Review

Role of Autophagy on Heavy Metal-Induced Renal Damage and the Protective Effects of Curcumin in Autophagy and Kidney Preservation

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Abstract: Curcumin is a hydrophobic polyphenol compound extracted from the rhizome of turmeric. The protective effect of curcumin on kidney damage in multiple experimental models has been widely described. Its protective effect is mainly associated with its antioxidant and anti-inflammatory properties, as well as with mitochondrial function maintenance. On the other hand, occupational or environmental exposure to heavy metals is a serious public health problem. For a long time, heavy metals-induced nephrotoxicity was mainly associated with reactive oxygen species overproduction and loss of endogenous antioxidant activity. However, recent studies have shown that in addition to oxidative stress, heavy metals also suppress the autophagy flux, enhancing cell damage. Thus, natural compounds with the ability to modulate and restore autophagy flux represent a promising new therapeutic strategy. Furthermore, it has been reported in other renal damage models that curcumin’s nephroprotective effects are related to its ability to regulate autophagic flow. The data indicate that curcumin modulates autophagy by classic signaling pathways (suppression of protein kinase B (Akt)/mammalian target of rapamycin (mTOR) and/or by stimulation of adenosine monophosphate-activated protein kinase (AMPK) and extracellular signal-dependent kinase (ERK) pathways). Moreover, it allows lysosomal function preservation, which is crucial for the later stage of autophagy. However, future studies of autophagy modulation by curcumin in heavy metals-induced autophagy flux impairment are still needed.

Keywords: heavy metal; autophagy; curcumin; kidney; autophagosome fusion

1. Introduction

Curcumin or diferuloylmethane (1,7-bis(4-hydroxy-3-methoxyphenyl)-16-heptadiene-3,5-dione) is a hydrophobic polyphenol extracted from the rhizome of Curcuma longa L., also known as turmeric [1], which is widely grown in the southern and south western tropical regions of Asia [2]. Curcumin is formed by two aromatic rings (with o-methoxy phenolic groups) linked to an α,β-unsaturated β-diketone moiety [3]. Curcumin can exist in tautomeric forms, keto and enol [4]. It has been widely used in traditional medicine due to its multiple antioxidant [5], antitumor [6], antiviral [7] and anti-inflammatory properties [8], among others [1,3]. Furthermore, studies have shown its protective effect in multiple diseases such as cancer [6], and neurological [9], metabolic [10], lung [11], liver [5], renal [12], and cardiovascular diseases [2]. In addition to its many therapeutic activities, curcumin has the advantage of its biosafety in animals as well as in healthy humans, even at high doses (12 g per day), without undesirable effects [13]. In addition, it has been shown that curcumin provides a nephroprotective effect against various xenobiotics, including heavy metals [12,14–17].
Heavy metals comprise elements like cadmium (Cd), lead (Pb), arsenic (As), mercury (Hg), and chromium (Cr), which are located along the Earth’s crust in various forms [18]. These elements are widely used in different anthropogenic activities, ranging from agriculture to industry [19]. Heavy metals can enter to the body by three routes: Intake (contaminated water or food), inhalation, and dermal contact [20]. Prolonged exposure (occupational or environmental) can cause serious problems in human health, especially in the kidneys, since they represent the major route of heavy metal excretion from the body [18]. Adverse effects of heavy metals have been usually associated with oxidative stress, which involves an overproduction of reactive oxygen species (ROS) and the loss of the endogenous antioxidant defense, as well as with mitochondrial dysfunction [18,21]. However, it was recently found that heavy metals can also impair autophagy flux [22–24], an evolutionarily-conserved self-digestive process, which is generally considered to be a cell survival mechanism [25]. In vitro and in vivo studies related to the participation of autophagy in kidney damage have described that autophagy helps to remove damaged mitochondria, avoiding ROS overproduction and cellular damage [26,27].

In this review we summarized the mechanisms involved in autophagy regulation by curcumin, as well heavy metals-induced autophagy flux impairment in the kidney. On the other hand, curcumin administration has been able to restore renal function in several models of kidney damage [8,14,28,29]. Furthermore, in hyperoxaluria, maleate and contrast-induced nephropathy, curcumin’s nephroprotection has been related to its ability to modulate the autophagy flux [17,30,31]. However, curcumin’s effect on heavy metals-induced autophagy flux impairment has not been explored yet, generating the opportunity for exploration in future studies.

2. Curcumin’s Antioxidant Effects and Mitochondrial Protection in Kidney Damage Models

Curcumin’s direct antioxidant effect derives from the presence of conjugated double bonds in its structure, allowing curcumin to act as an electron donor [32]. Therefore, it scavenges superoxide anion ($O_2^-$), hydroxyl radical (•OH), singlet oxygen ($^1O_2$), hydrogen peroxide ($H_2O_2$), nitric oxide (NO•), peroxynitrite (ONOO•), and peroxyl radical [33]. Moreover, curcumin also activates the antioxidant response element (ARE) by the kelch-like ECH-associating protein (Keap1)/nuclear factor erythroid 2-related factor 2 (Nrf2) system [34]. The protective effect of curcumin has been described in kidney damage models induced by ischemia/reperfusion [8], cisplatin [35], 5/6 nephrectomy [16], and heavy metals (Pb, Cd, Cr) [12,14,15], where its antioxidant activity is highlighted. In addition, it also preserves mitochondrial function [16,29,35], which also contributes to renal function preservation in kidney function. The anti-inflammatory effect of curcumin also plays a prominent role in acute and chronic kidney damage models [36–38]. In this regard, curcumin has an anti-fibrotic effect in glomerulonephritis model, which involves the reduction of transforming growth factor-$\beta_1$ and fibronectin production [36]. This effect is dependent on induction of heme oxygenase-1 (HO-1), a target of Nrf2. Following the same route, Ghosh et al. [37] in the 5/6 nephrectomy model, highlighted the effect of curcumin on the development of chronic renal failure, where inflammation plays an important role through tumor necrosis factor alpha (TNF$\alpha$) and the transcription factor nuclear kappa B (NF-kB).

In this model, curcumin partially suppressed the TNF$\alpha$-mediated NF-kB activity and avoided macrophage infiltration, as well as the functional (such as proteinuria, blood urea nitrogen, and plasma creatinine) and structural (tubular atrophy, hyperplasia, and glomerulosclerosis) alterations. Furthermore, in the renal ischemia/reperfusion model, curcumin attenuated interferon gamma (IFN$\gamma$) expression, while increasing the IL-10 levels [38]. On the other hand, in an in vitro and in vivo model of rhabdomyolysis, curcumin reduced renal damage associated with rhabdomyolysis. In addition to its antioxidant and anti-inflammatory effect, curcumin decreased ferroptosis-mediated cell death [39]. Interestingly, the inhibition of this type of autophagy associated to iron metabolism, is able to attenuate the function and structural alterations in the kidney. Finally, the protective effect of curcumin was also mediated by HO-1 [39].

3. Autophagy and Its Evaluation

Autophagy is a biological process that allows the preservation of cellular homeostasis by the removal of damaged macromolecules and/or organelles in response to a variety of stimuli [40]. Autophagy
consists of five steps: (1) Formation of the phagophore, a complex of Beclin-1, phosphatidylinositol 3-kinase/vacuolar protein sorting 34 (VPS34) and VPS15, (2) phagophore elongation and cargo recruitment, (3) autophagosome maturation, (4) fusion between the autophagosome and lysosome, and (5) autolysosome degradation [41,42]. In addition, formation of autophagosomes and autolysosomal degradation are essential stages to evaluate if the autophagy flux is functional. The usual hallmarks evaluated in these steps are the levels of: Microtubule-associated protein 1 light chain 3 (LC3-I) and its phosphatidyethanolamine form (LC3-II), as well as the ubiquitin binding protein p62, also called sequestosome 1. The LC3-II is an essential protein for elongation and closure of the phagophore, localized in the autophagosome membranes. Meanwhile, p62 is a receptor protein, which binds by its ubiquitin domain to the specific cargo, to anchor them to the LC3-II, present inside the autophagosome, to the subsequent cargo’s degradation in autophagolysosomes [43,44].

In relation to the signaling pathway, key regulators of autophagy include the class I phosphatidylinositol 3-kinase (PI3K) and adenosine monophosphate-activated protein kinase (AMPK) and the autophagy inhibitor the mammalian target of rapamycin (mTOR) [34]. mTOR activation is associated to the PI3K/protein kinase B (AKT)/p70 ribosomal protein S6 kinase (p70S6K) pathway and linked to growth factors. Thus, mTOR negatively regulates autophagy. Meanwhile, the AMPK kinase senses intracellular adenosine triphosphate levels and inhibits formation of the multiprotein kinase complex 1 of mTOR (mTORC1), leading to its dissociation from the Atg2/unc-51-like kinases complex (ULK) and to the dephosphorylation/activation of ULK1 and/or ULK2, which triggers autophagy’s initiation [45]. Likewise, mTORC1 is also regulated by tuberous sclerosis complex 1 (TSC1)/TSC2 and acts as a brake for this pathway, therefore modulating autophagy [42]. In this regard, Akt phosphorylates TSC2 and inactivates the TSC1/TSC2 complex, meanwhile its phosphorylation by AMPK has the opposite effect [42].

4. The Autophagy in Heavy Metals Kidney Damage

The role of autophagy in renal injury is still under debate [46]. It has been described in tubular epithelial cells that autophagy acts as a survival mechanism in multiple renal damage models, including cisplatin, cyclosporine, and ischemia [47]. However, its overactivation can be counterproductive, leading to autophagic cell death [48]. Therefore, due to the dual role of autophagy, factors such as the temporality and intensity of the stimulus have to be considered a priori, to determine the role of this process.

In relation to heavy metal-nephrotoxicity (as summarized in Table 1), the participation of autophagy in cadmium chloride (CdCl2)-induced damage in mice has been demonstrated [49]. It was shown that high doses of CdCl2 increase the LC3-II/LC3-I ratio and promote the formation of autophagosomes, which lead to autophagic cell death, instead of mitigating the renal damage [49]. This is consistent with the study of Shi et al. (2019), which revealed that prolonged exposure to CdCl2 increased apoptosis in chicken kidneys by c-Jun N-terminal kinase (JNK)-dependent autophagy [50]. In the same sense, Liu et al. (2017) found in rat proximal tubule (rPT) cells that exposure to CdCl2 increases LC3-II and Beclin-1 expression, as well as the number of autophagic vacuoles, in a dose-dependent manner [51].

Nevertheless, the increase in LC3-II expression and in autophagosome numbers are not irrefutable proof of autophagy flux. The lack of lysosomal degradation evidence represents a more reliable parameter to confirm the autophagy flux [52], since autophagosome accumulation can derive from the increase in autophagosome formation or the suppression of lysosomal degradation [53]. In this regard it was shown that exposure to a subtoxic dose (0.3 mg Cd/kg) for 5 days did not affect tubular or glomerular function in rat kidneys, although CdCl2 accumulation in the renal cortex was observed [54]. However, proximal convoluted tubule cells (PCT)-exposure to a low dose of CdCl2, showed significant morphological changes associated to autophagy, but not to apoptosis. Furthermore, autophagy upregulation derived from the binding of CdCl2 to the sulphhydryl groups of proteins, oxidative stress, and the endoplasmic reticulum stress-dependent autophagy induction [54]. Similarly, Liu et al. (2017) showed in rPT cells exposed to CdCl2, the anti-apoptotic effect of autophagy with rapamycin treatment, which decreased the apoptosis rate by Fas/Fas ligand (FasL) pathway inhibition, in contrast to 3-methyladenine (3-MA) treatment, an autophagy inhibitor, which increased the apoptosis rate [51].
Table 1. In vitro and in vivo effects of heavy metals on autophagy in kidney.

| Cell Line and Treatment | Effect | Cause/Mechanism | References |
|-------------------------|--------|-----------------|------------|
| rPT cells CdCl₂ (1.25–5 µM) for 12 h | ↑ LC3-II and p62 protein | ↓ Autophagosome-lysosome fusion | Ca²⁺ cytosolic overload impairment autophagic flux by autophagosome-lysosome fusion inhibition [23] |
| | ↑ p62 protein | ↑↑↑ LC3-II accumulation | Ubiquitinated proteins induced ER stress and autophagy activation as mechanisms to remove and detoxify the cell against CdCl₂ toxicity [54] |
| rPT cells Pb(NO₃)₂ (0.5 µM) for 12 h | ↑ LC3-II, p62, Beclin-1 and Atg5 proteins | ↑ Autophagosome accumulation | Autophagy inhibition is directly involved in Pb(NO₃)₂-induced apoptosis [55] |
| rPT cells PbNO₃ (0.5 µM) for 12 h | ↑ LC3-II, p62, p-mTOR, p-p70S6K and p-4E-BP1 protein | ↑ Autophagosome accumulation | Autophagic flux impairment by deregulation of AMPK/mTOR pathway [56] |
| rPT cells PbAc₂ (0.5 µM) for 12 h | ↑ LC3-II and p62 protein | ↑ Autophagosome accumulation | Autophagic flux impairment by lysosomal alkalization and LMP [22] |
Table 1. Cont.

| Cell Line and Treatment | Effect | Cause/Mechanism | References |
|-------------------------|--------|-----------------|------------|
| **In Vitro**             |        |                 |            |
| rTP and NRK-52E cells    | ↑ LC3-II, Beclin-1 and Atg5 proteins<br>↑ Autophagosome accumulation<br>↓ p62 protein<br>↑ p-ERK1/2 stimulated autophagy<br>↑ p-JNK1/2 and p-38 promoted apoptosis<br>- ERK inhibition (U0126):<br>↑ ER stress-induced autophagy<br>↓ Autophagy activation | p-ERK1/2-induced autophagy activation can suppress ER stress-mediated apoptosis | [57] |
| CdAc$_2$ (2.5–5 µM) for 12 h |        |                 |            |
| Male Kunming mice        | ↑ LC3-II/LC3-I ratio<br>↑ Autophagosome formation | ROS-mediated autophagy | [49] |
| CdCl$_2$ (0.4 mg/kg/day, i.p.) for 3 days |        |                 |            |
| Male ICR mice            | ↑ LC3-II and COX-2 protein<br>↓ p62 protein<br>- COX-2 inhibition (celecoxib) prevented:<br>Changes in LC3-II and p62 protein | COX-2 regulates CdCl$_2$-induced autophagy | [48] |
| CdCl$_2$ (0.2 mg to 5 mg/kg/day, i.p.) for 1 week |        |                 |            |
| Female BALB/c mice       | ↑ LC3-II/LC3-I, p62, SOCS3 and p-ERK/ERK protein<br>↑ SOCS3 mRNA, but IL-6 was unchanged<br>↓ p-STAT3/STAT3 protein | Impairment of autophagy flux resulted from ERK activation, by SOCS3 dependent-attenuation of IL-6/STAT3 activation | [24] |
| NaAsO$_2$ (12.5 mg/kg, s.c.) |        |                 |            |
| Female Wistar rats        | Glomerular and tubular functions not affected CdCl$_2$ accumulation on renal cortex<br>Structural changes in PCT cells associated to proliferation and autophagy | CdCl$_2$ accumulation generated ubiquitinated proteins, in order to remove them, autophagy was activated | [54] |
| CdCl$_2$ (0.3 mg/kg, i.p.) for 5 days |        |                 |            |
| Hyline Brown laying chicken treated with CdCl$_2$ (150 mg/kg in diet supplemented) for 90 days | ↑ Autophagic vesicles<br>↑ JNK, Beclin-1, Atg5, LC3-I and LC3-II protein and mRNA levels | CdCl$_2$ promotes JNK-dependent autophagy | [50] |

2-APB: 2-aminoethoxydiphenyl borate; 3-MA: 3-methyladenine; 4E-BP1: 4E-binding protein; Akt: protein kinase B; AMPK: adenosine monophosphate-activated protein kinase; Atg5: autophagy-related gene 5; ATP6V1A: vacuolar-ATPase subunit A; ATP6V1B1: vacuolar-ATPase subunit B1; ATP6V1B2: vacuolar-ATPase subunit B2; Baf: bafilomycin; Bax: Bcl-2-associated X protein; Bcl-2: B-cell lymphoma; COX-2: cyclooxygenase-2; ER: endoplasmic reticulum; ERK: extracellular signal-regulated kinase 1/2; FADD: Fas-associated death domain-containing protein; FasL: Fas ligand; IL-6: interleukin-6; JNK: c-Jun N-terminal kinase 1/2; LC3-I: microtubule-associated protein 1 light chain 3; LC3-II: LC3 conjugated with phosphatidylethanolamine; LMP: lysosomal membrane permeabilization; mTOR: mammalian target of rapamycin; p62: ubiquitin-binding protein p62; p70S6K: p70 ribosomal protein S6 kinase; PARP: poly(ADP-ribose) polymerase; PCT: proximal convoluted tubule; PI3K: phosphatidylinositol 3-kinase; rPT: rat proximal tubule; SOCS3: suppressor of cytokine signaling 3; STAT3: signal transducer and activator of transcription 3.
In support of this, Liu et al. [23] found in rPT cells treated with cadmium acetate (CdAc$_2$) an increase in LC3-II and p62 protein expression, whose expression is inversely correlated with autophagic activity. This suggests autophagy flux inhibition, which was associated with reduction of the autophagosome-lysosome fusion, as a consequence of the cellular fusion machinery depletion and cytosolic calcium increase.

It is important to highlight that cadmium is not the only heavy metal able to impair autophagy flux. In vitro studies in rPT cells treated with lead nitrate (PbNO$_3$) have shown the accumulation of LC3-II and p62 proteins [22,56]. Furthermore, autophagy inhibition increased the levels of cleaved caspase-3 and poly (ADP-ribose) polymerase (PARP), which evidenced its participation in lead-induced apoptosis in rPT cells [55]. The autophagy flux alterations in this condition were associated to autophagolysosome alkalization, as a consequence of the suppression of the two V-ATPase subunits (which hydrolyze ATP to pump protons into the lysosome lumen). Furthermore, lead induced lysosomal membrane permeabilization (LMP), allowing cathepsins-release to the cytoplasm, which induced caspases-mediated apoptosis [22]. The authors also demonstrated the deregulation of the AMPK/mTOR pathway and suggested its participation in the lead-induced impairment of autophagy flux [56].

Autophagy flux disruption has also been observed in arsenic models. Kimura et al. (2016) found in female mice administered with sodium arsenite (NaAsO$_2$), an accumulation of LC3-II and p62 proteins. The highlight of this study is that the participation of estrogens in the autophagic flux impairment was demonstrated, which affects more the females than the males. This phenomenon is associated with ERK overactivation, by SOCS3-dependent IL-6/STAT3 signaling pathway suppression [24].

About Cr (VI), its effect on autophagy in the kidneys is still unknown, but indirect evidence suggest that it may also inhibit the autophagy flux [58]. In rat kidney cortex and HK2 cells exposed to potassium dichromate (K$_2$Cr$_2$O$_7$), there was an increase in p-mTOR and in phosphorylated-p70 ribosomal protein S6 kinase (p-p70S6K) (mTOR target) levels, which suggests the inhibition of autophagy flux [58].

5. Protective Effects of Curcumin Related to Autophagy Regulation

Curcumin can modulate autophagy flux by different molecular mechanisms. In addition, studies have proposed that curcumin acts in different ways depending on its concentration: At low concentrations it functions as an antioxidant, at medium concentrations it also acts as an autophagy inductor, however it has been reported in an in vitro experiment in cancer cell that curcumin at high concentrations leads to cell death [1,4,59,60]. Although the mechanisms by which curcumin induces autophagy were initially described in cancer cells [61], they also occur in normal cells [25]. Nevertheless, their effect differ between both cell types: In normal cells, curcumin mainly induces autophagy [62] in different ways to promote cell survival (see detail in Table 2). Meanwhile, in cancer cells, curcumin induces autophagy over-activation leading to cell death [63,64].

Interestingly, the effects of curcumin administration on autophagic flow in kidney damage models have recently begun to be studied. In kidney diseases, curcumin stimulates autophagy activation at early times, while in neurological and cardiovascular diseases it prevents autophagy activation [9,65]. The main signaling pathway described for autophagy induction by curcumin involves the suppression of the PI3K/Akt/mTOR pathway [66], as well as the stimulation of the AMPK and ERK1/2 pathways [32]. In this respect, Wei et al. (2017) found in a model of advanced glycation end-product (AGE)-induced renal toxicity, that curcumin prevented apoptosis through PI3K/AKT pathway-dependent autophagy activation. In addition, the pretreatment with 3-MA, an autophagy inhibitor, corroborated the protective role of curcumin. Since 3-MA induced autophagy suppression, it stimulated AGE-induced apoptosis [67]. Importantly, curcumin alone did not enhance basal levels of autophagy-related proteins. Likewise, Hu et al. (2018) showed in human kidney cells (HK-2) exposed to H$_2$O$_2$, an autophagy flux reduction related to oxidative stress increase [66].

Furthermore, curcumin administration protected from damage in this model, which is attributable to its ability to restore the autophagy flux by the inhibition/dephosphorylation of the Akt/mTOR pathway. On the other hand, Yao et al. (2018) found in in vitro and in vivo models of diabetic cardiomyopathy,
that curcumin prevented apoptosis by restoring autophagy flux. Such process was mediated by JNK and AMPK/mTOR pathway activation. Furthermore, both kinases favored Bcl-2 phosphorylation, which disrupts its interaction with Beclin-1, an essential autophagy protein that initiates the autophagosome formation [68]. It is known that interaction of Beclin-1 with Bcl-2 does not allow the Beclin-1-binding to vacuolar protein sorting 34 (Vps34), meaning that Beclin-1-dependent autophagy is inhibited [69]. Stimulation of Beclin-1-dependent autophagy by curcumin has also been described in another injury type [70] as well as in various cancer cell lines [34].

In addition, curcumin not only participates in the early stage of autophagy, but also in the latter stage, such as in the fusion of autophagosomes and lysosomes, as well as in the degradation of autolysosomes [25,71]. Yan et al. (2018) found in mouse embryonic fibroblasts (MEF) cells exposed to curcumin, enhanced autophagic flux and lysosomal function (lysosomal acidification and cathepsin B activity) and an increased autophagosome-lysosome fusion. Lysosomal acidification by curcumin was achieved by upregulation of vacuolar V-ATPase gene expression [25]. Curcumin-induced autophagy activation was mediated by suppression of the AKT/mTOR pathway, although until now it is still under debate if the lysosomal function in autophagy depends on mTOR [25]. In support of this, Zhang et al. (2016) observed in Tsc<sup>−/−</sup> MEF cells, in which mTOR is constitutively active, that curcumin was unable to enhance the lysosomal function, suggesting mTOR-dependent lysosomal function. Likewise, it also showed a direct interaction between curcumin and the transcription factor EB (TFEB), which is key for the control of autophagy and lysosomal biogenesis, in MEF and HCT116 cells, which was evidenced by the increase of lysosomal acidification and enhanced cathepsin B activity [71]. In this sense, mTOR-dependent phosphorylation of TFEB (S211) is recognized by YWHA14-3-3 proteins that promote its cytoplasmic retention [72], but curcumin inhibits this phosphorylation, allowing TFEB translocation to the nucleus and enhancing its transcriptional activity [71]. Notably, curcumin is also effective in chronic kidney disease. A study in human kidney tubular epithelial cells (HKCs) found that curcumin exposure avoided the epithelial-to-mesenchymal transition through Akt/mTOR/p70S6K pathway inhibition [28]. In the same way, in the 5/6 nephrectomy-induced chronic kidney damage model, curcumin decreased the phosphorylation of mTOR and its targets (p70S6K and 4E-BP1) [73]. Although in both cases the study of autophagy was not the aim, the possible participation of this may be considered, since the Akt/mTOR pathway modulation is considered a classic mechanism for the regulation of autophagy [74,75]. However, more profound studies are necessary to provide clarification.

By contrast, other studies have also found that curcumin can inhibit autophagy in kidney damage, mainly in models of transition to chronic stage, where autophagy activation had harmful effects [30]. Li et al. (2019) described curcumin as an autophagy inhibitor in a model of hyperoxaluria-induced nephrolithiasis, which reduced the apoptosis rate. Furthermore, in a model using a contrast agent as a nephrotoxicity inducer, it was shown an increase in LC3-II expression in tubular epithelial cells, podocytes, mesangial cells, and macula dense. However, curcumin treatment decreased LC3-II levels and avoided autophagic cell death [17]. In support of this, in a maleate-induced nephrotoxicity model, it was found that curcumin’s protective effect was associated with decreased autophagy [31].

Taken together, these evidences suggest that the modulation of autophagy (stimulation or suppression) may be one of the mechanisms through which curcumin promotes cell survival. However, future studies of autophagy modulation by curcumin in heavy metals-induced autophagy flux impairment are still needed.

Summary of Heavy Metals-Induced Autophagy Flux Impairment in Renal Damage

Heavy metals-induced impairment of autophagy flux occurs in both, the early stage and the later stage (as we summarized in Figure 1). In the autophagy initiation steps, the inhibition may be associated with the stimulation of the Akt pathway (e.g., Cd) and/or with the inhibition of the AMPK pathway (e.g., Pb). Meanwhile, at the later stage, heavy metals impair the autophagosome-lysosome fusion and/or avoid lysosomal acidification. These alterations are closely related to lysosomal function loss.
Table 2. Curcumin effects on autophagy.

| Experimental Model     | Conditions                                                                 | Curcumin Effect                                                                 | References |
|------------------------|-----------------------------------------------------------------------------|---------------------------------------------------------------------------------|------------|
| HKC cells              | Cells were treated with TGF-β1 (5 ng/mL) alone or combination with curcumin (3.125–25 µM) for 72 h | ↓ Vimentin and α-SMA mRNA and protein levels  
↓ p-mTOR, p-Akt, p-p70S6K, and p-4E-BP1 protein  
↓ Epithelial-to-mesenchymal transition | [28] |
| MEF cells              | 20 µM curcumin for 12 h                                                    | ↑ LC3-II and ↓ p62 protein  
↑ Lysosomal acidification, enzyme activity of lysosomal cathepsin B and autophagosome-lysosome fusion  
↑ Atp6v1a, Atp6voc, and Atp6voe mRNA levels  
↓ p-Akt, p-mTOR and p70S6K protein | [25] |
| MEF and HTC116 cells   | 20 µM curcumin for 12 h                                                    | ↑ LC3-II protein  
↓ p62, p-Akt, p-mTOR, p70S6K, and p-TFEB protein  
↑ Lysosomal acidification and enzyme activities of lysosomal cathepsin B  
↑ Tfeb, Lamp1, and Atp6v1a mRNA levels  
↑ TFEB translocation and transcriptional activity  
In Tsc2−/− MEFs:  
Curcumin was unable to increase the lysosomal function | [71] |
| Wistar-albino rats     | Curcumin for 10 days (200 mg/kg/day, oral). On the 5th day contrast agent was administered | ↓ Kidney damage markers (Scr, BUN)  
↓ Lipid peroxidation  
↑ Antioxidant activity (SOD, CAT)  
↓ LC3-II expression in tubular epithelial cells, podocytes and mesangial cells in the glomeruli and the macula dense  
↓ Caspase-3 activation | [17] |
| NRK-52E cells          | Co-incubation of AGE (700 µg/mL) with curcumin (10 µM) for 48 h             | ↓ Apoptosis-related proteins (Bax, AIF and caspase-3)  
↑ LC3-II and Beclin-1 and p-Akt levels  
3-MA pretreatment:  
- Suppression of curcumin-induced apoptosis inhibition  
LY294002 (PI3K/AKT inhibitor) pretreatment:  
- Avoided the curcumin-induced autophagy activation | [67] |
**Table 2. Cont.**

| Experimental Model | Conditions | Curcumin Effect | References |
|--------------------|------------|-----------------|------------|
| Male C57BL/6 mice and H9c2 cells | STZ-induced diabetic mice were treated with curcumin (200 mg/kg/day in drinking water) for 3 months. Cells were co-incubated at high concentrations of D-glucose (30 mM) plus palmitate (0.1 mM) with curcumin for 24 to 36 h. | In both: ↓ Apoptosis-related proteins (cytochrome c and caspase-3 cleaved) ↑ Autophagic-like vesicle formation ↑ p-AMPK, p-JNK, p-Bcl-2, and Bim protein levels ↓ p-mTOR, p-p70S6K, and p-4E-BP1 protein levels In vitro: ↓ Interaction of Beclin1 with Bcl-2 and Bim 3-MA pretreatment: - Suppressed autophagic flux and increase apoptosis rate Inhibition of JNK- and AMPK-activation: - Abolished ability of curcumin to enhance LC3-II formation and to suppress apoptosis | [68] |
| Tubular epithelial (HK-2) cells and ICR male mice | Cells pretreated with H2O2 (200 µM) for 4 h followed by curcumin (20 µg/mL) incubation for 12 h. Curcumin (10 mg/kg, i.v.) was administered to animals in an ischemia (30 min)/reperfusion (24 h) model. | In vitro: ↓ Apoptosis (caspase-3 and -9) ↑ Autophagic-like vesicle formation and LC3-II protein ↓ p62, p-Akt and p-mTOR protein In vivo: ↓ Kidney damage markers (Scr, BUN) ↓ Tubular necrosis, luminal congestion, and pro-inflammatory cytokine (TNF-α and IL-6) | [66] |
| Male Sprague-Dawley rats | Rats that underwent 5/6 nephrectomy were administered with curcumin (75 mg/day, oral) for 6 weeks. | ↓ Kidney damage markers (Scr, BUN and proteinuria) ↓ Glomerular hypertrophy, tubular dilation, and fibrosis ↓ mTOR, p-mTOR, p-p70S6K, and p-4E-BP1 | [73] |

4E-BP1: 4E-binding protein; 3-MA: 3-methyladenine; AGE: advanced glycation end-product; AIF: apoptosis-inducing factor; Akt: protein kinase B; AMPK: adenosine monophosphate-activated protein kinase; α-SMA: alpha smooth muscle actin; Bax: Bcl-2-associated X protein; Bcl-2: B-cell lymphoma; BUN: blood urea nitrogen; CAT: catalase; HKC: human kidney tubular epithelial cells; IL-6: interleukin-6; JNK: c-Jun N-terminal kinase; LC3-I: microtubule-associated protein 1 light chain 3; LC3-II: LC3 conjugated with phosphatidylethanolamine; mTOR: mammalian target of rapamycin; p62: ubiquitin-binding protein p62; p70S6K: p70 ribosomal protein S6 kinase; Scr: serum creatinine; SOD: superoxide dismutase; STZ: streptozotocin; TFEB: transcription factor EB; TGF-β1: transforming growth factor beta; TNF-α: tumor necrosis factor alpha; TSC: tuberous sclerosis complex.
Therefore, its administration in heavy metals-induced renal damage could be a possible treatment strategy to reverse the autophagy impairment, thus contributing to the preservation of renal function.

6. Conclusions

Although several studies support the idea that curcumin can restore autophagy flux alterations in several models, aspects such as cellular type, stimulus intensity, and curcumin concentration must be considered.

Finally, it has been reported in other renal damage models that curcumin’s nephroprotective effects are related to its ability to regulate autophagy flow [17,67]. However, future studies of autophagy modulation by curcumin in heavy metals-induced autophagy flux impairment are still needed.

6. Conclusions

Curcumin administration has demonstrated the ability to restore autophagy flux balance. Therefore, its administration in heavy metals-induced renal damage could be a possible treatment strategy to reverse the autophagy impairment, thus contributing to the preservation of renal function.

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