

## Revisiting Hepatoprotective Natural Products from a Biological Point of View

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**Abstract** – Naturally occurring small molecules from plants, microorganisms, and animals allow the design of drugs that can be beneficial in virtually all kinds of human diseases. Liver diseases with diverse etiologies such as viral infection, chemical intoxication, and metabolic fat accumulation are one of the leading causes of human mortality. Unfortunately, however, there are few effective drugs available capable of stopping or reversing the progress of liver disease. Here, we discuss the current advances in developing hepatoprotective natural products for several arrays of liver disease pathogenesis.

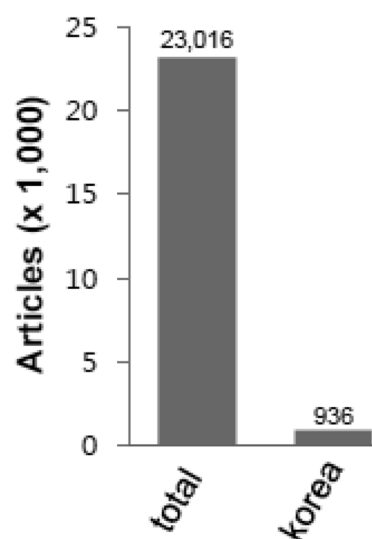
**Keywords** – The liver, Hepatoprotective, Natural Products, Apoptosis

### Introduction

Secondary metabolites from natural resources with a variety of structural diversity are known (Firn and Jones, 2003; Newman *et al.*, 2003). The occurrence of secondary metabolites in plants is considered to be a defense mechanism against environmental herbivores and microorganisms as well. It is now widely accepted that the metabolites are the result of a long-term evolutionary strategy of plants to inhibit or kill their enemies including competing plants.

From the dawn of human history, we have utilized the plants as food resources and medicinal purposes (Newman *et al.*, 2003). Treatment of liver-related diseases with natural products has been one branch of Oriental medicine. Now the isolation and functional characterization of hepatoprotective components has become a fruitful area of therapy (Chien *et al.*, 2011). Based on the historical literature and accumulated clinical experience, the search bioactive compounds or their mixtures to treat gastrointestinal diseases is a global pursuit. Regardless of the origins (synthetic or natural), however, few hepatoprotective agents are currently available. Ursodeoxycholic acid is an exception (Angulo, 2002).

The paucity of available compounds prompted us to review the status of research efforts in the context of the link between hepatoprotective natural products and the pathogenesis of hepatic diseases. A better understanding



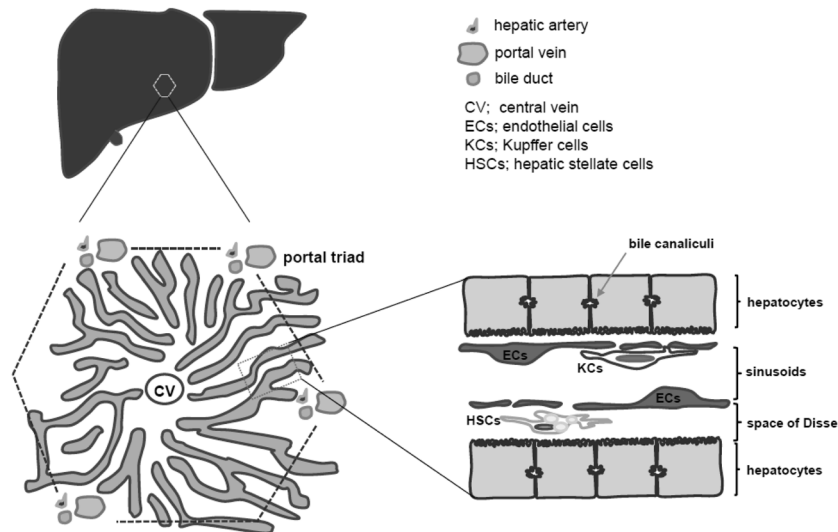
**Fig. 1.** The number of literatures on hepatoprotective phytochemicals contributed by Korean researchers.

concerning how disease pathogenesis is mechanistically inter-connected would provide better directions to mine out hepatoprotective materials. Involvement of natural killer cells in apoptotic event of hepatocyte damage provides an example (Jeong and Gao, 2008) and widens the window of therapeutic targets of immune-modulation.

As an approach, we conducted a literature search using the key words “hepatoprotective”, “liver protection and natural products”, and “phytochemicals”. The results were compiled according to the origin of the articles (Korean and non-Korean). As is clearly evident in Fig. 1, the contribution of Korean scholars is paucity. The need for

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**Fig. 2.** Hepatic blood flow and resident cells in the liver.

Korean-based mechanistic approaches toward the development of successful hepatoprotective materials from natural resources is indicated.

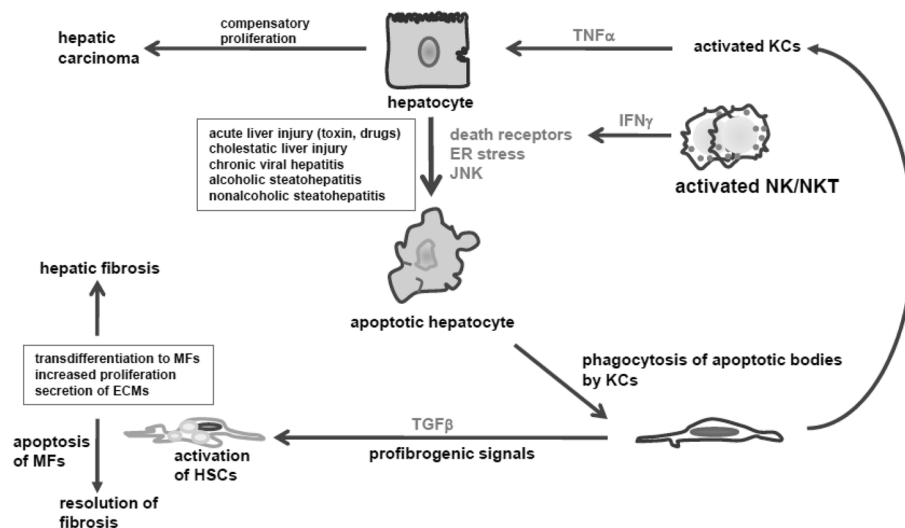
**Liver architecture and function** – The liver is the largest internal organ in the human body, having a wide range of functions that include detoxification, production of biochemicals for digestion, and metabolism (Zakim and Boyer, 2003). It is also a complex organ in terms of its pleiotropic functions and blood vessel structure. The circulatory network in the liver is very complicated. Blood gains access to the liver by two routes. Two-thirds of the blood enters the organ via the portal vein, which is partially deoxygenated since it is collected from components of the gastrointestinal tracts, such as stomach, intestine, gallbladder, spleen, and pancreas. The hepatic artery, in contrast, accounts for the remaining one-third of the oxygenated blood issuing from the heart. In this way, the liver receives 25% of the blood of cardiac output even though the liver mass accounts for only 2.5% of body weight (Fig. 2) (Rappaport, 1980). Both afferent systems merge at the sinusoidal bed, exiting the liver through the central vein. Therefore, the liver not only functions as a reservoir of blood volume but serves as a center for clearance and metabolism of endogenous or exogenous materials from various resources including food intake and toxins (Rappaport, 1980).

Parenchymal hepatocytes constitute about 70% of the total liver cell mass and 90% of the liver weight (Zakim and Boyer, 2003). As shown in Fig. 2, hepatocytes are only one to two cells thick and are separated from each other by capillary spaces called sinusoids. High permea-

bility of the sinusoids allows each hepatocyte to be in close contact with blood. Hepatic stellate cells (HSCs) reside in the space of Disse, which is located between the hepatocyte plate and the capillary endothelium (Fig. 2). The major function of HSCs of the normal liver is the storage of vitamin A. Kupffer cells (KCs) are liver macrophages that reside in the sinusoid. Bile produced by hepatocytes is secreted into thin channels called bile canaliculi. The canaliculi are drained by bile ducts (Fig. 2), which in turn merged into hepatic ducts that carry bile juice away from the liver to the small intestine. As a result, blood flows in the sinusoids whereas bile travels in the opposite direction.

### Pathophysiology of liver diseases

Hepatocytes are the most abundant cell type in the liver, and their apoptosis is prominent in virtually all forms of liver injury (Malhi and Gores, 2008). Liver injury encountered in clinical fields can be divided into acute and chronic types based upon the duration of the injury. In most cases, acute liver injury is reversible upon removal of the injurious insults, leading to a functional and structural restitution of the normal liver (Gressner *et al.*, 2007; Pan *et al.*, 2007). Hepatic fibrosis is a hallmark of chronic liver disease (Fig. 3), which eventually progresses to cirrhosis and/or hepatocellular carcinoma if it is not controlled properly. Chronic liver injuries reflect continuous cycles of acute liver injury and wound healing extended over time. Perpetuated cycles of the injury and healing result in fibrotic phenotypes of the liver, once



**Fig. 3.** Hepatocyte damage initiates the differential stage of liver diseases.

considered as one-way processes. With the paradigm shift that hepatic fibrosis may be reversible, inhibition and careful control of liver injury has become a therapeutic strategy aimed at preventing end-stage liver diseases (Gressner *et al.*, 2007; Pan *et al.*, 2007; Snowden and Fallowfield, 2011).

Traditionally, the secretion of transaminase enzymes from damaged hepatocytes has been utilized as the readout of liver injury. The levels of blood serum glutamic oxaloacetic transaminase (GOT, also called aspartate aminotransferase) and serum levels of glutamic pyruvic transaminase (GPT, also called alanine aminotransferase) are elevated with liver damage (Zakim and Boyer, 2003). Recently, a novel biomarker of hepatic damage has emerged, which is well-correlated with disease severity and response to therapeutic potential of candidate drugs. The M30 neoantigen is an enzymatic cleavage product of cytokeratin 18 that is generated upon apoptotic insults. It has been detected in virtually all liver diseases including cholestasis or cholangitis, and graft-versus host disease during liver transplantation and steatohepatitis (Malhi and Gores, 2008). Since the biomarker reflects hepatic apoptosis, it should prove to be a useful diagnostic tool for monitoring disease progress and therapy.

Various molecular players and subcellular organelles are unequivocally involved in apoptotic program in liver diseases (Fig. 2). A distinct feature of hepatocyte damage depending on the etiology is discussed elsewhere in detail (Malhi and Gores, 2008). In acetaminophen-mediated acute liver injury, for example, upregulation of death receptors and activation of its downstream signaling

pathway (Janus kinase, JNK) has been reported to be involved in liver cell death (Fig. 2) (Mitchell, 1998; Tujjos and Fontana, 2011). Mice deficient in death-receptor are resistant to the development of hepatic fibrosis upon bile-duct ligation, which also implies the role of apoptosis in disease progression (Gujral *et al.*, 2004). Consistent with the *in vivo* observation, caspase inhibition leads to attenuation of hepatocyte apoptosis and liver inflammation (Jeong and Gao, 2008). In a similar setting, however, triggering apoptosis in HSCs has been shown to decrease fibrosis (Siegmond *et al.*, 2005).

In response to hepatocyte death, resident KCs phagocytose the corpse and remove the dead cells while become activated, as shown in Fig. 2. Active KCs synthesize and secrete proinflammatory cytokines including tumor necrosis factor (TNF)- $\alpha$ , resulting in sustained hepatic damage (Siegmond *et al.*, 2005; Snowden and Fallowfield, 2011). Alternatively, growth promoting profibrogenic mitogens secreted by activated KCs stimulate HSCs that are otherwise normally quiescent. Activated HSCs grow rapidly and vigorously synthesize extracellular matrix proteins, which in turn occupy the space where the damaged hepatocytes are eliminated (Fig. 2). Even though different types of cells are able to differentiate to fibroblastic cells and can participate in hepatic fibrogenesis, these cells, including HSCs, seem to undergo apoptosis during the resolution stage of liver injury (Malhi and Gores, 2008). In conditions of ongoing liver damage, selective apoptosis in HSC has been proven to uncouple hepatocyte apoptosis from hepatic fibrogenesis (Gressner *et al.*, 2007). Conclusively, the above findings provide an important cue to handle liver damage; both

inhibition of hepatocyte apoptosis and promotion of HSC apoptosis may produce therapeutic benefits (Siegmund *et al.*, 2005).

### Natural product-derived components with hepatoprotective activity

From the brief overview of the pathogenesis in liver diseases, we may focus on various cell types and involved cellular processes such as apoptosis, immunomodulation, and differentiation programs to address the use of natural resources that contain bioactive constituents. Medicinal plants have been important resources for pharmaceutical drugs. As the demand for bioactive materials from natural resources increases, much effort has been invested to mine phytochemicals targeting human diseases including the liver. The versatility of natural remedies for the treatment of liver diseases has a long history (Firm and

Jones, 2003). Herbal medicines and their ingredients are still used all around the world. Known liver protective plants constituents are represented in Table 1; the constituents are structurally diverse and include flavonoids, coumarins, terpenoids, and xanthenes. Silymarin, a flavonolignan mixture extracted from *Silybum marianum*, has been a popular remedy for hepatic diseases. Baicalin (*Scutellaria baicalensis*), schizandrin B (*Schisandra chinensis*), tanshinones (*Salvia miltiorrhiza*), astragaloside (*Astragalus membranaceus*), glycyrrhizin (*Glycyrrhiza glabra*), and hypophyllanthin (*Phyllanthus niruri*) are potential candidates with hepatoprotective activity. Our lab has screened hepatoprotective and/or anti-fibrotic constituents from plant resources for the last two decades. The current review discusses latest accomplishment of isolated constituents with hepatoprotective activity, briefly summarized in Table 1.

**Table 1.** Classified hepatoprotective natural products from medicinal plants

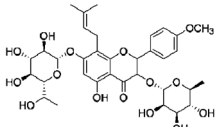
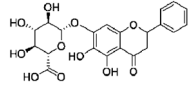
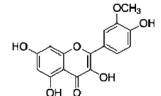
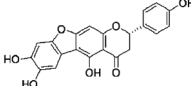
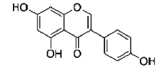
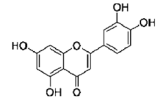
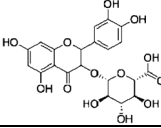
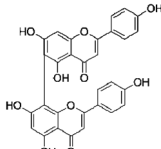
Class	Chemical Substances	Plant Source	Structure	References
	Icarrin and its glycoside	<i>Epimedium koreanum</i>		Lee (1995) Cho (1995)
	Baicalin	<i>Scutellaria baicalensis</i>		Wan (2008)
	Isorhamnetin	<i>Oenanthe javanica</i>		Lee (2008)
	Anastatin A, B	<i>Anantatica hierochuntica</i>		Yoshikawa (2003)
Flavonoid	Genistein	<i>Erycibe expansa</i>		Matsuda (2004)
	Luteolin	<i>Equisetum arvense</i>		Oh (2004)
	Quercetin-3O-β-D-glucuronopyranoside (methyl ester)	<i>Saururus chinensis</i>		Sung (1997)
	Agathisflavone	<i>Canarium manii</i>		Anand (1992)

Table 1. continued

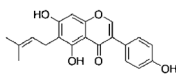
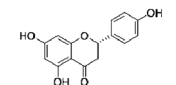
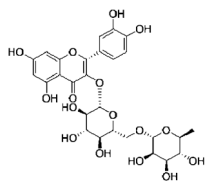
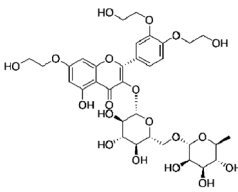
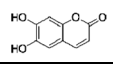
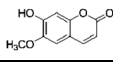
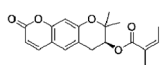
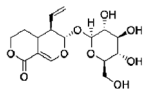
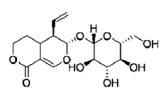
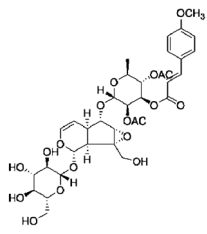
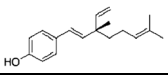
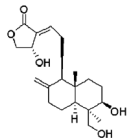
Class	Chemical Substances	Plant Source	Structure	References
Flavonoid	Wighteone	<i>Cudrania cochinchinensis</i>		Lin (1996)
	Naringenin	<i>Cudrania cochinchinensis</i>		Lin (1996)
	Rutin			Janbaz (2002)
	Troxeutin			Zhang (2009)
Coumarin	Kolaviron	<i>Garcinia kola</i>	Fraction containing Garcinia biflavonoids	Farombi (2009)
	Esculetin	<i>Cnidium monnieri</i>		Shin (2011)
	Scopoletin	<i>Solanum lyratum</i>		Kang (1998)
Iridoid	decursinol angelate	<i>Angelica gigas</i>		Yoo (2007)
	Genitopicroside	<i>Swertia japonica</i>		Hase (1997)
	Sweroside	<i>Swertia japonica</i>		Hase (1997)
	Scopolioside A	<i>Scrophularia koelzii</i>		Garg (1994)
	Picroliv	<i>Picrorhiza kurroa</i>		Visen (1991)
Monoterpenoid	(S)-bakuchiol	<i>Psoralea corylifolia</i>		Hyun (2001)
Diterpenoid	Andrographolide	<i>Andrographis paniculata</i>		Chander (1995)

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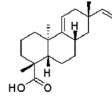
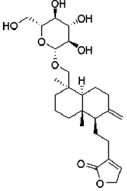
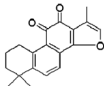
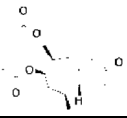
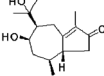
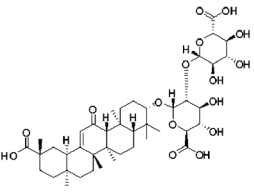
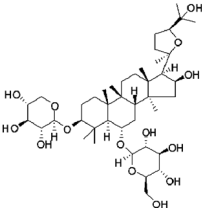
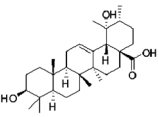
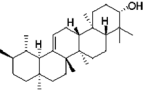
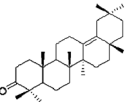
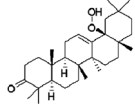
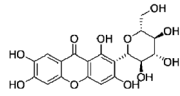
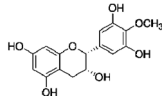
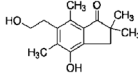
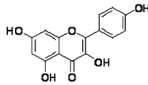
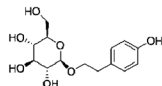
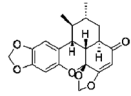
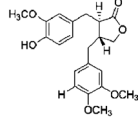
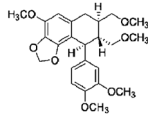
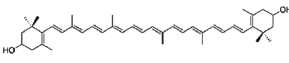
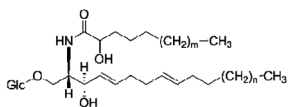
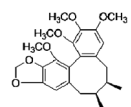
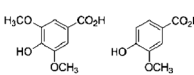
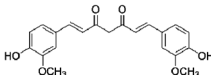
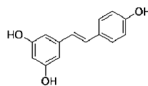

Class	Chemical Substances	Plant Source	Structure	References
	Acanthoic acid	<i>Acanthopanax koreanum</i>		Park (2004)
Diterpenoid	Neoandrographolide	<i>Andrographis paniculata</i>		Chander (1995)
	tanshinone	<i>Salvia miltiorrhiza</i>		Park (2009)
Sesquiterpenoid	Torilin	<i>Cnidium monnieri</i>		Oh (2002)
	Torilolone	<i>Cnidium monnieri</i>		Oh (2002)
	Glycyrrhizin	<i>Glycyrrhiza glabra</i>		Numazaki (1994)
	Astragaloside IV	<i>Astragalus membranaceus</i>		Liu (2009)
Triterpenoid	pomolic acid	<i>Astragalus membranaceus</i>		Liu (2009)
	Amyrin ( $\alpha$ -, $\beta$ -, $\gamma$ )	<i>Sedum sarmentosum</i> <i>Protium heptaphyllum</i>		Amin (1998) Oliveira (2005)
	$\gamma$ -amyrone	<i>Sedum sarmentosum</i>		Amin (1998)
	18 $\beta$ -hydroperoxy-olean, 12-en-3-one	<i>Sedum sarmentosum</i>		Amin (1998)

Table 1. continued

Class	Chemical Substances	Plant Source	Structure	References
	Mangiferin	<i>Salacia reticulata</i>		Yoshikawa (2002)
	4-O-methylgallocatechin	<i>Salacia reticulata</i>		Yoshikawa (2002)
Phenolic	Onitin	<i>Equisetum arvense</i>		Oh (2004)
	Kaempferol	<i>Rhodiola sachalinensis</i>		Song (2003)
	Salidroside	<i>Rhodiola sachalinensis</i>		Song (2003)
	Saquinone	<i>Saururus chinensis</i>		Sung (2000a) Sung (2000b)
Lignan	Arctigenin Traxillagenin 4'-Dimethyltraxillagenin	<i>Torreya nucifera</i>		Kim (2003)
	Hypophyllanthin	<i>Phyllanthus niruri</i>		Negi (2008)
	Cerebrosides	<i>Lycium chinensis</i>		Kim (1997b) Kim (1999) Kim (2000)
	Zeaxanthin Zeaxanthin dipalmitate	<i>Lycium chinensis</i>		Kim (1997c) Kim (1997a) Kim (2002)
	Schisandrin B	<i>Schisandra chinensis</i>		Tang (2003)
Msc	Syringic acid Vanillic acid	<i>Lentimula edodes</i>		Itoh (2010)
	Curcumin			Zhao (1996)
	Resveratrol	<i>Acer mono</i>		Yang (2005)
	Betaine	<i>Lycium chinensis</i>		Kim (1998)

### Conclusion: toward molecular targeting

Secondary metabolites originating from natural resources are synthesized by organisms under pressure to protect against environmental competitors or to ensure their reproductive success. Thus, they represent a population of molecules that can interact with multiple protein targets in nature (Evans *et al.*, 1988). Natural products, therefore, are developed and selected by molecular evolution to specifically bind these protein targets.

A representative example is rapamycin, which is a bacterial metabolite from soil Actinomycetes that show broad cellular effects via the binding to its protein target, a small (12 kDa) FK506-binding protein (FKBP12) and, in turn, the rapamycin-FKBP12 complex inhibits mammalian target of rapamycin (mTOR). In this manner, rapamycin has been utilized as a chemical tool to study the mTOR-related signaling pathways and serves to lead the design of potentially-active anti-cancer and anti-aging drugs.

Advances in molecular biology, cellular biology, proteomics and genomics dramatically increase in the number of molecular protein targets. High-throughput screening systems utilizing chemical libraries may be the first option to test and choose active materials to the molecular targets. Natural products also provide benefits in such efforts since they have unique chemical diversity found in their structures. In addition, new technologies are available to overcome the traditional bottlenecks in the drug discovery process from natural origins. However, the natural product library is composed of crude or semi-pure compound mixtures. This heterogeneity surely adds another layer of complexity to screening system with natural products.

As mentioned above, natural products represent a valuable source with diverse chemical structures for drug development. Current technical advances that we witness in assay methods, chemical separation and structure elucidations lower the inherent difficulties in handling mixtures of these complex molecules.

In the context of the candidate liver molecules to target, the ample attempts that modify a cellular process, autophagy, by small molecules provides a cogent example. Autophagy is a newly-introduced cellular process in a variety of clinical and basic research fields. Some gastroenterologists have already noted that autophagy appears in the liver under basic and pathophysiological processes. Autophagy is a cellular process in which cells degrade their own organelles and unfolded protein aggregates through lysosomal engagement. Accumulating

evidence indicates that the autophagic process is involved in a wide range of cell physiological activities ranging from homeostatic stress response to inflammation to aging. Moreover, it actively participates in disease progress including neurodegenerative diseases, tumorigenesis, and inflammatory bowel disease. Although its importance in biological aspects is recognized, the link to the etiology of liver disease is relatively unexplored. Hidvegi *et al.* (2010) recently investigated how the well-known compound, cabamazepine, stimulates the autophagic process to successfully clear the toxic effects of misfolded  $\alpha$ 1-antitrypsin protein in cells and in a mouse model of associated hepatic fibrosis. Similarly, silibin has been explored with respect to its capacity to activate and induce autophagic cell death, by which it can kill fibrosarcoma cells. These observations imply that autophagy may have a role in hepatic pathophysiology. Research focusing on defining the underlying mechanisms of disease process often lead to new treatment strategies. Rather than correcting the genetic mutations, targeting the autophagy responsible for the elimination of unfolded protein aggregates or damaged organelles would provide an excellent example of the new concept of therapy and drug development.

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