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Endoplasmic Reticulum Stress Signaling in Mammalian Oocytes and Embryos: Life in the Balance

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Abstract

Mammalian oocytes and embryos are exquisitely sensitive to a wide range of insults related to physical stress, chemical exposure, and exposures to adverse maternal nutrition or health status. Although cells manifest specific responses to various stressors, many of these stressors intersect at the endoplasmic reticulum, where disruptions in protein folding and production of reactive oxygen species initiate downstream signaling events. These signals modulate mRNA translation and gene transcription, leading to recovery, activation of autophagy, or with severe and prolonged stress, apoptosis. ER stress signaling has recently come to the fore as a major contributor to embryo demise. Accordingly, agents that modulate or inhibit ER stress signaling have yielded beneficial effects on embryo survival and long-term developmental potential. We review here the mechanisms of ER stress signaling, their connections to mammalian oocytes and embryos, and the promising indications that interventions in this pathway may provide new opportunities for improving mammalian reproduction and health.

1. INTRODUCTION

The maturing oocyte and early mammalian embryos are notable for their unique cellular physiologies and unique mechanisms of developmental regulation. Oocytes and early embryos lack many of the mechanisms that exist in somatic cells to perform basic metabolic and homeostatic functions, such as free radical scavengers, ion transporters, and osmoregulatory mechanisms. Oocytes and embryos also undergo unique cellular events not seen in somatic cells. For example, fertilization results in massive calcium release and extensive changes to the cell membrane. Meiotic cell cycle progression leads to asymmetric cell division, with attendant mechanisms that must position and orient the meiotic spindle appropriately. The cell cycle of the early cleavage stage embryo is unique in that DNA replication and cytokinesis occur in the absence of substantial cell growth. Oocyte maturation encompasses global transcriptional repression, so that maturing oocytes and early embryos rely predominantly on post-transcriptional mechanisms to sustain and modify protein content of the cell and to execute key developmental transitions.

These unique characteristics of maturing oocytes and early embryos create unique challenges. Indeed, these unique challenges may underlie the relative sensitivity of these cells to exogenous insults. Although the early mammalian embryo is often noted for its apparent plasticity, enabling it to compensate for dramatic perturbations such as cell extirpation, the maturing oocyte and early embryo are quite sensitive to exogenous stresses. It is becoming increasingly apparent that insults to oocytes and early embryos underlie long-term phenotypic alterations observed during both fetal and post-natal life (Latham et al., 2012). The simplest interpretation of these observations is that oocytes and early embryos can undergo physiological adaptations to environmental perturbations, and that these adaptations likely involve epigenetic changes that permanently modify cellular properties by establishing abnormal genome programming.

Such adaptations highlight the fascinating interplay between the environment and developmental biology, particularly the sensitivity of early embryonic genomes undergoing early developmental programming processes. However, such adaptations to environmental stress are only possible when the oocyte or embryo survives the insult.

This chapter focuses on the role of unfolded protein response (UPR) and endoplasmic reticulum stress signaling (ERSS) in the responses of oocytes and embryos to environmental stress, the unique consequences that ERSS may have in oocytes and early embryos, and the potential for novel approaches to manage ERSS in enhancing oocyte and embryo quality and survival. The latter possibility stands at the frontier of modern mammalian embryology, and offers many exciting new possibilities for enhancing clinical and applied outcomes in humans and other mammalian species.

2. OVERVIEW OF UPR AND ERSS

Sensing and responding to exogenous stress is a vital part of cellular physiology. It has become increasingly apparent that one of the key mechanisms of initiating cellular response to a variety of exogenous stressors resides in the endoplasmic reticulum (ER). Secreted proteins and membrane-associated proteins are synthesized in the ER, and must then undergo proper folding, glycosylation, and disulfide bond formation in order to generate functional proteins. A quality control mechanism that detects and eliminates incorrectly processed or unprocessed proteins is thus vital to overall cellular functioning, including cell division, homeostasis, functional responses and cell-cell interactions, and differentiation.

The unfolded protein response fills this need (Bernales et al., 2006). But UPR also fills a much greater role in the cell by providing an indirect means of detecting and responding to stress, because many exogenous stressors negatively impact the ER environment and protein processing (Fig. 1), for example by altering amino acid availability affecting rates of protein synthesis, carbon substrate availability for glycosylation, Ca²⁺ concentration required for proper folding, cellular redox state related to disulfide bond formation and macromolecular oxidation states, ATP availability for biosynthesis, protein denaturation, lipid availability for protein lipidation, and rates of protein trafficking and secretion.

Disruptions in any of these protein-processing steps by any of a wide variety of stressors leads to accumulation of unfolded or incorrectly folded proteins in the ER. Whereas under

normal conditions unfolded and incorrectly folded proteins are removed from the ER by the ER associated protein degradation machinery (ERAD) and targeted for degradation in the cytosol by the ubiquitin-proteasome pathway, any exogenous stress that increases the abundance of such defective proteins in the ER initiates a series of responses that lead either to cellular recovery or cell death (Fig. 1). This sequence is initiated by the transfer of glucose regulated protein 78 (GRP78 a.k.a. HSPA5, BIP) from three primary responder ER membrane proteins eukaryotic translation initiation factor 2A (EIF2A) kinase 3 (PERK), activating transcription factor 6 (ATF6), and endoplasmic reticulum-to-nucleus signaling protein (ERN, a.k.a. IRE1)(Fig. 1, blue boxes) to the accumulating unfolded proteins. Disruption of GRP78 leads to preimplantation arrest, blastocyst failure to hatch, cell division defects and apoptosis in the inner cell mass (Luo et al., 2006), attesting to the importance of ERSS in the early embryo. Once PERK, ATF6 and/or IRE1 are activated, they initiate an early adaptive response to unfolded proteins that, if the stress is short-lived, can facilitate clearance of the unfolded proteins and cell survival—specifically, inhibition of protein translation via EIF2A phosphorylation to reduce the rate of unfolded protein production, transcriptional induction and increased expression of chaperones to facilitate protein folding and negatively feed back to the primary responders, and increased expression of ERAD components to enhance clearance from the ER (Fig. 1, green arrows and boxes). Certain stressors can lead to translational inhibition via alternative EIF2A kinases, thereby augmenting the basic ERSS response. Early ATF4 expression can activate autophagy and enhance cell survival.

With prolonged stress, however, additional responses are initiated, including continued accumulation of ATF4 and ATF6 α (p50), leading to transcriptional induction of CHOP, transcriptional induction of JNK/JUN (v-JUN oncogene homolog and kinase) signaling, and activation of p38MAPK14 (mitogen-activated kinase 14) and p53, all of which promote cell apoptosis (Fig. 1 red arrows and boxes). Apoptosis is driven by both mitochondrial dependent and independent pathways (Logue et al., 2013; Minamino and Kitakaze, 2010). IRE1 mediated activation of XBP1 (X-box binding protein) mRNA splicing to produce functional (short form) XBP1 can either promote autophagy or activate transcriptional responses leading to apoptosis via JNK/JUN. Prolonged or increased induction of autophagy, however, can lead to apoptosis rather than survival. Consequently the balance between ERSS and autophagy is a key determinant of survival, and ERSS can diminish to cell survival in conjunction with autophagy induction. One other possible consequence of ERSS is induction of caspase-12, which distinguishes apoptosis deriving from ERSS from other causes of apoptosis (Nakagawa et al., 2000).

3. INTERACTIONS WITH OTHER SIGNALING PATHWAYS

The above summary of ERSS (Fig. 1) addresses a limited set of pathways and interactions that lie at the core of ERSS, mediating a common set of responses to diverse stressors. But the ERSS response is integrated into overall cellular physiology so that it interacts with many other pathways, and can in turn be modulated by these other pathways. A detailed summary of all such interactions is not possible here, but particular interactions are worth mentioning due to potential relevance to oocytes and embryos.

3.1 Interaction with TOR pathway

One key pathway that interacts with ERSS is the TOR (Target of Rapamycin) signaling pathway. This pathway is relevant particularly in the context of effects of diabetes, hyperglycemia, insulin resistance, and insulin signaling. TOR signaling inhibits autophagy under normal conditions but TOR promotes apoptosis during ER stress, and inhibition of TOR promotes autophagy and cell survival under ER stress (Kapuy et al., 2014). ERSS can negatively affect TOR signaling (Qin et al., 2010). The nature of the agent causing ER stress and the duration of the stress affect outcome (autophagy vs. apoptosis) and the activation state of TOR contributes to this choice (Kapuy et al., 2014). Amino acid deprivation, for example, may activate ERSS whilst biasing a cell toward apoptosis by activating TOR. Activation of TOR signaling can promote phosphorylation of the translation inhibitors EIF4EBP1 and 2 (eukaryotic translation initiation factor 4E binding proteins 1 and 2) (Jansson et al., 2012) and other translation factors (e.g., ribosomal protein S6) to promote translation of certain mRNAs that increase translation of particular mRNA classes. Multiple signals linked to nutrient signaling operate through controlling EIF2A phosphorylation, including PI3/AKT (*v*-AKT oncogene homolog), TOR, GCN2 (general control non-derepressible 2, a.k.a. EIF2AK4), AMPK (AMP protein kinase), and GSK3 β (glycogen synthase kinase 3 beta), and these same signals can directly or indirectly modulate TOR activity, so that diverse signals may modulate the translation response to TOR signaling. AMPK can diminish ERSS downstream events (Salvado et al., 2013; Terai et al., 2005; Tirupathi Pichiah et al., 2011) but can also activate ERSS (Lin et al., 2014b; Yang et al., 2013). PPAR β/δ (peroxisome proliferator activated receptors) inhibit events downstream of ERSS by AMPK activation (Salvado et al., 2014). Transcriptional responses to TOR signaling affecting genes related to amino acid transport, lipid metabolism, nucleotide metabolism, and protein synthesis can also be affected by other regulators such as hypoxia inducible factor HIF1A (Jansson et al., 2012). The overall effect is that the status of TOR activation and signaling in a cell is modulated by many factors. Whether cells arrest growth and either survive (autophagy) or undergo apoptosis under ER stress, and the magnitude and duration of ER stress tolerated before initiating apoptosis vary in this context of TOR signaling and interacting pathways such as AMPK signaling.

3.2 Interaction with SRC pathway

Another interacting pathway is the cell growth, proliferation, migration and differentiation regulator SRC (*v*-SRC oncogene homolog). ER stress induced by thapsigargin or tunicamycin activates SRC kinase (Moon et al., 2014; Yu and Kim, 2010), and this has been implicated in significant cellular responses such as epithelial-to-mesenchymal transition (Ulianich et al., 2008) and other changes (Yu and Kim, 2010). ERSS inhibitors (e.g., salubrinal, guanabenz) reduce SRC activation (Wan et al., 2014). Conversely, overexpression of the SRC homology domain adapter protein NCK1 reduces ERSS mediated EIF2A phosphorylation by inhibiting PERK and recruiting phosphatases (Kebache et al., 2004; Latreille and Larose, 2006).

3.3 Interaction with NRF2 pathway

Another key stress response pathway is linked to ERSS, as PERK activates NRF2 (NFE2 related factor 2) (Cullinan and Diehl, 2004; Cullinan et al., 2003). NRF2 activation leads to increased glutathione levels and buffering of reactive oxygen species (Cullinan and Diehl, 2004) providing a valuable connection between unfolded protein and oxidative stress responses. As oxidized lipids also induce ERSS (Chen et al., 2013; Garbin et al., 2014), activation of NRF2 may also help protect from this source of oxidative stress. Interestingly, reactive oxygen species can be generated during ER stress via oxidative protein folding in the ER (Wang et al., 2014b), so that activation of NRF2-mediated buffering of reactive oxygen species by PERK may minimize damage arising from protein oxidation. Additionally, fatty acid oxidation in the ER inhibits oxidative protein folding, and inhibition of fatty acid oxidation can reduce ERSS (Tyra et al., 2012).

3.4 Interaction with NF κ B pathway

There is also extensive interaction between ERSS and the NF κ B (nuclear factor kappa B) pathway. NF κ B is involved in many processes such as cell adhesion, proliferation and differentiation, cell morphology changes, apoptosis, and inflammation (Pahl and Baeuerle, 1995; Prell et al., 2014). Activation of the NF κ B pathway by ERSS can arise with some selectivity (e.g., accumulated protein, oxidative stress and calcium chelation, but not unfolded protein in ERSS) (Pahl and Baeuerle, 1995, 1996). NF κ B pathway activation is mediated by IRE1 and TRAF2 (Kaneko et al., 2003; Mauro et al., 2006) or by inhibition of production of the I κ B α (NF κ B inhibitor) (Deng et al., 2004). Other signaling pathways and kinases [e.g., GSK3 β and CK2 (casein kinase type II); (Piazza et al., 2013)] connect to the NF κ B pathway and modulate cell survival. NF κ B can protect cells from stress by modulating reactive oxygen species generation (Mauro et al., 2006). Additionally, NF κ B and TOR may cooperate to initiate unfolded protein aggregation in the cytoplasm to protect cells from ER stress (Liu et al., 2012).

3.5 Interaction with p53 pathway

P53 acting downstream of p38/MAPK14 activation, can contribute to the EIF2A and EIF4E phosphorylation (Jiang et al., 2014). P53 can also contribute to cell cycle arrest and apoptosis. ERSS can inhibit p53-mediated apoptosis via GSK3b signaling and de-stabilize p53 via synoviolin (an E3 ubiquitin ligase used in ERAD, ER associated degradation) during mild ER stress, but severe stress leads to p53 activation and stabilization (Yamasaki et al., 2007).

One lesson that emerges from the foregoing considerations is that ERSS is complex and is an integral part of cellular responses to many different stressors, and to many different signaling pathways or combinations of pathways. Thus, ERSS can operate in parallel with multiple signaling mechanisms in different situations, and the overall outcome in terms of autophagy, cell survival, or cell apoptosis depends on the additive effects on downstream effectors.

4. CHEMICAL ACTIVATORS AND INHIBITORS OF ER STRESS

As is clear above, ERSS is complex, and moreover intersects with many key regulatory pathways in the cell. As a result, many compounds have been identified that directly or indirectly regulate ERSS. Some of these are described in Table 1. Activators of ER stress include agents that inhibit protein glycosylation or export from the ER, induce oxidative stress, affect mitochondrial function, diminish ER calcium stores, activate PERK, increase NO production, or disrupt GRP78 binding. Some agents have multiple effects in cells. Importantly, some agents impact interacting pathways, further attesting to the close integration of ERSS with other cellular processes. One interesting class of agents that inhibit ER stress comprises inhibitors of EIF2A dephosphorylation. EIF2A phosphorylation is one aspect of the early ER stress response. Continued protein translation contributes to later events such as ATF4 mRNA translation and protein synthesis following transcriptional induction. Continued protein synthesis may slow the recovery from ER stress. The inhibitors of EIF2A dephosphorylation appear to inhibit ER stress by reducing the rate of protein synthesis to minimize the genesis of unfolded proteins, as well as by inhibiting protein synthesis-dependent aspects of ERSS that lead to apoptosis. Activators/inhibitors of autophagy can have opposing effects, due to the relationship between short- and long-term autophagy induction and ERSS. One other feature of ERSS-modulating agents is that many are derived from natural sources. A variety of natural compounds function as antioxidants or can induce ROS generation. Others can inhibit the proteasome, inhibit TOR signaling, or modulate ER calcium stores.

5. RELATIONSHIP OF ERSS PATHWAYS TO OOCYTE MATURATION AND EARLY EMBRYOGENESIS

The foregoing discussion of ERSS highlights the complexity of the process. Indeed, ERSS occupies a nodal regulatory role in the cell, serving to integrate a myriad of environmental cues and other cellular signaling pathways into a network of events that coordinate numerous events in diverse cellular organelles, including the nucleus, mitochondria, the ER, and the Golgi, and coordinating diverse stress responses with a variety of cell functions including transit through the cell cycle, growth and metabolism.

In the multicellular organism, the ERSS achieves an overall objective of either repairing or eliminating damaged cells. In the oocyte and fertilized zygote, however, cell death eliminates the entire organism; in the cleaving embryo cell death may severely compromise embryo viability. Reduced rates of cell division may likewise lead to more long-term effects.

Additionally, the development of high-quality oocytes and the elaboration of high-quality embryos are linked to correct dialog between the developing oocyte, the supporting cumulus cells and other cells within the follicle (Hao et al., 2014; Matzuk et al., 2002). Thus, stressors that impact follicle cells can also indirectly affect oocyte and embryo viability. Successful mammalian development also relies on successful implantation and placentation and efficient maternal-fetal exchange of materials. Stressors that impinge on formation and function of the placenta can thus exert severe detrimental effects on developmental outcomes. The following sections review what has been discovered with respect to ERSS in

mammalian oocytes, embryos, and follicle cells, and what measures may be taken to enhance reproductive outcomes. It is noted that there is an extensive body of literature on ERSS in trophoblast and placenta, describing the relationships between inflammation, ERSS, maternal health state, and pregnancy outcomes. Though too extensive to cover in this chapter, those studies provide additional insight into the link between ERSS and mammalian reproduction.

6. ACTIVATION AND INHIBITION OF ER STRESS IN OOCYTES AND EMBRYOS

Many *in vivo* and *in vitro* stressors impact oocyte and preimplantation embryo health and developmental potential, and long-term developmental outcomes. These include heat, DNA damage or DNA damaging agents, osmotic stress and availability of organic osmolytes, oxygen and oxidative stress, hyperglycemia and carbon substrate availability, hyperlipidemia and oxidized lipids, calcium ionophores, cytokines, amino acid deprivation, insulin signaling, and serum components. The overall effects of these stresses range from delayed cleavage to developmental arrest at the 2-cell stage, to significant fetal and post-natal effects, such as intrauterine growth restriction, post-natal compensatory growth, abnormal developmental programming of metabolism, hypertension, diabetes, obesity, and potential epigenetic changes that may even be transmitted transgenerationally. Such responses underlie some of what are now recognized as developmental origins of adult disease (DOADs) (Latham *et al.*, 2012). Immediate embryo responses to these stressors can be multi-pronged, invoking multiple parallel signaling pathways as discussed above; these can include p38/MAPK14 activation, AMPK activation, DNA damage response, TOR, and GSK3 β . ERSS occupies a central role in mediating overall cellular responses and cell survival/death, and thus may be linked to the long-term effects on developmental programming and developmental origins of disease. As such, it is worthwhile to summarize activation of ERSS in oocytes and embryos subsequent to a range of different insults.

6.1 Temperature

Maternal heat stress impairs cleavage in mouse embryos *in vivo*, with most embryos arresting at the 2-cell stage (Bellve, 1973), and severe heat stress can inhibit oocyte maturation (Kim *et al.*, 2002). Similar effects are seen in other mammals (Edwards and Hansen, 1996; Isom *et al.*, 2007; Makarevich *et al.*, 2007; Payton *et al.*, 2004). Heat stress is associated with increased reactive oxygen species and apoptosis (Paula-Lopes and Hansen, 2002; Sakatani *et al.*, 2004; Tseng *et al.*, 2006). A variety of exogenous factors such as antioxidants, melatonin, retinol, IGF1 and sphingosine-1-phosphate can protect embryos from heat stress (Cebrian-Serrano *et al.*, 2013; Lawrence *et al.*, 2004; Namekawa *et al.*, 2010; Roth and Hansen, 2004; Sakatani *et al.*, 2007). Heat stress induces changes in gene expression in surviving embryos (Hickman *et al.*, 2013). Cold stress associated with cryopreservation, particularly slow freezing can also affect embryo development and gene regulation (Larman *et al.*, 2011; Saenz-de-Juano *et al.*, 2012). Prolonged exposure to room temperature can inhibit cleavage and alter the Golgi complex in early embryos (Hegele-Hartung *et al.*, 1991), and can degrade meiotic spindle structure in ovulated oocytes (Van

der Elst et al., 1988). In addition to generating ROS, negative effects on intracellular structures could impact ER function and lead to ERSS.

6.2 Osmotic stress

Osmotic stress has profound effects on early embryos, including induction of 2-cell arrest in mouse embryos (Wang et al., 2011), a response that is genotype-dependent and that underlies the classical 2-cell arrest of outbred mouse lines in sub-optimum culture conditions. Early embryos lack some of the osmoregulatory mechanisms that operate in somatic cells, and rely instead on amino acids as internal organic osmolytes (Baltz and Tartia, 2010; Petronini et al., 1992; Steeves and Baltz, 2005). Availability of amino acids in the culture medium can ameliorate stress responses, such as translational repression, and facilitate survival (Petronini et al., 1992; Richards et al., 2010). Thus, amino acid starvation may not only affect protein synthesis and other metabolic processes, but may also compromise osmoregulatory ability in the early embryo. Hyperosmolarity induces activation of p38/MAPK14 and JNK signaling, and induction of HSP70.1 (Baltz and Tartia, 2010; Fiorenza et al., 2004; Xie et al., 2013). Interestingly, organic osmolytes can also reduce negative effects of other stressors (Zander-Fox et al., 2013). Osmotic stress affects ER to Golgi trafficking (Jiang and Storrie, 2005; Lee and Linstedt, 1999) and may affect calcium sequestration, which may activate ERSS via calreticulin involvement (Jiang and Storrie, 2005; Samak et al., 2011; Wang et al., 2012a). Thus, specific cellular responses to cell shrinking/swelling under changes in external osmolarity can be accompanied by ERSS and associated downstream events.

Sorbitol supplementation (25 μ M) of mouse embryo culture medium to create hyperosmolarity decreases blastocyst formation and induces ERSS, including XBP1s expression and increased nuclear localization, as does tunicamycin treatment at 1–2 μ g/ml (Zhang et al., 2012). Higher concentrations of tunicamycin (5 μ g/ml) and sorbitol (50mM) arrest embryos at the 2-cell stage and increase cytoplasmic but not nuclear XBP-1 staining (Zhang et al., 2012). The higher concentrations applied to blastocysts increase apoptosis. All of these ERSS mediated responses are inhibited by treatment with tauroursodeoxycholic acid (TUDCA), which improves embryo development in vitro (Abraham et al., 2012; Kim et al., 2012; Zhang et al., 2012).

6.3 pH stress

Alkalinized medium during the zygote stage can alter post-natal growth in mice (Banrezes et al., 2011). Zygotes and embryos prior to compaction have lesser abilities to regulate intracellular pH, and treatment with ammonium may contribute to inter-blastomere heterogeneity and stress-mediated effects on gene transcription, cell lineage commitment and cell survival (Brison et al., 2014; Zander et al., 2006). Exposure to acid to reduce intracellular pH by less than 0.2 pH units for either brief period or a prolonged period impairs blastocyst formation in mouse embryos; exposure of zygotes does not reduce implantation of these blastocysts but alters fetal growth, indicating the exquisite sensitivity of zygotes to mild stress and the impact on developmental programming of later development (Zander-Fox et al., 2010). Importantly, early embryos lack active mechanisms for alleviating acid loads, and can only alleviate alkaline loads by sodium-independent

bicarbonate/chloride exchange (Baltz et al., 1991a, b), making them susceptible to perturbations in pH. Because pH can affect protein folding, calcium signaling and sequestration, and other processes, ERSS may contribute to effects on viability and long-term developmental programming following pH insults.

6.4 Maternal nutrition and physiology, and nutrient availability in vitro

The preceding sections touched upon the connections between glucose availability, lipid exposure, oxidative stress and the ERSS in affecting oocyte quality and embryo development. Other aspects of maternal nutrition and physiology, and in vitro availability of nutrients also contribute to embryonic ERSS activation. Maternal hyperglycemia and insulin resistance are significant factors. Maternal insulin resistance is harmful to embryos, due to downregulation of IGF1R (insulin like growth factor 1 receptor) and reduced glucose uptake (Louden et al., 2008). Nutrient availability is sensed in part by activation of AMPK in response to elevated AMP/ATP ratio, and AMPK inhibits TOR activity (Rehman et al., 2014). Activation of AMPK in blastocysts rescues them from the effects of maternal insulin resistance (Louden *et al.*, 2008), implicating TOR signaling in the detrimental effects of nutrient stress on early embryos. TOR may enhance amino acid transport in placenta (Jansson *et al.*, 2012). Maternal dietary protein intake is also important. Low protein diets during oocyte maturation and preimplantation development can perturb embryo metabolism, alter epigenetic information, and contribute to post-natal abnormalities (Fleming et al., 2011; Kwong et al., 2006; Kwong et al., 2007; Mitchell et al., 2009; Watkins et al., 2010, 2008a, 2008b; Williams-Wyss et al., 2014). The stress of in vitro embryo growth can perturb normal cellular physiology. Availability of nutrients in the medium such as amino acids and vitamins can ameliorate negative effects of culture (Lane and Gardner, 1998). Such effects may be mediated in part by modulating ERSS.

6.5 Oxidative stress, oxygen availability, and glucose availability

Oocyte development is sensitive to oxidative stress in the ovary (Devine et al., 2012). In embryos, oxidative stress can be problematic as early embryos may not express all the same factors that help protect somatic cells from reactive oxygen species (ROS) (Johnson and Nasr-Esfahani, 1994; Nasr-Esfahani and Johnson, 1992). Genes related to oxidative stress response (and ERSS) are expressed in immature and mature oocytes and in embryos, but may be subject to translational control as well as induction in response to stress (El Mouatassim et al., 1999).

Oxidative stress can arise for many reasons. Oxidative phosphorylation in mitochondria generates ROS. There is an intimate connection between ER stress and oxidative stress. ERSS itself can generate ROS (Landau et al., 2013). Oxidative stress then impedes correct protein folding and transport, and calcium homeostasis, and can trigger ERSS (Malhotra and Kaufman, 2007). Embryo culture in elevated (atmospheric) oxygen levels, elevated glucose, and other factors that alter oxidative phosphorylation and mitochondrial function can induce oxidative stress. Environmental toxins induce oxidative stress in embryos (Chu et al., 2013). Transient exposure to elevated glucose concentrations (or older media formulations with higher glucose concentrations) in vitro can increase ROS generation, and alter gene expression and energy metabolism (Cagnone et al., 2012; Karja et al., 2006; Rinaudo et al.,

2006), as can exposure to maternal hyperglycemia (Moley, 2001; Moley et al., 1998). Maternal hyperglycemia degrades oocyte and preimplantation embryo quality, impacts meiosis, and alters mitochondria in embryos (Wang and Moley, 2010; Wyman et al., 2008). These conditions can contribute to diabetes, growth abnormalities, and other long-term effects in progeny (Wyman *et al.*, 2008). Responses to other maternal stressors that may compromise oocyte quality and early embryogenesis can be accompanied by oxidative stress and increase in ROS production (Koyama et al., 2012; Lian et al., 2013; Ozawa et al., 2002). Embryo glutathione levels change during development and are altered in vitro (Gardiner and Reed, 1994). Oxidative stress in embryos can lead to DNA damage (Takahashi et al., 2000). Paraquat, an inducer of oxidative stress and ER stress, inhibits preimplantation development (Hausburg et al., 2005), highlighting the importance of ERSS in early development.

The benefits of reduced oxygen availability during embryo culture have long been recognized (Rinaudo et al., 2006; Umaoka et al., 1991, 1992), as have the beneficial effects of adding antioxidants or glutathione to the culture medium, particularly in overcoming cleavage stage developmental arrest (Hu et al., 2012; Lee et al., 2000; Leese, 2012; Legge and Sellens, 1991; Nonogaki et al., 1991; Takahashi, 2012; Umaoka et al., 1991). Antioxidants can improve oocyte quality and enhance embryo development (Huang et al., 2013; Ideta et al., 2012; Orsi and Leese, 2001; Takeo et al., 2014; Wang et al., 2014a) and also protect against peroxide-induced damage (Yu et al., 2014) and other stress induced damage (Cebrian-Serrano *et al.*, 2013; Koyama *et al.*, 2012; Sakatani *et al.*, 2007) in embryos and oocytes (Devine *et al.*, 2012). Additionally, reduced glucose availability in vitro has long been recognized as beneficial to embryos (Chatot et al., 1989; Leese, 2012), although glucose starvation can actually increase ROS production and is not beneficial (Jansen et al., 2009). Other maternal stressors that may compromise oocyte quality and early embryogenesis can be redressed by antioxidant treatments (Lian *et al.*, 2013). Growth factors may also improve development after oxidative stress (Kurzawa et al., 2002).

Embryos respond to oxidative stress in part via a p66(Shc)-dependent pathway (Betts et al., 2014; Favetta et al., 2007; Ren et al., 2014). ERSS and oxidative stress response often accompany each other, ERSS can mediate response to oxidative stressors (Xu et al., 2012), and treatments that alleviate one often alleviate both. Separate cellular responses to oxidative stress and ER stress combine to dictate developmental outcomes (Yoon et al., 2014b). Interestingly, the mouse zygote apoptotic response to oxidative stress resides in the cytoplasm (Liu and Keefe, 2000), consistent with the absence of early gene transcription. Overall, the observations to date indicate that many different stressors are accompanied by oxidative stress and ROS production, and that the response to these stressors arises from a combination of oxidative stress response involving p66 and activation of ERSS, and that inhibition of ERSS as well as suppression of ROS can enhance embryo viability.

6.6 Lipids

Fatty acid oxidation (β -oxidation) is essential for oocyte maturation, providing an important energy source and availability of acetyl-CoA (Dunning et al., 2014; Paczkowski et al., 2014). Inhibiting fatty acid oxidation in oocytes impairs maturation (Downs et al., 2009), and stimulating fatty acid oxidation reduces glucose metabolism, increases lipid stores, and

improves embryo development after fertilization (Johnson and Nasr-Esfahani, 1994; Paczkowski et al., 2014). Changing the serum lipid profile can change lipid availability in the ovarian follicle and affect embryogenesis after fertilization (Johnson and Nasr-Esfahani, 1994). Dietary fatty acids can enhance oocyte maturation and quality (Moallem et al., 2013). But excess circulating lipids and excess amounts of non-esterified fatty acids can increase ROS and negatively affect oocyte quality and embryo development (Leroy et al., 2005; Shehab-El-Deen et al., 2009). Palmitic acid can induce ERSS (Jung et al., 2012). Elevated levels of some polyunsaturated fatty acids in vivo can adversely affect oocyte quality and embryo development (Wakefield et al., 2008). Exposure of embryos in vitro to elevated serum lipid negatively impacts pluripotency gene expression in bovine embryos, but does not reduce blastocyst formation (Cagnone and Sirard, 2014). A high fat diet can exert subtle changes in DNA methylation and gene expression in oocytes of the treated mother, and also in the oocytes and tissues of their offspring (Ge et al., 2014). Maternal obesity negatively affects clinical outcome in human in vitro fertilization, and also degrades oocyte and embryo developmental in mice, and this is associated with abnormalities in fetal growth and changes in oocyte mitochondrial properties (Grindler and Moley, 2013; Jungheim et al., 2010; Luzzo et al., 2012). Oxidized oil and lipids are detrimental and can induce ER stress (Guo et al., 2013; Larroque-Cardoso et al., 2013; Otsuki et al., 2007), and oxidized low density lipoprotein is associated with ER stress in adipocytes (Chen et al., 2013). Hypercholesterolemia can induce ER stress in some tissues (Cai et al., 2013). These latter observations indicate that excess lipid availability may activate ERSS and possibly oxidative stress response in oocytes, and thereafter negatively affect embryogenesis. Potential negative effects of excess lipid availability may combine with negative effects of hyperglycemia in maternal diabetes, obesity, and altered nutrition status.

6.7 Cytokines

Inflammatory cytokines can activate SRC signaling, which interacts with the ERSS pathway. This can be attenuated with inhibitors of ERSS (Wan et al., 2014). Interleukins can activate ERSS, which can be pro-survival, and can also influence cellular responses to other stimuli, such as activators of NF κ B signaling (Lee et al., 2014). Additionally, ERSS can activate interleukin and cytokine expression in paracrine interactions (Meares et al., 2014). However, IL-10 deficiency combined with NADPH oxidase deficiency can induce ERSS (Tretton et al., 2014). Cytokines such as CSF1 and GM-CSF positively affect preimplantation development in mice and improve in vitro survival, and CSF1 deficiency can impair pre- and post-implantation development (Bhatnagar et al., 1995; Karagenc et al., 2005; Robertson et al., 2001; Sjoblom et al., 2005). The beneficial effects of CSF1 and GM-CSF are mediated in part by suppressing ERSS (Chin et al., 2009).

6.8 Shear stress

Shear stress during embryo handling can induce a stress response. Mouse embryo pipetting generates shear stress that is ameliorated by the presence of the zona pellucida (Xie et al., 2006). The stress activated protein kinases MAPK8 (also known as SAPK or JNK1) is activated by prolonged shear stress, and more so in hatched embryos (Xie et al., 2007). Transient stress may not lead to negative effects on the embryo, but prolonged or repeated handling or handling without the zona pellucida may be more deleterious (Xie *et al.*, 2007).

6.9 Autophagy

Induction of autophagy as opposed to apoptosis leads to cell survival. Fertilization is accompanied by increase in autophagic activity and mutation of autophagy genes impedes early development (Song et al., 2012). Induction of autophagy in bovine embryos by rapamycin enhances embryo development in vitro whereas inhibition of autophagy with 3-methyladenine reduces development, and treatment with TUDCA ameliorated the effects of autophagy inhibition (Song *et al.*, 2012). Thus, increased ER stress in early embryos may diminish the beneficial effects of autophagy and tip the scale in favor of apoptosis.

6.10 Electrofusion

One study in cloning by somatic cell nuclear transfer revealed a potential effect of electrofusion in the induction of ERSS in bovine oocytes and early embryos, including activation of ERSS related genes (Song et al., 2011). Electrofusion yielded embryos with fewer cells and different nuclear remodeling changes as compared to embryos produced by Sendai virus fusion. The negative effects of electrofusion could be prevented with TUDCA treatment to inhibit ERSS. ERSS induction by electrofusion could be arise following calcium depletion in the ER. It is noted, however, that whereas electrofusion in the bovine activates the oocyte, this does not occur in the mouse; hence it is has not been determined to what degree electrofusion or electrical pulses will elicit the ERSS in different species or using different parameters across protocols.

7. UNIQUE CONSIDERATIONS FOR ERSS IN OOCYTES AND EMBRYOS-DEVELOPMENTAL OUTCOMES

Induction of ER stress in mammalian oocytes and embryos may conflict with normal developmental processes. The progression of oocytes and early cleavage stage embryos is controlled in large part by the timely recruitment, translation and degradation of maternal mRNAs in the ooplasm. Thus, although utilizing ERSS to cope with stress may help maintain cell and embryo survival, this may come at the cost of a disruption in the normal temporal coordination of events regulated by post-transcriptional mechanisms. Additionally, activation of apoptotic mechanisms in the early embryo may eliminate damaged cells that cannot recover from severe or prolonged stress, and although mammalian embryos have considerable regulative capacities, the loss of a substantial proportion of cells can greatly reduce developmental potential.

Another challenge in mammalian embryogenesis relates to genome integrity. Embryos may begin life with DNA damage in either or both parental genomes, and this DNA damage needs to be repaired. Activation of ERSS can interfere with DNA repair (Yamamori et al., 2013; Yasui et al., 2014). The “quiet embryo hypothesis” posits that a low metabolic rate, low glucose consumption, low glycolysis, low amino acid turnover, and low oxidative phosphorylation from the zygote to the morula stage limits ROS production and maximizes viability (Baumann et al., 2007; Leese, 2002). Stress activates AMPK, which in turn increases glucose transport and glycolysis, but which also leads to changes in gene regulation. High amino acid turnover correlates with increased DNA damage (Sturmey et al., 2009). Stress and excess exogenous nutrients can thus combine to increase metabolism

and amino acid turnover (Leese, 2012), impede the repair of DNA damage, and alter gene regulation, leading either to embryo demise or long-term changes in developmental phenotype. One other link of ERSS to genome integrity is that XBP1 is observed associated abundantly in the oocyte germinal vesicle, then abundantly with the meiotic spindle in MI oocytes, but less so on spindles of MII stage oocytes, then in the cytoplasm of 1-cell embryos, and subsequently in the nuclei of 2-cell and 4-cell embryos (Zhang *et al.*, 2012). This dynamic regulation of XBP1 localization may reflect a role in chromatin regulation during meiosis and cleavage as well as periods of greater or lesser capacity for ERSS-induced DNA repair (Tao *et al.*, 2011).

Another unique aspect of early mammalian embryos relates to mitochondrial biology. Early mammalian embryos display reduced rates of oxygen consumption, which increases with cavitation, and much of the oxygen consumed by blastocysts can be via non-mitochondrial processes (Leese, 2012). Mature oocytes and early embryos display mitochondria with an inert appearance, which take on a more active appearance as development progresses, becoming elongated with more transverse cristae (Sathananthan and Trounson, 2000). Stressors such as toxins, hyperglycemia, and maternal high fat diet can induce ERSS, generate defective mitochondria in the oocyte and embryo, increase ROS, disrupt AMPK activity, and possibly inhibit the removal of defective mitochondria by mitophagy (Grindler and Moley, 2013; Yuzefovych *et al.*, 2013). This suggests that stressors that activate ERSS may not only alter embryo metabolism but may also disrupt the normal pattern of mitochondrial biogenesis/elimination to ensure healthy embryonic cells.

Early embryogenesis is also a period of extensive chromatin remodeling and susceptibility to factors that may interfere with correct developmental programming of the genome. The ERSS encompasses multiple components of transcriptional response, including modifications of histone acetylation, histone methylation, and DNA methylation, which in turn regulate diverse genes including genes related to ER stress response, genes related to the ERAD process, and other stress regulatory genes such as NRF2 (Baumeister *et al.*, 2009; Chen *et al.*, 2014; Han *et al.*, 2013; Martin *et al.*, 2014; Nakajima and Kitamura, 2013; Tao *et al.*, 2011). Additionally, histone deacetylases can modify transcription factor activity and other proteins (Fazi *et al.*, 2009; Kimura *et al.*, 2012; Palsamy *et al.*, 2014; Sato *et al.*, 2014). For example, HDAC4 sequesters ATF4 in the cytoplasm (Zhang *et al.*, 2014a). Thus, ERSS may exert a special penalty on early embryonic cells by modifying the developmental programming, and this may be one aspect of how diverse stressors can exert long-term effects on progeny phenotype.

Another special consideration for mammalian embryos related to ERSS is the wide range of manipulations to which embryos may be subjected during *in vitro* experimentation, assisted reproduction, or genome manipulation. Nutrient makeup of culture media, shear stress and embryo pipetting or handling, thermal and osmotic stress, exposure to atmospheric oxygen level, and electrofusion are common aspects of *in vitro* oocyte and embryo handling. Additionally, maternal health status, maternal serum carbohydrate and lipid content, and maternal toxin exposure also impact oocyte and embryo quality and potential responses to these oocyte/embryo handling parameters. Thus, the combination of all of these parameters

will work additively to determine the extent, severity, and duration of ERSS activation, and whether ERSS is activated in concert with other stress responses.

All of this ultimately impacts the ability of the embryo to eliminate unfolded protein, repair DNA damage, eliminate damaged mitochondria, undergo autophagic processes to repair cellular damage, execute normal transcriptional programs, execute the required maternal mRNA translational program, and survive with a sufficient number of viable cells to sustain normal long-term development. What emerges from all of these considerations is that, while mammalian embryos are remarkably resilient and can tolerate a range of exogenous stresses without dying, such regulative responses come at a potentially high cost of aberrant phenotypes lasting into adult life.

8. CONCLUDING REMARKS

The above considerations suggest that some portion of adult disease and health disorders might be preventable by minimizing insults to the oocyte and embryo *in vivo*, and also by taking precautions to minimize *in vitro* conditions that may activate ERSS. Indeed, exciting recent developments illustrate that embryo viability can be improved by managing embryonic ERSS. This includes managing the REDOX state of embryonic cells, using inhibitors of ERSS, using histone deacetylase inhibitors to modulate ERSS, promoting autophagy, and in some species avoiding procedures that deplete ER calcium stores (Song et al., 2011; Song et al., 2012; Song et al., 2014; Yoon et al., 2014b; Zhang et al., 2012). The specific interventions will need to be optimized according to species, genotype, and possibly maternal health status and specific *in vitro* culture conditions. One exciting prospect will be to discover whether *in vitro* manipulation of ERSS in oocytes and embryos might provide a way to repair damage or improve oocyte and embryo quality that may have been compromised by maternal age or adverse maternal health or nutrition states. Additionally, the need for managing ER stress in the oocyte and early embryo may extend to post-implantation stages. Recent studies reveal links between ER stress, placenta or trophoblast cell defects such as inflammation and preeclampsia, and adverse outcomes such as pregnancy loss, intrauterine growth restriction, and low birth weight (Jain et al., 2012; Kawakami et al., 2014; Lian et al., 2011; Liu et al., 2011; Redman and Sargent, 2009; Sankaralingam et al., 2006; Shi et al., 2012; Wang et al., 2012b; Yung et al., 2008; Yung et al., 2012). An overall strategy of minimizing the likelihood of activating ERSS in oocytes, early embryos, and the placenta should be beneficial for enhancing reproductive efficiencies clinically, but may also enhance overall reproductive health and reduce the incidence of adult disorders and diseases that have their origins in early embryonic stress. Further study of the benefits of avoiding exposures of oocytes, embryos, and placenta to exogenous stressors, and providing exogenous factors that inhibit ERSS (e.g., antioxidants or other compounds) before conception, during clinical and applied reproduction procedures, and during pregnancy, may yield many beneficial discoveries.

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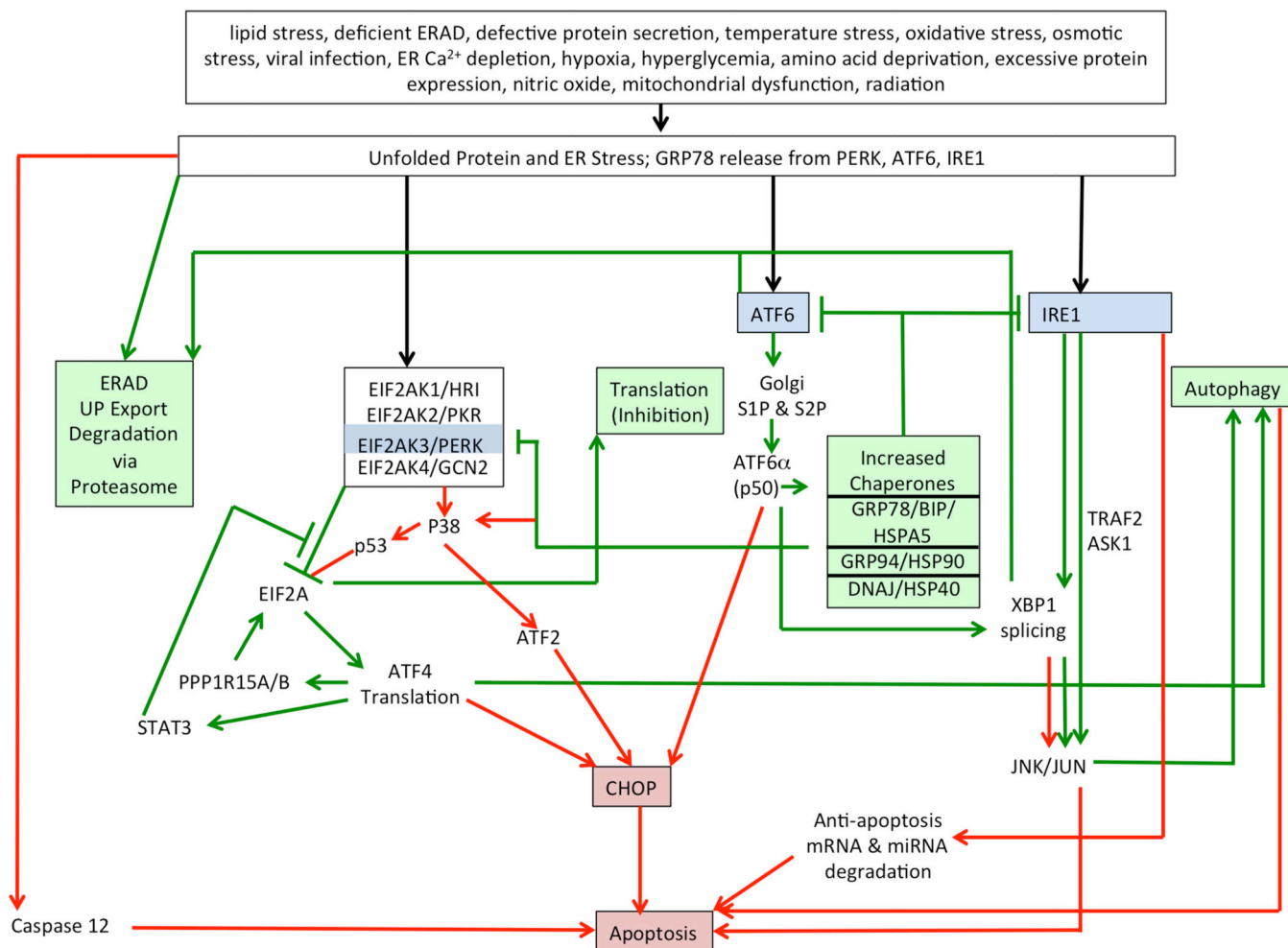


Figure 1. Summary of unfolded protein response and endoplasmic reticulum stress signaling pathways. Stress mediated by diverse agents (top box) causes accumulation of unfolded protein, which then bind to GRP78/BIP/HSPA5, releasing the three primary transducers (blue boxes). Green boxes and green lines/arrows designate pathway components that promote survival and recovery. Red boxes and red lines/arrows designate pathway components that promote apoptosis when stress is too severe or prolonged to allow survival.

Table 1

Chemical Activators and Inhibitors of ER Stress Response and Cell Death

Agent	Probable Modes of Action	Comments	Reference
Activators			
2-deoxy-D glucose	activation of AMPK as stress sensor, inhibit autophagy, ATP lowering, oxidative stress		Xi et al. 2011; Xi et al. 2013; Saez et al. 2014; Wang et al. 2014
2,4-Dichlorophenol	EIF2A de phosphorylation		Zhang et al. 2014
Ampelopsin	ROS generation	falvanol from <i>Ampelopsis grossedentata</i>	Wang et al. 2014; Zhou et al. 2014
Asymmetric dimethylarginine	inhibits nitric oxide production		Hong et al. 2014
bichalcone analog TSWU-CD4	Ca ER depletion, suppression PI3/Akt signaling		Lin et al. 2014
Brefeldin A	Inhibits protein transport from ER to Golgi		Rao et al. 2001
Calcium ionophores	Ca depletion		Rao et al. 2001
CCT020312	Activates PERK		Stockwell et al. 2012
Celastrol, triterpene	Ca release		Matos et al. 2014; Yoon et al. 2014
Clofoctol	unknown	Gram+ bacterial antibiotic	Wang et al. 2014
CuSO4	Disrupt GRP78 protein binding, induces GRP78 recomparmentalization		Qian et al. 2005; Matos et al. 2014
Flouride	intracisternal granule accumulation		Ito, Nakagawa et al. 2009,
Gadolinium	effects on Ca stores		Feng et al. 2011
H2O2	oxidative stress		Suyama et al. 2014
Homocysteine	reductive stress		Outinen et al. 1998; Yang et al. 2014
Indium, Indium tin oxide	oxidative stress		Brun et al. 2014
Indomethacin	oxphos uncoupler, cyclooxygenase inhibitor, Ca channel blockade		Tsutsumi et al. 2004; Narabayashi, Ito et al. 2014
Lead	Disrupt GRP78 protein binding, induces GRP78 recomparmentalization		Qian and Tiffany-Castiglioni 2003; Qian et al. 2005; Shinkai et al. 2010
L-NAME	nitric oxide diminishment		Shen et al. 2014
Neferine	oxidative stress	alkaloid from lotus seed embryo	Yoon et al. 2013
Nitric oxide	PPAR-delta		Cheang et al., 2014
Palm itate	ER Ca depletiobn, accumulation unfolded proteins		Laybutt et al. 2007; Cnop et al. 2010; Park et al. 2014
Paraquat			Hausberg et al., 2005
Quinones	Inhibit proteasome	derived from xenobiotics and endogenous molecules	Xiong et al. 2014
Sarsasapogenin	oxidative stress	from the Chinese medical herb <i>Anemarrhena asphodeloides</i> Bunge; a steroidal sapogenin; anti-diabetic	Shen et al. 2013; Shen et al. 2013
SNX-2112	HSP90 inhibition		Wang et al. 2014
Thapsigargin	SERCA inhibition, Ca depletion in ER		Humeres et al. 2014; Zhang et al. 2014

Agent	Probable Modes of Action	Comments	Reference
Tunicamycin	GlcNAc phosphotransferase Inhibitor; blocks N-linked glycosylation		Perez-Martin et al. 2014; Saez et al. 2014; Zhang et al. 2014
Inhibitors			
4-(2-aminoethyl)benzenesulfonyl fluoride (AEBSF)	serine protease inhibitor		Okada et al. 2003
4-phenylbutyric acid PBA	chemical chaperone		Cadavez et al. 2014; Humeres et al. 2014; Yao et al. 2014
Baicalin	nitric oxide synthase induction		Shen et al. 2014
Betaine	inhibits hyperhomocysteinemia		Ji and Kaplowitz 2003
Cajaninstilbene acid	antioxidant	a natural stilbene isolated from <i>Cajanus cajan</i> leaves	Liu et al. 2014
Emodin	Ca regulation		Wu et al. 2014
Exendin-4	GLP1R agonist		Younce et al. 2013
Glucagon like peptide 1	activation of SERCA2	effective against hyperglycemia and lipid induced ERSS	Cnop et al. 2010; Younce et al. 2013
Guanabenz	inhibits EIF2A dephosphorylation		Wan et al. 2014
Honiokol	GRP78 inducer		Kudo et al. 2008
Imipramine	works via Sigma1 receptor which is a ER chaperone that modulates Ca release		Ono et al. 2012
Metallothionein	antioxidant		Guo et al. 2009
Metformin	increases nitric oxide bioavailability, activates AMPK		Cheang et al. 2014
Metyrapone	inhibits TOR to promote autophagy and inhibit apoptosis		Kapuy et al. 2014
N-acetylcysteine	antioxidant		Yang et al. 2014
Naltrexone	opioid receptor antagonist; works via Sigma1 receptor which is a ER chaperone that modulates Ca release		Moslehi et al. 2014
Nitric oxide	PPAR-delta		Cheang et al. 2014; Narabayashi et al. 2014
Rapamycin	inhibits TOR to promote autophagy and inhibit apoptosis	bacterial macrolide	Li et al. 2014
Salubrinal	inhibits EIF2A dephosphorylation		Boyce et al. 2005; Kapuy et al. 2014
sodium tanshinone IIA sulfonate	reduces reactive oxygen species in irradiated cells		Gu et al. 2014
tauroursodeoxycholic acid (TUDCA)	ER chaperone properties; prevents BAX transport to mitochondria		Rivard et al. 2007; Cadavez et al. 2014; Yan et al. 2014
Trolox	antioxidant	analog of vitamin E	Brito et al. 2014