further dissociated by gently pulling up and down

## Materials and Methods

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with a 1ml pipette tip to obtain a single cell suspension. Subsequently,  $9 \times 10^4$  single mES cells/ml were seeded onto MEF feeder cell layers in fresh LIF-pLpro pre-warmed medium.

To generate EBs mES cells were enzymatically dissociated as described and seeded at a density of  $3 \times 10^6$  cells/ml into siliconized spinner flasks containing 125 ml of CCE differentiation medium (figure 2.1B). The speed of the magnetic stirrer system was adjusted to 20 rotations per min. After 4 rotation rounds the direction of rotation was changed to avoid EB agglomeration. After 24h, 125ml CCE differentiation medium was added to give a final volume of 250ml. Every day 125 ml of culture medium was exchanged for fresh medium. After 3 days of cell culture in spinner flasks EBs were transferred into 100mm bacteriological plates to be used in experiments on cardiovascular differentiation (figure 2.1C).



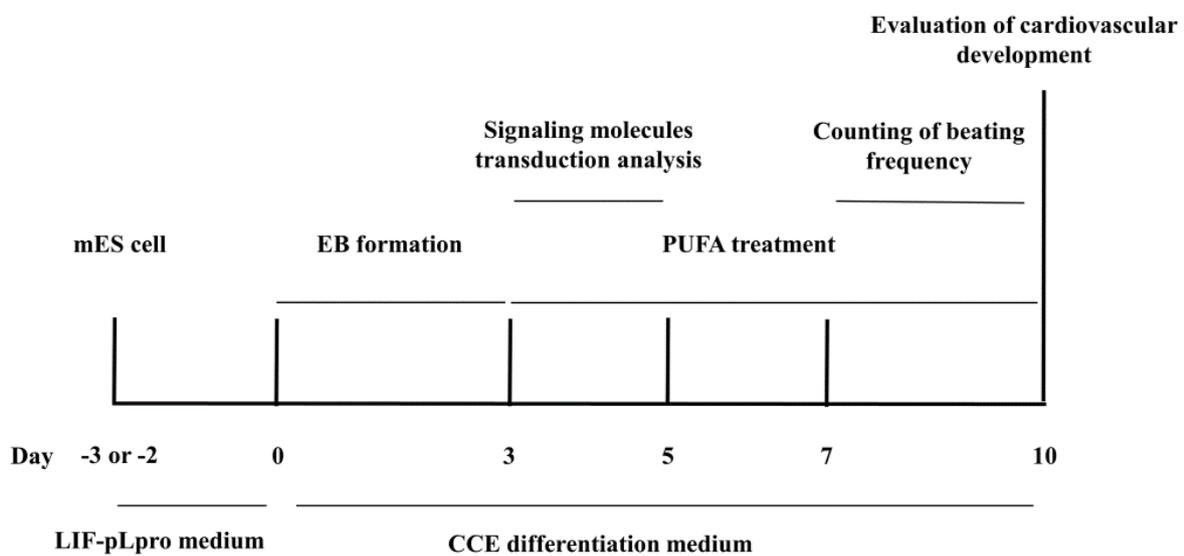
**Figure 2.1: EB formation from differentiating mES cells.** (A): Transmission image of undifferentiated ES cells cultivated on mitotically inactivated MEF feeder layers. (B): Spinner flask culture for EBs derived from mES cells. (C): Transmission image of EBs. The scale bars represent 200 $\mu$ m.

### 2.2.1.5 Protocol of PUFA treatment

For each, experiment approximately 25-30 EBs (3-day-old) were transferred to 60mm bacteriological tissue culture dishes. EBs were treated with series of physiological concentrations of LA as well as EPA, ranging from 1 to 100 $\mu$ M or were treated with appropriate concentrations of DMSO (vehicle). Petri dishes with EBs were placed on the shaker inside the incubator. The cell culture medium supplemented with PUFAs was changed every 24h. For investigation of signaling events upon PUFA treatment 5-day-old

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EBs were collected. For immunofluorescence or protein extraction, EBs were treated with PUFAs until day 10 of cell culture. The beating frequency of spontaneously beating cardiomyocyte clusters (foci) within EBs as well as the number of contracting EBs were recorded from day 7 to day 10 of cell culture using an inverted light microscope (Zeiss Axiovert 40 C), which was connected with a microscope heating stage TRZ 3700 and equipped with a 5x/0.12 objective (figure 2.2).



**Figure 2.2: Schematic overview on the experimental setup**

### 2.2.1.6 Freezing mES cells

mES cells were frozen upon confluency of 70-80%. Briefly the mES cells were washed with PBS and incubated with 2ml pre-warmed trypsin/EDTA for 2 min at 37 °C. The cells were dissociated by gently pipetting. Subsequently the suspension of mES was transferred into Falcon tubes containing 20ml CCE differentiation medium to stop enzymatic digestion. Then the tube was centrifuged for 5 min at 209g. The supernatant was aspirated, and the pellet was cooled down on the ice (+4°C), 1ml of the freezing medium was added slowly and re-suspended gently. Finally, the suspension was transferred to pre-cooled

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1.8ml CryoPur tubes and stored at -80 °C overnight. The next day frozen mES cells were transferred to a liquid nitrogen tank for long-term storage.

### 2.2.2 Immunocytochemistry

EBs grown in suspension culture or cultivated on coverslips were collected, washed with PBS, and fixed in ice-cold methanol for 20 min at -20 °C. Subsequently, methanol was removed and the samples were washed 3-4 times with 0.01% PBST buffer. For permeabilization EBs were incubated in permeabilization buffer containing 1% PBS-Triton X-100 for 10 min at room temperature (RT). Then EBs were washed 3-4 times with 0.01% PBST buffer and incubated in blocking buffer containing 10% non-fat dry milk at RT for 60 min. EBs were incubated at RT for 120 min or overnight with primary antibodies, which were freshly prepared. Anti-CD31 was diluted with 10% non-fat dry milk and anti- $\alpha$ -actinin with 10% FCS. Primary antibodies were diluted 1:100. After incubation with primary antibodies EBs were washed 3-4 times with 0.01% PBST buffer and incubated with secondary antibody, which was prepared in blocking buffer, and incubated in a dark place at RT for 60 min. After that, the secondary antibody was removed, and EBs were washed 4-5 times with 0.01% PBST buffer, and kept in the same buffer at +4°C till immunofluorescence analysis. For long-term storage tissues were fixed and sealed in Fluoromount-G<sup>®</sup> solution. (Table 10).

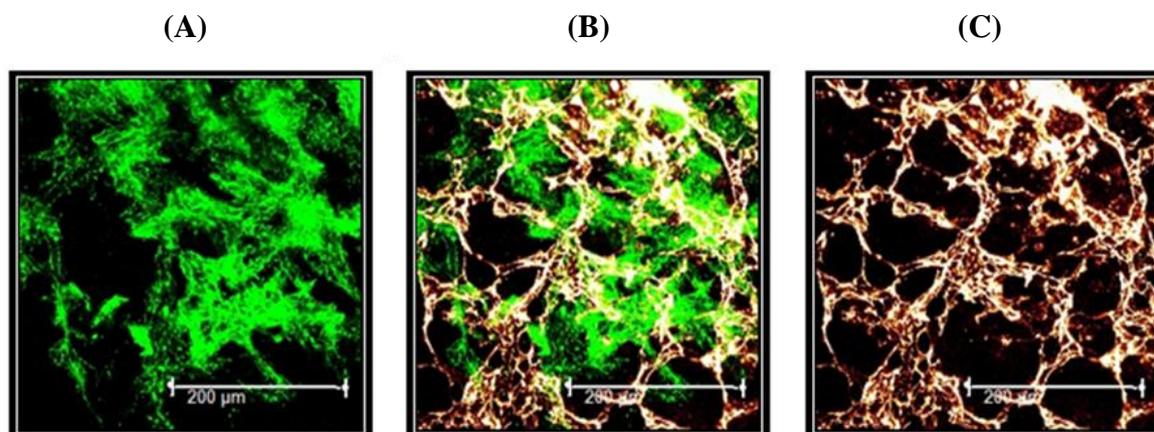
**Table 10: Immunocytochemistry protocol for staining of cardiomyocytes and endothelial cells within EBs**

Steps	Specimen	Duration	Solutions and reagents	Temperature
Washing	Endothelial cells	~1 min	1xPBS	RT
	Cardiomyocytes			
Fixation	Endothelial cells	$\geq 20$ min	ice-cold 100% methanol	-20 °C
	Cardiomyocytes			
Permeabilization	Endothelial cells	10 min	1% PBST	RT
	Cardiomyocytes			
Blocking	Endothelial cells	60 min	10% nonfat- dry milk	RT
	Cardiomyocytes		10% FCS	

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Primary antibody	Endothelial cells	60 min	Anti-CD31 diluted in 10% non-fat dry milk 1:100	+4°C
	Cardiomyocytes	overnight	Anti- $\alpha$ -actinin diluted in 10% FCS 1:100	
Secondary antibody	Endothelial cells	60 min	Diluted in 10% non-fat dry milk 1:100	RT
	Cardiomyocytes	60 min	Diluted in 10% FCS 1:100	RT

Fluorescence recordings were performed by means of a confocal laser scanning setup (Leica SP2 AOBS) connected to an inverted microscope. The confocal setup was equipped with a 5 milliwatt (mW) helium/neon laser, single excitation 633 nm, and an argon-ion laser, single excitation 488 nm. Emission was recorded using the band-pass filter BP 515-550 and long-pass LP 650 filter sets, respectively. For the analysis, a 10 $\times$ , numerical aperture (NA) 0.5 objective was used. The pinhole settings of the confocal setup were adjusted to give a full-width half maximum of 5 $\mu$ m. For the quantification of capillary areas within EBs an optical sectioning routine based on CLSM was used. Images (512  $\times$  512 pixels) were acquired from CD31-stained EBs using the extended depth of focus algorithm of the confocal setup. In brief, 10 full frame images, separated by a distance of 10 $\mu$ m in z-direction, were recorded. The obtained images included information of the capillary area and spatial organization of CD31-positive cell structures in a tissue slice 100 $\mu$ m thick. The acquired images were processed to generate a single in-focus image projection of vascular structures in the scanned tissue slice. By use of an image analysis software (Metamorph), the branching points of vascular structures within the three-dimensional projection were identified and counted in relation to the size of the respective EB. Moreover, other settings of CLSM were used to identify  $\alpha$ -actinin-positive cardiac cell areas (figure 2.3).



**Figure 2.3: Cardiovascular differentiation of mES cells.** (A): Cardiac cell structure stained with an antibody directed against  $\alpha$ -actinin. (C): Vascular cell structures labeled with an antibody directed against vascular CD31. (B): Overlay image of cardiac (green) and vascular (white) cell structures. The scales bar in A, B, and C represents 200 $\mu$ m.

### 2.2.3 Measurement of ROS generation

Intracellular ROS generation was quantified using H<sub>2</sub>DCF-DA, which is the most widely used cell-permeable dye for analysis of the intracellular redox state. Intracellularly, H<sub>2</sub>DCF-DA is hydrolyzed by cellular esterases to yield the DCFH carboxylate anion, which is trapped in the cells. DCFH is rapidly oxidized into the fluorescent product (DCF) 2, 7-dichlorofluorescein by intracellular ROS. To measure intracellular ROS, the EBs were incubated with 20 $\mu$ M H<sub>2</sub>DCF-DA for 30 min in the dark on a shaker at 37 °C, 5% CO<sub>2</sub>. After staining EBs were washed once with fresh serum-free Iscove's basal medium and immediately transferred into the slide chamber and monitored using CLSM. Fluorescence excitation was performed using the 488nm line of an argon-ion laser, and emission was recorded at 515-550nm (Bartsch et al., 2011; Sharifpanah et al., 2016).

### 2.2.4 Measurement of NO generation

To evaluate the intracellular NO concentration within EBs, the EBs were loaded with 10 $\mu$ M of the cell-permeable specific fluorescent NO indicator DAF-FM DA and incubated for 30 min on a shaker in the dark at 37 °C, 5% CO<sub>2</sub>. Subsequently, the EBs were washed with serum-free Iscove's basal medium and further incubated for 30 min. The EBs were

## Materials and Methods

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transferred to the slide chamber which was fixed on the stage of the CLSM. DAF fluorescence was excited at 488nm and emission recorded at 515-550nm (Mascheck et al., 2015; Sharifpanah et al., 2019).

### 2.2.5 Recording of the intracellular calcium concentration

To measure intracellular calcium transients, single cells were loaded with the cell-permeable dye Fluo-4-AM. Initially, 3-day-old EBs were enzymatically dissociated by using PBS containing 2 mg/ml of collagenase B for 30 min at 37 °C. Single cells were seeded on (0.1%) gelatin-coated coverslips in 24-well cell culture plates and incubated in CCE differentiation medium at 37 °C, 5% CO<sub>2</sub> and relative humidity of 95%. After 24h, single cells were attached, and were then incubated in serum-free Iscove's basal medium containing 5µM Fluo-4-AM at 37 °C, 5% CO<sub>2</sub> for 30 min. Subsequently, the coverslips were transferred to an incubation slide chamber containing fresh serum-free Iscove's Basal Medium, and then transferred to the stage of the CLSM. Cells were superfused with PUFAs after 300 s of recording. For fluorescence excitation the 488 nm a wavelength of an argon laser was used. Emission was recorded at 500–550nm. Fluorescence images were acquired every 3 s (Mascheck et al., 2015; Sharifpanah et al., 2016).

### 2.2.6 Western blot assay

Western blotting is the most common and widely used assay to detect protein expression qualitatively and quantitatively in biological specimen such as cells or tissues. Western blotting was performed according to a previously published protocol (Ali et al., 2018).

#### 2.2.6.1 Protein extraction

After the EBs were prepared, they were collected in 1.5ml micro-tubes, the medium was aspirated and EBs were washed with ice-cold PBS to remove the rest of cell culture medium. Subsequently 100µl of protein extraction buffer (lysis-buffer) was added, which was prepared freshly. The EBs were grinded with a pellet pestle in the 1.5ml micro-tube, incubated for 20 min on ice, then the cells were homogenized again. Subsequently, the micro-tube was centrifuged in a pre-cooled centrifuge for 10 min at +4 °C at 35395g.

## Materials and Methods

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Finally, the supernatant was transferred to another 0.5ml micro-tube and stored at -80 °C till use.

### 2.2.6.2 Lowry protein assay

Lowry protein assay (Lowry et al., 1951) is a biochemical assay, which is used to quantify protein levels in biological samples. This method is dependent on color change of the solution, which can be measured by colorimetric techniques. The sample was prepared by mix-up of solution D, solution E, DW, and protein sample in the ratio 1000:100:100:10, respectively. In parallel, a concentration series of BSA protein standards was prepared. The mixed solutions were incubated for 10 min at RT, then 200µl of solution F was added to 1000µl of the mixed solutions, and then incubated for an additional 30 min at 37 °C. The solutions were centrifuged for 30 min at RT and 35395g. Finally, 200µl of solutions were transferred to a 96-well ELISA plate and absorbance measured by using an ELISA reader (Tecan Infinite m200 Microplate reader) and a wavelength of 578nm.

### 2.2.6.3 Gel electrophoresis

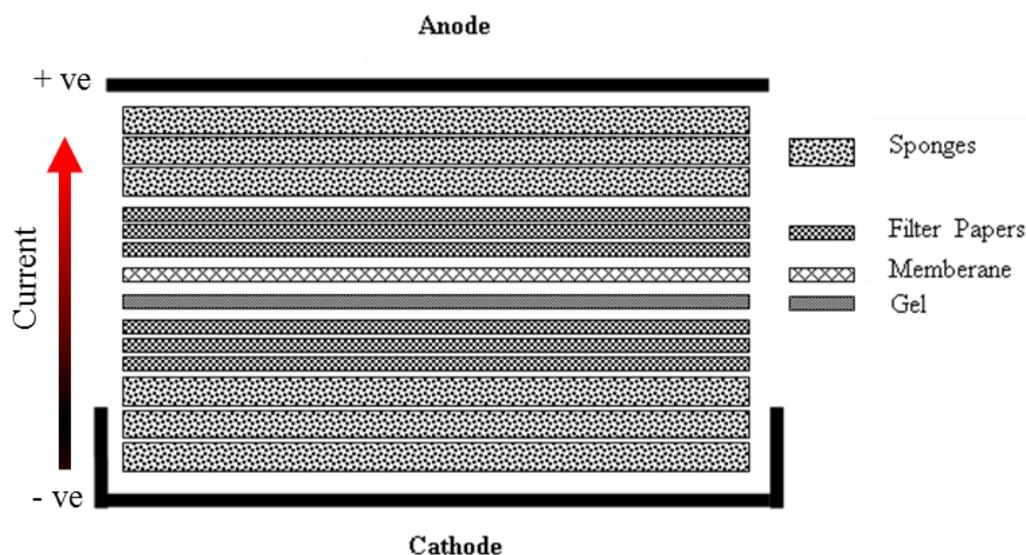
Protein electrophoresis refers to the migration of charged proteins in polyacrylamide gels in an electric field. This leads to protein separation depending on the respective molecular weight. In the present study semi-quantitative western blot was performed. An amount of 20µg of the protein sample was mixed with 1xNuPAGE LDS buffer and 1xNuPAGE reducing agent. Subsequently, the protein sample was denatured by heating for 10 min at 70 °C. The protein samples were loaded into the pockets of 4% stacking PAGER EX polyacrylamide gels, which were mounted in the electrophoresis chamber. A volume of 500µl antioxidant solution was added to the electrophoresis chamber, which was filled with 0.67x of ProSieve EX running buffer. The chamber was connected with the power supply, and 200 voltage (V) and 120 milliampere (mA) were applied for 45-55min.

### 2.2.6.4 Protein transfer from the gel to the PVDF membrane

After completion of gel, electrophoresis proteins were transferred to PVDF membranes with a pore size of 0.2µM. PVDF membranes were wetted for 1 min with methanol, washed with DW, and then transferred to 1xProSieve EX transfer buffer. Additionally,

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filter papers and sponges were soaked in transfer buffer. Subsequently, the transfer sandwich was prepared as illustrated below (figure 2.4). The transfer sandwich was inserted in the blotting chamber (Xcell II Blot module). The chamber was connected with the power supply, which was adjusted at 25V and 375mA for 60-90 min.



**Figure 2.4: Example of sandwich preparation for western blot analysis.**

### 2.2.6.5 Ponceau-S staining.

Ponceau-S staining is utilized to detect successful protein transfer onto the PVDF membrane. The membrane was soaked in the Ponceau-S staining solution for 5 min to visualize protein bands.

### 2.2.6.6 Protein detection

In this step, the target protein bands were detected on the membrane by immune-staining with specific antibodies. Ponceau-S staining solution was removed by washing the membrane with 0.1% TBST for blocking nonspecific protein binding. The membrane was incubated in blocking buffer (5% nonfat- dry milk in 0.1% TBS-Tween) at RT for 60 min on a shaker. After that, the membrane was incubated on a shaker overnight in primary antibody solution at +4 °C, and then for 60 min at RT with secondary antibody. The primary and secondary antibodies were previously diluted with 5% BSA in ratio as

## Materials and Methods

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described in 2-1-9. Between incubation with primary and secondary antibody the membranes were washed at least 3 times for 5 min with 0.1% TBS-Tween (Table 11).

**Table 11: Summary of the antibody staining protocol on PVDF membranes**

Step	Solution	Duration	Temperature
Washing	0.1% TBS- Tween	3x5 min	RT
Blocking	5% nonfat- dry milk	60 min	RT
Primary antibody	Diluted in 5% BSA	Overnight	+4 °C
Secondary antibody	Diluted in 5% BSA	60 min	RT

### 2.2.6.7 ECL detection

ECL detection is a method depending on an enzymatic reaction between horseradish peroxidase (HRP)-labelled secondary antibody and substrate, which consists of a stable peroxide solution and an enhanced luminol solution. This reaction generates a light signal which can be detected using a chemiluminescence imaging system. The PVDF membrane with labelled proteins was soaked in 10ml of freshly prepared ECL detection solution for ~1 min with gentle agitation. Subsequently, the membrane was wrapped in a plastic container, filled with ECL detection solution, and placed into the chemiluminescence imaging system. The luminescence light signal of the membranes was recorded under darkroom conditions, and saved as tagged image file format (TIFF).

### 2.2.6.8 Densitometry image analysis

The image was analyzed using the densitometry. Densitometric signal strength depends on the grey level intensity of protein bands in images. The signal intensity of protein expression in images was converted into a digital value by use of the Java software Image J. The value for relative specific protein expression was normalized with the internal loading control, which is an endogenous protein and is unaffected by experimental conditions or treatments.

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### 2.2.6.9 Stripping and reprobing

Stripping refers to removal of the primary and secondary antibody from the western blot membrane by using the stripping buffer. This method was used for investigating more than one protein on the same blot membrane. The membrane was soaked in restore western blot stripping buffer for 20 min at RT with gently shaking. Subsequently, the membrane was washed with 0.1% TBS-Tween three times for 5 min. After this procedure, the membrane is ready for re-staining with another antibody by the same steps as mentioned in 2-2-6-6.

### 2.2.7 Statistical analysis

For statistical analysis, PRISM statistics software (GraphPad Software, San Diego, CA) was used. Data are given as mean  $\pm$  standard deviation (SD), with  $n$  denoting the number of experiments performed with independent ES cell cultures. In each experiment, at least 20 culture objects were analyzed unless otherwise indicated. Student's t-test for unpaired data and one-way analysis of variance (ANOVA) (Tukey or Dunnett multiple comparison tests) were applied as appropriate for statistical analysis. A value of  $P \leq 0.05$  (\*) or (#) was considered significant.

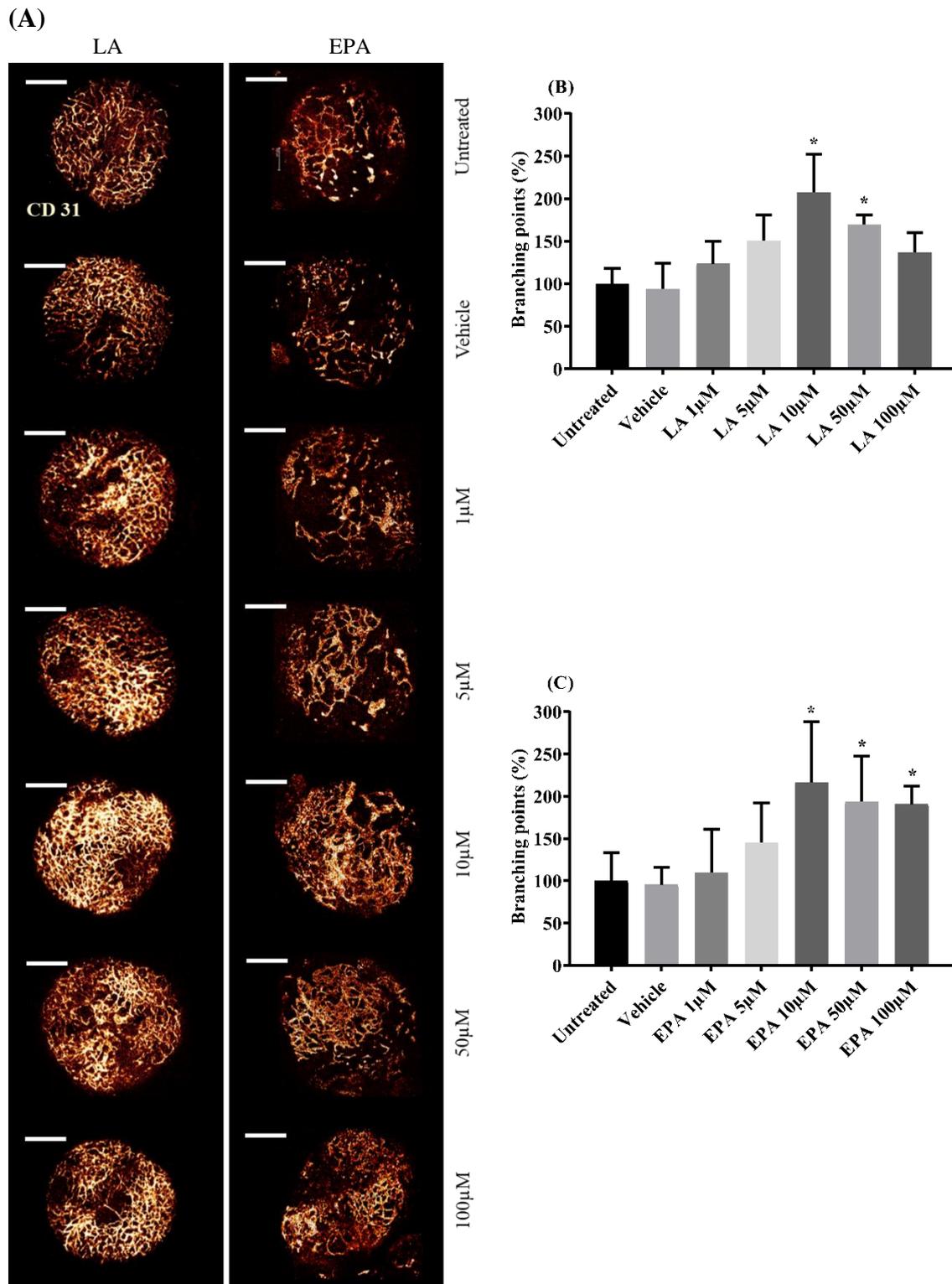
### 3. Results

#### 3.1 Vasculogenesis upon PUFA treatment in EBs

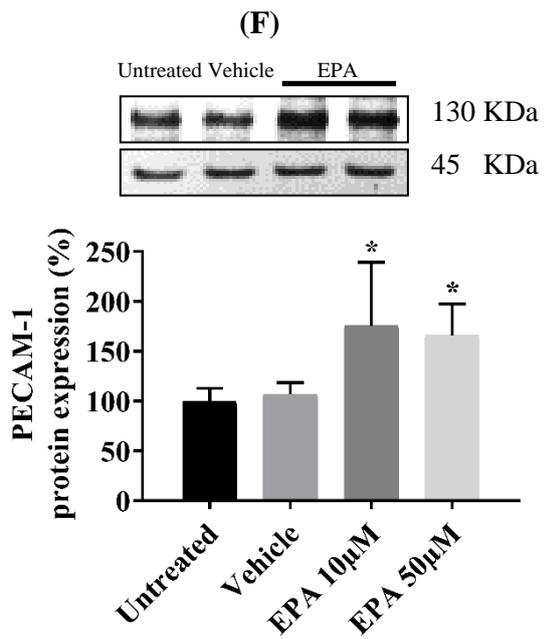
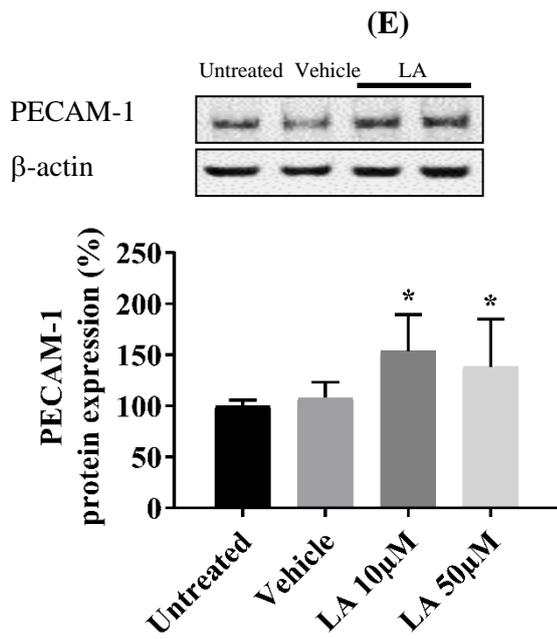
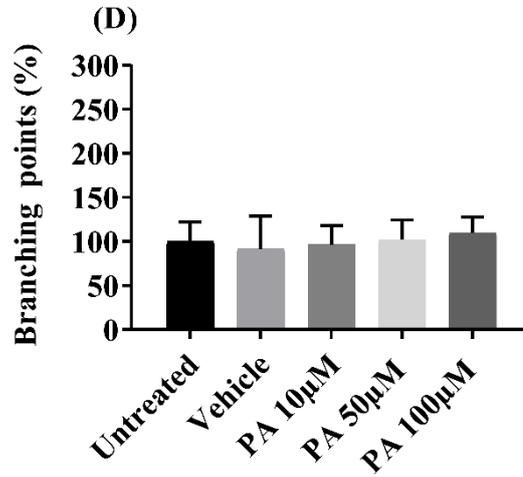
##### 3.1.1 LA and EPA stimulated vasculogenesis in ES cell-derived EBs

Vasculogenesis refers to the differentiation of endothelial cells from angioblasts and endothelial precursor cells, in addition to the *de novo* formation of a primitive vascular network (Kolte et al., 2016). According to our working hypothesis, PUFAs may regulate ES cell differentiation towards the cardiovascular lineage. The effect of PUFAs and PA on vascular sprout formation was evaluated. EBs were formed in spinner flask culture, and were transferred on day 3 of cell culture to bacteriological plates. EBs were incubated with either EPA or LA in concentrations ranging from 1 to 100 $\mu$ M, and vascular structure formation as well as protein expression of the vascular markers PECAM-1 and VE-Cadherin were investigated. EPA as well as LA treatment dose-dependent significantly increased vascular structure formation (figure 3.1A–C) and the expression of vascular markers (figure 3.1E–H). In contrast, treatment with PA (10, 50, 100 $\mu$ M) did not affect vasculogenesis as compared with the untreated and the vehicle control (figure 3.1D). Optimum effects on vasculogenesis were achieved with 10–50 $\mu$ M EPA and LA. Moreover, increased expression of FLK-1, which is an early marker of vascular progenitor cells, was observed upon either EPA or LA treatment (figure 3.1I, J), suggesting stimulation of vasculogenesis in ES cells upon incubation with PUFAs.

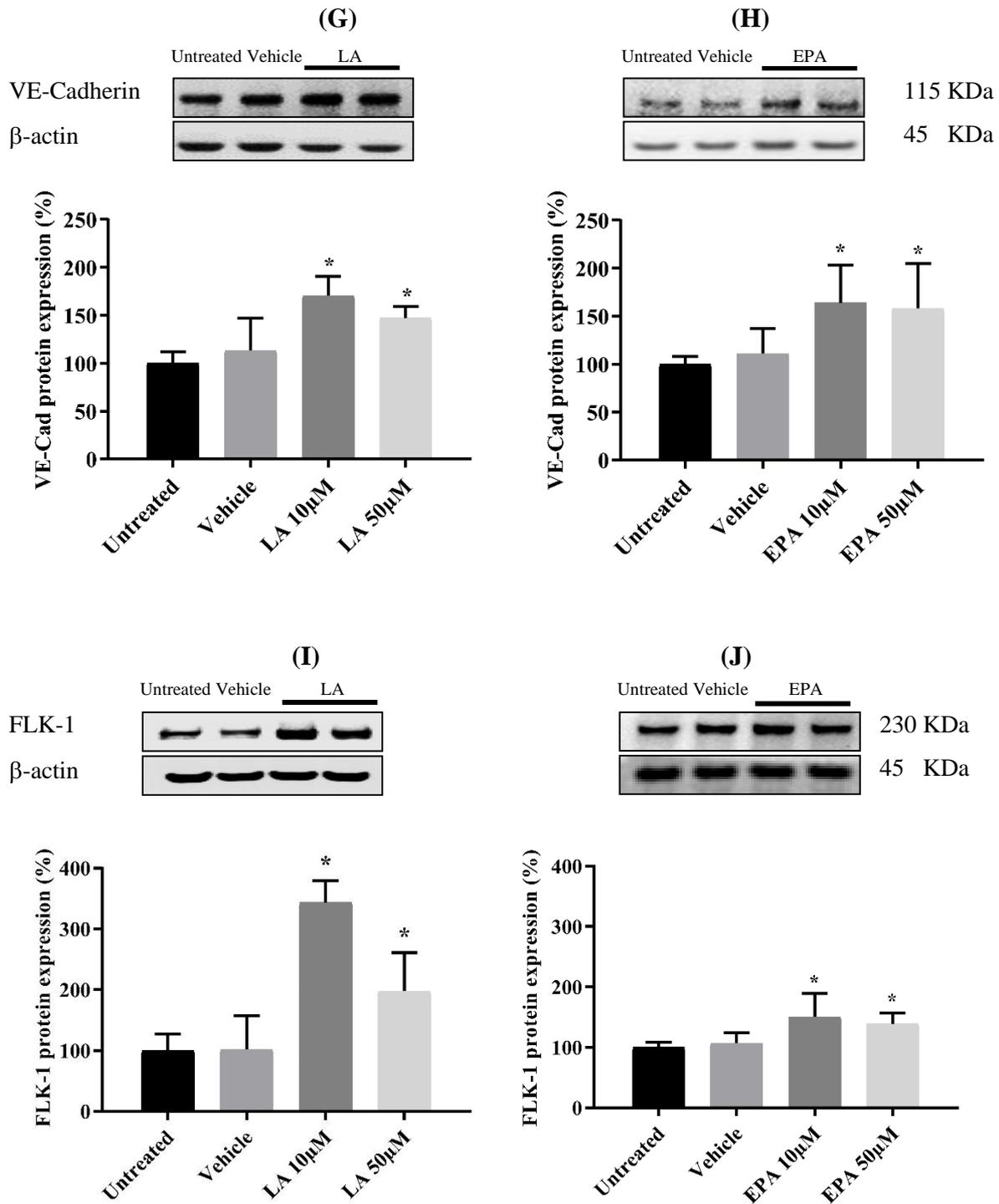
## Results



## Results



## Results



**Figure 3.1: Stimulation of vasculogenesis and vascular marker expression by EPA and LA.** ES cells differentiating within EBs were treated from day 3 to 10 of cell culture with increasing concentrations (1–100µM) of either EPA or LA, remained untreated or were treated with vehicle (DMSO). Vascular structures were stained with an anti-PECAM-1 antibody, and branching points were analyzed by computer-assisted image

## Results

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analysis. (A) Representative PECAM-1 labeled EBs. The bar represents 300 $\mu$ m. (B–D) Quantitative presentation of branching points (%) following incubation with different concentrations of (B) LA (n = 3), (C) EPA (n = 6) or (D) PA (n = 3). (E and F) Western blot analysis of PECAM-1 expression upon treatment with either (E) 10 or 50 $\mu$ M LA (n = 9) or (F) EPA (n = 7). (G and H) Western blot analysis of VE-Cadherin expression upon treatment with either (G) 10 or 50 $\mu$ M LA (n = 4) or (H) EPA (n = 7). (I and J) Western blot analysis of fetal liver kinase-1 (FLK-1) expression upon treatment with either (I) 10 or 50 $\mu$ M LA (n = 4) or (J) EPA (n = 6). \*P  $\leq$  .05, significantly different to the vehicle and control.

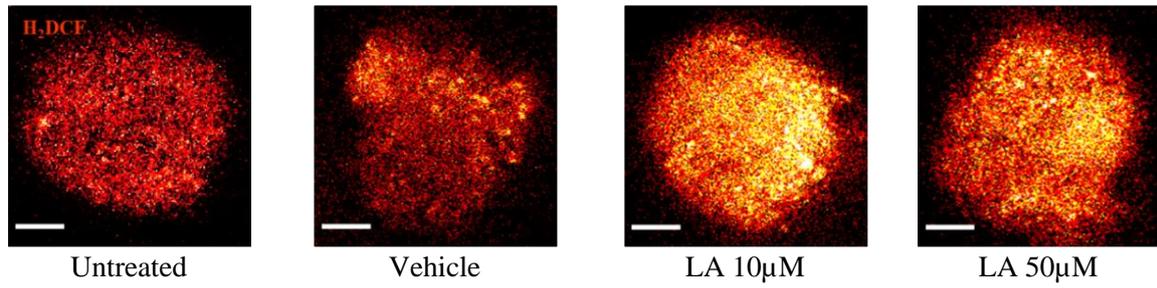
### 3.1.2 PUFAs-induced signaling pathway elements in ES cells.

#### 3.1.2.1 Generation of ROS and NO upon treatment of ES cells with PUFAs

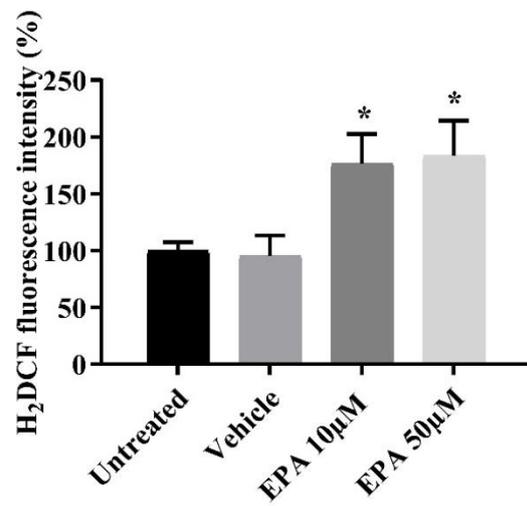
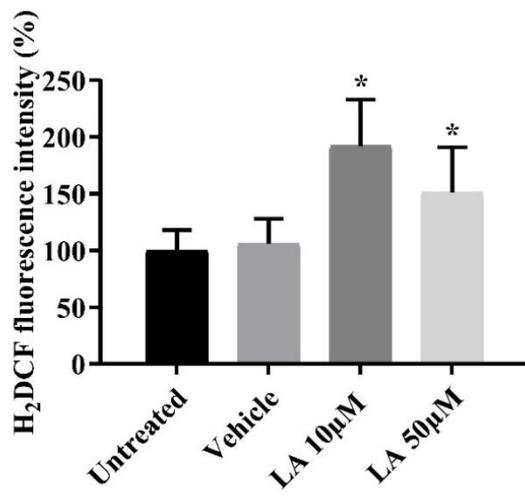
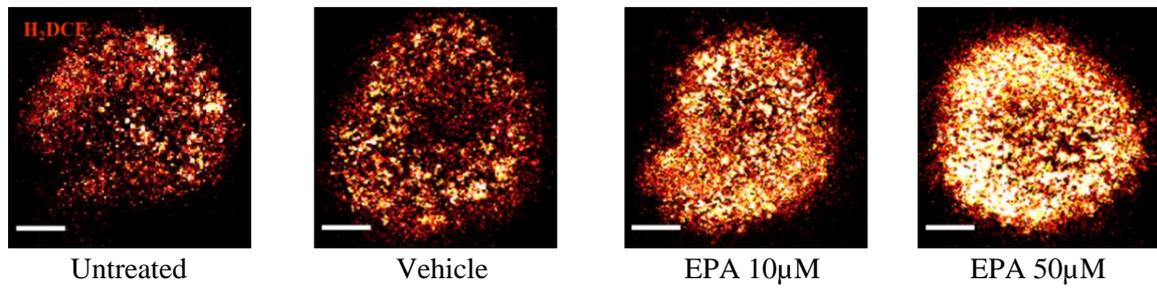
Previous studies of us and others have demonstrated that cardiovascular differentiation of ES cells is dependent on ROS and NO, acting as signaling molecules in intracellular signal transduction pathways (Ali et al., 2018; Cencioni et al., 2018; Sharifpanah et al., 2016). We therefore assessed ROS generation using the ROS-sensitive indicator H<sub>2</sub>DCF-DA as well as the NO-sensitive indicator DAF-FM for ROS and NO determination, respectively. Moreover, activation of eNOS was evaluated upon treatment with either EPA or LA. It was observed that treatment with EPA and LA (10 and 50 $\mu$ M) increased ROS generation (figure 3.2A, B). Investigation of the time course of ROS generation by LA revealed that ROS production occurred within few min after PUFA administration and remained on an elevated plateau for at least 24h (figure 3.2C). Coadministration of EPA and LA with the free radical scavengers Trolox (100 $\mu$ M) and NMPG (100 $\mu$ M) completely abolished the stimulation of vascular structure formation obtained upon PUFA treatment (figure 3.2D-F). Moreover, the stimulation of vasculogenesis by PUFAs was significantly inhibited in presence of the NADPH oxidase inhibitor VAS2870 (50 $\mu$ M; figure 3.2D–F). PUFA treatment of differentiating ES cells likewise increased NO generation (figure 3.3A–C) and increased eNOS phosphorylation at residue Ser1177 within few min (figure 3.3D, E). The phosphorylation of eNOS as well as vascular structure formation were completely blunted in presence of the NOS inhibitor L-NAME (figure 3.3F, G), indicating the importance of NO in parallel to ROS for the induction of vasculogenesis by PUFAs.

## Results

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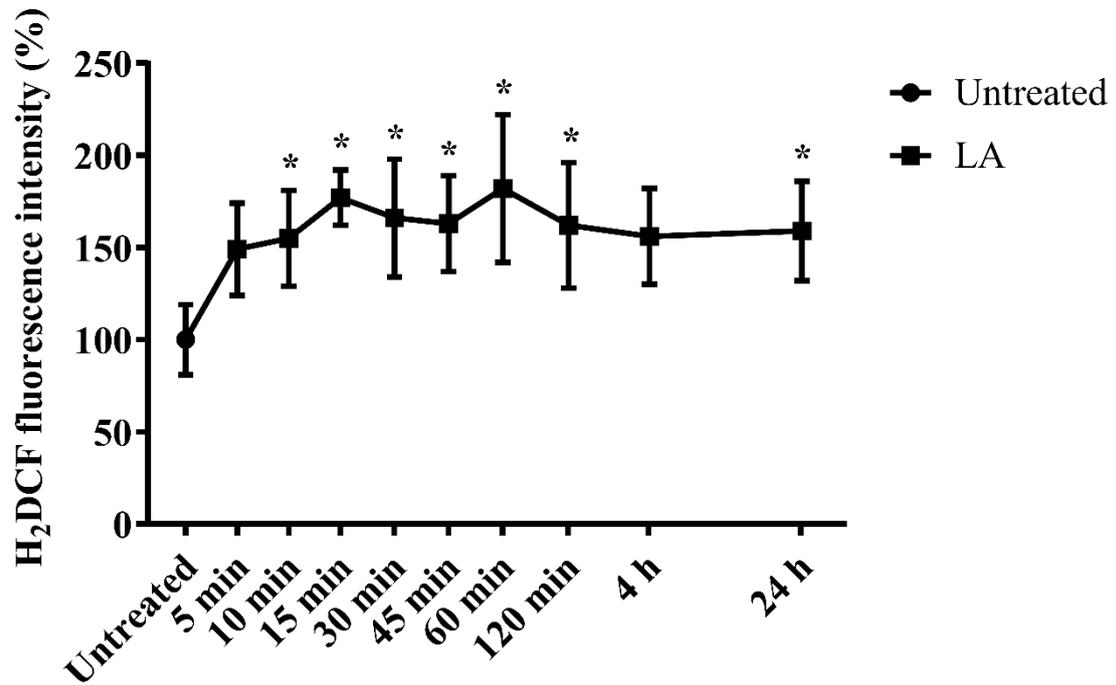
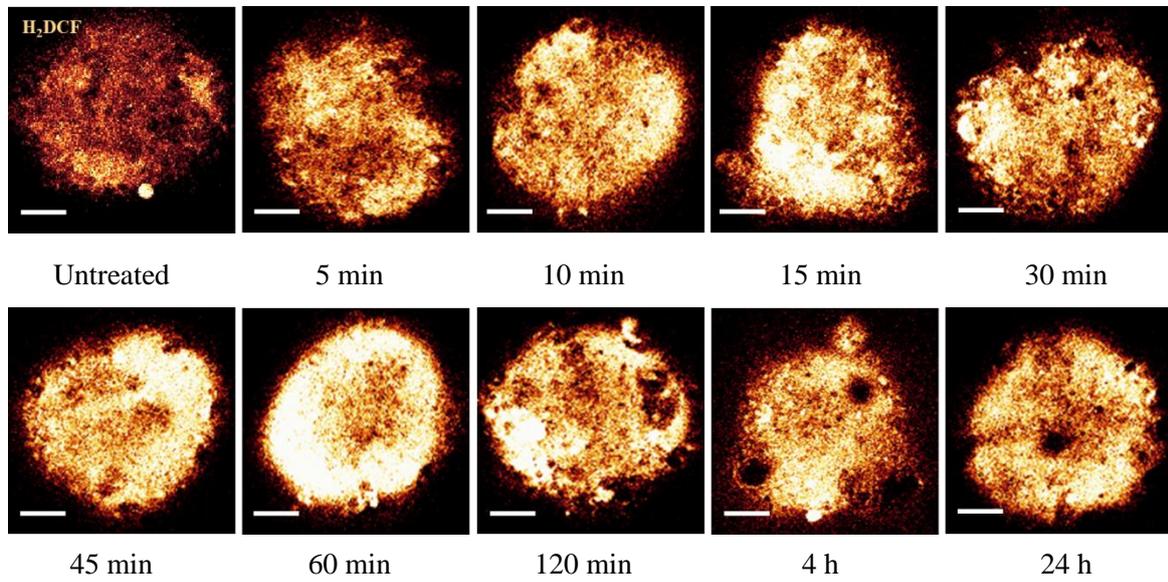


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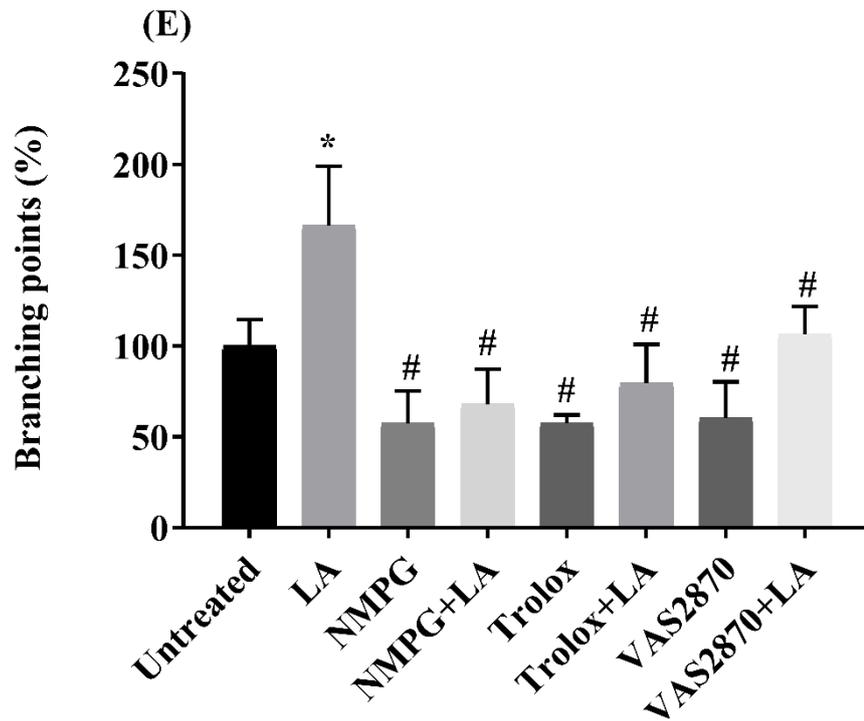
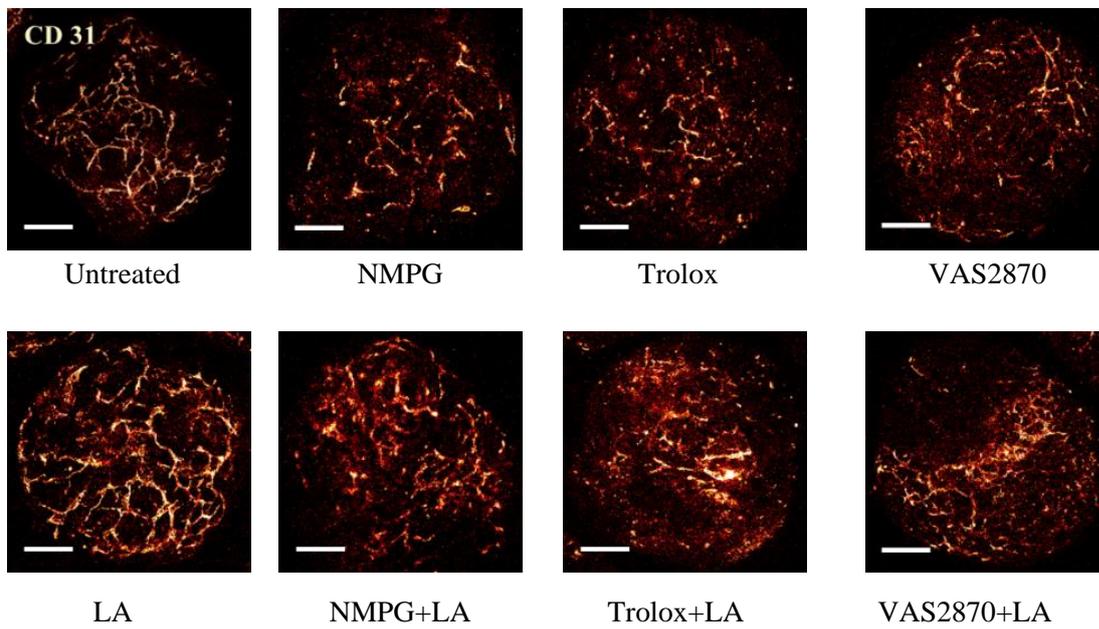
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(C)

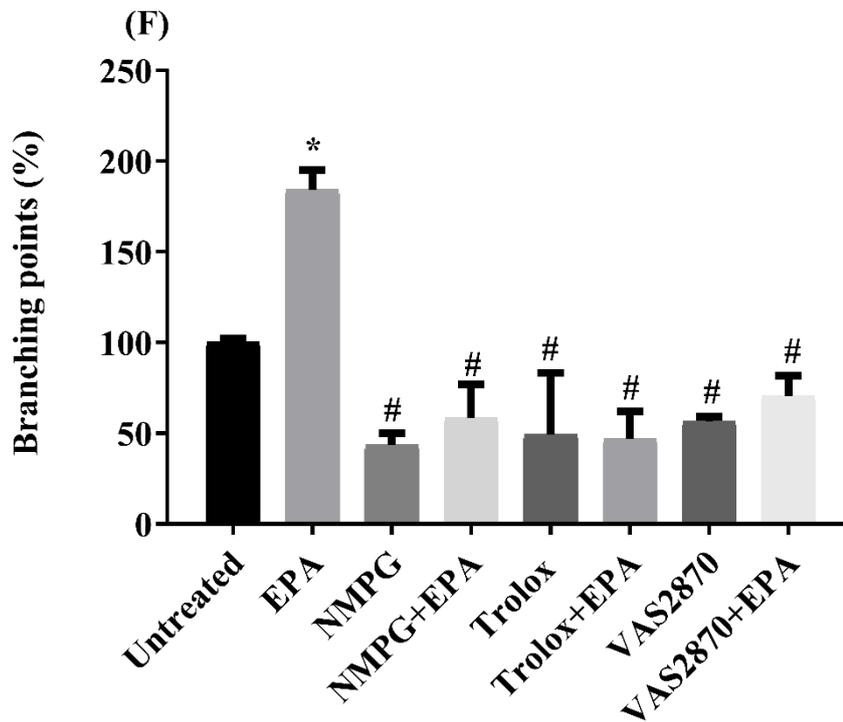


## Results

(D)

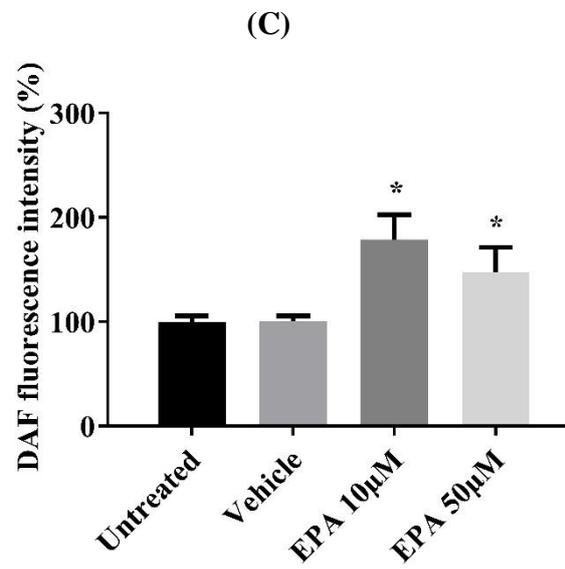
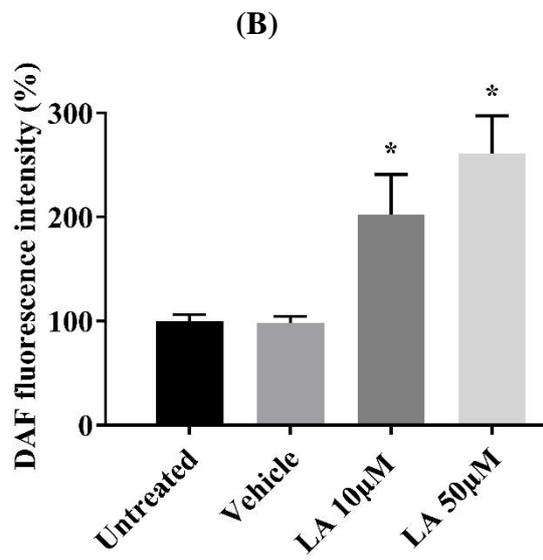
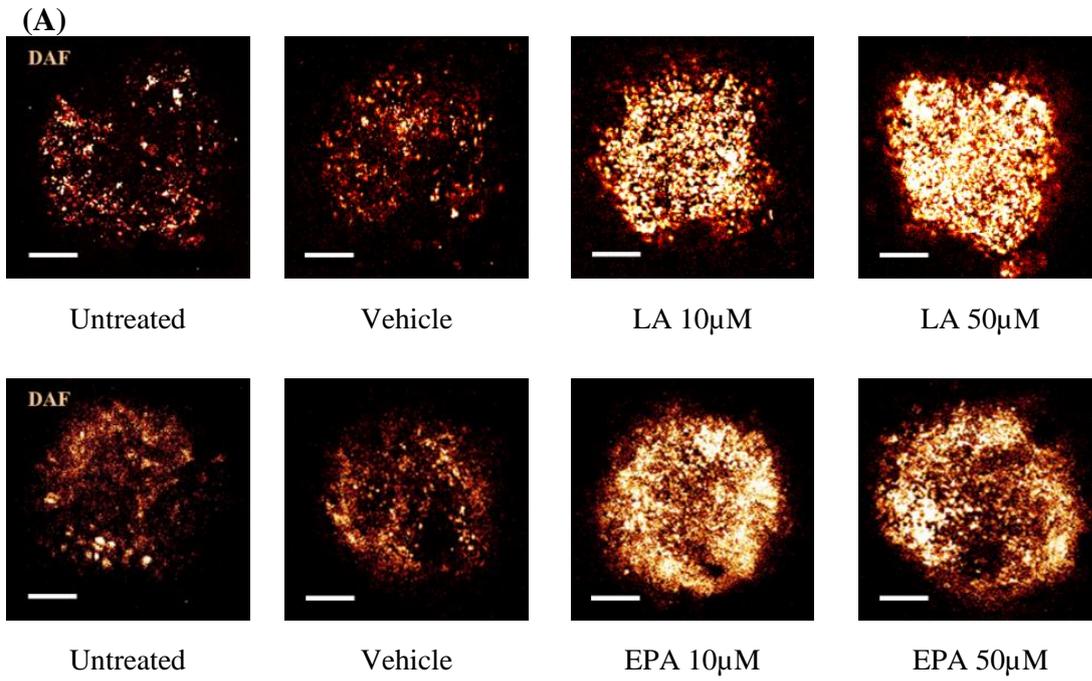


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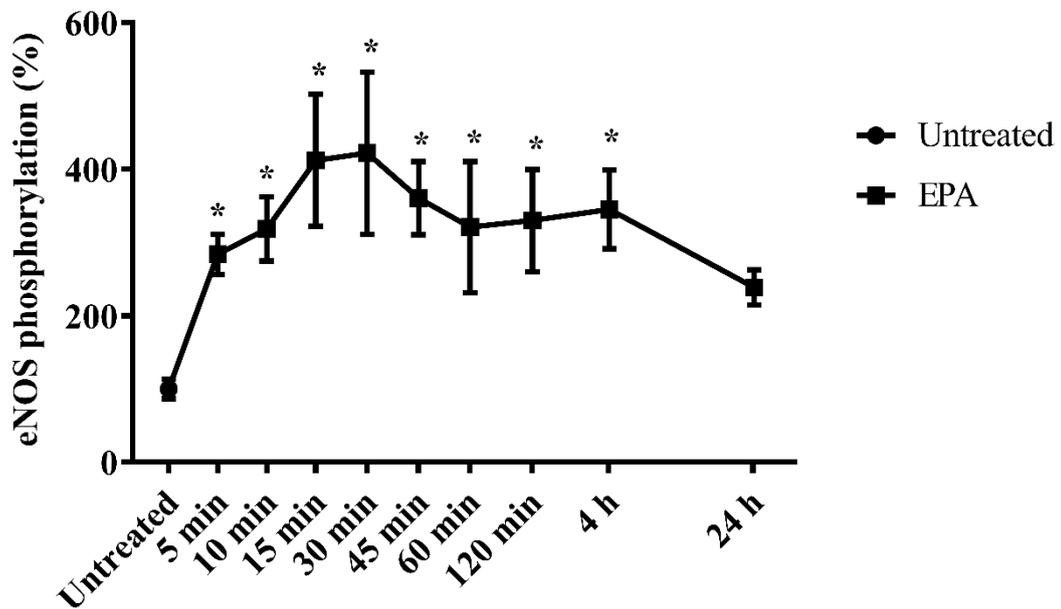
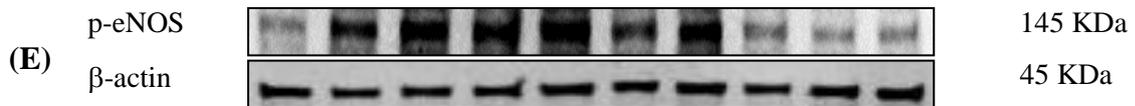
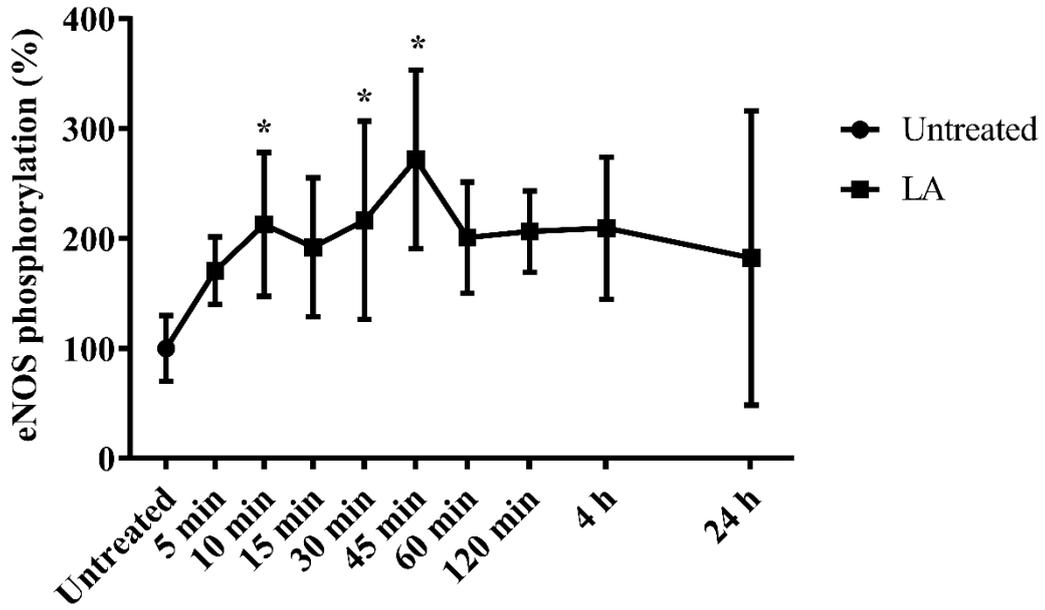
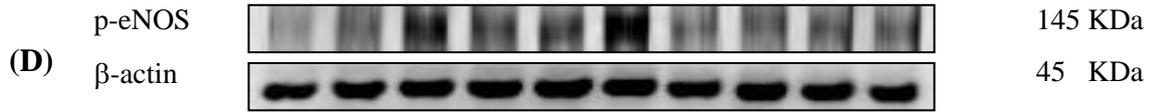


**Figure 3.2: Generation of ROS in EPA as well as LA treated differentiating ES cells and effects of free radical scavengers and NADPH oxidase inhibition on vasculogenesis.** (A and B) EBs (4-day-old) were treated with either 10 or 50 $\mu$ M LA (A) or EPA (B) or vehicle (DMSO), and ROS generation was assessed 24h thereafter using H<sub>2</sub>DCF as fluorescent ROS indicator (n = 3). Shown are representative DCF-labeled EBs. The bar represents 200 $\mu$ m. (C) Time course of ROS generation after different min of LA (50 $\mu$ M) incubation (n = 3). (D–F) Effect of the free radical scavengers NMPG (100 $\mu$ M) and Trolox (100 $\mu$ M) or the NADPH oxidase inhibitor VAS2870 (50 $\mu$ M) on vascular branch formation. (D) Representative EBs stained with an antibody against CD31. The bar represents 200 $\mu$ m. (E and F) Quantitative analysis of vascular branching points (%) following treatment with (E) LA (50 $\mu$ M; n=3) or (F) EPA (n=3). \*p  $\leq$  .05, significantly different to the untreated control, #p  $\leq$  .05, significantly different to the PUFA treated sample.

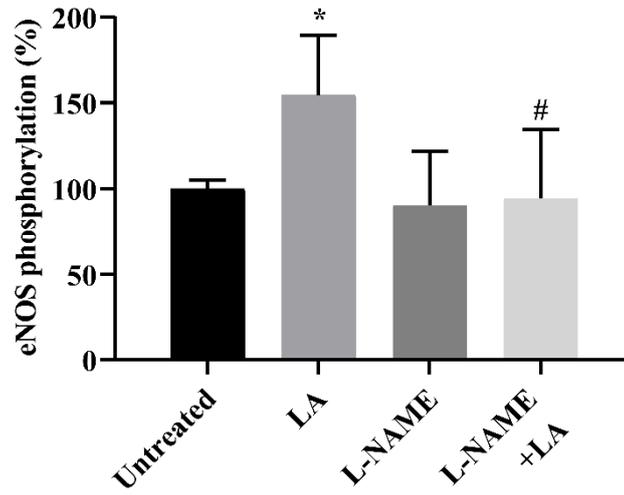
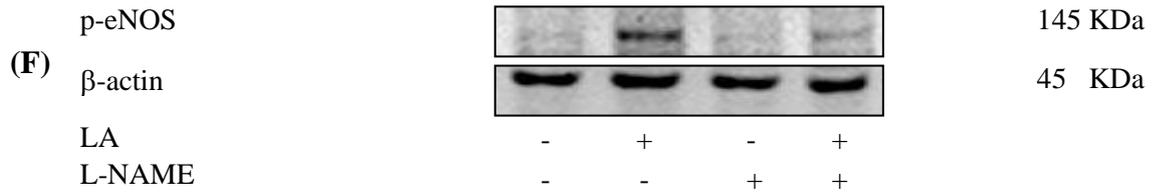
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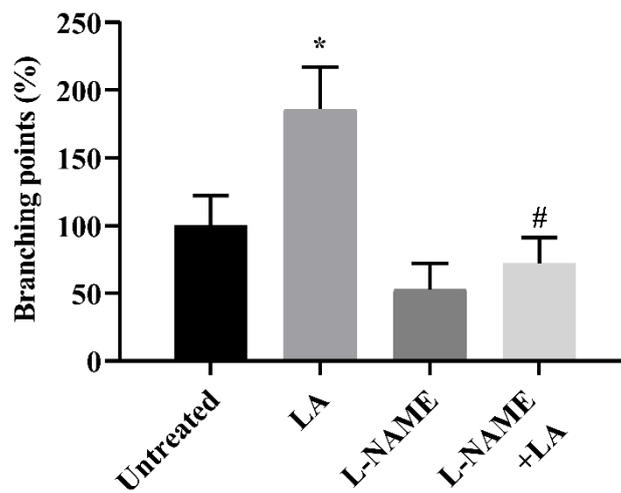
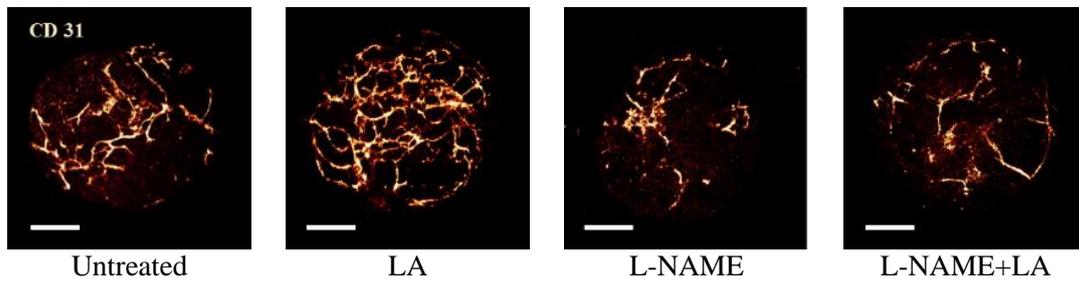
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## Results



(G)



## Results

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**Figure 3.3: Generation of NO in EPA as well as LA treated differentiating ES cells and effect of the NOS inhibitor L-NAME on vasculogenesis.** (A–C) EBs (4-day-old) were treated with either (A and B) 10 or 50 $\mu$ M LA or (A and C) EPA or vehicle (DMSO), and NO generation was assessed 24h thereafter using DAF-FM as fluorescent NO indicator. In (A) representative EBs labeled with DAF-FM are shown which were treated with either 10 or 50 $\mu$ M LA and EPA, respectively. The bar represents 200 $\mu$ m. (B and C) Quantification of DAF fluorescence (%) following treatment with (B) LA (n=3) and (C) EPA (n = 4). (D and E) Western blot analysis of eNOS phosphorylation at different time points of (D) LA (50 $\mu$ M; n=6) and (E) EPA (50 $\mu$ M; n=3) application. (F) Effect of the NOS inhibitor L-NAME on eNOS phosphorylation following treatment of EBs with LA (50 $\mu$ M) as evaluated by western blot analysis (n=6). (G) Effect of L-NAME on vasculogenesis (PECAM-1 staining) elicited upon treatment of differentiating ES cells with LA (50 $\mu$ M; n=3). \*P  $\leq$  05, significantly different to the untreated control, #P  $\leq$  .05, significantly different to the PUFA treated sample.

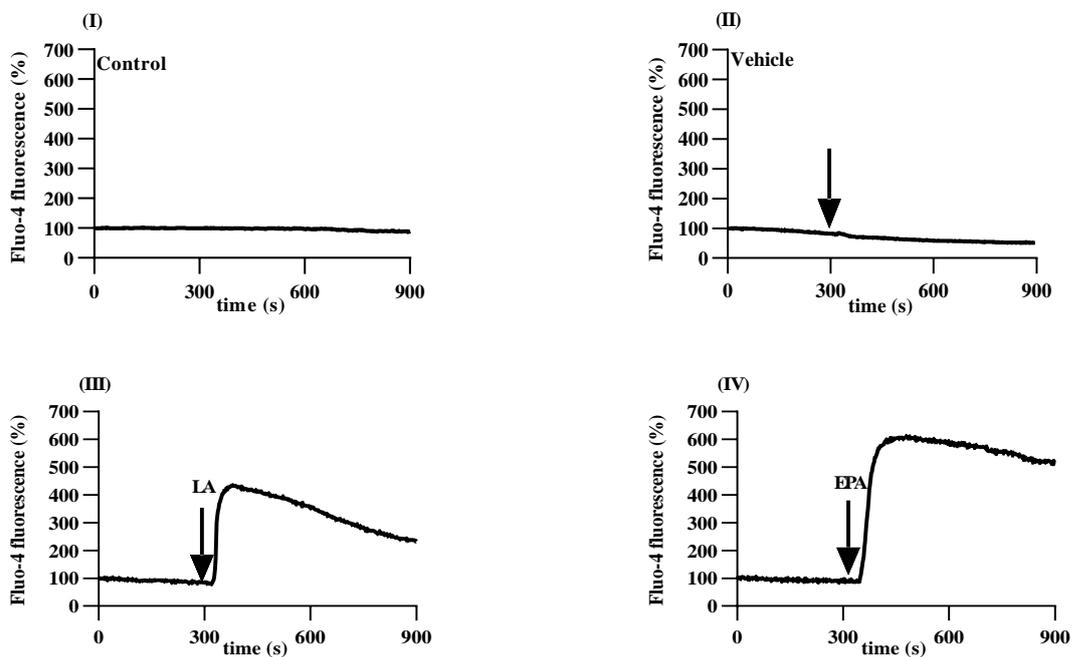
### 3.1.2.2 Release of intracellular calcium upon PUFA treatment

$\Omega$ -3 as well as  $\omega$ -6 PUFAs have been previously shown to raise intracellular calcium in different cells (Carrillo et al., 2011; Kim et al., 2015; Min Hee Kim et al., 2009; Moberaten et al., 2013). To investigate whether changes in intracellular calcium concentrations relay PUFA treatment to the cellular messengers ROS and NO as well as to vasculogenesis of ES cells, intracellular calcium changes were assessed by Fluo-4 microfluorometry. It was evidenced that both EPA and LA (50 $\mu$ M) treatment transiently raised intracellular calcium concentrations (figures 3. 4A, III, IV). In contrast, there was no noticeable change in the intracellular Ca<sup>2+</sup> level in the untreated group as well as the vehicle (DMSO) control (figures 3.4A, I, II). Pretreatment for 30 min with the intracellular calcium chelator BAPTA-AM (10 $\mu$ M) completely abolished the calcium response observed with PUFAs, indicating that calcium was released from intracellular stores (figures 3.4A, VII, VIII). CD36 imports fatty acids inside cells and is a member of the class B scavenger receptor family of cell surface proteins (Glatz & Luiken, 2018). Since PUFA entry via CD36 may initiate the intracellular calcium response, it was investigated whether the CD36 inhibitor (SSO; 200 $\mu$ M) would affect PUFA-induced calcium transients. Indeed, the increase in intracellular calcium was associated to PUFA entry into the cell via CD36 since the calcium response was significantly reduced by SSO (figures 3.4A, V, VI). When cells were pre-incubated for 24h with a blocking antibody against CD36, the calcium response was decreased in amplitude at an antibody concentration of 2 $\mu$ g/ml (figures 3.4A, IX, X) and

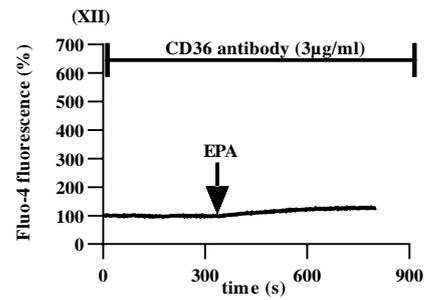
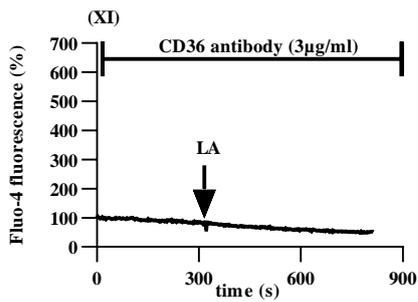
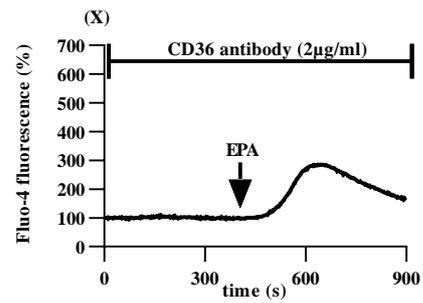
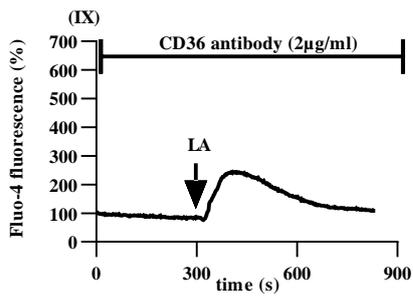
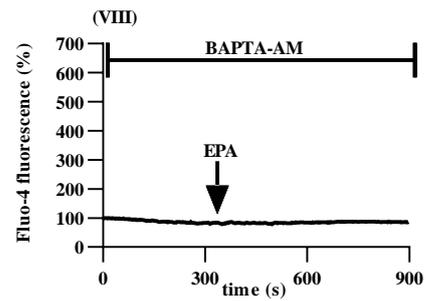
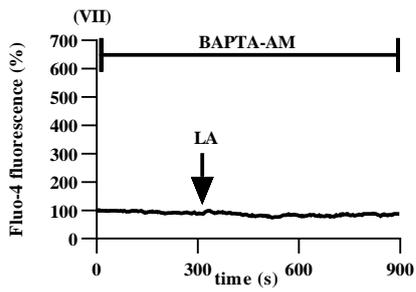
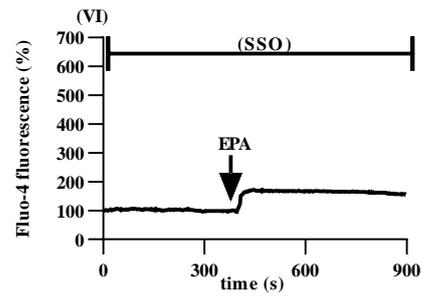
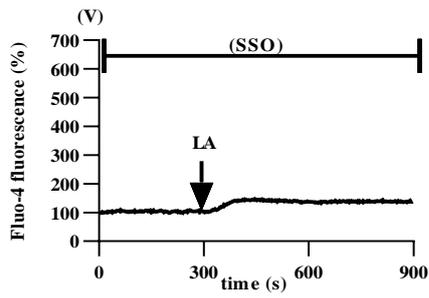
## Results

totally abolished at 3 $\mu$ g/ml (figures 3.4A, XI, XII). Furthermore, to find out, whether ROS generation required intracellular Ca<sup>2+</sup>, ROS generation upon PUFAs treatment was assessed under conditions of intracellular Ca<sup>2+</sup> chelation. 5-day-old EBs were pre-treated for 2h with 10 $\mu$ M of the intracellular calcium chelator BAPTA-AM, and in parallel loaded either in the absence or in the presence (50 $\mu$ M) of PUFAs with the ROS indicator (H<sub>2</sub>DCF-DA). The data show that the chelator BAPTA-AM (10 $\mu$ M) significantly blocked ROS generation stimulated by PUFAs. In other words, the enhancement of ROS generation by PUFAs requires Ca<sup>2+</sup> release from intracellular stores (figures 3.4B, C). Moreover, the intracellular calcium response upon PUFA treatment was necessary for the vasculogenesis process since pre-incubation with BAPTA-AM significantly inhibited vascular structure formation following treatment of differentiating ES cells with either EPA (50 $\mu$ M) or LA (50 $\mu$ M; figure 3.4D, E).

(A)

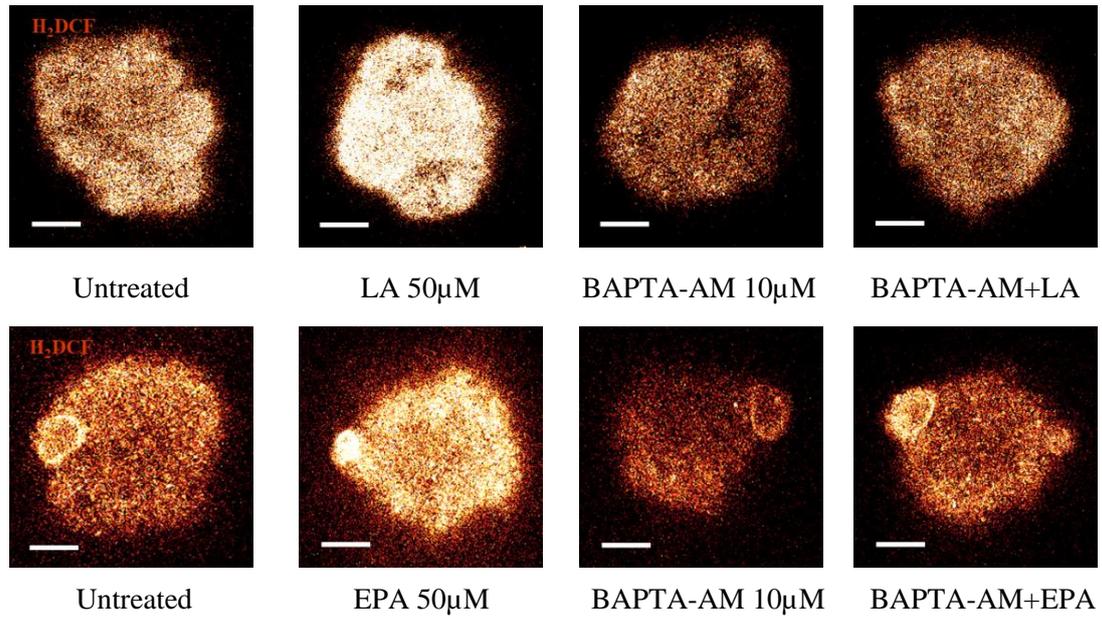


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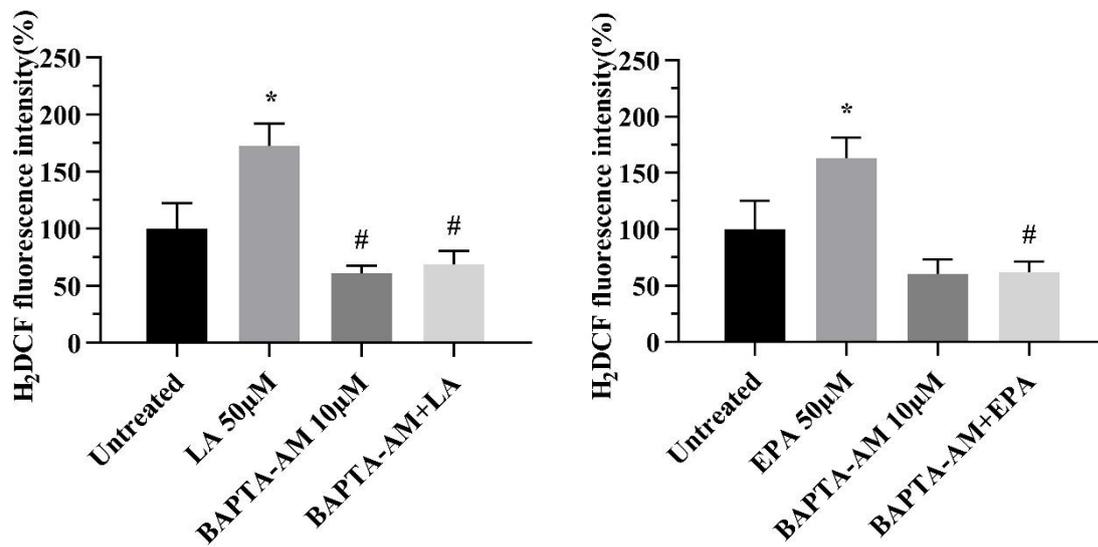


## Results

(B)

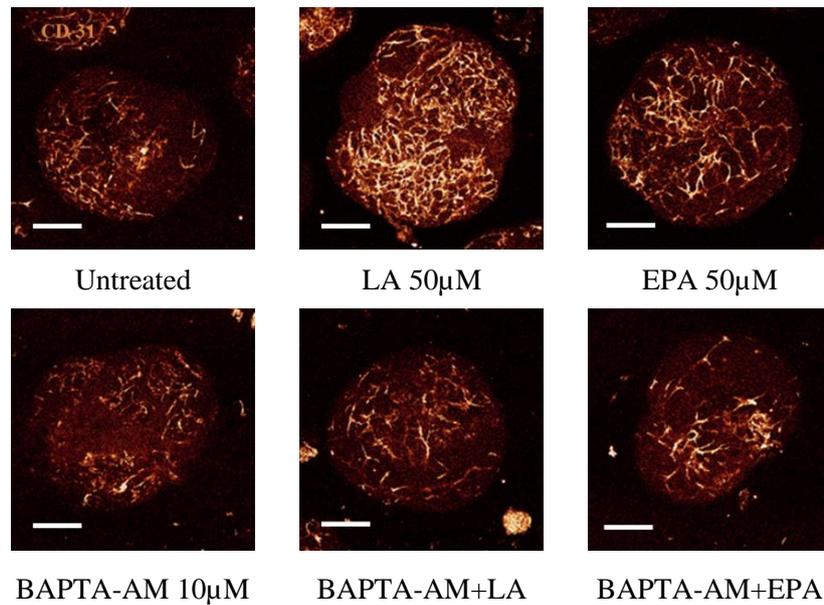


(C)

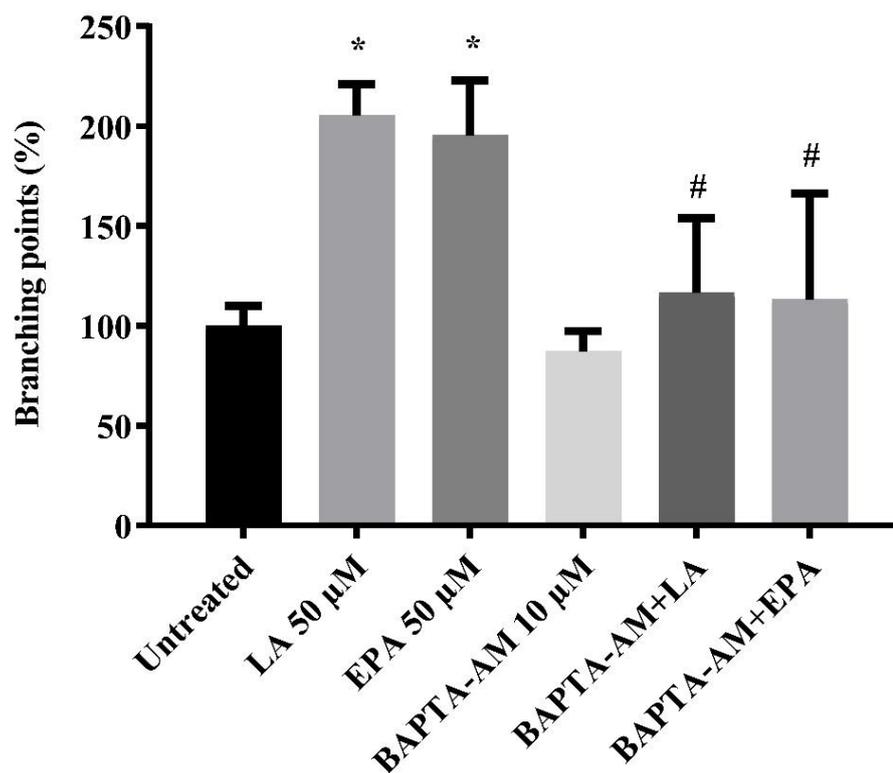


## Results

(D)



(E)



**Figure 3.4: Release of intracellular calcium upon treatment of differentiating ES cells with EPA and LA.** Intracellular calcium was monitored by Fluo-4 microfluorometry in single cells enzymatically dissociated from 3-day-old EBs. (AI) control (n = 4), (AII) vehicle (DMSO; n = 3), (AIII) LA (50 $\mu$ M; n = 3), (AIV) EPA (50 $\mu$ M; n = 3), (AV) LA

## Results

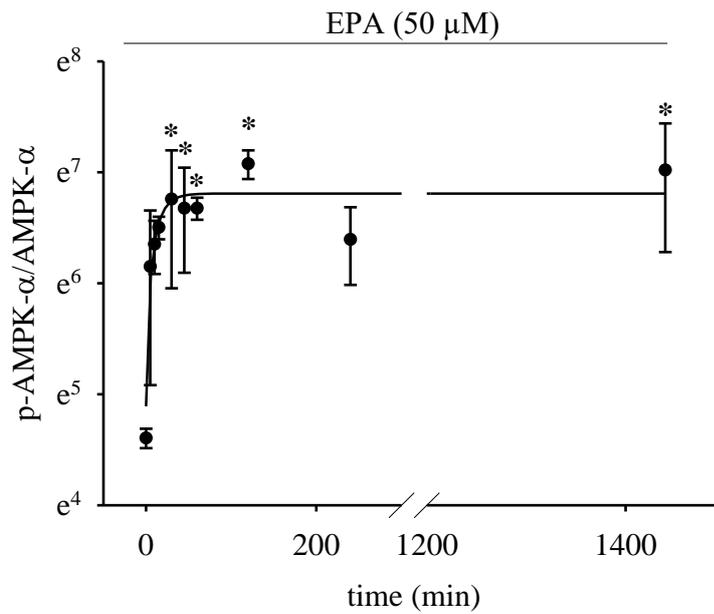
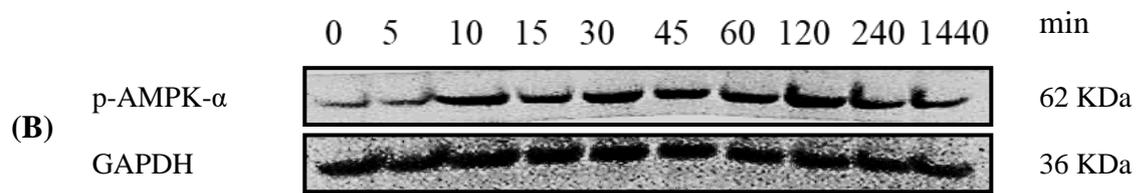
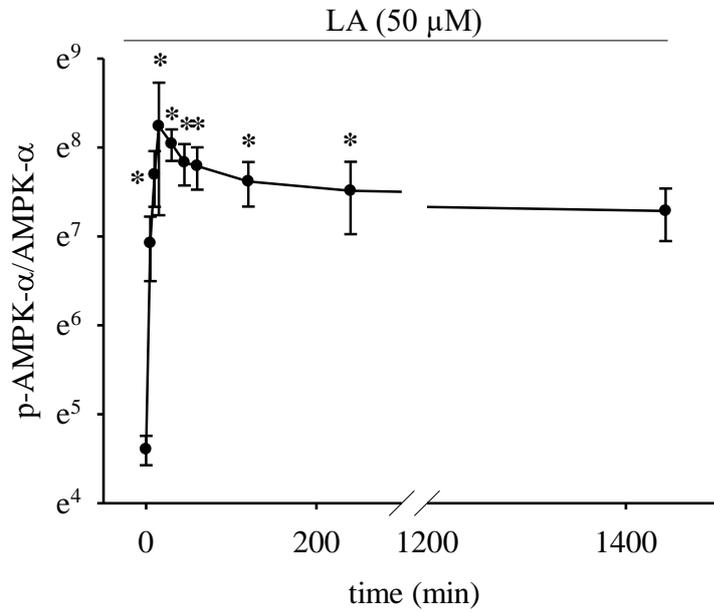
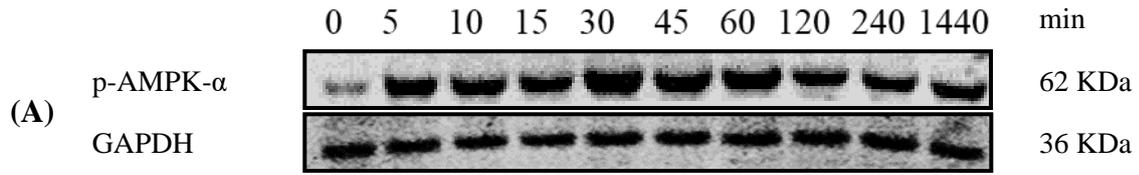
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(50 $\mu$ M) in presence of SSO (200 $\mu$ M; n = 3), (AVI) EPA (50 $\mu$ M) in presence of SSO (200 $\mu$ M; n = 3), (AVII) LA (50 $\mu$ M) in presence of BAPTA-AM (10 $\mu$ M; n = 3), (AVIII) EPA (50 $\mu$ M) in presence of BAPTA-AM (10 $\mu$ M; n = 3), (AIX) LA (50 $\mu$ M), (AX) EPA (50 $\mu$ M) in presence of CD36 antibody (2 $\mu$ g/ml; n = 3), (AXI) LA (50 $\mu$ M), (AXII) EPA (50 $\mu$ M) in the presence of CD36 antibody (3 $\mu$ g/ml; n = 3). (B and C) Inhibition of PUFA-induced ROS generation by intracellular calcium chelation. (B) Representative images of the ROS indicator H<sub>2</sub>DCF-DA, DCF fluorescence intensity was recorded by CLSM. Scale bars represent 200 $\mu$ m, (C) Bar charts show quantification of fluorescence intensity (%) of means  $\pm$  SD of LA (n = 4) and EPA (n = 3) experiments. (D and E) Inhibition of PUFA-induced vasculogenesis by intracellular calcium chelation. EBs were treated from day 3 to 10 of differentiation with either EPA (50 $\mu$ M) or LA (50 $\mu$ M) either in absence or presence of BAPTA-AM (10 $\mu$ M). Vascular structures were visualized by PECAM-1 immunostaining. In (D) representative EBs labeled with an antibody against PECAM-1 are shown. The bar represents 200 $\mu$ m. The quantitative data on vascular branch formation (branching points (%)) (E) were assessed by computer-assisted image analysis of PECAM-1 positive vascular structures (n = 3). \*P  $\leq$  .05, significantly different to the untreated control, #P  $\leq$  .05, significantly different to the PUFA treated sample.

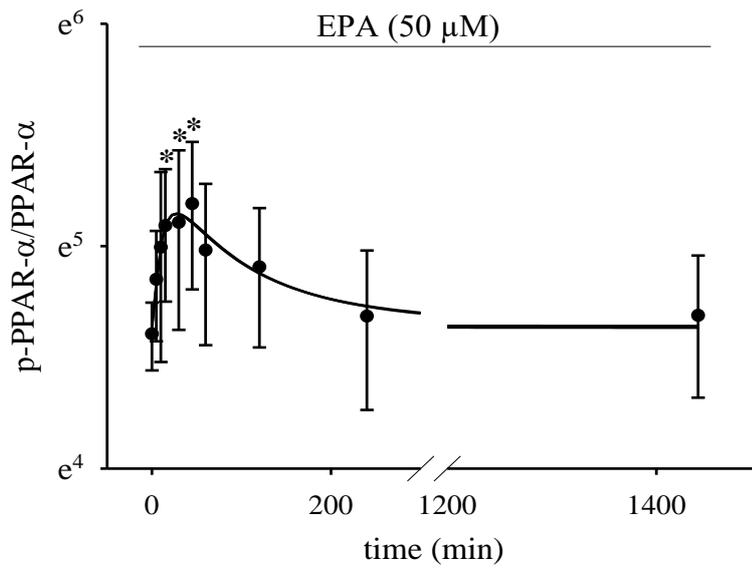
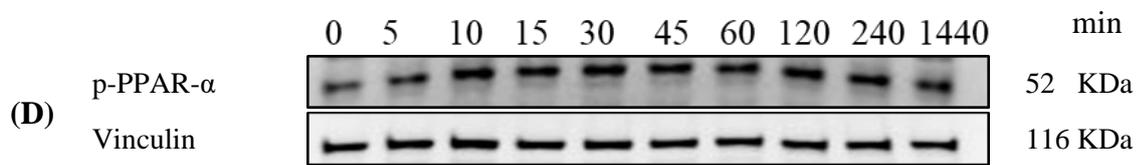
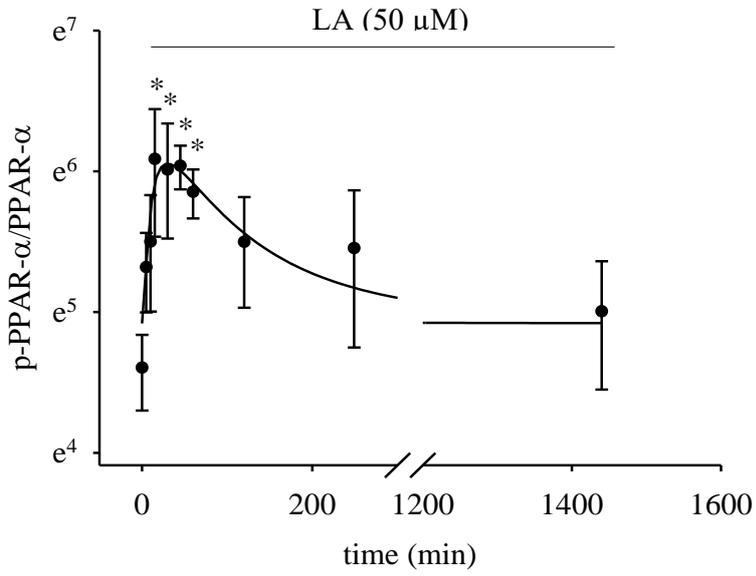
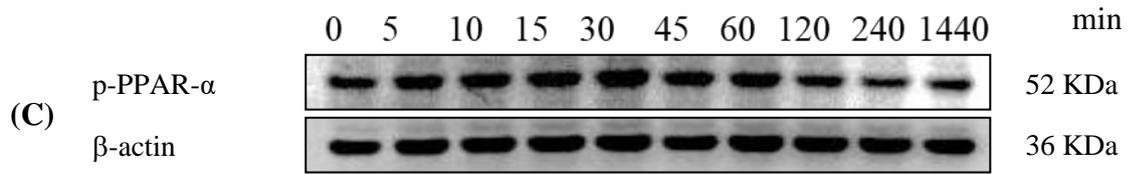
### 3.1.2.3 Activation of AMPK- $\alpha$ and PPAR- $\alpha$ upon PUFA treatment of ES cells

PUFAs regulate energy metabolism as ligands of PPARs. The transcription factor PPAR- $\alpha$  is essential for metabolic adaptation to starvation by inducing genes for  $\beta$ -oxidation and ketogenesis (Nakamura et al., 2014). In cooperation with AMPK, PPAR- $\alpha$  activates fatty acid oxidation in order to switch the cellular energy supply from glycolysis to fatty acid utilization (Grabacka et al., 2013). To investigate, whether LA and EPA activate PPAR- $\alpha$  and AMPK- $\alpha$ , phospho-specific antibodies were applied. Indeed, both LA (50 $\mu$ M) and EPA (50 $\mu$ M) increased phosphorylation of AMPK- $\alpha$  at residue Thr172, starting already 5 min after PUFA application (figures 3.5A, B), whereas total of protein expression remained unchanged (data not shown). Moreover, activation of PPAR- $\alpha$  at residue S12 was observed with comparable time kinetics (figure 3.5C, D). To identify the location of activated PPAR- $\alpha$ , cells enzymatically dissociated from 4-day-old EBs and plated on coverslips were incubated for 60 min with either EPA (50 $\mu$ M) or LA (50 $\mu$ M), and stained with a phospho-specific anti-PPAR- $\alpha$  antibody. The data clearly showed that phospho-PPAR- $\alpha$  exclusively localized to cell nuclei of all cells under inspection. This effect could be abolished upon pre-incubation (2h) with the PPAR- $\alpha$  inhibitor GW6471 (5 $\mu$ M), which drives the displacement of coactivators from PPAR- $\alpha$  and promotes the recruitment of co-repressor proteins like nuclear co-repressor (H Eric Xu et al., 2002) (figure 3.5E, F).

## Results

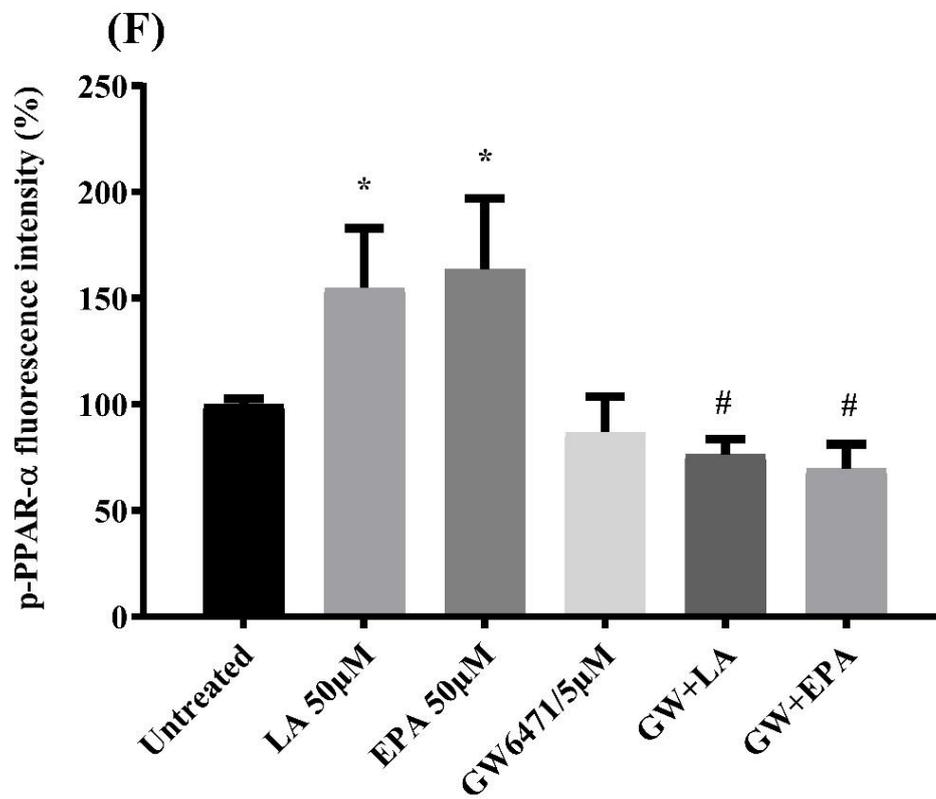
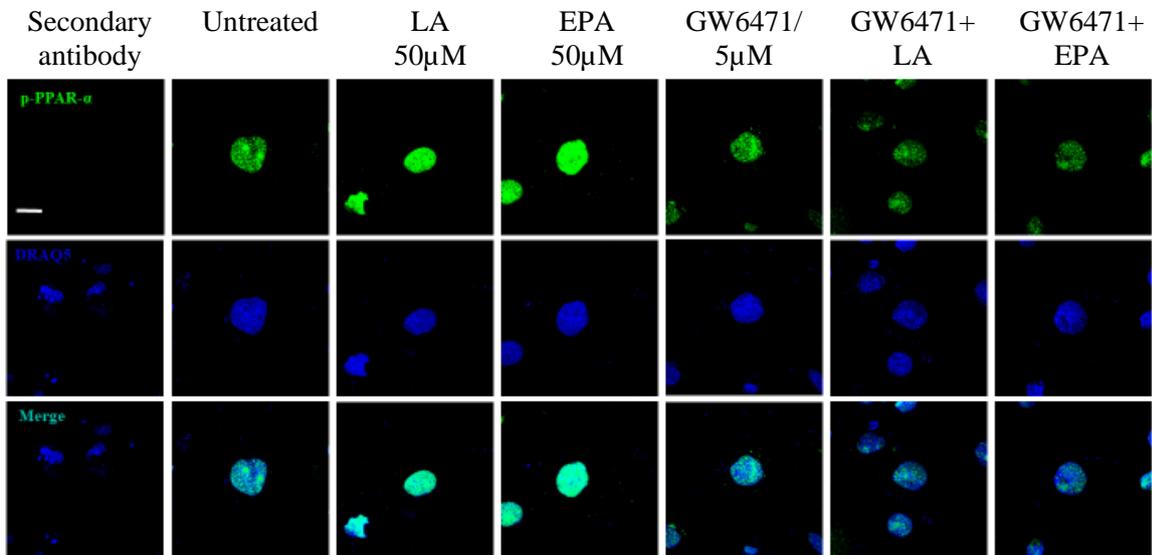


## Results



## Results

(E)



**Figure 3.5: Time course of AMPK -  $\alpha$  and PPAR -  $\alpha$  activation upon treatment of differentiating ES cells with PUFAs.** Five-day-old EBs were treated with 50 $\mu$ M of either LA or EPA, and protein was collected at different time points and subjected to western blot analysis. (A and B) AMPK- $\alpha$  phosphorylation following treatment with either (A) LA or (B) EPA. The upper panels show representative western blots stained with a phospho-specific antibody against AMPK- $\alpha$ . The lower panels show the means  $\pm$  SD of (n = 4) experiments for LA and EPA, respectively, corrected for total AMPK- $\alpha$  protein expression (p-AMPK- $\alpha$ /AMPK- $\alpha$ ). (C and D) PPAR- $\alpha$  phosphorylation following treatment with either LA (C) or EPA (D). The upper panels show representative western blots stained with a phospho-specific antibody against PPAR- $\alpha$ . The lower panels show the means  $\pm$  SD of (n = 8) experiments for LA and (n = 10) for EPA, respectively, corrected for total PPAR- $\alpha$  protein expression (p-PPAR- $\alpha$ /PPAR- $\alpha$ ). \*P  $\leq$  .05, significantly different to time 0 min. (E and F) Nuclear PPAR- $\alpha$  activation in single cells enzymatically dissociated from 3-day-old EBs upon treatment for 1h with either LA (50 $\mu$ M) or EPA (50 $\mu$ M) or pretreatment (2h) with GW6471 (5 $\mu$ M). (E) Representative cells which were labeled with a phospho-specific antibody against PPAR- $\alpha$  (upper row) or only secondary antibody, co-stained with the nuclear cell marker DRAQ5 (middle row) and merged (lower row). The bar represents 15 $\mu$ m. (F) Bar chart showing the means  $\pm$  SD of nuclear p-PPAR- $\alpha$  expression upon PUFA treatment in absence and presence of GW6471 (5 $\mu$ M) (n = 3) \*P  $\leq$  .05, significantly different to the untreated control and #P  $\leq$  .05, significantly different to the PUFA treated sample.

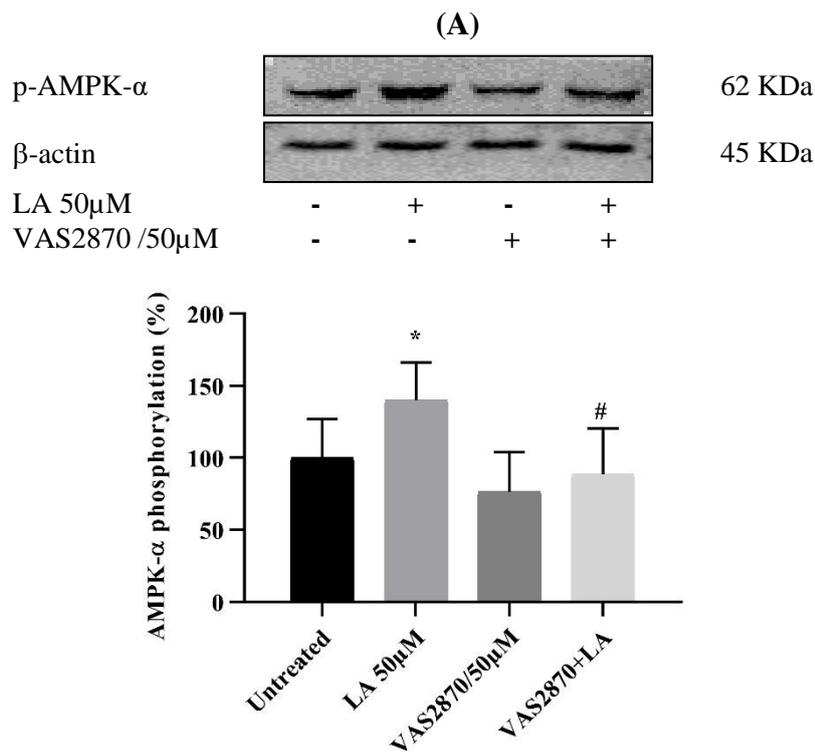
### 3.1.2.4 Effect of calcium chelation as well as NADPH oxidase and eNOS inhibition on AMPK- $\alpha$ activation upon PUFA treatment of ES cells

The data of the present study demonstrate that treatment of differentiating ES cell with  $\omega$ -3 and  $\omega$ -6 PUFAs raised intracellular calcium, ROS and NO. To investigate whether the observed activation of AMPK- $\alpha$  is related to these intracellular messengers, experiments were carried out, where NADPH oxidase was inhibited with VAS2870 (50 $\mu$ M), intracellular calcium was chelated with BAPTA-AM (10 $\mu$ M), and eNOS was inhibited with L-NAME (100 $\mu$ M). The data demonstrate that inhibition of NADPH oxidase by VAS2870 totally abolished AMPK- $\alpha$  activation by LA (50 $\mu$ M) (figure 3.6A). Moreover, AMPK- $\alpha$  activation by LA (50 $\mu$ M) and EPA (50 $\mu$ M) was inhibited upon pre-incubation with L-NAME (figures 3.6D, E) as well as with BAPTA (figures 3.6B, C).

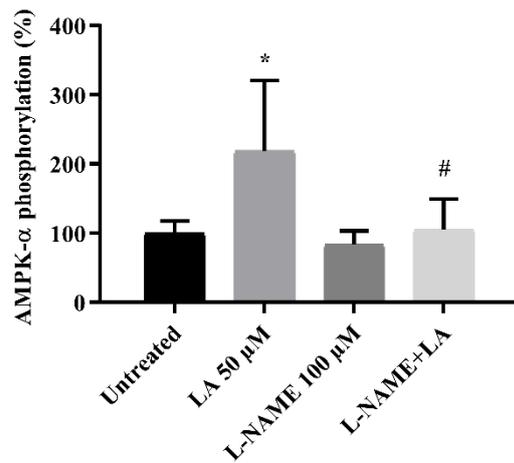
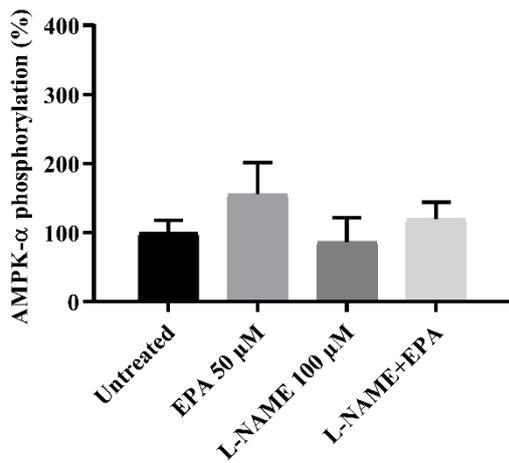
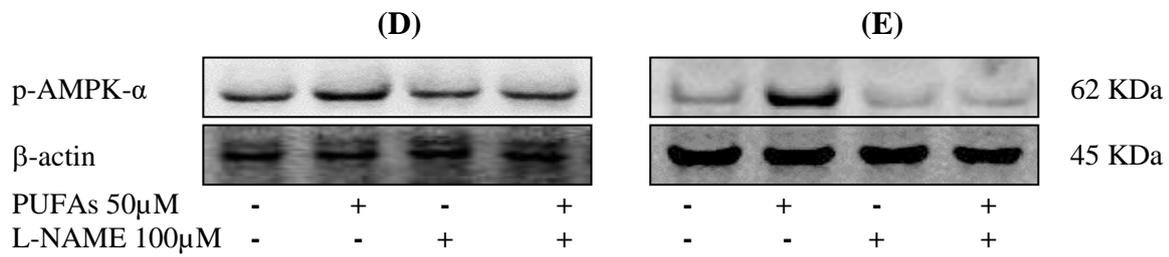
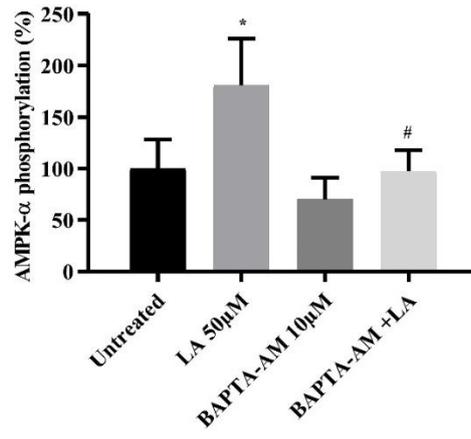
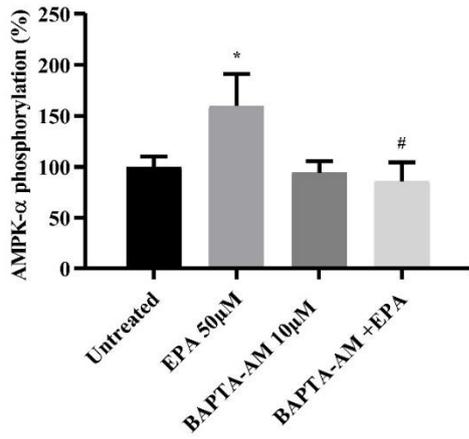
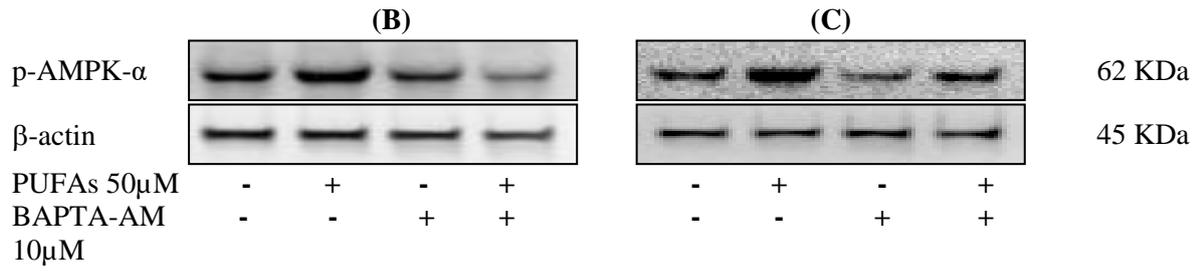
## Results

### 3.1.2.5 Abrogation of PUFA-mediated vasculogenesis upon inhibition of PPAR- $\alpha$

If PUFAs are stimulating vasculogenesis of ES cells via AMPK- $\alpha$  and downstream of PPAR- $\alpha$ , inhibition of the latter should abolish this effect. To validate our assumption we performed experiments, where differentiating ES cells were co-incubated with the PPAR- $\alpha$  inhibitor GW6471. Our data demonstrate that upon co-incubation of EPA (50 $\mu$ M) and LA (50 $\mu$ M) with GW6471 (5 $\mu$ M) the stimulation of vasculogenesis, as evaluated by assessment of vascular branching points, was completely abolished (figure 3.6F, G). This finding clearly underscores the impact of PPAR- $\alpha$  for PUFA-induced vasculogenesis of ES cells (see figure 4.1).

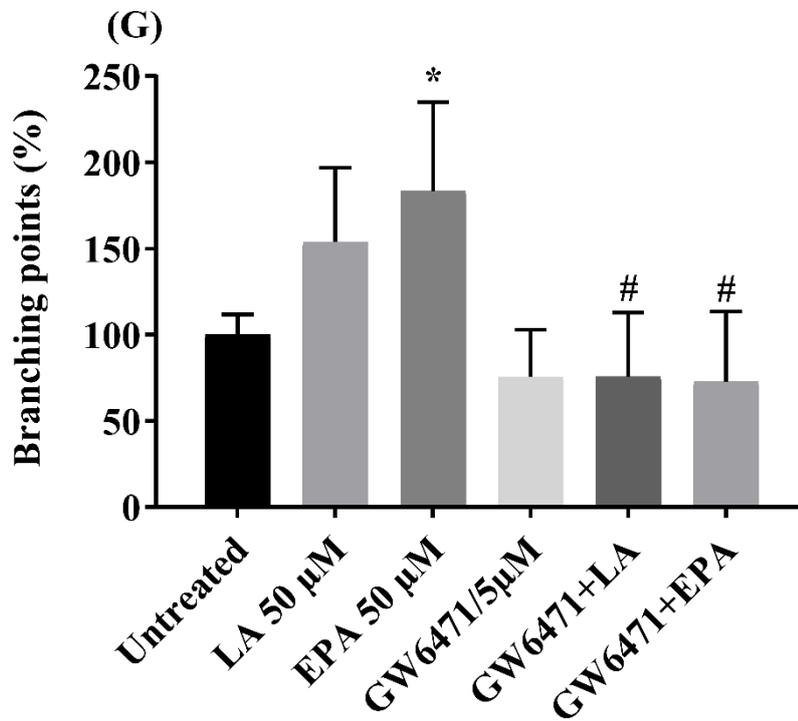
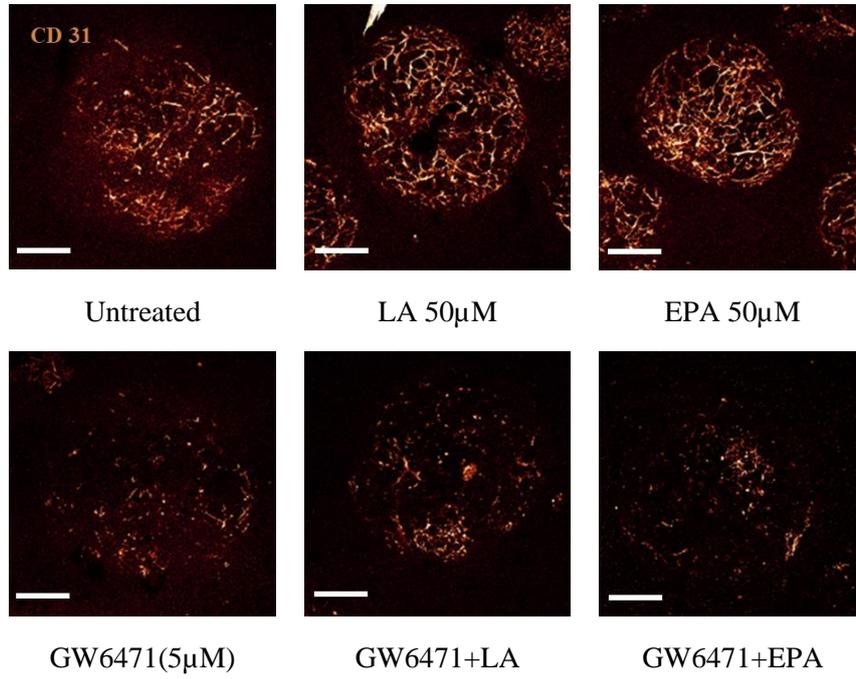


## Results



## Results

(F)



## Results

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**Figure 3.6: Inhibition of PUFA-induced AMPK- $\alpha$  activation by interference with calcium, ROS and NO signaling.** Five-day-old EBs were treated with 50 $\mu$ M of either EPA or LA and protein was collected 30 min thereafter and subjected to western blot analysis. Pretreatment with VAS2870 (50 $\mu$ M), BAPTA-AM (10 $\mu$ M), L-NAME (100 $\mu$ M), and GW6471 (5 $\mu$ M) was done 2 h before addition of PUFAs. (A) LA-induced AMPK- $\alpha$  activation was abolished by the NADPH oxidase inhibitor VAS2870 (n = 7). (B and D) EPA, and (C and E) LA-induced AMPK- $\alpha$  activation was abolished by the NOS inhibitor L-NAME (n = 3 for EPA and n = 5 for LA), and the calcium chelator BAPTA-AM (n = 4 for EPA and LA, respectively). Shown are representative western blots as well as bar charts showing the means  $\pm$  SD of the indicated number of experiments. (F and G) Inhibition of PUFA-induced vasculogenesis by the PPAR- $\alpha$  inhibitor GW6471. The images in (F) show representative EBs treated from day 3 to 10 of differentiation with PUFAs, either in absence or presence of GW6471 (the bar represents 200 $\mu$ m). The bar chart in (G) shows the means  $\pm$  SD of n = 6 experiments. \*P  $\leq$  .05, significantly different to the untreated control, #P  $\leq$  .05, significantly different to the PUFA treated sample.

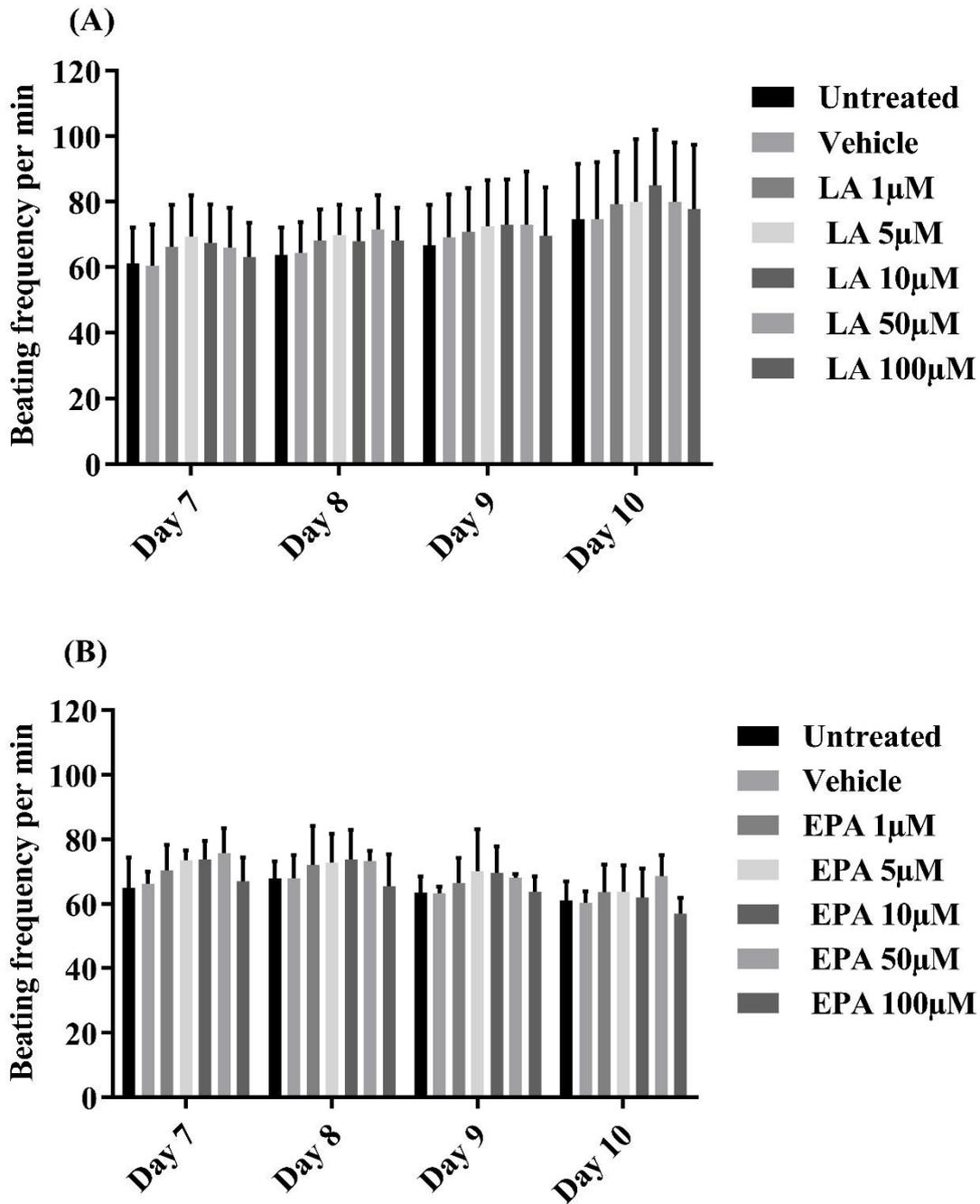
### **3.2 Effect of PUFAs on cardiomyogenesis and cardiac cell function of ES cell-derived EBs**

Under normal physiological states, fatty acid oxidation is fundamental for adult cardiomyocytes and prevails over the oxidation of other nutrients (Gaspar et al., 2014). Comparably to adult cardiac cells, cardiomyocytes derived from mESCs display a gene expression signature, which correlates with fatty acid metabolism and glycolysis (Sander et al., 2013). To investigate optimal concentrations of PUFAs to induce cardiomyogenesis in ES cells, 3-day-old EBs either outgrown on coverslips or in suspension culture were treated with different concentrations of LA and EPA versus untreated and vehicle groups (DMSO).

#### **3.2.1 Effect of PUFAs on the beating function of cardiac cells**

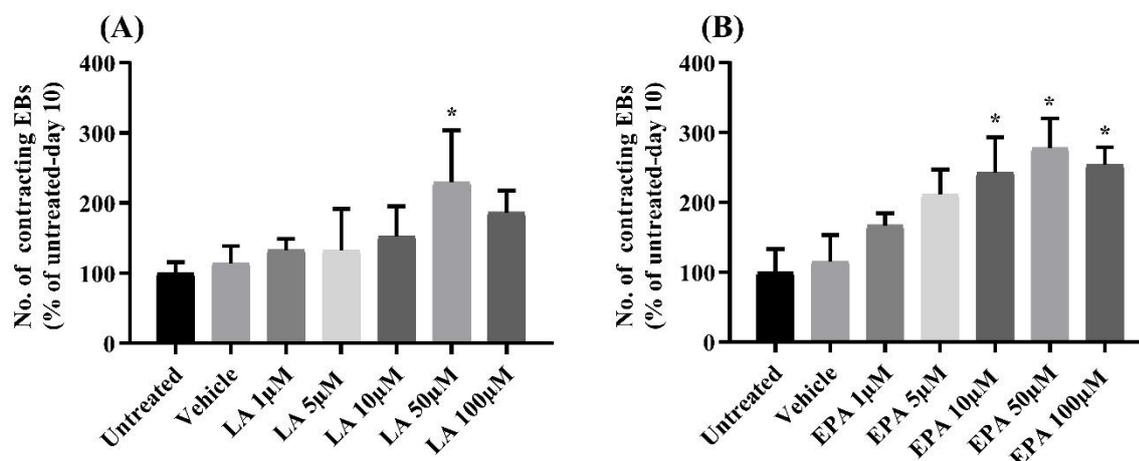
Spontaneous contractions are one of the fundamental functions of cardiomyocytes. To investigate, whether PUFAs accelerated the beating frequency of cardiomyocytes derived from ES cells or increase the number of contracting EBs, they were incubated with different concentrations of PUFAs (1, 5, 10, 50, and 100 $\mu$ M) from day 3 till day 10. In absence of PUFAs, spontaneous beating of cardiac cells clusters started on day 6 or 7. Therefore, beating frequency was evaluated by visual inspection on day 7 for each PUFA concentration in comparison with untreated and vehicle groups until day 10 of cell culture. The results showed that LA and EPA accelerated the beating frequency per min for cardiac contracting areas (foci), which was, however, non-significant in time and dose-dependent manner (figures 3.7A, B). In further experiments the number of contracting EBs was counted by microscopic inspection on day 10 of differentiation time upon PUFA treatment versus the control and vehicle group. The results showed that LA and EPA significantly increased the number of contracting EBs in dose-dependent manner (figures 3.8A, B), indicating stimulation of cardiomyogenesis upon PUFA treatment.

## Results



**Figure 3.7: Effect of PUFAs on beating frequency of EBs differentiated from mES cells.** (A, B) Beating frequency of cardiac cells clusters; 3-days-old EBs were incubated for 7 days with either LA (A) ( $n = 6$ ) or EPA (B) ( $n = 3$ ), and beating frequency per min was counted from day 7 to day 10 of cell culture. Shown are the means  $\pm$  SD.

## Results



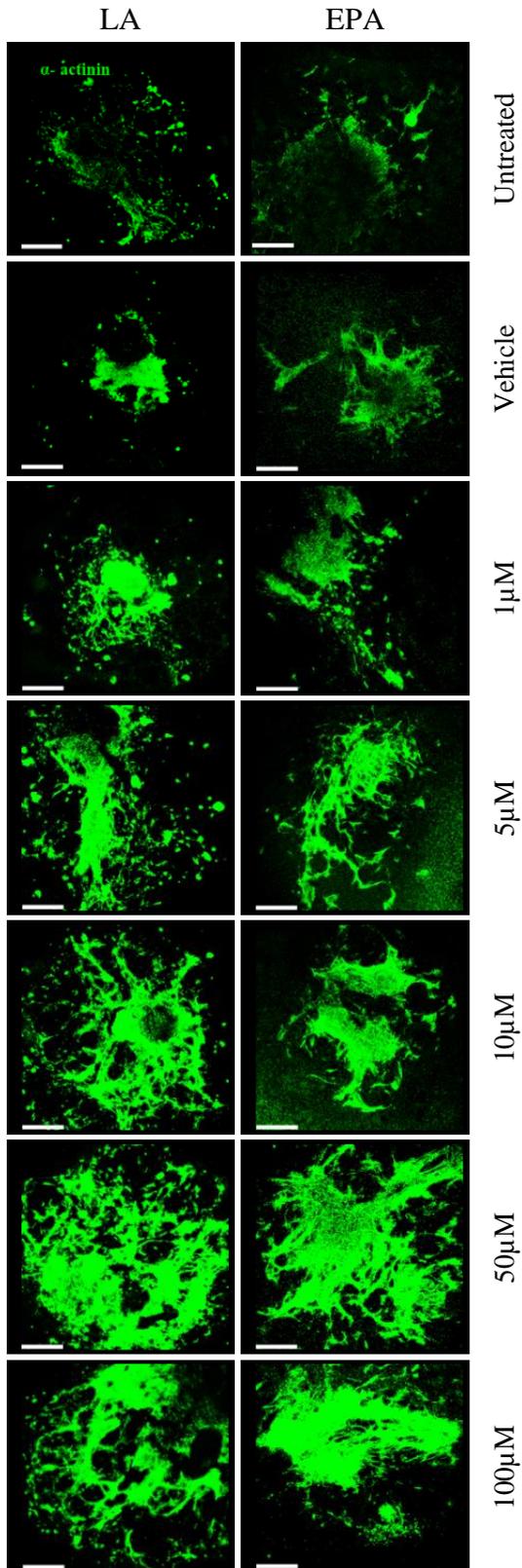
**Figure 3.8: Effect of different concentrations of PUFAs on the number of spontaneously contracting EBs.** (A, B) Bar charts show numbers (%) of spontaneously beating EBs (10-day-old) derived from ES cells. EBs were grown in presence of a series of concentrations of either LA (A) or EPA (B). Shown are the means  $\pm$  SD of (A) ( $n = 3$ ; LA) and (B) ( $n = 3$ ; EPA). \* $P \leq 0.05$ , statistically significant as compared with the untreated group.

### 3.2.2 Effect of PUFAs on the size of spontaneously contracting cardiac areas

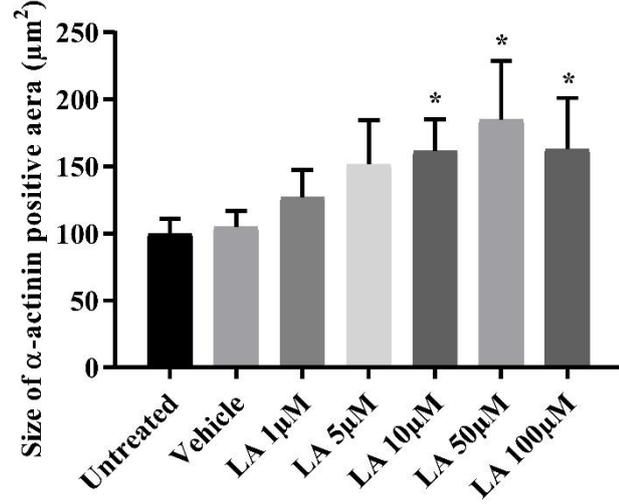
To further investigate stimulation of cardiomyogenesis by PUFAs, the size of beating cardiac clusters following PUFA exposure was assessed. To achieve this aim EBs were labeled with a monoclonal mouse anti-sarcomeric  $\alpha$ -actinin antibody. The  $\alpha$ -actinin (actin-binding protein) is associated with the Z-lines in cardiomyocytes (Djinovic-Carugo et al., 1999) (figure 3.9A). Our data demonstrated that PUFAs significantly increased the size of  $\alpha$ -actinin-positive cardiac cell clusters in a dose-dependent manner, suggesting that PUFAs increased the number of cardiomyocytes differentiated from ES cells (figures 3.9B, C).

## Results

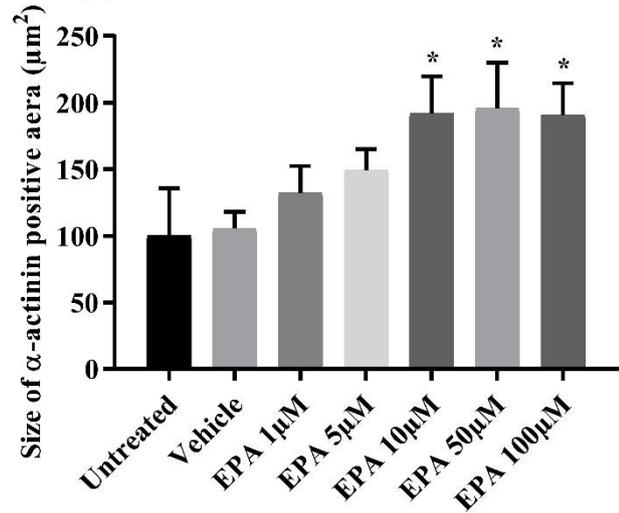
(A)



(B)



(C)

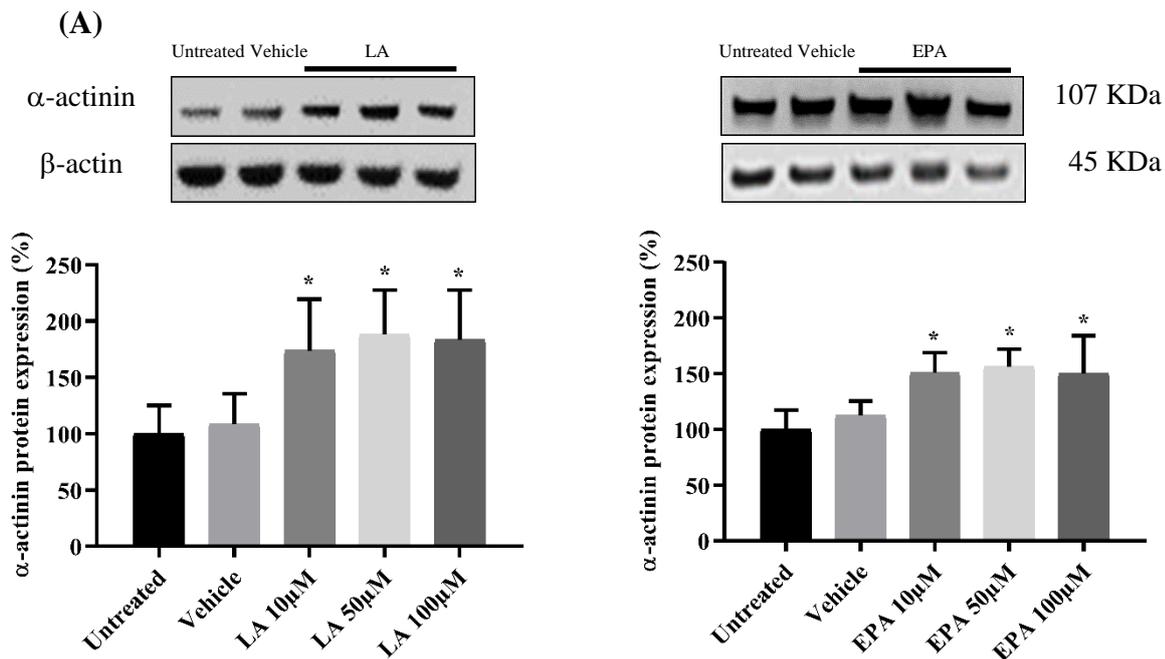


## Results

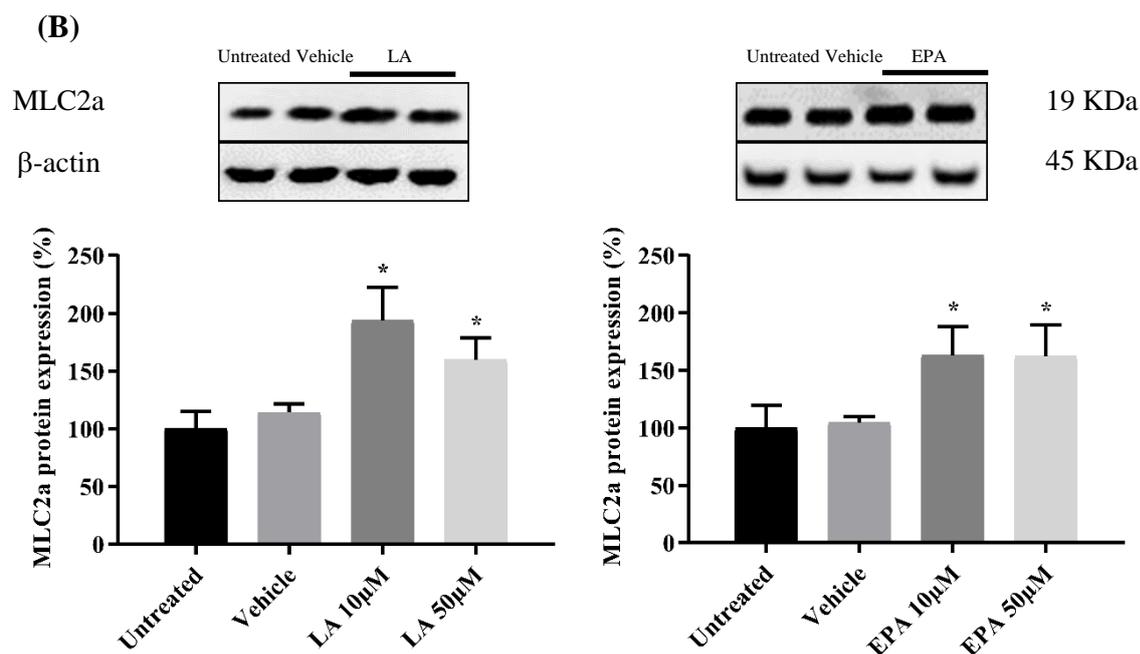
**Figure 3.9: Effect of PUFAs on the size of  $\alpha$ -actinin-positive cardiac cells clusters differentiated from ES cells.** (A, B) LA, (A, C) EPA. Shown are representative images of cardiac areas, which were stained with an antibody against  $\alpha$ -actinin (green) after 10 days of cell culture. The scale bars represent 200 $\mu$ m. (B, C) Bar charts show quantification of the size of  $\alpha$ -actinin-positive cardiac cell areas ( $\mu$ m<sup>2</sup>). Shown are the means  $\pm$  SD for LA (n = 4) and EPA (n = 3). \*P  $\leq$  0.05, statistically significant as compared with the untreated group.

### 3.2.3 Cardiac structural protein expression upon PUFA treatment

To analyse the effects of PUFAs on cardiomyogenesis in more details, Western blot analysis was performed to investigate cardiac-structural protein expression under PUFA treatment conditions, i.e.,  $\alpha$ -actinin (figure 3.10A) and MLC2a (figure 3.10B). As shown in figure 3.10, PUFAs upregulated  $\alpha$ -actinin (A) and MLC2a (B) protein expression in a dose-dependent manner. The optimal PUFA concentration was 50 $\mu$ M versus the untreated and vehicle (DMSO) group, while 100 $\mu$ M exerted slightly adverse effects.



## Results



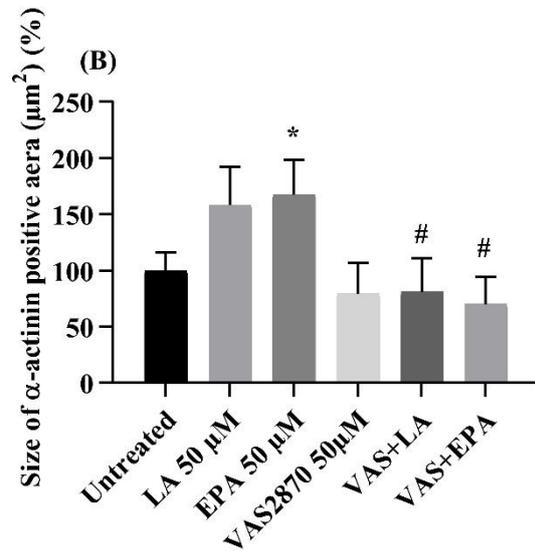
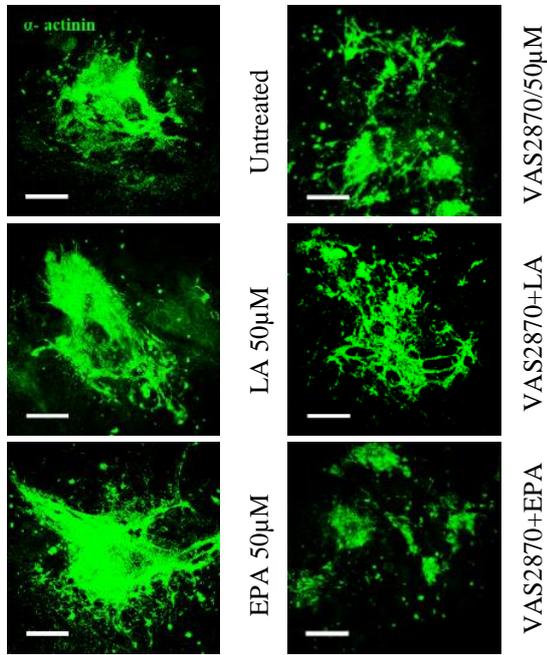
**Figure 3.10: Effect of PUFAs on cardiac structural protein expression.** (A, B) Upper panels, western blot analysis of  $\alpha$ -actinin (A) and MLC2a (B) protein expression upon PUFA treatment.  $\beta$ -actin was used as internal loading control. (A, B) The bar charts show quantitative data of cardiac protein expression upon PUFA treatment in different concentrations conditions. Shown are the means  $\pm$  SD of 4-independent experiments for (A) (LA) and (EPA) as well as (B) (LA) and (EPA). \* $P \leq 0.05$ , statistically significant as compared with the untreated group.

### 3.2.4 Involvement of NADPH oxidase and PPAR- $\alpha$ in PUFA-induced cardiomyogenesis of ES cells

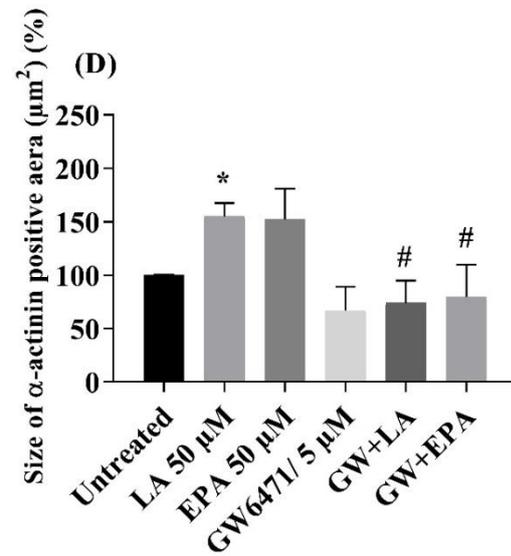
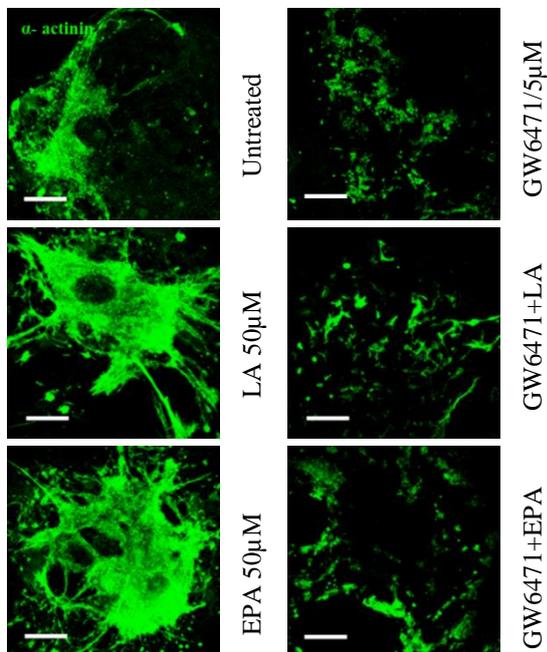
To investigate the hypothesis that PUFAs stimulate cardiomyogenesis from ES cells via NADPH oxidase and PPAR- $\alpha$ , EBs were pre-incubated from day 3 to 10 of cell culture either with the NADPH oxidase inhibitor VAS2870 (50 $\mu$ M) or with the PPAR- $\alpha$  antagonist GW6471 (5 $\mu$ M) in presence or absence of either LA (50 $\mu$ M) or EPA (50 $\mu$ M). Then, the size of cardiac areas ( $\alpha$ -actinin-positive area) and beating frequency were evaluated. The results show that the NADPH oxidase inhibitor VAS2870 or PPAR- $\alpha$  antagonist GW6471 significantly reduced the PUFAs-induced differentiation of cardiac cells from ES cells as well as the increase in beating frequency observed upon PUFA treatment (figure 3.11A-E).

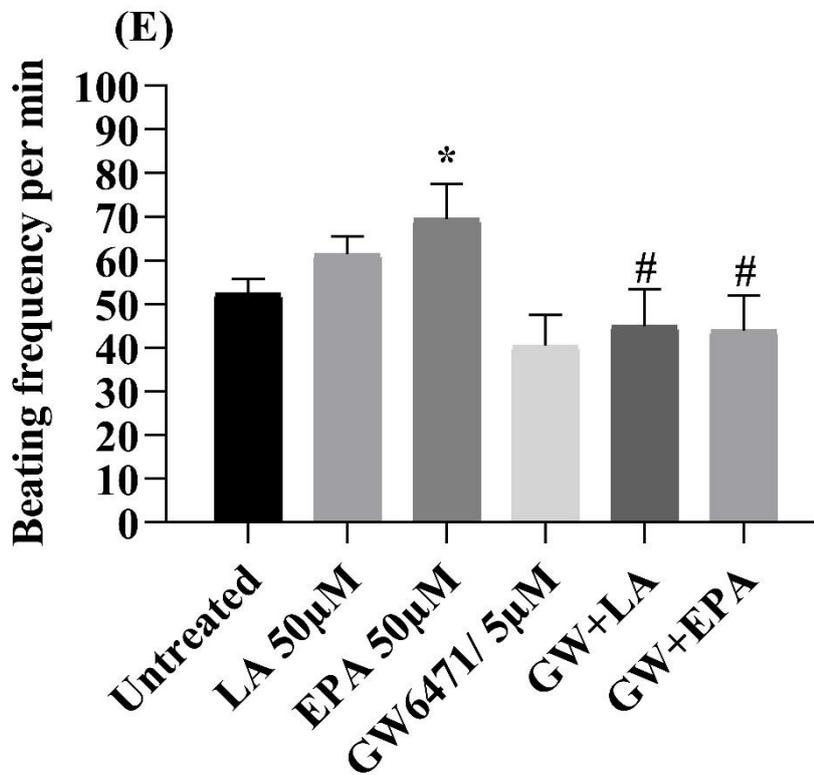
## Results

(A)



(C)





**Figure 3.11: Effect of NADPH oxidase inhibition by VAS2870 and PPAR- $\alpha$  inhibition by GW6471 on PUFA-induced cardiomyogenesis.** EBs were incubated from day 3 to 10 of differentiation with PUFAs, either in absence or presence of VAS2870 (50µM) or GW6471 (5µM). (A and C) Representative images of EBs cultivated on coverslips and labeled with an antibody against  $\alpha$ -actinin (green) after 10 days of cell culture. The scale bars represent 200µm. (B and D) show bar charts representing the means  $\pm$  SD of independent experiments for VAS2870 (50µM; n=3) and GW6471 (5µM; n=3). (E) The beating frequency per min was counted on day 10 of cell culture in absence or presence of GW6471 (5µM) condition (n=6). \*P  $\leq$  0.05, statistically significant as compared with the untreated group. #P  $\leq$  .05, significantly different to the PUFA treated sample.

### 4. Discussion

Objective of the present study was to investigate vasculogenesis and cardiomyogenesis of mES cells exposed to either EPA as  $\omega$ -3 PUFA, LA as  $\omega$ -6 PUFA or PA as a saturated fatty acid. Most previous studies have focused on small chemical molecules to stimulate stem cell differentiation. Only a few studies have examined the impact of the endogenous metabolism on cardiovascular differentiation (Yanes et al., 2010), although a number of studies have demonstrated that PUFAs exert distinct effects on the proliferation and differentiation of stem cells originating from different sources (see table 12).

Recently an *American Heart Association* presidential advisory on dietary fats and cardiovascular disease (CVD) discussed and suggested that lower intake of saturated fat and higher intake of polyunsaturated and monounsaturated fat is associated with lower rates of CVD and cardiovascular mortality (Rimm et al., 2018). In contrast, a recent analysis of published clinical trials suggested that increasing EPA and DHA has little or no effect on mortality or cardiovascular health (Abdelhamid et al., 2018). The main beneficial health effects have been mainly attributed to PUFAs. ALA is found in plant oils, EPA, and DHA are originating in marine algae and phytoplankton as well as in the ascending food chain in marine crustaceans and marine fish. The lead substance for  $\omega$ -6 PUFAs is LA, which is produced in many nuts and seeds and is a precursor of AA (Sokoła-Wysoczańska et al., 2018). Mammals are unable to produce  $\omega$ -3 fatty acids, but can ingest the shorter-chain  $\omega$ -3 fatty acid ALA through diet and use it to synthesize the essential  $\omega$ -3 PUFAs EPA and DHA. After uptake into cells, PUFAs are metabolized by COXs, LOXs, and CYP to generate bioactive eicosanoids, including prostanoids, LTs, HETEs, and EETs (Hu et al., 2018). CYP enzymes are membrane-bound, heme-containing terminal oxidases that are part of a multi-enzyme complex, which also includes a FAD/FMN-containing NADPH CYP reductase and cytochrome b5 (Hu et al., 2018).

#### 4.1 Stimulation of vasculogenesis by PUFAs treatment

In the present study, the effects of EPA as a lead substance for  $\omega$ -3 and LA for  $\omega$ -6 PUFAs were investigated. It was demonstrated that EPA as well as LA stimulated vasculogenesis

## Discussion

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in a comparable concentration range, that is, between 1 and 100 $\mu$ M and maximum effects at 10 and 50 $\mu$ M. Previously reported range of physiological concentrations of EPA and LA in human serum amount to 14–100 $\mu$ M and 2,270–3,850 $\mu$ M, respectively (Dumancas et al., 2010). Interestingly the saturated fatty acid PA did not show any effect on vasculogenesis of ES cells, suggesting that the unsaturated double bond chemical structure of fatty acids may be a decisive determinant for their biological function. Moreover, this finding supports the notion, that dietary intake of PUFAs may support vasculogenesis and vascular health in comparison to saturated fatty acids like PA.

EPA and LA treatment increased ROS as well as NO generation (see figure 4.1), which was abolished in presence of the NADPH oxidase inhibitor VAS2870 and the NOS inhibitor L-NAME. Moreover, PUFA-stimulated vasculogenesis was blunted upon inhibition of ROS and NO generation by using pharmacologic NADPH oxidase and eNOS inhibitors. These data corroborate previous studies of us showing that the stimulation of vasculogenesis in ES cells by AA was dependent on ROS generation (Huang et al., 2016). It is well reported that upon chemical reaction of ROS with NO, peroxynitrite is formed which-in high concentrations is generally regarded as very reactive and toxic substance and causes endothelial dysfunction (Kanaan & Harper, 2017). In low, physiologic concentrations, ROS/reactive nitrogen species (ROS/RNS) species including hydrogen peroxide, superoxide, and peroxynitrite are second messenger molecules, which involve in functional oxidative and nitrosative modification of proteins (Hansen et al., 2016; Sauer et al., 2001). In this respect, previous studies have demonstrated that ROS and NO are critical determinants in cardiovascular differentiation of stem cells (Bekhite et al., 2016; Gentile et al., 2013; Huang et al., 2010; Mazrouei et al., 2015; Sharifpanah et al., 2016).

Notably, peroxynitrite in low physiologic concentrations may be proangiogenic and act as intracellular messenger. This has been evidenced in studies, where peroxynitrite increased vascular endothelial growth factor (VEGF) expression in vascular endothelial cells (Platt et al., 2005), mediated VEGF-dependent angiogenic function in brain microvascular endothelial cells by c-src and MT1-matrix metalloproteinase activation (Prakash et al., 2012), and stimulated cell endothelial cell migration by focal adhesion kinase activation as well as low molecular weight protein tyrosine phosphatase S-glutathionylation (Abdelsaid

## Discussion

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& El-Remessy, 2012). It is well known that  $\omega$ -3 and  $\omega$ -6 PUFAs are metabolized in distinct intracellular pathways. AA is the substrate for the series 2 PGs, prostacyclins, TXs, and LTs (LTB<sub>4</sub>, series B<sub>4</sub> LT); EPA is the substrate for the series 3 prostanoids and LTB<sub>5</sub> (series B<sub>5</sub> LT) (Benatti et al., 2004). The redox biology of PUFAs is complex and not sufficiently investigated. Generally, it is assumed that  $\omega$ -3 PUFAs are anti-oxidative and anti-inflammatory, whereas  $\omega$ -6 PUFAs exert pro-oxidative and pro-inflammatory effects (Sokoła-Wysoczańska et al., 2018). However, this notion was recently challenged since a number of studies reported that also  $\omega$ -3 PUFAs moderately raise ROS in several cell types like endothelial cells (Okada et al., 2017), vascular smooth muscle (Crnkovic et al., 2012) and skeletal muscle cells (da Silva et al., 2016). Interestingly, ROS arising from PUFAs have been frequently noticed in tumour cells, for example, colon cancer cells (Pettersen et al., 2016) as well as lung and melanoma cancer cells (Zajdel et al., 2013; Zajdel et al., 2015). This is of importance since aggressive cancer cells mimic early-development stem cells, and properties of ES cells are retained in cancer cells (Ratajczak et al., 2018).

The intracellular signaling cascade resulting in ROS/RNS generation is not sufficiently investigated. The initial step should be the uptake of PUFAs into the cell. The cellular import of PUFAs is accomplished by CD36, which is a member of the class B scavenger receptor family of cell surface proteins. Inside the cell, PUFAs may elicit intracellular calcium responses. This has been shown for LA in pancreatic  $\beta$  cells (Zhao et al., 2013), bovine neutrophils (Mena et al., 2013), HeLa cells (Figueroa et al., 2013) and mES cells (Min Hee Kim et al., 2009). EPA raised intracellular calcium in several cell types including endothelial cells (Kim et al., 2015; Okuda et al., 1994; Omura et al., 2001; Wu et al., 2018), human colon epithelial cells (Kim et al., 2015) and vascular smooth muscle cells (Engler et al., 1999). Intracellular calcium responses elicited by EPA in endothelial cells and SMCs were independent of NO generation (Okuda et al., 1997), which may indicate that the calcium response is upstream of NO generation. Calcium dependence of eNOS is well documented in the scientific literature (Vanhoutte et al., 2016). In the present study EPA as well as LA transiently raised intracellular calcium (see figure 4.1) which was totally inhibited upon pre-incubation with the calcium chelator BAPTA-AM and significantly reduced by the CD36 inhibitor SSO as well as by a blocking antibody against

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CD36, suggesting that PUFAs enter cells via CD36 and release calcium from intracellular stores. The intracellular calcium response is apparently a prerequisite for ROS generation since BAPTA-AM abolished the effect. Moreover, ROS/NO are necessary for the vasculogenic effects of PUFAs since the NADPH oxidase inhibitor VAS2870 as well as the NOS inhibitor L-NAME abolished the differentiation of blood vessel-like structures upon PUFA treatment.

PUFAs have been previously shown to activate the metabolic sensor AMPK, which senses ATP deficiency and blocks anabolic processes (Shackelford & Shaw, 2009). Upon activation, AMPK inhibits glycolysis by phosphorylation of multiple forms of phosphofructo-2-kinase and by mammalian target of rapamycin inhibition, and in parallel initiates energy supply in form of fatty acid oxidation (Grabacka et al., 2013). The activity of AMPK is upstream of PPAR- $\alpha$  which is responsible for fat mobilization during fasting and activates mitochondrial and peroxisomal fatty acid  $\beta$ - and  $\omega$ -oxidation and ketogenesis, simultaneously inhibiting glycolysis and fatty acid synthesis (Grabacka et al., 2013). The data of the present study show that EPA as well as LA activated AMPK- $\alpha$  and PPAR- $\alpha$ . Previous studies have evidenced that  $\omega$ -3 PUFAs increased fatty acid oxidation in AMPK-dependent manner to generate ATP, thereby inhibiting unnecessary pathways, such as fatty acid synthesis (Lyons & Roche, 2018). Our data show that AMPK- $\alpha$  activation was dependent on PUFA-induced intracellular calcium, ROS and NO changes, since activation was abolished upon calcium chelation as well as inhibition of NADPH oxidase and NOS. This is in line with previous studies of us, demonstrating that PPAR- $\alpha$  agonists, such as WY14643, GW7647, and ciprofibrate stimulated cardiomyogenesis of ES cells by a mechanism involving ROS generation (Sharifpanah et al., 2008). Consequently, the PPAR- $\alpha$  inhibitor GW6471 inhibited vasculogenesis elicited by EPA and LA in the present study.

### **4.2 Effect of LA and EPA on cardiomyogenesis of ES cells and cardiac cell function**

Several reports have shown that one of the significant causes of morbidity and mortality worldwide is CVD. It is responsible for 17 million cases of deaths every year that represent about 31% of all global deaths such as ischemic heart disease (IHD), heart failure (HF) and

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myocardial infarction (MI). It is well known that the regenerative capacity of the heart is limited, which necessitates to differentiate and cultivate cardiac cells under cell culture conditions for cell therapy approaches of diseased hearts. The most promising source of cells for cardiomyoplasty are ES cells. Cardiomyocytes differentiated from pluripotent ES cells are currently used in the development of tools for in vitro generation of cardiac tissues for disease modelling, drug testing, and illustrating developmental processes (du Pré et al., 2013; Madonna et al., 2019; Park & Yoon, 2018; Shen et al., 2017; World Health Organization, 2014). Moreover, cardiac cell therapy holds the potential to enhance healing of ischemic heart, repopulate injured myocardium and recover cardiac function (Der Sarkissian et al., 2017). Reaching these aims requires the improvement of efficient protocols, which induce and direct cardiomyocyte differentiation from ES cells to achieve sufficient cardiac cell numbers for cell therapy. It is therefore necessary to perform studies, which focus on exogenous and endogenous signaling pathways to differentiate ES cells towards cardiovascular lineages. Previous studies have outlined the importance of PUFAs such as LA and EPA to improve coronary heart disease (CHD) as well as CVD. Farvid and his colleagues have demonstrated that higher dietary consumption of LA is correlated with a lower risk of (CHD). They therefore recommended to replace the use of saturated fatty acid by PUFAs in the diet (Farvid et al., 2014). Moreover,  $\omega$ -3 and  $\omega$ -6 PUFAs, and their metabolites are crucial elements of cell structure and function that could affect stem cell fate. The available evidence demonstrates that  $\omega$ -6 and  $\omega$ -3 PUFAs and their metabolites can perform through multiple mechanisms to contribute to the proliferation and differentiation of several stem cell types (Table. 12) (Kang et al., 2014). These and our own observations suggest that PUFAs could activate distinct signaling networks to advance the cardiomyocyte differentiation of ES cells.

Fatty acids include the major metabolites for energy supply in cardiomyocytes (Lopaschuk et al., 2010). They also contribute to the regulation of cardiac-specific gene expression as well as activity of ion channels that are taking charge of heart beating (S. S. Chan et al., 2009; Leifert et al., 1999; Xiao et al., 2001). Energy metabolism is a fundamental aspect of cardiomyocyte biology. The heart is a metabolic omnivore, and the adult heart picks the substrate best suited for each condition. Fatty acid oxidation is considered the most

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important source of ATP production, and is preferred in order to achieve the high energy requirement of the contracting myocardium (Malandraki-Miller et al., 2018; Ulmer et al., 2018).

The data of the present study demonstrated that LA and EPA stimulated cardiomyocyte differentiation from ES cells. The beating frequency of cardiomyocytes-derived from ES cells under LA and EPA conditions was accelerated, indicating improvement of cardiac cell function. Recently it has been demonstrated that PUFA analogues vary in their selectivity for human voltage-gated ion channels involved in the ventricular action potential (Bohannon et al., 2020). Selectivity for a specific ion channel to broadly modulating cardiac ion channels was evidenced for all three ion channel families, i.e. voltage-gated sodium channel ( $\text{Nav}$ ), voltage-gated calcium channel ( $\text{Cav}$ ), and voltage-gated potassium channel ( $\text{Kv}$ ), which generate the cardiac action potential. Other authors showed that PUFAs induced activation  $\text{Kv}7.1$  and  $\text{Kv}7.1\text{-KCNE1}$  channels in cardiac cells.  $\text{Kv}7.1\text{-KCNE1}$  channels produce the slow-delayed rectifying  $\text{K}^+$  current,  $I_{Ks}$ , which is important during the repolarization phase of the cardiac action potential (Liin et al., 2018; Thompson et al., 2021). These data support the notion that PUFAs exert anti-arrhythmic properties which may be related to improvement of gap-junctional coupling via connexin 43 ( $\text{Cx}43$ ) (Benova et al., 2019). The anti-arrhythmic properties of PUFAs were confirmed in a recent clinical study which showed that  $\omega$ -3 PUFA supplementation of the diet improved cardiac function in children with premature ventricular contractions (Oner et al., 2018). Another important finding of the present study was that PUFAs dose-dependent increased the number of contracting EBs in a specific time window of differentiation, i.e. day 7-10 of cell culture which indicates an increase in the number of differentiated cardiomyocytes from ES cells. Prior studies have reported the importance of this time window in cardiomyogenesis of ES cells (Bartsch et al., 2011; Boheler et al., 2002). PUFAs dose-dependent increased the  $\alpha$ -actinin-positive area and enhanced cardiac protein expression such as  $\alpha$ -actinin and  $\text{MLC}2a$ . In accordance with the present results, previous studies have demonstrated that a low dose ( $10\mu\text{M}$ ) of EPA promoted ES cell differentiation toward cardiomyocytes, increased the cell area of contracting cardiomyocytes and the expression of cardiac markers such as  $\text{Nkx}2.5$ ,  $\text{MEF}2\text{C}$ ,  $\text{MYH}6$ ,

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TNNT2, and Cx43 in EBs derived from mES cells (Shabani et al., 2015). The same group reported that preincubation of ES cell-derived cardiac cells with  $\omega$ -3 PUFAs and ascorbic acid improved the regenerative capacity and reduced fibrosis in a rat infarction model (Shabani et al., 2019).

Since data of the present study showed that stimulation of vascular differentiation by PUFAs was mediated by ROS, we investigated whether the same effect would underlie cardiomyogenesis. Our data demonstrated that PUFA-induced cardiomyogenesis was blunted in presence of the ROS inhibitor NADPH oxidase inhibitor VAS2870. This observation corroborates previous research from our group and others which demonstrated that intracellular ROS are implicated in cardiomyocyte differentiation from ES cells via different ROS-involving signaling cascades (Bartsch et al., 2011; Buggisch et al., 2007; Crespo et al., 2010; Murray et al., 2013; Sauer et al., 2004; Sauer et al., 2000; Schmelter et al., 2006; Wo et al., 2008).

### **4.3 Impact of PUFAs on cardiac cell maturation**

Besides stimulating the differentiation, proliferation and function of cardiac cells, PUFAs may also regulate maturation of cardiomyocytes. Recently it has been outlined that the energy metabolism may be involved in the regulation of maturation of cardiomyocytes derived from pluripotent stem cells with unexpected effects on electrical and mechanical function, and it has been discussed that PUFAs may activate multiple signaling networks leading to improved maturation of cardiac progenitor cells from the fetal phenotype (Karbassi et al., 2020). Cardiomyocytes differentiated from pluripotent stem cells are significantly immature compared to adult cardiomyocytes and display a fetal-like phenotype (Feric & Radisic, 2016; Yang et al., 2014). The immature properties of cardiomyocyte limit their efficacy in disease modeling and therapeutic purposes. Although some progress has been performed to achieve more mature human pluripotent stem cell-derived cardiomyocytes (hPSC-CMs), there is enough space for improvement because of the marked postnatal cardiac metabolism changes associated to cardiac cell maturation (Makinde et al., 1998; Yang et al., 2019). Maturation refers to the end state of heart development and primes the organ for powerful, effective and persistent pumping during

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lifespan. It is characterized by structural, gene expression, metabolic and functional specializations in terminally differentiated cardiomyocytes as the heart transits from fetal to adult states. Due to maturation defects in pluripotent stem cell–derived cardiomyocytes, their use in myocardial regeneration and their possible contribution to cardiac disease is so far limited. There are measurable parameters that distinguish cardiomyocyte maturation, such as myofibril maturation, changes in electrophysiology and  $\text{Ca}^{2+}$  metabolism, and others (Guo & Pu, 2020). Yang and his colleagues have mentioned that metabolic substrate shift from glucose to fatty acids is an essential feature of postnatal cardiomyocyte maturation. They suggested that applying fatty acids would improve hPSC-CM maturation. They designed a cocktail based on breastfed human infant serum (Hardell & Walldius, 1980) and breast milk (Gibson & Kneebone, 1981). Their study evidenced that palmitate, oleate and linoleate could induce hPSC-CM hypertrophic growth, force generation, calcium dynamics, action potential upstroke velocity and oxidative capacity and that these molecules activate multiple signaling pathways (Yang et al., 2019). Knight et al., (2021) have exhibited a modified method to create further mature human induced pluripotent stem cell-derived cardiomyocytes (hiPSC-CMs) by seeding cells in a medium supplemented with fatty acids as an important energy source and plating these cells on micropatterned growth surfaces. They noted that cells displayed elongation, enhanced sarcomeric maturity, metabolic gene expression, mitochondrial fatty acid uptake and cardiolipin maturation under these conditions. The isolated myofibrils from these cells showed that mechanical performance more similarly resembled those from adult human ventricular tissue, particularly in terms of force generation. These mature hiPSC-CMs also respond well to hypertrophic stimulation, producing an adult-like hypertrophic response and alterations in myofibril relaxation similar to human hypertrophic cardiomyopathy (HCM) tissue (Knight et al., 2021). In addition, Zhang and his colleagues have demonstrated that hPSC-CM are dependent upon exogenous sources of lipids for metabolic maturation (Zhang et al., 2020). The mechanism of PUFAs to enhance cardiomyocyte maturation is not sufficiently investigated. In mature cardiomyocytes, CD36 and fatty acid transport proteins (FATPs) facilitate fatty acid uptake to produce ATP by mitochondrial oxidative phosphorylation utilized to drive cardiac contraction (Steinbusch et al., 2011). Moreover, the expression of

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many key genes such as fatty acid oxidation and CD36 increased upon maturation treatment in hiPSC-CMs (Gentillon et al., 2019). In the heart, CD36 not only facilitates long-chain fatty acid uptake, but also regulates other cellular activities such as intracellular  $\text{Ca}^{2+}$  signaling (Pietka et al., 2012) and AMPK activation (Samovski et al., 2015). Notably, the data of the present study demonstrated that intracellular  $\text{Ca}^{2+}$  elicited by PUFAs was inhibited by the pharmacological CD36 inhibitor SSO as well as by neutralizing antibodies against CD36, thus relating PUFA import with intracellular  $\text{Ca}^{2+}$  signaling cascades and presumably cardiac maturation processes.

A further major player in cardiac cell maturation is PPAR- $\alpha$ . PPARs (comprising of PPAR- $\alpha$ ,  $\beta$  and  $\gamma$ ) are nuclear receptors that perform as ligand-dependent transcription factors, which are activated by different types of fatty acids. PPARs have important roles for various metabolic and cellular processes such as fatty acid  $\beta$ -oxidation and adipogenesis (lipid homeostasis). Moreover, PPARs perform an essential function during placental, embryonal, and fetal development and regulatory processes related to aging, comprising oxidative stress, inflammation, and neuroprotection (Echeverría et al., 2016). Recently, proteomic analysis of hES cell-derived, fetal and adult ventricular cardiomyocytes demonstrated that protein expression related to PPAR- $\alpha$  signalling cascades were differentially regulated in ES cell-derived and fetal cardiomyocytes as compared to adult cardiac cells. Stimulation of PPAR- $\alpha$  with WY-14643 resulted in increased fatty oxidative enzyme level, hyperpolarized mitochondrial membrane potential and induced a more organized morphology (Poon et al., 2015). Previous studies of us reported that agonists of PPAR- $\alpha$  increased cardiac differentiation from ES cells (Sharifpanah et al., 2008), which was related to elevation in the expression of cardiac-specific transcription factors and proteins. In the present study it was evidenced that EPA and LA activated PPAR- $\alpha$  in cell nuclei of differentiating cardiac cells which was inhibited in presence of the PPAR- $\alpha$  antagonist GW6471. This observation indicated that in addition to the activation of genes encoding contractile proteins, myofibrillogenesis is supported by enzymes involved in fatty acid metabolism (Kraft et al., 2006). In the experiments of the present study co-incubation of LA and EPA with the PPAR- $\alpha$  antagonist GW6471 abolished PUFA-induced cardiomyogenesis. Ding and colleagues (2007) who

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demonstrated that activation of PPAR- $\alpha$  is a pre-requisite for cardiomyocyte differentiation of murine ES cells. PPAR- $\alpha$  activation is dependent on either pharmacological ligands like fibrates or natural ligands, e.g. fatty acids (Ding et al., 2007). Previous transcriptomic analyses performed in mice demonstrated that single oral dose of synthetic triglycerides composed of one single fatty acid stimulated expression of cardiac genes which were regulated in a PPAR $\alpha$ -dependent manner, thus underscoring the importance of PPAR $\alpha$  in mediating transcriptional regulation by fatty acids in the heart (Georgiadi et al., 2012). Recently, research was undertaken to promote hiPSC-CM maturation by combining hypoxia inducible factor 1- $\alpha$  (HIF-1 $\alpha$ ) inhibition with molecules that target key pathways involved in the energy metabolism. In this study hiPSC-CMs were treated with a HIF-1 $\alpha$  inhibitor alone or in combination with the PPAR- $\alpha$  agonist WY-14643 and the postnatal factors triiodothyronine hormone T3, insulin-like growth factor-1 (IGF-1) and dexamethasone (Gentillon et al., 2019). The treatment resulted stimulated fatty acid oxidation, increased mitochondrial number and cardiac cell maturation thus supporting notion of the present study that stimulation of fatty acid oxidation by PUFAs supports cardiac cell maturation via signaling pathways, which involve the energy sensor AMPK- $\alpha$  and PPAR- $\alpha$ -mediated signaling cascades.

Stem cell pluripotency does not represent a single defined metabolic state. Metabolic signatures are highly characteristic for a cell and may contribute to the fate of the cell (Ellen Kreipke et al., 2016). Cellular metabolism is changing between naïve (early stage) and primed (late stage) ES cells, representing preimplantation and postimplantation embryos, respectively. The transition from naïve to primed ES cells has been shown to be accompanied by a distinct metabolic switch from bivalent (fatty acid oxidation/glycolytic) to highly glycolytic state which adopts the cells to the hypoxic condition of the early postimplantation embryo and protects the embryo from oxidative stress (Ellen Kreipke et al., 2016). During the following differentiation and maturation processes a second metabolic switch towards oxidative phosphorylation occurs. Then highly respiring mitochondria fulfill the dramatically augmenting energy needs during shaping of the cardiovascular system and organ formation. This physiological “metabolic inflammation”

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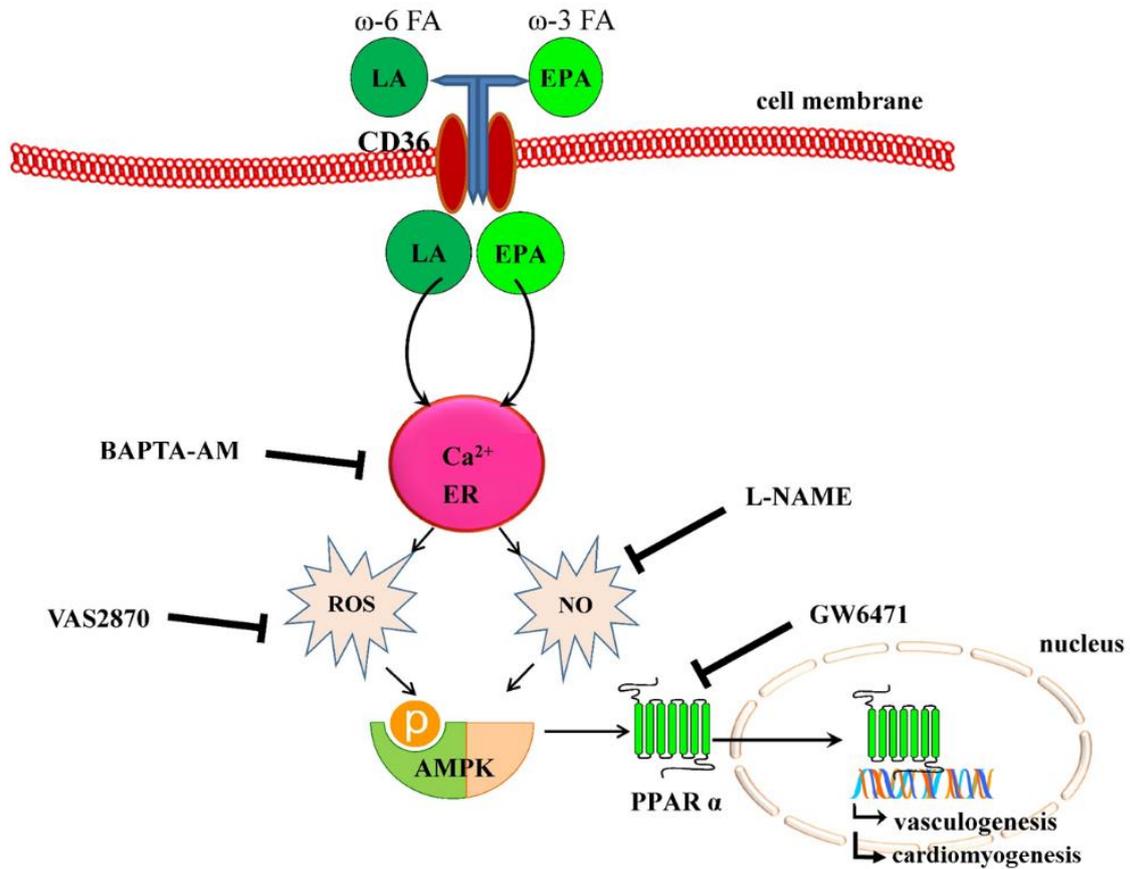
requires ROS/RNS as well as PUFA-derived eicosanoids as intracellular messengers to initiate differentiation and maturation of progenitor cells in the cardiovascular system.

**Table 12. The effects of PUFAs and their metabolic products on the proliferation and differentiation from various stem cell types.** The table was modified and updated with permission of Kang and his colleagues (Kang et al., 2014).

Fatty acids/ Metabolites	Stem cell models	Effects	Potential pathways	References
LA	ES cells	Increase proliferation	Ca21/PKC, PI3K/Akt and MAPKs	(Min Hee Kim et al., 2009)
		Promote differentiation	Ca <sup>2+</sup> , ROS, NO, AMPK- $\alpha$ and PPAR- $\alpha$	(Taha et al., 2020)
AA	ES cells	Promote differentiation	LT signaling pathway and 5-LOX-activating protein (FLAP)	(Huang et al., 2016)
	C2C12 cells	By prolonging proliferation and preventing differentiation, PUFAs control the transition cells from proliferation to differentiation.	Wnt signaling pathway	(Huang et al., 2016; Risha et al., 2021)
	Mesenchymal stem cells	Promote differentiation	Unknown	(Kan et al., 2007)
ALA	Embryonic neural stem cells	Increase proliferation Promote differentiation	Notch1, hes1 and Ki-67	(Mahmoudi et al., 2019)
EPA	ES cells	Promote differentiation	Unknown	(Shabani et al., 2015)
			Ca <sup>2+</sup> , ROS, NO, AMPK- $\alpha$ and PPAR- $\alpha$	(Taha et al., 2020)
DHA	Neural stem cells	Promote differentiation, increase proliferation	RXR, PPAR, dendritic spine-related genes and basic helix-loop-helix	(He et al., 2009; Kan et al., 2007; Katakura et al., 2009; Sakamoto et al., 2007)
	Mesenchymal stem cells	Promote osteogenic differentiation	Enhanced Akt activation at the plasma membrane	(Levental et al., 2017)
		Promote neuronal differentiation	Unknown	(Kan et al., 2007)
C2C12 cells	By prolonging proliferation and preventing differentiation, PUFAs control the transition cells from proliferation to differentiation.	Wnt signaling pathway	(Risha et al., 2021)	
PGE1	Hematopoietic stem cells	Inhibit proliferation	Unknown	(Gidáli & Fehér, 1977; Kurland et al., 1978; Motomura & Dexter, 1980)
PGE2	Hematopoietic stem cells	Increase proliferation, inhibit apoptosis, promote differentiation to dendritic cells	CXCR4, survivin, and Wnt	(Chen et al., 2007; Goessling et al., 2009; Pelus et al., 1979)

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	ES cells	Increase proliferation, inhibit apoptosis	PI3K/Akt	(Hou et al., 2013; SP Yun et al., 2009)
	Human umbilical cord blood-derived mesenchymal stem cells	Increase proliferation	Epac1/Rap1/Akt, and $\alpha$ -arrestin-1/JNK	(Jang et al., 2012; Yun et al., 2011)
	Neural stem cells	Increase proliferation	Unknown	(Jung et al., 2006; Sasaki et al., 2003)
	Bone marrow-derived cells	Promote endothelial	AMPK	(Zhu et al., 2011)
	Tendon stem cells	Promote osteogenic differentiation	PI3K/Akt	(Liu et al., 2013)
$\Delta$ 12,14-PGJ2	ES cells	inhibit proliferation	JAK-STAT	(Rajasingh & Bright, 2006)
	Neural stem cells	Biphasic regulation of proliferation	Unknown	(Katura et al., 2009, 2010)
LTB4	Neural stem cells	Biphasic regulation of proliferation, promote differentiation to neurons	Hedgehog	(Bijlsma et al., 2008; Wada et al., 2006)
	Hematopoietic stem cells	Increase proliferation, inhibit apoptosis	Unknown	(Chung et al., 2005)
LTD4	ES cells	Increase proliferation	STAT3, calcineurin, PI3K/Akt, and GSK 3b/b-catenin	(Kim et al., 2010)
LXA4	Neural stem cells	Directly regulates proliferation and differentiation to neurons	EGFR, cyclin E, p27, and caspase 8	(Wada et al., 2006)
NPD1	ES cells	Promote neuronal and cardiac differentiation	Unknown	(Yanes et al., 2010)
TXA2	Adipose tissue-derived mesenchymal stem cells	Increase proliferation, promote differentiation to smooth-muscle-like cells	ERK, p38 MAPK	(Mi Ra Kim et al., 2009; Doo Hee Yun et al., 2009)



**Figure 4.1: Schematic view of the stimulation of cardiomyogenesis and vasculogenesis in ES cells by PUFAs.** LA and EPA are entering the cell through CD36 and release intracellular calcium from intracellular stores. Calcium stimulates ROS from NADPH oxidase and NO from eNOS, which in concert activate AMPK- $\alpha$ . Downstream of AMPK, PPAR- $\alpha$  is phosphorylated in the cell nucleus and initiates vasculogenic and cardiomyogenic genes. Figure adapted with permission from (Taha et al., 2020); copyright John Wiley and Sons.

## Summary

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### 5-Summary

PUFAs and their metabolites may influence differentiation processes of stem cells. The aim of the current study was to determine the effects of LA as  $\omega$ -6-PUFA and EPA as  $\omega$ -3-PUFA on vasculogenesis and cardiomyogenesis of ES cells.

LA and EPA increased vascular structure formation and protein expression of the endothelial-specific markers FLK-1, PECAM-1, and VE-Cadherin, whereas the saturated fatty acid PA was without effect. LA and EPA increased ROS and NO, activated eNOS and raised intracellular calcium. The calcium response was inhibited by the intracellular calcium chelator BAPTA-AM and SSO, which is an antagonist of CD36, the scavenger receptor for fatty acid uptake. Comparable effects were achieved with CD36-neutralizing antibodies.

In a dose-dependent manner, LA and EPA enhanced cardiomyocyte differentiation and function. LA and EPA accelerated beating frequency of cardiac cells cluster (foci), increased the percentage of contracting EBs, the area of  $\alpha$ -actinin-positive cardiac cells (beating area), and protein expression of the cardiac-specific markers  $\alpha$ -actinin and MLC2a.

Prevention of ROS generation by radical scavengers or the NADPH oxidase inhibitor VAS2870 blunted vasculogenesis and cardiomyogenesis of ES cells. Moreover, inhibition of eNOS by L-NAME abolished vasculogenesis of ES cells.

PUFAs stimulated AMPK- $\alpha$  as well as PPAR- $\alpha$ . AMPK- $\alpha$  activation was abolished by calcium chelation as well as inhibition of ROS and NO generation. Moreover, PUFA-induced vasculogenesis and cardiomyogenesis were blunted by the PPAR- $\alpha$  inhibitor GW6471.

In conclusion, the data demonstrated that, LA as well as EPA, enhance cardiomyogenesis and vasculogenesis by mechanisms involving ROS, NO and changes in intracellular calcium, which are locked to the energy sensors AMPK- $\alpha$  and PPAR- $\alpha$ .

### 6- Zusammenfassung

Mehrfach ungesättigten Fettsäuren (PUFAs) beeinflussen möglicherweise Differenzierungsprozesse von Stammzellen. In der vorliegenden Studie wurde die Bedeutung von Linolsäure (LA), als Beispiel einer  $\omega$ -6-PUFA, und Eicosapentaensäure (EPA), als Beispiel einer  $\omega$ -3-PUFA, für die Vaskulogenese und Kardiomyogenese embryonaler Stammzellen der Maus untersucht.

LA und EPA stimulierten die Bildung reaktiver Sauerstoffspezies (ROS) und Stickmonoxid (NO), aktivierten die endotheliale NO-Synthase (eNOS) und erhöhten das intrazelluläre Calcium. Die Calciumantwort wurde durch den intrazellulären Calciumchelator BAPTA-AM und Sulfo-N-Succinimidyl Oleat (SSO), einen Antagonisten von CD36, dem Scavenger-Rezeptor für die Aufnahme von Fettsäuren, inhibiert. Vergleichbare Effekte wurden durch einen neutralisierenden Antikörper gegen CD36 erzielt.

LA und EPA stimulierten Dosis-abhängig die Kardiomyozyten-Differenzierung und Funktion. LA and EPA beschleunigten die Kontraktionsfrequenz der kontraktiven kardialen Areale, erhöhten die Anzahl kontrahierender EBs, die Fläche der kardialen Zellareale sowie die Proteinexpression der kardialen Marker  $\alpha$ -Actinin and MLC2a. LA und EPA erhöhten außerdem die Bildung vaskulärer Strukturen und die Protein-Expression der Endothel-spezifischen Marker FLK-1, PECAM-1 und VE-Cadherin. Im Gegensatz zu LA und EPA konnte keine Stimulation der Vaskulogenese durch die gesättigte Fettsäure PA festgestellt werden.

Unterdrückung der ROS-Erzeugung durch Radikalfänger oder den NADPH-Oxidase-Inhibitor VAS2870 inhibierte die PUFA-vermittelte Vaskulogenese und Kardiomyogenese von ES-Zellen. Darüber hinaus führte die Inhibition der eNOS durch N( $\omega$ )-Nitro-L-Argininmethylester (L-NAME) zur Hemmung der Vaskulogenese von ES-Zellen.

PUFAs stimulierten die AMP-aktivierte Proteinkinase (AMPK- $\alpha$ ) sowie den Peroxisom-Proliferator-aktivierten Rezeptor- $\alpha$  (PPAR- $\alpha$ ). Die PUFA-vermittelte AMPK- $\alpha$ -Aktivierung wurde durch Chelation von Calcium sowie die Hemmung der ROS und NO

## Summary

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inhibiert. Weiterhin wurde die PUFA-induzierte Kardiomyogenese und Vaskulogenese der Stammzellen durch den PPAR- $\alpha$ -Inhibitor GW6471 verhindert.

Zusammenfassend zeigen die Daten der vorliegenden Studie, dass die PUFAs LA und EPA die Kardiomyogenese und Vaskulogenese embryonaler Stammzellen durch Signalkaskaden stimulieren, die von einer Erhöhung der Signalmoleküle Calcium, ROS und NO begleitet, und mit einer Aktivierung der Energiesensoren AMPK- $\alpha$  und PPAR- $\alpha$  verknüpft sind.

## List of abbreviations

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### 7. List of abbreviations

<b>°C</b>	Degrees Celsius
<b>µl</b>	Microliter
<b>µs</b>	Microsecond
<b>8-EETs</b>	8-epoxyeicosatrienoic acids
<b>AA</b>	Arachidonic acid
<b>ADP</b>	adenosine diphosphate
<b>AICAR</b>	5-aminoimidazole-4-carboxamide riboside
<b>AKT</b>	Protein kinase B
<b>ALA</b>	α-linolenic acid
<b>AMP</b>	adenosine monophosphate
<b>AMPK</b>	5' adenosine monophosphate-activated protein kinase
<b>ANOVA</b>	Analysis of variance
<b>ATP</b>	Adenosine triphosphate
<b>BMP</b>	Bone morphogenetic protein
<b>BSA</b>	Bovine serum albumins
<b>Cat. No.</b>	Catalog number
<b>Cav</b>	Voltage-gated calcium channel
<b>CC BY 2.0</b>	Creative Commons Attribution 2.0 Generic
<b>CC BY 4.0</b>	Creative Commons Attribution 4.0 International
<b>CC-BY-NC-ND</b>	Creative Commons Attribution Non-Commercial No Derivatives
<b>CD31</b>	Cluster of differentiation 31
<b>cGMP</b>	Cyclic guanosine monophosphate
<b>CHD</b>	Coronary heart disease
<b>CLSM</b>	Confocal laser scanning microscope
<b>COX</b>	Cyclooxygenases
<b>CPN</b>	Core pluripotency network
<b>CVD</b>	Cardiovascular disease
<b>Cx43</b>	Connexin 43
<b>CYP</b>	Cytochrome P450
<b>D5D</b>	Delta-5 desaturase
<b>D6D</b>	Delta-6 desaturase
<b>DAF-FM DA</b>	4-amino-5-methylamino-2',7'-difluorofluorescein diacetate
<b>DGLA</b>	Dihomo-gamma-linolenic acid
<b>DHA</b>	Docosahexaenoic acid
<b>DMEM</b>	Dulbecco's modified Eagle medium
<b>DMSO</b>	Dimethyl sulfoxide
<b>DNA</b>	Ddeoxyribonucleic acid
<b>DPA</b>	Docosapentaenoic acid
<b>DRAQ5</b>	1,5-bis(2-(di-methylamino)ethylamino)-4,8-dihydroxyanthracene-9,10-dione
<b>DUOX</b>	Dual oxidase
<b>DW</b>	Distilled water
<b>EB</b>	Embryoid body
<b>ECL</b>	Enhanced chemiluminescence
<b>EDTA</b>	Ethylenediaminetetraacetic acid
<b>EETs</b>	Epoxyeicosatrienoic acids
<b>EFA</b> s	Essential fatty acids
<b>EGFR</b>	Epidermal growth factor receptor

## List of abbreviations

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<b>ELOVLs</b>	Elongation of very long-chain fatty acids
<b>eNOS / NOS III</b>	Endothelial nitric oxide synthase
<b>EPA</b>	Eicosapentaenoic acid
<b>ER</b>	Endoplasmic reticulum
<b>ERK</b>	Extracellular signal-regulated kinase
<b>ES cells</b>	Embryonic stem cells
<b>Esrrb</b>	Estrogen-related receptor beta
<b>ETA</b>	Eicosatetraenoic acid
<b>ETC</b>	Electron transport chain
<b>FADS</b>	Fatty acid desaturase
<b>FATPs</b>	Fatty acid transport proteins
<b>FCS</b>	Fetal calf serum
<b>FGF</b>	Fibroblast growth factor
<b>FLAP</b>	5-LOX-activating protein
<b>Flk-1</b>	Fetal liver kinase-1
<b>Fluo-4-AM</b>	4-(6-Acetoxy-methoxy-2,7-difluoro-3-oxo-9-xanthonyl)-4'-methyl-2,2'-(ethylenedioxy)dianiline-N,N,N',N'-tetraacetic acid tetrakis (acetoxymethyl) ester
<b>g</b>	Gram
<b>g</b>	Gravity 1g = 9,81m/s <sup>2</sup>
<b>GAPDH</b>	Glyceraldehyde-3-phosphate dehydrogenase
<b>GLA</b>	Gamma-linolenic acid
<b>h</b>	Hour
<b>H<sub>2</sub>DCF-DA</b>	2',7'-dichlorodihydrofluorescein diacetate
<b>HCl</b>	Hydrochloric acid
<b>HCM</b>	Hypertrophic cardiomyopathy
<b>HDL</b>	High-density lipoprotein
<b>hESCs</b>	Human embryonic stem cells
<b>HETEs</b>	5-hydroxyeicosatetraenoic acids
<b>HF</b>	Heart failure
<b>HIF-1<math>\alpha</math></b>	Hypoxia inducible factor 1- $\alpha$
<b>hiPSC-CMs</b>	Human induced pluripotent stem cell-derived cardiomyocytes
<b>hPSC-CMs</b>	Human pluripotent stem cell-derived cardiomyocytes
<b>HRP</b>	Horseradish peroxidase
<b>ICM</b>	Inner cells mass
<b>IGF-1</b>	Insulin-like growth factor-1
<b>IgG</b>	Immunoglobulin G
<b>IHD</b>	Ischemic heart disease
<b>IL-6</b>	Interleukin-6
<b>iNOS/ NOS II</b>	Inducible nitric oxide synthase
<b>IP3</b>	Inositol trisphosphate
<b>iPSCs</b>	Induced pluripotent stem cells
<b>JNK</b>	c-Jun N-terminal kinase
<b>KDa</b>	Kilodaltons
<b>Klf4</b>	Kkruppel-like factor 4
<b>Kv</b>	Voltage-gated potassium channel
<b>LA</b>	Linoleic acid
<b>LDL</b>	Low-density lipoprotein
<b>LIF</b>	Leukemia inhibitory factor

## List of abbreviations

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<b>LIF-pLpro</b>	LIF-Plasmocin-Prophylactic
<b>LIMP2</b>	Lysosomal integral membrane protein II
<b>L-NAME</b>	N $\omega$ -Nitro-L-arginine methyl ester
<b>LOX</b>	Lipoxygenases
<b>LTs</b>	Leukotrienes
<b>mA</b>	Milliampere
<b>MAPK</b>	Mitogen-activated protein kinase
<b>MEF2C</b>	Myocyte enhancer factor 2C
<b>MEFs</b>	Mouse embryonic fibroblasts
<b>mES cells</b>	Mmouse embryonic stem cells
<b>mg</b>	Milligram
<b>MI</b>	Myocardial infarction
<b>min</b>	Minute
<b>ml</b>	Milliliter
<b>MLC2a</b>	Myosin regulatory light chain 2, atrial isoform
<b>mm</b>	millimeter
<b>mM</b>	Millimolar
<b>MUFAs</b>	Monounsaturated fatty acids
<b>mW</b>	Milliwatts
<b>NA</b>	Numerical aperture
<b>MW</b>	Molecular weight
<b>NADPH</b>	Nicotinamide adenine dinucleotide phosphate
<b>Nanog</b>	Nanog homeobox protein
<b>Nav</b>	Voltage-gated sodium channel
<b>NEAA</b>	Non-essential amino acids
<b>nm</b>	Nanometers
<b>NMPG</b>	N-(2-mercaptopropionyl)-glycine
<b>nNOS/ NOS I</b>	Neuronal nitric oxide synthase
<b>NO</b>	Nitric oxide
<b>NOS</b>	Nitric oxide synthase
<b>NPD1</b>	Neuroprotectin D1
<b>Oct4</b>	Octamer-binding transcription factor 4
<b>PA</b>	Palmitic acid
<b>p-AMPK-<math>\alpha</math></b>	Phosphorylation of AMP-activated protein kinase- $\alpha$ -subunits
<b>PBS</b>	Phosphate buffered saline
<b>PD1</b>	protectins D1
<b>PECAM-1</b>	Platelet endothelial cell adhesion molecule-1
<b>p-eNOS</b>	Phosphorylation of Endothelial Nitric-oxide Synthase
<b>PGRN</b>	Pluripotency gene regulatory network
<b>PGs</b>	Prostaglandins
<b>PI3K</b>	Phosphatidylinositol-3-kinase
<b>PKC</b>	Protein kinase C
<b>PKG-1</b>	Protein kinase G
<b>PPAR-<math>\alpha</math></b>	Peroxisome proliferator-activated receptor alpha
<b>p-PPAR-<math>\alpha</math></b>	Phosphorylation of peroxisome proliferator-activated receptor $\alpha$ alpha
<b>PUFAs</b>	Polyunsaturated fatty acids
<b>RIPA</b>	Radioimmunoprecipitation assay
<b>RNS</b>	Reactive nitrogen species
<b>ROS</b>	Reactive oxygen species

## List of abbreviations

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<b>RT</b>	Room temperature
<b>Rvs</b>	Resolvins
<b>RyRs</b>	Ryanodine receptors
<b>s</b>	Second
<b>SCARB3</b>	Scavenger receptor class B member 3
<b>SDA</b>	Stearidonic acid
<b>SDS</b>	Sodium dodecyl sulfate
<b>Ser</b>	Serine
<b>sGC</b>	Soluble guanylate cyclase
<b>SMCs</b>	Smooth muscle cells
<b>SOD</b>	Superoxide dismutase
<b>SOX2</b>	Sex-determining region Y/Box 2
<b>SR</b>	Sarcoplasmic reticulum
<b>SR-B1</b>	Scavenger receptor-B1
<b>SSO</b>	Sulfo-N-succinimidyl oleate
<b>STAT3</b>	Signal transducer and activator of transcription 3
<b>TBST</b>	Tris-buffered saline with 0.1% Tween
<b>Tbx3</b>	T-box transcription factor 3
<b>Thr</b>	Threonine
<b>TIFF</b>	Tagged image file format
<b>TXs</b>	Thromboxanes
<b>V</b>	Voltage
<b>v/v</b>	volume/volume
<b>VE-Cadherin</b>	Vascular endothelial cadherin
<b>VEGF</b>	Vascular endothelial growth factor
<b>VEGFRs</b>	Vascular endothelial growth factor receptors
<b>VLCFAs</b>	Very long-chain fatty acids
<b>w/v</b>	Wweight/volume
<b>XO</b>	Xanthine oxidase
<b>Δ12,14-PGJ2</b>	15-Deoxy-delta(12,14)-prostaglandin J(2)
<b>μm</b>	Micrometer
<b>μM</b>	Micromolar
<b>ω-3</b>	Omega-3
<b>ω-6</b>	Omega-6

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## Publications

### Parts of this thesis have been published in:

- **Taha, A.**, Sharifpanah, F., Wartenberg, M., & Sauer, H. (2020). Omega-3 and Omega-6 polyunsaturated fatty acids stimulate vascular differentiation of mouse embryonic stem cells. *Journal of Cellular Physiology* 235(10), 7094-7106. doi: 10.1002/jcp.29606.
- Ali, E. H., Sharifpanah, F., **Taha, A.**, Tsang, S. Y., Wartenberg, M., & Sauer, H. (2018). The Milk Thistle (Silybum marianum) Compound Silibinin Inhibits Cardiomyogenesis of Embryonic Stem Cells by Interfering with Angiotensin II Signaling. *Stem cells International* 2018:9215792. doi: 10.1155/2018/9215792.

### Poster Presentations

- **Amer Taha, Fatemeh Sharifpanah, Maria Wartenberg & Heinrich Sauer.** Omega-6 and Omega-3 polyunsaturated fatty acids stimulate vascular differentiation of mouse embryonic stem cells. Excellence Cluster Cardio-Pulmonary System ECCPS Retreat 2016. Hotel Dolce Bad Nauheim 07th 08th Jul. 2016. Germany.
- **Amer Taha, Fatemeh Sharifpanah, Maria Wartenberg & Heinrich Sauer.** Omega-6 and Omega-3 polyunsaturated fatty acids stimulate vascular differentiation of mouse embryonic stem cells. Joint International Meeting in vascular Biology 26th - 28th Sept. 2016. Goethe University Hospital, Frankfurt. Germany.
- **Amer Taha, Fatemeh Sharifpanah, Yu-Han Huang, Maria Wartenberg & Heinrich Sauer.** Omega-6 and Omega-3 polyunsaturated fatty acids stimulate vasculogenesis and cardiomyogenesis of embryonic stem cells. The annual meeting of the German Physiology Society (DPG) 16th – 18th Mar. 2017. The University of Greifswald, Institute of Physiology, Greifswald. Germany.
- **Amer Taha, Fatemeh Sharifpanah, Yu-Han Huang, Maria Wartenberg & Heinrich Sauer.** Omega-6 and Omega-3 polyunsaturated fatty acids stimulate vasculogenesis and cardiomyogenesis of embryonic stem cells. ECCPS Symposium Poster Presentation-Area Cellular Plasticity and Heterogeneity 7th – 8th June 2017. Bad Nauheim, Germany.

**Declaration**

I declare that I have completed this dissertation single-handedly without the unauthorized help of a second party and only with the assistance acknowledged therein. I have appropriately acknowledged and referenced all text passages that are derived literally from or are based on the content of published or unpublished work of others, and all information that relates to verbal communications. I have abided by the principles of good scientific conduct laid down in the charter of the Justus Liebig University of Giessen in carrying out the investigations described in the dissertation.

**Amer Hussein Taha**

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