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Targeting disease through novel pathways of apoptosis and autophagy

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Abstract

Introduction—Apoptosis and autophagy impact cell death in multiple systems of the body. Development of new therapeutic strategies that target these processes must address their complex role during developmental cell growth as well as during the modulation of toxic cellular environments.

Areas covered—Novel signaling pathways involving Wnt1-inducible signaling pathway protein 1 (WISP1), phosphoinositide 3-kinase (PI3K), protein kinase B (Akt), β -catenin and mammalian target of rapamycin (mTOR) govern apoptotic and autophagic pathways during oxidant stress that affect the course of a broad spectrum of disease entities including Alzheimer's disease, Parkinson's disease, myocardial injury, skeletal system trauma, immune system dysfunction and cancer progression. Implications of potential biological and clinical outcome for these signaling pathways are presented.

Expert opinion—The CCN family member WISP1 and its intimate relationship with canonical and non-canonical *wingless* signaling pathways of PI3K, Akt1, β -catenin and mTOR offer an exciting approach for governing the pathways of apoptosis and autophagy especially in clinical disorders that are currently without effective treatments. Future studies that can elucidate the intricate role of these cytoprotective pathways during apoptosis and autophagy can further the successful translation and development of these cellular targets into robust and safe clinical therapeutic strategies.

Keywords

β -catenin; Akt; Alzheimer's disease; apoptosis; autophagy; Beclin 1; cancer; cardiovascular disease; caspase; CCN family; diabetes mellitus; erythropoietin; forkhead transcription factors; FoxO; glycogen synthase kinase-3 β ; mammalian target of rapamycin; neurodegenerative disease; oxidative stress; Parkinson's disease; phosphoinositide 3-kinase; programmed cell death; Wnt1-inducible signaling pathway protein 1

1. Introduction

The demise of a cell can occur through a number of pathways that involve apoptosis (type I cell death), autophagy (type II cell death) and necrosis (type III cell death). Apoptosis is

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considered to be one variant of programmed cell death (PCD) that is a directed process that involves reduction in cell size, chromatin condensation and nuclear DNA fragmentation [1]. Autophagy represents another form of PCD that is highly regulated to reduce and recycle cell components into smaller units through lysosomes that leads to the degradation of cellular organelles [2]. Other forms of PCD can include anoikis that describes apoptotic death to cells that lose contact with the extracellular matrix and cornification that is used to maintain the integrity of the epidermis through the replacement of cellular components with keratin. Although Carl Vogt in 1842 initially described the process of apoptosis, it was not until several years later, such as in 1965, that investigators began to take a greater interest in distinguishing apoptotic cell death from necrotic cell death with the work of John Kerr [3,4]. Early descriptions of autophagy surfaced approximately during the same time period with the identification of several of the essential components of autophagy that includes [5] autophagic vacuoles [6]. In contrast to the processes of PCD that may be beneficial such as during the development of an organism, necrosis is an unexpected and premature death of cells as a result of exposure to environmental toxins or trauma that is usually not considered to be beneficial [7].

In disorders involving multiple disease entities and several systems of the body such as the nervous system, immune system, cardiac system, skeletal system and vascular system, PCD that is associated with apoptosis or autophagy has recently been shown to lead to cell demise. As a result, many proposed treatments for the development of new therapeutic strategies seek to modulate apoptosis or autophagy to either prevent cell injury or foster cellular repair and regeneration. However, a fine balance within the processes of apoptosis and autophagy exists since activation of these pathways is sometimes necessary for developmental cell growth or protection against toxic environments. As a result, development of new avenues of treatment must address these challenges to enhance clinical efficacy and limit potential disability or death. Here, the authors discuss an integrated series of novel signaling pathways that involve Wnt1-inducible signaling pathway protein 1 (WISP1), phosphoinositide 3-kinase (PI3K), protein kinase B (Akt), β -catenin and mammalian target of rapamycin (mTOR) signaling for future clinical development that are increasingly being identified as primary targets against apoptotic or autophagic cell death.

2. Cell injury through oxidative stress, apoptosis and autophagy

Reactive oxygen species (ROS) are formed through agents such as superoxide free radicals, hydrogen peroxide, singlet oxygen, nitric oxide (NO) and peroxynitrite and lead to oxidative stress [7,8]. Under normal physiological conditions, ROS exist at non-detrimental levels and are kept in check by endogenous antioxidant systems that include superoxide dismutase, catalase, glutathione peroxidase and vitamins C, D, E and K [9–11]. Yet, early studies suggested that exposure to elevated oxygen, a high metabolic rate and ROS could reduce lifespan expectations [12]. Once produced, ROS affect mitochondrial function, DNA integrity and protein folding leading to cellular injury. More current studies have now linked oxygen free radical production to DNA damage in diabetic patients [13], mitochondrial injury and aging mechanisms [14] and nutritional impairment [15]. Oxidative stress can affect multiple systems of the body and result in cardiac disease [16], vasculature dysfunction [17,18], immune-mediated disorders [19–21], aging [22], cerebral ischemia [23,24], diabetic complications [25,26] and cognitive loss [27]. Ultimately, oxidative stress that has pathological consequences can lead to cell death through apoptosis or autophagy (Figure 1).

Two primary components of apoptosis consist of an early phase that involves the exposure of membrane phosphatidyl-serine (PS) residues and a late phase that involves the destruction of genomic DNA (Figure 1) [7]. The early phase can tag cells with the membrane

externalization of PS residues and essentially function as a 'homing device' for inflammatory cells to engulf and remove injured cells [28,29]. For this to occur such as during periods of oxidative stress, inflammatory cells increase their expression of the membrane phosphatidylserine receptor (PSR) with a concurrent increase in the proliferation and activation of the inflammatory cells. It is important to note that as a vital therapeutic component, modulation of inflammatory cell activation may be required since removal of temporarily injured cells expressing membrane PS residues may sometimes result in the loss of otherwise functional cells [30,31]. As a potential therapeutic strategy, treatment agents can target the reversal of membrane PS externalization on temporarily injured cells with a simultaneous blockade of the PSR receptor on inflammatory cells to limit their activation and proliferation during apoptosis. Externalization of membrane PS residues can result from a broad array of toxic injuries that result in oxidative stress, such as during oxygen radical exposure, infection, ischemia, vascular clot formation and β -amyloid deposition [32–34].

The second or later phase of apoptosis consists of the cleavage of genomic DNA (Figure 1) [35–37]. Once initiated, genomic DNA degradation usually is not a reversible step. Enzymes responsible for DNA degradation include the acidic cation-independent endonuclease (DNase II), cyclophilins and the 97 kDa magnesium-dependent endonuclease. Three separate endonuclease activities also exist in the nervous system that include a constitutive acidic cation-independent endonuclease, a constitutive calcium/magnesium-dependent endonuclease and an inducible magnesium-dependent endonuclease [1].

During apoptotic cell injury, caspases are activated [38]. Caspases are usually synthesized as inactive zymogens that are proteolytically cleaved into subunits at the onset of apoptosis and function as active caspases after reconstitution to molecular heterodimers. Caspases are composed of three domains including an N-terminal pro-domain, a large subunit and a small subunit. As a result of their activation sequence, caspases are classified as either initiator caspases (also known as apical caspases) or effector caspases. An initiator caspase cleaves and subsequently activates an effector caspase. The apoptotic-associated caspases include initiator caspases, such as caspase 2, 8, 9 and 10, that activate downstream effector caspases leading to an amplification of cascade activity. The initiator caspases consist of long N-terminal pro-domains that contain caspase recruitment domains (CARDs) in caspase 2 and caspase 9 or death effector domains (DEDs) in caspase 8 and caspase 10. The effector caspases consist of caspase 1, 3, 6 and 7 that function to directly cleave crucial cellular protein substrates that result in cell death. The effector caspases contain either short pro-domains or are absent of pro-domains [39].

Activation of caspases can occur through two distinct pathways, namely the extrinsic and intrinsic pathways [1,39]. The extrinsic pathway is initiated by death receptor activation at the cell surface resulting in the recruitment and activation of the initiator caspase 8 or caspase 10 on apoptotic stimuli. The intracellular death domain of death receptors, such as the TNF superfamily Fas/CD95/Apo-1, can bind to extracellular ligands and lead to an intracellular death-inducing signaling complex following recruitment of adaptor molecules, such as the Fas-associated death domain (FADD). FADD recruits caspase 8 and 10 through its DED domain to result in the activation of caspase 8 and 10. Activation of caspase 8 can subsequently lead to caspase 3 activation. In addition, active caspase 8 can cleave BH3-only protein Bid, a pro-apoptotic member of the Bcl-2 family and result in truncated Bid (tBid) that promotes cytochrome *c* release through Bax resulting in the subsequent activation of executioner caspases. In regards to activity in the intrinsic caspase pathway, this involves mitochondrial membrane depolarization that is associated with the release of cytochrome *c* and the subsequent activation of caspase 9 followed by activation of caspase 3. The process is regulated by the Bcl-2 subfamily BH3-only proteins including Bid, Bad, Bim, Bmf, Puma and Noxa, which are normally located in cellular compartments other than mitochondria.

The translocation of these proteins to mitochondria delivers an apoptotic signal through the interaction with Bax, a multiple Bcl-2 homology domain containing protein, to promote permeabilization of the outer mitochondrial membrane and the release of cytochrome *c* that then binds to apoptotic protease-activating factor-1 (Apaf-1). Apaf-1 consists of three different domains that include CARD, repeats of tryptophan and aspartate residues (WD-40 repeats) and a nucleotide-binding domain CED-4. Binding of cytochrome *c* to Apaf-1 results in the removal of the WD-40 domain, masking the CED-4 and CARDS, and leads to the oligomerization of Apaf-1 with the requirement of dATP/ATP. The oligomerization of Apaf-1 promotes the allosteric activation of caspase 9 by forming the Apaf-1 apoptosome. Caspase 9 can subsequently activate caspase 3 as well as caspase 1 through the intermediary caspase 8. Activation of caspase 1 and caspase 3 not only leads to DNA fragmentation, but also results in membrane PS exposure [40,41].

Apoptotic injury is believed to be a significant contributor to the pathogenesis of a variety of disorders. For example, in the brains of patients with Alzheimer's disease, apoptotic DNA fragmentation [42] and caspase activation has been observed [43]. Experimental models of Alzheimer's disease also have identified apoptotic proteins in the brain [44]. Apoptotic neuronal nuclei and caspase 3 has been identified in the postmortem nigra of Parkinson's disease patients, suggesting that neuronal loss during Parkinson's disease is a result of apoptosis [45]. During cardiac transplant rejection, both early apoptotic membrane PS exposure and later caspase 3 activation has been reported [46]. Furthermore, injury to cells of the immune system, such as microglia, is believed to be mediated through apoptotic mechanisms [47,48].

Autophagy has three different categories termed microautophagy, macroautophagy and chaperone-mediated autophagy [49]. In general, autophagy allows cells to recycle cytoplasmic components, remove defective organelles and maintain important cytoskeletal structures during development, cell differentiation and tissue remodeling [50]. Macroautophagy includes the bulk degradation of cytoplasmic material and the sequestration of the cytoplasmic protein and organelles into autophagosomes. The autophagosomes fuse with lysosomes for degradation and reuse by essential cellular processes [51]. In most descriptions of autophagy, macroautophagy is usually depicted. Microautophagy is the sequestration of cytoplasmic components by invagination of the lysosomal membrane. Vesicles subsequently formed are transferred to the lumen of the lysosomes for digestion. In chaperone-mediated autophagy, the cytoplasmic component is delivered by cytosolic chaperones to the receptors on the lysosomal membranes and the cellular organelle is translocated across lysosomal membranes into the lumen.

Induction of autophagy can occur through many environmental stimuli but is frequently described during oxidative stress and nutrient depletion (Figure 1). In some circumstances, activation of autophagy has been reported to be potentially cytoprotective such as during neurodegenerative disorders [52,53] and in the setting of prion protein-mediated neurotoxicity [54]. For example, in Parkinson's disease, autophagy has been associated with the processing of the protein α -synuclein. Mutation of α -synuclein and accumulation of wild-type α -synuclein in dopaminergic neurons have been associated with progression of Parkinson's disease [55]. Of the three types of autophagy, both chaperone-mediated autophagy and macroautophagy are involved in the degradation of α -synuclein. Chaperone-mediated autophagy appears to be more critical for the clearance of aberrant α -synuclein in neurons since inhibition of this autophagic pathway leads to accumulation of high molecular weight and detergent insoluble α -synuclein. This leads to neurotoxicity and further inhibition of chaperone-mediated autophagy [55]. Mutant α -synuclein, which is poorly internalized into lysosomes, is degraded by macroautophagy. As a result, activation of

autophagic pathway protects against neurodegeneration and α -synuclein in Parkinson's disease [52].

However in a number of scenarios, autophagy may be detrimental to cell survival. During cellular ischemia, autophagy results in cell death in cerebral astrocytes [56], spinal cord motor neurons [57] and cortical neurons [58]. Autophagic cell death also occurs during growth factor deprivation in Purkinje neurons [59] and in sympathetic neurons [60]. Other toxins such as glutamate, potassium deprivation and staurosporine lead to autophagy [59]. Interestingly, the pathways of autophagy and apoptosis may be closely aligned. For example, methamphetamine leads to cell death not only through apoptosis, but also through autophagy by inhibiting the disassociation of the Bcl-2/Beclin 1 complex [61]. Bcl-2/Bcl-x_L is an antiapoptotic protein and a protein that blocks autophagy through its inhibitory interaction with Beclin 1 (Figure 1) [62]. In other circumstances, autophagy and apoptosis may have opposing roles. Some studies report that progression of apoptosis may conversely require the inhibition of autophagy [63–65]. Other work also suggests that some pathways may function as a switch between autophagy that is associated with cell survival and autophagy that is associated with cell death. For example, Draper, the *Drosophila melanogaster* ortholog of the *Caenorhabditis elegans* engulfment receptor CED-1, is required for autophagy during cell death in *Drosophila* salivary glands. Knockdown of Draper has been shown to prevent autophagy in dying salivary glands [66]. However, in the fat body in which autophagy is associated with cell survival, Draper knockdown does not prevent starvation-induced autophagy, suggesting that Draper can have dual roles during autophagy and may be one factor that can distinguish between autophagy-associated cell death and autophagy-associated cell survival [66].

3. The CCN family member WISP1

Initially identified as a downstream target of the *wingless* Wnt1 signaling pathway and in the mouse mammary epithelial cell line C57MG transformed by Wnt1, WISP1 was later associated with neoplastic growth in the gastrointestinal tract [67]. WISP1 is expressed in several tissues including the epithelium, heart, kidney, lung, pancreas, placenta, ovaries, small intestine, spleen and brain. WISP1 is a member of the CCN family of proteins and is also known as CCN4. The CCN family of proteins is defined by the first three members of the family that include cysteine-rich protein 61, connective tissue growth factor and nephroblastoma overexpressed gene and consists of six secreted extracellular matrix-associated proteins. Each family member contains four cysteine-rich modular domains that include insulin-like growth factor-binding domain, thrombospondin domain, von Willebrand factor type C module and C-terminal cysteine knot-like domain. This connective tissue growth factor protein family has multiple cellular functions that include skeletal system development, vascular repair, cellular survival and extracellular matrix growth. In regards to the extracellular matrix, WISP1 can bind to leucine-rich proteoglycans that may affect the ability of other cells to anchor to the extracellular matrix [68].

WISP1 has been shown to have elevated expression during events such as cardiac ischemia [69], neuronal exposure to oxidative stress [70,71], lung epithelial damage [72] and cellular repair of fractured bone [73,74]. The increased expression of WISP1 may be suggestive of a reparative and regenerative process that is controlled by this CCN family member. Furthermore, WISP1 is a target of the *wingless* pathway Wnt1, a cysteine-rich glycosylated protein that controls neuronal development, angiogenesis, tumorigenesis and stem cell proliferation (Figure 2) [75,76]. Wnt1 is upregulated during cortical injury [23], on endothelial cell [35,77] and pancreatic cell [78] exposure to elevated glucose [35,77], during spinal cord injury [79], in reactive central nervous system astrocytes [80], in settings of intestinal inflammation [81], during vascular cell aging [82] and during neurodegenerative

disease [83]. Wnt1 prevents apoptotic cell injury during stroke injury to the brain [23], protects against ethanol toxicity in osteoblasts [84], blocks cell loss in dopaminergic neurons in models of Parkinson's disease [85], limits vascular injury during experimental diabetes [35,77] and maintains microglial cell survival during A β exposure [21,34,86]. By contrast, inhibition or loss of Wnt1 signaling can lead to apoptosis [76,86–88] as well as the progression of autophagy [89].

In regards to WISP1, early work highlighted the ability of WISP1 to prevent p53-mediated DNA damage and apoptosis in kidney fibroblasts [90]. Recent studies have demonstrated both a proliferative and protective role against apoptotic cell injury for WISP1. WISP1 may promote cardiac remodeling after myocardial infarction [69], stimulate lung tissue repair [72], lead to cardiomyocyte proliferation [91] and assist with vascular smooth muscle growth [92]. WISP1 may be necessary to prevent cell death during bone fractures [73,74], to limit doxorubicin-induced cardiomyocyte death [93] and block oxygen--glucose deprivation injury in primary neuronal cells [70,71].

4. Cytoprotective canonical and non-canonical pathways of PI3K, Akt and β -catenin

Although WISP1 is a target of Wnt1, WISP1 utilizes cytoprotective pathways that are sometimes exclusive of the traditional *wingless* canonical and non-canonical signaling of Wnt1 (Figure 2). For example, WISP1 phosphorylates and activates Akt under conditions of cardiac tissue injury, oxidative stress, neuronal degeneration and vascular smooth muscle proliferation [70,90,92,93]. WISP1 protects against apoptotic injury through PI3K and Akt pathways. WISP1 has been shown to rely on PI3K and Akt to provide cytoprotection in renal fibroblasts [90], neurons [70,71] and cardiomyocytes [69,92,93]. WISP1 ultimately modulates apoptotic pathways of Bad, glycogen synthase kinase-3 β (GSK-3 β), Bim, Bcl-x_L, mitochondrial membrane permeability, cytochrome *c* release and caspase activation to prevent cell injury [70]. PI3K and Akt are critical pathways to foster cellular proliferation and block apoptotic injury. During apoptotic injury, the PI3K/Akt pathway can enhance glial cell survival, maintain the integrity of endothelial cells, prevent neurodegeneration, promote cardioprotection and provide tolerance against oxidative stress [94]. Given the proliferative effects of the PI3K/Akt pathway, inhibition of this pathway may be desirable under some circumstances, such as to control tumor growth and promote apoptosis. Experimental strategies targeted to block activation of the PI3K/Akt pathway can suppress medulloblastoma growth [95], reduce colorectal cancer growth [96], increase radiosensitivity in tumors [97] and be beneficial to patients with gynecological malignancies [98]. In addition, inhibition of the PI3K/Akt pathway also can target tumor growth through the induction of autophagy. In oral squamous carcinoma cell lines, application of the agent erufosine that blocks Akt activity leads to cell death through autophagy [99]. Similar results of Akt inhibition that lead to autophagic cell death have been reported in ovarian cancer with other treatments [100].

In regards to the canonical pathways of Wnt1 [75,76], WISP1 controls the post-translational phosphorylation of β -catenin as well as the cellular trafficking of this protein (Figure 2). WISP1 can block phosphorylation of β -catenin in neurons that may be mediated through the inhibition of GSK-3 β [71]. GSK-3 β inhibition is known to prevent β -catenin phosphorylation [75]. WISP1 also can block GSK-3 β activity in other cell systems such as cardiac cells [70,93]. During the inhibition of GSK-3 β , β -catenin is not phosphorylated, ubiquitinated or degraded and therefore can translocate to the nucleus to initiate 'antiapoptotic' pathways and prevent cellular apoptosis [88,101]. In addition to modulating the post-translational phosphorylation of β -catenin [82], WISP1 also controls the subcellular trafficking of β -catenin. In neurons [71], osteoclasts [102], vascular cells [82] and

cardiomyocytes [93], WISP1 can increase nuclear expression of β β -catenin. In neurons, WISP1 through a PI3K-mediated pathway promotes the translocation of β -catenin from the cytoplasm of neurons to the nucleus that can allow for the transcription and eventual translation of pathways that can limit apoptotic cell death [71]. WISP1 requires β -catenin to limit neuronal cell injury during oxidative stress, since inhibition of β -catenin activity blocks neuroprotection against apoptosis by WISP1 [71]. The ability of WISP1 to control the phosphorylation and cellular trafficking of β -catenin also appears to be necessary for WISP1 to autoregulate its own expression. WISP1 expression is governed by β -catenin activity and WISP1 regulates its own expression through the ability of WISP1 to control β -catenin phosphorylation and nuclear translocation [71]. Although β -catenin also may function to limit cell injury through the blockade of autophagy [103,104], the ability of WISP1 to control β -catenin activity does not appear to alter autophagy progression, at least in primary neurons during oxidative stress [71].

5. Downstream mTOR signaling during apoptosis and autophagy

Given that PI3K and Akt are central pathways in WISP1 signaling and can determine the onset and progression of apoptosis and autophagy, consideration of intimately linked downstream pathways, such as mTOR, may offer new directions for the modulation of apoptosis and autophagy in multiple disorders (Figure 2). mTOR signaling is controlled through two protein complexes [105–107]. mTOR Complex 1 (mTORC1) uses the regulatory-associated protein of mTOR (Raptor) protein to allow mTORC1 to bind to its substrates. mTORC1 consists 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). If 4EBP1 is not phosphorylated, it can block protein translation by binding to eukaryotic translation initiation factor 4 epsilon (eIF4E) through the eukaryotic translation initiation factor 4 gamma (eIF4G). 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 p70S6K by mTORC1 leads to mRNA biogenesis, translation of ribosomal proteins and cellular proliferation. PRAS40 is an inhibitory protein and can block the binding of the mTORC1 substrates p70S6K and 4EBP1 to Raptor. mTORC2 is similar to mTORC1 in that it also is composed of mTOR, mLST8 and Deptor. mTORC2 is also different from mTORC1 since it contains Rictor as a component rather than Raptor. mTORC2 also 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 increases the activity of mTORC2. mTORC2 controls actin cytoskeleton organization, cell growth, endothelial cell survival and migration and cell cycle progression. Interestingly, mTORC2 can control the activity of Akt. mTORC2 phosphorylates Akt to lead to its activation. mTORC2 also controls protein kinase C (PKC), P-Rex1, P-Rex2, Rho GTPases and Rho signaling pathways that control cell-to-cell contact [108,109].

Both apoptosis and autophagy can be modulated by mTOR. Activation of mTOR signaling is usually protective during apoptosis. Oxidative stress has been shown to prevent mTOR kinase activity and results in apoptotic cell death in neuronal cells [110]. Inflammatory cells also can sustain apoptotic injury during oxidative stress if deprived of mTOR activation [86,111]. In addition, apoptotic cell death such as in dopaminergic neurons can be blocked during application of agents that increase mTOR activity [112]. During periods of serum deprivation that prevent mTOR activation, insulin has been shown to be unable to rescue cells from apoptotic injury unless mTOR activity is restored [113]. Other growth factors similar to insulin, such as erythropoietin (EPO), also have been shown to be dependent on

mTOR activation for cytoprotection against apoptosis [47,86,114]. However, under other circumstances, inhibition of mTOR may increase cell survival and block apoptosis. For example, during Alzheimer's disease, post-mitotic neurons that attempt to enter the cell cycle do not replicate, but ultimately succumb to apoptotic cell death [115,116]. In work that examines amyloid oligomer exposure, neurons can be prevented from entering the cell cycle during the inhibition of mTOR [117].

mTOR controls apoptotic cell death through 4EBP1 and p70S6K. Activation of p70S6K by mTOR blocks apoptosis through pathways that can increase 'anti-apoptotic' Bcl-2/Bcl-x_L expression and inactivate the 'pro-apoptotic' protein BAD [118]. If mTOR is not active, 4EBP1 binds to eIF4E to result in the translation of pro-apoptotic proteins and cell death [119]. Prevention of apoptotic cell death by mTOR is dependent on Akt. mTOR requires Akt activation to protect endothelial cells against apoptosis [120] and mediate protection through the inactivation of forkhead transcription factors, such as FoxO3a [35,120]. Akt also functions to modulate apoptosis with mTOR through the inhibition of PRAS40. If PRAS40 activity is not prevented, activation of apoptotic pathways can ensue [121]. Phosphorylation of PRAS40 by Akt can block the activity of this substrate and lead to its dissociation from mTORC1 and binding to cytoplasmic 14-3-3 proteins [122].

Similar to studies with mTOR and apoptosis, activation of mTOR is not consistently cytoprotective during autophagy. It is possible that the degree of mTOR activation may be a significant variable. For example, during the early phases of autophagy, mTOR activity can be inhibited [123]. Reactivation of mTOR appears necessary to continue with autophagy, but increased mTOR activity can ultimately block autophagy [123]. In regards to cytoprotection, mTOR activation can prevent neurodegeneration during oxidative stress-mediated autophagy in dopamine neurons [112] and loss of mTOR activity results in autophagic cell death [100]. mTOR activation also appears vital to block autophagy in the cardiac system, limit forkhead transcription FoxO3a activity and prevent cardiac atrophy and dysfunction [124]. During normal physiology, mTOR activity may be required to regulate autophagy, since loss of mTOR activity leads to cardiomegaly and decreased cardiac contractility [125]. However, some chronic disease processes in the nervous or vascular systems may benefit from inhibition of mTOR to allow the progression of autophagy, as suggested in some models of Alzheimer's disease [53]. Furthermore, the benefits of exercise may require a brief inactivation of mTOR for autophagic pathways to proceed [126].

6. Conclusion

Apoptosis (type I cell death) and autophagy (type II cell death) play a significant role in cell death for multiple disease entities that can include the nervous system, cardiovascular systems, skeletal system and immune system. As a result, development of novel therapeutic strategies that can target these processes are viewed with great excitement. The CCN family member WISP1 and its intimate relationship with canonical and noncanonical *wingless* signaling pathways offer a novel approach for modulating the pathways of apoptosis and autophagy. WISP1 can have increased expression in a variety of cells during injury that can involve cardiomyocytes, neurons, bone cells and lung cells, suggesting that WISP1 pathways are necessary for tissue repair and re-growth. WISP1 oversees critical cell survival and proliferation cell mechanisms that involve PI3K, Akt1 and β -catenin. Ultimately, these pathways can converge on mTOR signaling that can either repress or promote the induction of apoptosis or autophagy during normal physiology, acute illness or chronic degenerative disorders. Given the intimate relationship WISP1 holds with apoptotic and autophagic pathways that can affect multiple systems throughout the body, WISP1 and its signaling

pathways offer promising and novel avenues for the future development of therapeutic strategies, especially for disorders that are currently without effective treatments.

7. Expert opinion

Central to the onset and progression of multiple disease entities are the pathways leading to cell demise through apoptosis or autophagy. Under conditions of toxic insults such as exposure to oxidative stress, activation of apoptosis and autophagy may be instrumental in leading to neurodegeneration, immune system dysfunction or cardiac disease. Yet, apoptotic and autophagic pathways may sometimes be necessary for normal physiological function and balance. Furthermore, prevention of cell death in some disease processes may require the initiation and progression of apoptosis or autophagy. For example, proper early neural development of the nervous system in a variety of organisms requires active apoptotic pathways that require Wnt signaling as well as other pathways [127,128]. Activation of autophagy also may be required during oxidant stress injury to the brain to prevent neuronal cell loss [22].

On the converse side, it is clear that in other disorders, inhibition of apoptosis and autophagy is necessary to prevent disease progression. The presence of apoptotic proteins in the brain may be a significant contributor to the progression of Alzheimer's disease as well as alter immune system response [8,44]. During acute cerebral ischemia, the onset and progression of autophagy may ultimately increase brain infarction size [58]. Furthermore, cell death becomes more elaborate with evidence that apoptosis and autophagy may synergistically lead to cell death under some circumstances [61], but under other scenarios, apoptosis may oppose the progression of autophagy to result in cell death. For example, apoptosis can block autophagy by increasing caspase-mediated cleavage of Beclin 1 [64].

Given the complexities of cell death that are controlled by apoptosis and autophagy and the variable relationship between these processes of PCD, new therapeutic strategies must address a number of critical concerns. For example, why does it appear that processes such as autophagy appear to be protective during chronic disorders, such as neurodegeneration [52,53], but detrimental during acute injury such as cellular ischemia [56–58]? Are there cellular pathways that differ between acute and chronic disorders that can alter the modulation of apoptosis or autophagy to affect cellular survival? Are there also specific genetic pathways that can be altered by different toxic environments that determine the role of PCD pathways during disease? New work suggests that in some organisms this may be the case. Draper is required for autophagy during cell death in *Drosophila* salivary glands, but Draper does not appear to have a role with autophagy during starvation [66].

Future strategies for new treatments also must address the role of novel new targets, such as WISP1, PI3K, Akt, β -catenin and mTOR, during apoptotic and autophagic cell death. Although each of these pathways in combination or independently can be highly effective in controlling cell death during apoptosis or autophagy, several unanswered questions remain. For example, WISP1 can effectively limit injury in bone cells [73,74], cardiac cells [69], lung tissue [72] and neurons [70,71] during oxidative stress through blockade of apoptotic pathways. However, autophagy appears to have little or no role with WISP1 cytoprotection or cell death [71] even though pathways that are responsible for WISP1 cytoprotection such as PI3K, Akt and β -catenin are closely tied to modulating autophagic cell death [71,99]. Can cytoprotective pathways such as WISP1 alter endogenous cellular PCD pathways to have one PCD pathway become dominant over another? In addition, are specific cell types, such as neurons, more susceptible to one type of PCD over another dependent on age, toxin exposure or other variables? The specific pathways that are activated during cell injury also may influence biological outcome. In some disorders, mTOR activation is considered

beneficial. mTOR signaling may prevent insulin resistance during diabetes mellitus [129] and lessen the toxicity of β -amyloid in the brain [86,130]. However, other parameters such as duration of activity with mTOR may influence outcome. Acute activation of mTOR yields cardioprotection [131], but long-term mTOR activity may lead to vasculopathy [132]. As a result, WISP1, PI3K, Akt, β -catenin and mTOR offer significant excitement and potential to target apoptotic and autophagic pathways in multiple disorders throughout the body. Yet, it is vital that well-focused future investigations provide essential knowledge of the intricate role these cellular targets play during complex PCD processes that involve apoptosis and autophagy for the successful translation and development of these pathways into robust and safe clinical treatment strategies.

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Article Highlights

- Programmed cell death that is associated with apoptosis or autophagy can lead to pathological consequences in multiple disease entities through oxidative stress that can involve the nervous system, immune system, cardiac system, skeletal system, and vascular system
- Apoptosis leads to cell death through an early phase that involves the exposure of membrane phosphatidylserine residues and a late phase that involves the destruction of genomic DNA
- Autophagy recycles cytoplasmic components, removes defective organelles, and maintains important cytoskeletal structures, but can also determine whether a cell will survive a toxic insult
- Apoptosis and autophagy can work in unison or in an opposing fashion to modulate cell survival in multiple systems of the body
- WISP1, a target of the *wingless* Wnt1 pathway, is cytoprotective in a variety of tissues that include cells of the nervous system, cardiovascular system, renal system, and musculoskeletal system
- WISP1 and its integrated pathways of PI 3-K, Akt, and β -catenin as well as the downstream signaling of mTOR govern apoptosis and autophagy during normal physiology as well as during cell injury

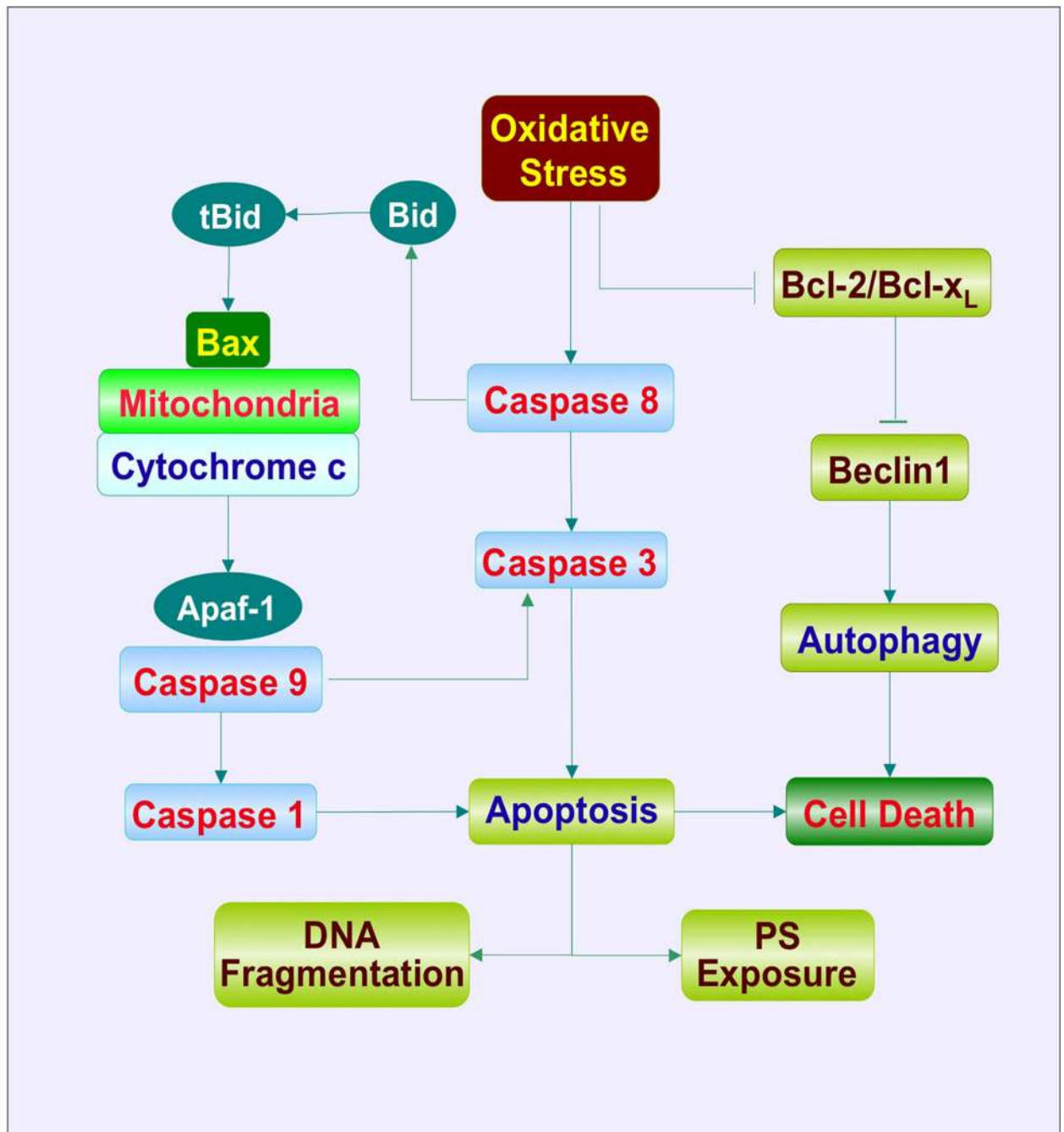


Figure 1. Signaling pathways of apoptosis and autophagy during oxidative stress

Oxidative stress can lead to the activation of initiator caspases, such as caspase 8, which can directly activate caspase 3 and cleave BH3 domain only protein Bid. The resultant truncated Bid (tBid) promotes the release of cytochrome *c* from mitochondria through Bax.

Cytochrome *c* interacts with apoptotic protease-activating factor-1 (Apaf-1) resulting in the oligomerization of Apaf-1 and the subsequent activation of caspase 9. Activated caspase 9 can activate caspase 3 as well as caspase 1 leading to apoptotic DNA fragmentation and phosphatidylserine (PS) exposure. In response to oxidative stress, expression of the anti-apoptotic protein Bcl-2/Bcl-x_L is down-regulated. This process releases the autophagy-related protein Beclin 1 from inhibitory binding by Bcl-2/Bcl-x_L and initiates autophagic

cell death. In addition, Bcl-2/Bcl-x_L also antagonizes Bax-mediated mitochondrial release of cytochrome *c* to prevent the induction of caspase activation and apoptosis.

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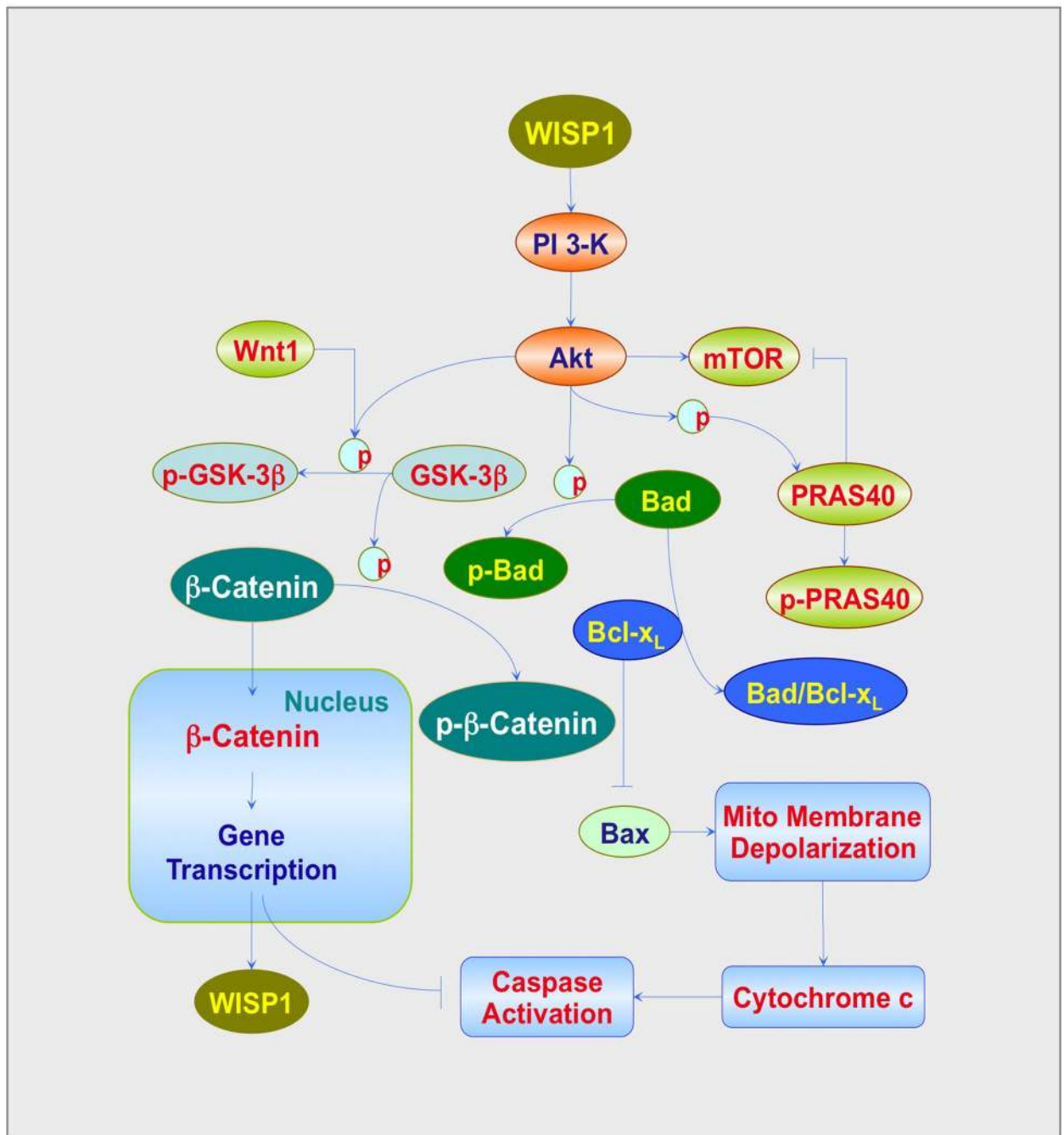


Figure 2. Pathways of WISP1, PI3K, Akt, β-catenin and mTOR that control cell fate
WISP1 results in the activation of phosphoinositide 3-kinase (PI3K) and Akt (protein kinase B). Following activation, Akt can phosphorylate glycogen synthase kinase-3β (GSK-3β). The phosphorylated (p) GSK-3β loses its ability to phosphorylate β-catenin, allowing β-catenin to enter into the nucleus and promoting gene transcription that fosters WISP1 expression and leads to ‘anti-apoptotic’ protein production. Activation of Akt also phosphorylates the pro-apoptotic protein Bad. This process releases Bcl-x_L from the binding to Bad to prevent Bax-mediated mitochondrial (Mito) membrane depolarization, cytochrome *c* release and subsequent caspase activation. In addition, activation of the mammalian target of rapamycin (mTOR) is intimately involved in cell apoptosis and autophagy. Proline-rich

Akt substrate 40 kDa (PRAS40) is one target of Akt phosphorylation. PRAS40 is an inhibitory protein of mTOR and its phosphorylation by Akt results in the loss of its ability to prevent mTOR activation.

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