

Closing in on the link between apoptosis and autophagy

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Abstract

While there is a clear connection between apoptosis and autophagy, the mechanisms that regulate the interaction have been difficult to identify. The initial clue to the link was the observation that Bcl-2 was located at the endoplasmic reticulum (ER), where it could prevent some forms of apoptosis and also bind to the autophagy regulatory protein Beclin-I. However, both of these enigmatic observations have been united with the discovery of the nutrient-deprivation autophagy factor-I (NAF-I) protein. As an ER-localized protein that enhances the interaction of Bcl-2 and Beclin-I and that also binds to the pro-apoptotic protein Bik, NAF-I is perfectly placed to be a central regulator of the switch between autophagy and apoptosis.

Introduction and context

Over the past several years, extensive work has been done to elucidate the process whereby members of the Bcl-2 protein family at the mitochondrial outer membrane regulate the decision phase that commits cells to apoptosis. The current consensus is that the pro-apoptotic Bcl-2 family members Bax (Bcl-2-associated X protein) and Bak (Bcl-2 homologous antagonist/killer) exist as inactive monomers. After exposure of the cell to an apoptotic stimulus, these proteins change their conformation in the mitochondrial outer membrane, form oligomers, and permeabilize the membrane, allowing the contents of the inter-membrane space to escape. The released proteins initiate the downstream caspase cascade that executes apoptosis. The two other subclasses of the Bcl-2 family (the Bcl-2 homology region 3 (BH3)-only proteins and the antiapoptotic members) have distinct roles in this process: BH3-only proteins activate Bax, Bak, or both, whereas the anti-apoptotic members such as Bcl-2 (B-cell lymphoma 2), Bcl-XL (B-cell lymphoma-extra large), and Mcl-1 (induced myeloid leukemia cell differentiation protein) inhibit the BH3-only proteins as well as Bax and Bak.

However, all three subclasses of the Bcl-2 family are also found at the ER. The critical event that initiates apoptosis

at the ER is more contentious than the action of the mitochondrial outer membrane. This is complicated by observations suggesting that Bcl-2 family proteins regulate both caspase-dependent apoptosis and autophagy (an independent cell death/survival pathway initiated at the ER). The laboratory run by Gordon Shore at McGill University, Montreal, has been a leader in investigating the consequences of the Bcl-2 family localization to the ER. They have previously shown that an ER-specific BH3-only family member, Bik, can bind to Bcl-2 located at the ER and cause apoptosis [1]. They also identified a Bcl-2associated protein that links Bcl-2 to autophagy. Significantly, a homozygous mutation in this protein that disrupts interaction with Bcl-2 was recently identified in three pedigrees with Wolfram syndrome, a severe neurodegenerative disorder [2].

Major recent advances

By using a form of Bcl-2 specifically targeted to the ER (Bcl2-cb5) [3], the nutrient-deprivation autophagy factor-1 (NAF-1) protein was demonstrated to be required for Bcl-2 to antagonize the autophagy regulatory protein Beclin-1 [4]. Crosslinking and subsequent mass spectrometry were used to identify proteins that both bound to Bcl2-cb5 and were displaced by Bik. The most

abundant protein found in this search was a previously recognized small (15 kD) type 1 transmembrane protein containing a cytosolic iron-sulfur binding cluster designated NAF-1 [4]. When analyzed by immunofluorescence, NAF-1 seems to be localized to the ER. The data obtained using a variety of approaches including binding to glutathione S-transferase fusion proteins and immunoprecipitation of transfected cells, as well as the original crosslinking experiments, make it highly likely that the interaction between NAF-1 and Bcl-2 at the ER is direct, although this has yet to be formally proven. Importantly, decreasing NAF-1 expression by silencing RNA did not affect apoptosis initiated by Bik binding to Bcl-2. NAF-1 knockdown does enhance both starvation-induced autophagy as well as Bik-dependent autophagy, mediated by the displacement of Beclin-1 binding to Bcl-2. Beclin-1 is an authentic BH3-only binding protein [5] that, when released from Bcl-2, participates in a multi-component complex that initiates the formation of autophagosome formation at the ER [6,7].

In co-immunoprecipitation assays, NAF-1 was also shown to be present in a complex containing Bcl-2 and the inositol triphosphate receptor (IP3R). This interaction may be functionally relevant because the effect of Bcl-2 binding to IP3R on intraluminal calcium stores, a mechanism elegantly demonstrated by Clark Distelhorst's group [8], is also inhibited by knocking down NAF-1 expression.

Future directions

These important findings shed light on the regulation of both apoptosis and autophagy at the ER by Bcl-2. Functionally, the interaction between Beclin-1 and Bcl-2 is enhanced by NAF-1, indeed, the data from Chang et al. [4] strongly suggest that NAF-1 binds directly to Bcl-2. It is unclear whether or not NAF-1 also binds directly to Beclin-1 and therefore acts as an adaptor protein for the Bcl-2-Beclin-1 interaction, or alternatively whether NAF-1 binding alters the conformation of the Bcl-2 BH3 binding region such that it binds to Beclin-1 more tightly. There is already support for the theory that binding of BH3-only proteins to Bcl-2 at the ER alters Bcl-2 conformation and function [9]. If proven to be true, this would be the first instance where a non-BH3-only protein would be shown to have a similar effect to Bid (BH3 interacting domain death agonist) or Bim (Bcl-2-like protein 11). This would mean that NAF-1 would function as a specific allosteric regulator, as it does not seem to alter the binding of Bcl-2 to whichever BH3-only protein is displaced by Bik to initiate apoptosis.

Although not yet proposed in the literature, the presence of NAF-1 in Bcl-2–IP3R complexes has intriguing implications for both the composition of the Bcl-2 complex at the

ER and the potential involvement of NAF-1 in regulating autophagy via IP3R. The functional importance of this interaction is highlighted by the recent observation that baseline activity of IP3R is required to mediate low-level calcium release from the ER, where it is taken up by the mitochondrial calcium uniporter [10]. In this organelle, it is required to turn on pyruvate dehydrogenase activity, mediating a switch from glycolysis to the Krebs acid cycle thereby enhancing oxidative phosphorylation. When this process is interrupted at any of the steps between calcium release and ATP generation, energy deprivation is sensed and macro-autophagy is initiated in an mTOR (mammalian target of rapamycin)-independent fashion. Thus, by facilitating the binding of Bcl-2 to both Beclin and IP3R, NAF-1 may mediate the control of two separate arms of autophagic cell death or survival.

Therefore, NAF-1 may act as a switch between autophagy and apoptosis at the ER. It is intriguing to note that the other mutated gene associated with Wolfram's syndrome also encodes a transmembrane ER protein that regulates calcium homeostasis, suggesting that this is an important pathway for neuronal survival [11]. Investigating how NAF-1 interaction with Bcl-2 modulates binding of partners through different binding domains will be essential to understanding how this switch is regulated. It will also be important to determine whether all these proteins are present in the same complex or whether some interactions are mutually exclusive. Finally, it will be interesting to find out whether regulation is achieved by different complexes (if they exist) competing with each other for limiting factors.

Abbreviations

Bak, Bcl-2 homologous antagonist/killer; Bax, Bcl-2-associated X protein; Bcl-2, B-cell lymphoma 2; BH3, Bcl-2 homology region 3; ER, endoplasmic reticulum; IP3R, inositol triphosphate receptor; NAF-1, nutrient-deprivation autophagy factor-1.

Competing interests

The authors declare that they have no competing interests.

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