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Cardiovascular Disease and mTOR Signaling

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Abstract

The cell signaling pathways of the mammalian target of rapamycin (mTOR) are broad in nature, but are tightly integrated through the protein complexes of mTORC1 and mTORC2. Although both complexes share some similar subcomponents, mTORC1 is primarily associated with the regulatory protein Raptor while mTORC2 relies upon Rictor. Pathways of mTOR that partner with Wnt as well as growth factor signaling are vital for endothelial and cardiomyocyte growth. In mature differentiated endothelial cells and cardiac cells, mTOR activation regulates both apoptotic and autophagic pathways during oxidative stress that can be dependent upon the activation of protein kinase B (Akt). These protective pathways of mTOR can promote angiogenesis and limit acute cell death to foster cardiac repair and tissue regeneration. However, under some conditions, blockade of mTOR pathways may be necessary to limit vasculopathy and promote microcirculatory flow. Future work that further elucidates the vital regulatory pathways of mTOR can offer new therapeutic insights for the treatment of cardiovascular diseases.

Keywords

Akt; cardiac; endothelial; erythropoietin; TORC1; TORC2; Wnt; wingless

Introduction

The signaling pathways of the mammalian target of rapamycin (mTOR) are closely linked to the formation of two different protein complexes (Chong et al. 2010; Hwang and Kim 2011) (Figure 1). The first complex, mTOR Complex 1 (mTORC1), relies upon the regulatory-associated protein of mTOR (Raptor) protein to enable mTORC1 to bind to its substrates. mTORC1 is also composed of the proline rich Akt substrate 40 kDa (PRAS40), the DEP domain-containing mTOR interacting protein (Deptor), and the mammalian lethal with Sec13 protein 8 (mLST8). mTORC1 controls the serine/threonine kinase ribosomal protein p70S6K and the eukaryotic initiation factor 4E-binding protein 1 (4EBP1). When 4EBP1 is hypophosphorylated, it can block protein translation by binding to eukaryotic translation initiation factor 4 epsilon (eIF4E) through the eukaryotic translation initiation factor 4

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gamma (eIF4G), a protein that shepherds mRNA to the ribosome. Phosphorylation of 4EBP1 by mTORC1 leads to the dissociation of 4EBP1 from eIF4E to allow eIF4G to begin mRNA translation. mTORC1 phosphorylation also increases the kinase activity of p70S6K. Phosphorylation of the p70S6K serine/threonine kinase by mTORC1 results in mRNA biogenesis, translation of ribosomal proteins, and cell growth. In addition, PRAS40 can block the binding of the mTORC1 substrates p70S6K and 4EBP1 to Raptor (Chong et al. 2010; Hwang and Kim 2011).

The second mTOR complex, mTORC2, is similar to mTORC1 in that it also is composed of mTOR, mLST8, and Deptor (Figure 1). However, mTORC2 has Rictor as a component rather than Raptor and associates with the mammalian stress-activated protein kinase interacting protein (mSIN1) and protein observed with Rictor-1 (Protor-1). Rictor is not sensitive to rapamycin and promotes the activity of mTORC2. mTORC2 controls actin cytoskeleton organization, cell size, endothelial cell survival and migration, and cell cycle progression. One target of mTORC2 is protein kinase B (Akt). Rictor allows mTORC2 to phosphorylate Akt at Ser⁴⁷³ to lead to its activation and to facilitate threonine³⁰⁸ phosphorylation by phosphoinositide-dependent kinase 1 (PDK1). mTORC2 also regulates protein kinase C (PKC), P-Rex1, P-Rex2, Rho GTPases, and Rho signaling pathways that control cell to cell contact (Chong et al. 2010; Hwang and Kim 2011).

Stem cell proliferation

Expression of mTOR occurs in many systems of the body that include the cardiac, pulmonary, immune, reproductive, and gastrointestinal systems (Chong et al. 2010; Hwang and Kim 2011). As a result, mTOR can have a significant role in cell development with stem cell regulation (Table 1). In endothelial cells, mTOR may be necessary for endothelial progenitor cell development since inhibition of mTOR pathways with rapamycin lead to endothelial progenitor cell death that may result from inhibiting growth factor signaling (Miriuka et al. 2006). Growth factors, such as erythropoietin (EPO), rely upon mTOR pathway signaling. EPO controls angiogenesis, endothelial survival, and cardiomyocyte protection (Ammar et al. 2011; Maiese et al. 2005). EPO requires mTOR signaling for microglia survival during oxidative stress (Shang et al. 2011) and for osteoblastogenesis and osteoclastogenesis (Kim et al. 2012). However, in hematopoietic stem cells, mTOR may be associated with aging since mTOR activity is increased in the hematopoietic stem cells of older mice (Chen et al. 2009).

In relation to cardiac tissue, human embryonic stem cell-derived cardiomyocyte growth may be dependent upon mTOR, since pharmacological loss of mTOR can limit stem cell growth (Foldes et al. 2011). In general, it is believed that mTOR may have an important role for the proliferation and differentiation of embryonic stem cells. Deletion of the C-terminal six amino acids of mTOR that control kinase activity leads to a decrease in cell size and limits the proliferation of embryonic stem cells (Murakami et al. 2004). In addition, ablation of mouse mTOR gene results in the early lethality and arrest of embryonic stem cell proliferation (Gangloff et al. 2004). The mTOR pathway also has an important role in the maintenance of pluripotency and differentiation of cells. As a downstream target of mTOR, p70S6K is a critical factor for protein translational control. Expression of constitutively active p70S6K or siRNA-mediated knockdown of both tuberous sclerosis complex tuberin (TSC2) and Rictor to increase p70S6K activation leads to the differentiation of human embryonic stem cells (Easley et al. 2010). The activity of mTOR is also essential for the long-term renewal of human embryonic stem cells, since inhibition of mTOR impairs pluripotency, prevents cell proliferation, and enhances mesoderm and endoderm activities in embryonic stem cells (Zhou et al. 2009).

Stem cell proliferation and homeostasis through mTOR may require partnering with other pathways, such as the wingless pathway. Wnt proteins and their signaling pathways are vital to multiple cell process that involve stem cell proliferation, cell development, and cellular survival (Maiese et al. 2008). The Wnt pathway can increase the activity of mTOR through glycogen synthase kinase-3 β (GSK-3 β) (Chong et al. 2010). GSK-3 β phosphorylates TSC2 on serine 1341 in combination with the AMP activated protein kinase (AMPK) phosphorylation of TSC2 on serine 1345 that results in the inhibition of mTOR activity. Wnt proteins counteract mTOR inhibition and foster mTOR activity by inhibiting GSK-3 β through phosphorylation. In hematopoietic stem cells, the balance between Wnt and GSK-3 β activation controls self renewal and lineage commitment (Huang et al. 2009).

Apoptosis, autophagy, and oxidative stress

Oxidative stress has a vital role in the onset of cardiovascular injury and can affect multiple related systems in the body that affect metabolic homeostasis, ischemic disease, and cardiopulmonary function (Maiese et al. 2010). The release of reactive oxygen species (ROS) leads to oxidative stress and promotes both cell injury as well as aging pathways. ROS can be generated in excessive quantities through different sources such as superoxide free radicals, hydrogen peroxide, singlet oxygen, nitric oxide (NO), and peroxynitrite. Under normal physiological conditions, ROS are scavenged by endogenous antioxidant systems that include superoxide dismutase, glutathione peroxidase, catalase, and vitamins that include C, D, E, and K.

Both apoptosis and autophagy may result in cardiac injury through oxidative stress (Figure 1). Apoptotic pathways can lead to cardiac failure, cardiomyocyte injury, cardiac metabolic disease, and reperfusion injury (Ammar et al. 2011; Das et al. 2011). Apoptosis has at least two phases that involve the early exposure of membrane phosphatidylserine (PS) residues and the subsequent destruction of genomic DNA (Chong et al. 2005). Externalization of membrane PS residues occur initially during cellular apoptosis and are a signal for the phagocytosis of cells. Apoptotic membrane PS exposure occurs in a broad array of cell that include cardiac, vascular, and inflammatory cells. The loss of membrane phospholipid asymmetry leads to the exposure of membrane PS residues on the cell surface and assists inflammatory cells to identify injured cells for phagocytosis. Exposure of membrane PS residues also affects the cardiovascular system since membrane PS externalization also occurs on platelets and has been associated with clot formation in the vascular system (Popescu et al. 2010).

Regulation of apoptosis by mTOR is mediated through 4EBP1 and p70S6K (Table 1). Activation of p70S6K by mTOR blocks apoptosis through pathways that can increase "antiapoptotic" Bcl-2/Bcl-x_L expression and inactivate the "pro-apoptotic" protein BAD (Pastor et al. 2009). In the absence of mTOR activity, 4EBP1 also binds to eIF4E to result in the translation of pro-apoptotic proteins and cardiac injury (Zhang et al. 2010). Blockade of apoptosis by mTOR also relies upon the serine-threonine kinase Akt. Akt activation leads to enhanced cell growth and cardiac protection during oxidative stress, endothelial cell injury, metabolic disease, and cardiac hypertrophy (Maiese et al. 2010). mTOR has been shown to require Akt activation to protect endothelial cells against apoptosis (Dormond et al. 2007) and mediate protection through the inactivation of forkhead transcription factors, such as FoxO3a (Chong et al. 2011; Dormond et al. 2007). Akt also functions to modulate apoptosis with mTOR through the inhibition of PRAS40 which can lead to the activation of apoptotic pathways (Thedieck et al. 2007). Phosphorylation of PRAS40 by Akt can inhibit the activity of this substrate and lead to its dissociation from mTORC1 and binding to cytoplasmic 14-3-3 proteins (Nascimento et al. 2010).

Autophagy is a process in which cells are able to recycle cytoplasmic components and dispose of defective organelles. This allows the early destruction of organelles and other cytoplasmic components but the preservation of cytoskeletal structures that is in contrast to processes that occur during apoptosis. Yet, autophagy can lead to cell death, such as during acute ischemic injury (Xin et al. 2011). In cardiomyocytes, cardiotoxicity by kinase inhibitor agents can be blocked during gene silencing of proteins responsible for autophagosome formation that involve Beclin 1 (autophagy-related gene 6 (Atg6)) (Zhao et al. 2010). However, autophagic degradation during normal physiology may be necessary with mTOR modulation to some degree since loss of Vps34, a known regulator of autophagy, can depress mTOR activation and result in cardiomegaly and decreased cardiac contractility (Jaber et al. 2012). In addition, benefits of exercise may require a brief inaction of mTOR for autophagic pathways to proceed (Ogura et al. 2011). In other circumstances, mTOR activation appears vital to block autophagy, possibly limit forkhead transcription FoxO3a activity, and prevent cardiac atrophy and dysfunction (Schips et al. 2011) (Table 1).

Angiogenesis, ischemic Injury, and cardioprotection

Angiogenesis, the process of new capillary formation, is an important component of cardiac tissue protection and regeneration that can be regulated by mTOR (Maiese et al. 2009). Inhibition of mTOR signaling, such as during the exposure to cigarette smoke, leads to a sequence of events with elevated matrix metalloproteinase-1, the blockade of tissue inhibitor of metalloproteases-3 that, and subsequent impaired angiogenesis (Lemaitre et al. 2011). Loss of mTOR activity also blocks endothelial proliferation and angiogenesis (Humar et al. 2002) as well as the proliferation of endothelial progenitor cells (Miriuka et al. 2006). Without effective angiogenesis following cardiac injury, tissue repair and regeneration may be limited or not occur at all (Table 1).

In regards to ischemic cardiac injury, activation of mTOR can be protective against oxidant ischemic/reperfusion injury in cardiomyocytes (Hernandez et al. 2011). Cardiac protection by mTOR may require the inhibition of GSK-3 β through Wnt signaling (Vigneron et al. 2011), similar to other survival pathways that are both GSK-3 β and Wnt dependent (Shang et al. 2012) (Figure 1). During periods of ischemic post-conditioning, mTOR is necessary to prevent apoptotic cardiac death (Wagner et al. 2010). In studies that block the phosphorylation and activation of p70S6K, cardiac infarct area and scar tissue are increased, further illustrating a cardioprotective role for the mTOR signaling pathway (Lajoie et al. 2009). It is important to note that prolonged activation of mTOR may have detrimental consequences to the cardiac system. Under these circumstances, inhibition of mTOR may be desirable. For example, chronic activation of mTOR can lead to vascular dysfunction. (Popescu et al. 2010) Inhibition of mTOR for prolonged treatment has been shown to reduce vasculopathy (Mancini et al. 2003) and to improve coronary flow through the microcirculation following cardiac transplantation (Sinha et al. 2008).

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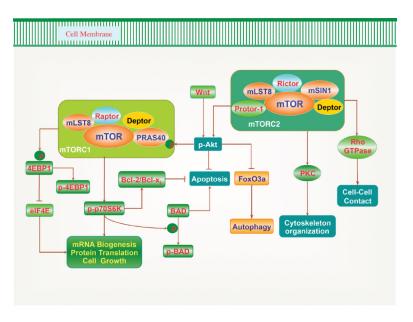


Figure 1. Cellular signaling of mTOR

The mammalian target of rapamycin (mTOR) functions through two complexes, mTORC1 and mTORC2. The components of mTORC1 include the catalytic multiple protein mTOR, regulatory-associated protein of mTOR (Raptor), proline rich Akt substrate 40 kDa (PRAS40), mammalian lethal with Sec13 protein 8 (mLST8), and DEP-domain-containing mTOR-interacting protein (Deptor). Once active, mTORC1 phosphorylates its two downstream targets, p70 ribosomal S6 kinase (p70S6K) and eukaryotic initiation factor 4E (eIF4E)-binding protein 1 (4EBP1), to regulate mRNA biogenesis, translation, and cell growth. p70S6K also promotes the phosphorylation of pro-apoptotic protein BAD and enhances the expression of anti-apoptotic protein Bcl-2/Bcl-x_L. In addition to mTOR, mSLT8, and Deptor, mTORC2 also contains rapamycin-insensitive companion of mTOR (Rictor), mammalian stress-activated protein kinase interacting protein (mSIN1) and protein observed with Rictor-1 (Protor-1). mTORC2 primarily regulates cytoskeleton organization of the cell through its major downstream targets, protein kinase B (Akt) and protein kinase C (PKC). mTORC2 can also activate Rho GTPases and control cell to cell contact via Rho signaling pathways. Wnt can lead to Akt activation. Subsequently, Akt can directly phosphorylate PRAS40 to inhibit apoptosis and control autophagy through forkhead transcription factors, such as FoxO3a.

Table 1
Regulatory Roles of mTOR in Cardiovascular System

Targets	Biological function of mTOR
Stem Cells	Regulates proliferation/differentiation of embryonic stem cells Maintains the pluripotency of embryonic stem cells Promotes long-term renewal of embryonic stem cells Associates with aging of hematopoietic stem cells Mediates erythropoietin induced osteoblastogenesis and osteoclastogenesis Promotes endothelial progenitor cell development
Programmed Cell Death	Modulates apoptosis and autophagic cell death
Angiogenesis	Promotes the proliferation of endothelial progenitor cells and endothelial cells Prevents the induction of matrix metalloproteinases Promotes angiogenesis
Cardiovascular Disease	Protects cardiomyocytes against ischemia/reperfusion Inhibits autophagy in cardiomyocytes to prevent cardiac atrophy Prolonged activation leads to vasculopathy