Aripiprazole Cytotoxicity Coincides with Activation of the Unfolded Protein Response in Human Hepatic Cells

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Received December 8, 2019; accepted June 8, 2020

ABSTRACT
Schizophrenia is a mental disease that results in decreased life expectancy and well-being by promoting obesity and sedentary lifestyles. Schizophrenia is treated by antipsychotics. Although the second-generation antipsychotics (SGA), Olanzapine and Aripiprazole, are more effective in treating schizophrenia, they display a higher risk of metabolic side effects, mostly by development of diabetes and insulin resistance, weight gain, and dyslipidemia. Endoplasmic reticulum (ER) stress is induced when ER homeostasis of lipid biosynthesis and protein folding is impaired. This leads to the activation of the unfolded protein response (UPR), a signaling cascade that aims to restore ER homeostasis or initiate cell death. Chronic conditions of ER stress in the liver are associated with diabetes and perturbed lipid metabolism. These metabolic dysfunctions resemble the pharmacological side effects of SGAs. We therefore investigated whether SGAs promote the UPR in human and mouse hepatocytes. We observed full-fledged activation of ER stress by Aripiprazole not by Olanzapine. This occurred at low micromolar concentrations and to variable intensities in different cell types, such as hepatocellular carcinoma, melanoma, and glioblastoma. Mechanistically, Aripiprazole caused depletion of ER calcium, leading to activation of inositol-requiring enzyme 1 (IRE1) and protein kinase R (PKR)-like endoplasmic reticulum kinase (PERK), two major transducers of the UPR. Cells underwent apoptosis with Aripiprazole treatment, which coincided with UPR induction, and this effect was reduced by adding glutathione without affecting UPR itself. Deletion of IRE1 from HepG2, a human liver cancer cell line, protected cells from Aripiprazole toxicity. Our study reveals for the first time a cytotoxic effect of Aripiprazole that involves the induction of ER stress.

SIGNIFICANCE STATEMENT
The antischizophrenic drug Aripiprazole exerts cytotoxic properties at high concentrations. This study shows that this cytotoxicity is associated with the induction of endoplasmic reticulum (ER) stress and IRE1 activation, mechanisms involved in diet-induced obesity. Aripiprazole induced ER stress and calcium mobilization from the ER in human and mouse hepatocytes. Our study highlights a new mechanism of Aripiprazole that is not related to its effect on dopamine signaling.

Introduction
Schizophrenia is a worldwide mental disease that results in decreased life expectancy associated with obesity and sedentary lifestyle. It affects around 0.3%–0.7% of the population at young ages, although in certain cases, it can appear in a later stage of life (van Os and Kapur, 2009). Schizophrenia typically occurs during late adolescence and early adulthood and is characterized by abnormal social behavior. Common symptoms are false beliefs, unclear or confused thinking, and hearing imaginary voices. The causes of schizophrenia are assumed to be a combination of genetic and environmental factors (Owen et al., 2016). From a genetic point of view, the
heritability is around 80%, although environmental factors such as living environment, drug abuse, and prenatal stress also contribute to its etiopathogenesis (van Os and Kapur, 2009). Aripiprazole (Ari) and Olanzapine (Ola), drugs of the latest generation of antipsychotics, referred to as second-generation antischizophrenics (SGAs), have shown high efficacy to treat this mental disorder. However, they display a greater risk of metabolic side effects (i.e., weight gain, dyslipidemia, and type 2 diabetes). The mechanisms responsible for these severe metabolic disturbances are still unknown. Some studies suggest that hypothalamic dopamine antagonism or disruption of hypothalamic regulation of glucose levels, along with anticholinergic-induced inhibition of insulin secretion, could play a role (Panariello et al., 2011). However, peripheral effects of the drugs that are unrelated to their dopamine antagonism or effects in the central nervous system were also invoked.

The endoplasmic reticulum (ER) is the entry point into the secretory system in which proteins acquire their folded state and assemble into complexes. ER stress occurs when the amount of protein entering the ER exceeds its folding capacity. This imbalance induces an adaptive reaction collectively termed the unfolded protein response (UPR) (Frakes and Dillin, 2017). The mammalian UPR is transduced by three major ER sensors, IRE1, PERK, and activating transcription factor 6 (ATF6), that undergo activation under ER stress conditions. When unfolded proteins accumulate at the ER, the chaperone 78-kDa glucose-regulated protein (GRP78) dissociates from the sensors and allows their activation (Amin-Wetzel et al., 2017). IRE1 and PERK are activated by autophosphorylation, although ATF6 traffics to the Golgi. Activated IRE1 splices the mRNA of XBP1 in a noncanonical fashion, yielding the potent transcription factor–spliced X box binding protein 1 (XBP1) (Yoshida et al., 2001). Activated PERK phosphorylates eukaryotic translation initiation factor 2α (eIF2α), which attenuates the proliferation of global protein synthesis. At the same time, selective translation of certain mRNA molecules such as ATF4 is induced. ATF6 is proteolytically fragmented in the Golgi, yielding an active transcription factor (Ye et al., 2000). Whereas acute ER stress leads to apoptosis, chronic sublethal ER stress causes cellular adaptation and resistance to apoptosis, characterized by modulation of mRNA localization and stability (Rutkowski et al., 2006). The UPR profoundly affects cellular homeostasis, regulating cell cycle, apoptosis, metabolism, and autophagy. Studies in animal models have demonstrated that extended activation of the UPR plays an important role in chronic metabolic diseases, including obesity, insulin resistance, and diabetes (Ozcan et al., 2004). Specifically, obesity may be a consequence of chronic ER stress, particularly in the liver and adipose tissues. Part of the mechanism is the role of UPR in promoting inflammatory conditions (Liu et al., 2016). In this regard, all three main branches of the UPR mediate cell autonomous proinflammatory transcriptional programs, which are mainly governed by transcription factors such as nuclear factor kappa-light-chain-enhancer of activated B cells (NF-kB) and AP-1 (Hotamisligil and Erbay, 2008; Verfaillie et al., 2013). The signaling pathways involved in the UPR and inflammation are interconnected through various mechanisms, including the production of reactive oxygen species (ROS) and the release of calcium from the ER (Zhang and Kaufman, 2008) as well as direct stimulation of the secretion of proinflammatory cytokines. Interestingly, free fatty acids, which are elevated in obesity, have the potential to induce ER stress in various cells, including adipocytes and hepatocytes (Jiao et al., 2011; Cnop et al., 2012; Pardo et al., 2015).

Because of the connection of ER stress and obesity, we explored whether SGA drugs induce ER stress. We observed full-fledged activation of the UPR at low micromolar concentrations of Ari but not of Ola in hepatic cells. This was associated with cell death and was rescued in part by inhibition of IRE1 or the addition of reduced glutathione. Our data indicate a previously unknown pharmacological activity of Ari in inducing ER stress conditions and UPR signaling. It should be noted that the concentrations that induce the UPR in cultured cells are approximately 25-fold higher than the pharmacokinetic Cmax of Ari under its regular dosing regimen, suggesting that this pharmacodynamic property may be related more to Ari over dosage toxicity. However, the exact exposure of the liver to Ari, particularly after oral administration, is not known.

### Material and Methods

#### Experimental Procedures

**Cell Lines and Culture Conditions.** HepG2, Hep3B, melanoma (Mel526), and glioblastoma cells were cultured in high-glucose Dulbecco’s Modified Eagle medium (Sigma-Aldrich) supplemented with 10% fetal bovine serum (Invitrogen), 2 mM of L-glutamine (Biologic Industries, Israel), 1% penicillin-streptomycin solution (Biologic Industries), and 1 mM of sodium pyruvate (Biologic Industries). Cells were incubated at 37°C under 5% CO₂.

**Chemical Reagents.** Aripiprazole (129722-12-9; Acros Organics), Olanzapine (035M4781V; Sigma-Aldrich), Rifampicin (AS470650; Apollo), Z-VAD(Fmk)-FMK (0542537-4; Cayman chemical), N-acetyl-L-cysteine (616-91-1; Sigma-Aldrich), BAPTA-AM (B6769; ThermoFisher), EGTA (67-42-5; Sigma-Aldrich), STF-083010 (1A213103; Tocris Bioscience), Glutathione reduced ethyl ester (92614-59-0; Sigma-Aldrich), propidium iodide (25535-16-4; Sigma-Aldrich), GSK2606414 (5107; Tocris), Thapsigargin (67526-95-8; Fermontek), Tunicamycin (11089-65-9; Fermontek), and DMSO (440075; Biosolve) were used.

**Western Blotting.** Cells were either trypsinized or directly harvested by cell scrapping, centrifuged at 3000 relative centrifugal force (rcf) for 5 minutes, and washed twice in cold PBS. For cell lysis, RIPA buffer supplemented with protease and phosphatase inhibitors was added in a volume of about four times the cells pellet and then vibrated for 20 minutes at 4°C. Lysates were cleared by centrifugation at 20,000 rcf for 30 minutes at 4°C. Protein content was measured by BCA. Reduced Laemmli sample buffer (5 ×) was added, boiled for 5 minutes at 95°C, and loaded on SDS-PAGE. After SDS-PAGE, gels were blotted onto Polyvinylidene Fluoride (PVDF) membranes by using BioRad PowerPacTM. Blots were blocked in 10% skim milk in Tris-buffered saline/Tween 20 buffer for 1 hour at room temperature. The following primary antibodies were used: rabbit anti-IRE1 (1:1000) (3294S; Cell Signaling); rabbit anti-phosphorylated IRE1 (S724) (1:1000) (ab124945; Abcam); mouse anti-puromycin antibody, IRE1 (1:1000) (3294S; Cell Signaling); rabbit anti-phosphorylated IRE1 (S724) (1:1000) (ab124945; Abcam); mouse anti-puromycin antibody, clone 12D10 (1:1000) (MABE343; Millipore); and polyclonal rabbit anti-PERK (1:1000) (5683; Cell Signaling). The following primary antibodies were used: rabbit anti-PERK (1:1000) (5683; Cell Signaling); rabbit anti-phosphorylated PERK (S703) (1:1000) (ab124945; Abcam); mouse anti-PERK (1:1000) (5683; Cell Signaling); rabbit anti-PKR (1:1000) (5679; Cell Signaling); rabbit anti-IRE1 (1:1000) (3294S; Cell Signaling); rabbit anti-phosphorylated IRE1 (S724) (1:1000) (ab124945; Abcam); mouse anti-puromycin antibody, clone 12D10 (1:1000) (MABE343; Millipore); and polyclonal rabbit anti-p70 (1:1000) (provided by Dr. Ariel Stanhill, Open University, Israel). Secondary horseradish peroxidase–conjugated goat anti-rabbit and anti-mouse (Jackson ImmunoResearch, West Grove, PA) were used. Blots were developed with Bio-Rad ChemiDoc XR. Quantification was done by densitometry using Image Laboratory software.

### Quantitative Real-Time Polymerase Chain Reaction and Analysis of XBP1 mRNA Splicing

Total RNA was isolated by using TRI-reagent (Bio-Rad Laboratories, Berkeley, CA). cDNA was synthesized by using 1 μg of RNA with qScript cDNA Synthesis KIT (Quanta Biosciences, Beverly, MA) according to the manufacturer’s instructions. Bio-Rad iQ universal SYBR Green Supermix was used for quantitative polymerase...
chain reaction (PCR) analyses. Actin was used as an endogenous housekeeping gene for PCR quantification. Analysis was performed on CFX Connect Real-Time PCR Detection System (Bio-Rad), using the Bio-Rad CFX manager 3.1 software. The following primers were used: CYP3A4 forward, 5′-TGAAGTTCATCTGCAGACCAG-3′, and reverse, 5′-AGTCTGTTCTGCTTCTGAGTG-3′; and Actin forward, 5′-CCA ACTAATCCCCAGATCATCTACA-3′, and reverse, 5′-ACATGC GATATGCTGCA-3′. Reverse-transcription PCR (RT-PCR) was used for the detection of XBP1 mRNA splicing, using 5′-Red Load Taq (LAROVA) with forward, 5′-CCTCTGAAATTGAGAACCAG-3′, and reverse, 5′-CCAAAAGGATATACGACTCGG-3′.

Puromycin Labeling for Protein Synthesis Evaluation

Cells were harvested, centrifuged, and resuspended in 3 ml of fresh medium. A portion of 1 ml was analyzed for cell counts, and the remaining 2 ml were pulsed for 10 minutes at 37°C to detect puromycin incorporation. After the addition of puromycin, cells were immediately incubated at 37°C under mild shaking, and ice-cold PBS was added to a volume of 40 ml to terminate the protein labeling. Cells were washed twice with cold PBS. 10 × 10⁶ cells, according to the count analysis performed, were then lysed in 200 μl of 1× reduced Laemmli sample buffer preheated to 70°C. Equal volumes were loaded on 12% SDS-PAGE and probed with an antipuromycin antibody after blotting.

Live Cell Calcium Imaging

Cells were spotted onto glass coverslips coated with Poly D-Lysine and allowed to settle down for 2.5 hours at 37°C and 5% CO₂. Cells were loaded with 2.5 μM Fura-2-AM dissolved in Ringer’s solution (150mM NaCl, 2.5mM KCl, 1.8mM CaCl₂, 1mM MgSO₄, 10mM HEPES, and 10mM D-glucose), adjusted to pH 7.4 with NaOH and supplemented with Pluronic F-127 acid (0.02%), for 1 hour at room temperature. After the addition of puromycin, cells were immediately incubated at 37°C under mild shaking, and ice-cold PBS was added to a volume of 40 ml to terminate the protein labeling. Cells were washed twice with cold PBS. 10 × 10⁶ cells, according to the count analysis performed, were then lysed in 200 μl of 1× reduced Laemmli sample buffer preheated to 70°C. Equal volumes were loaded on 12% SDS-PAGE and probed with an antipuromycin antibody after blotting.

Generation of Knockout Cells Using CRISPR/Cas9

Guide RNA (gRNA) was designed according to Doench et al. (2016) and cloned into PX459 vector according to the Zhang laboratory protocols. HepG2 cells were transfected by using Transit 2020 (3 μl per 1 μg DNA; Mirus Bio, Madison, WI) and then underwent puromycin selection (2 μg/ml) for 48 hours followed by single-cell cloning by limiting dilution. Clones were screened by immunoblotting for the target protein. gRNA sequences for human PERK were 5′-GAATATACCGAAGTTCAAAG-3′; and reverse, 5′-CCAAAAGGATATACGACTCGG-3′.

Flow Cytometry

Cells were harvested, centrifuged, washed twice in PBS, and filtered through a 100-μm strainer directly to FACS tubes. Analyses were performed with Cytolux cytometer (Beckman Coulter, Brea, CA) using the CytExpert software.

Propidium Iodide Staining

To evaluate the percentage of dead cells using flow cytometry analysis, 1 μl of propidium iodide (1 mg/ml) was added to the cells before the analysis. Equal number of events were recorded, and PI-positive cells were gated and considered as dead cells.

Isolation and Culture of Primary Mouse Hepatocytes

Mice received health care according to Spanish and European legislation. All animal experimentation was approved by the Spanish National Research Council (CSIC) and Comunidad de Madrid Animal Care and Use Committees. Primary mouse hepatocytes were isolated from 8- to 10-week-old male C57Bl6 mice by perfusion with collagenase as described (Benveniste et al., 1988). Cells were seeded on 6- or 12-well collagen IV precoated plates (Corning, New York, NY) and cultured in media containing Dulbecco’s Modified Eagle medium and Ham’s F-12 medium (1:1) supplemented with heat-inactivated 10% FBS, 2 mM glutamine, 15 mM glucose, and 1 mM sodium pyruvate (attachment media). Cells were maintained in this medium for 24 hours. Then, the medium was replaced with fresh media containing the indicated drugs.

Oil Red-O and Hematoxylin and Eosin Staining

Hematoxylin and Eosin and Oil Red-O staining was performed as previously reported (Erez et al., 2017).

Results

Aripiprazole at Low Micromolar Concentration Induces ER Stress. The peak plasma concentration of Aripiprazole in treated patients typically reaches a 1- to 2-μM concentration under chronic regimen (Nagai et al., 2017). However, the corresponding concentrations of the drug in the liver can be much higher. For example, Ola levels in the liver of rats are 30-fold higher than that of the serum levels (Aravagiri et al., 1999). Although we have not found documentation on Aripiprazole concentrations in the liver or serum in rodents, we assume that high micromolar to low millimolar concentrations may represent a reasonable exposure to the liver at the clinical dosage regimen. Treatment of the human hepatocellular carcinoma (HCC) cell line HepG2 with increasing concentrations of Aripiprazole induced IRE1 hyperphosphorylated at concentrations higher than 20 μM (Fig. 1, A and B). The sarco/endoplasmic reticulum Ca²⁺-ATPase (SERCA) inhibitor thapsigargin (Tg) was used as a positive control for UPR induction. Moreover, the total level of IRE1 was induced by Aripiprazole, a typical effect of ER stress (Shemorry et al., 2019). We detected phosphorylated IRE1 as early as 4 hours after Aripiprazole addition (Fig. 1C). This was accompanied by the splicing of XBP1 mRNA (Fig. 1A, lower panel). Immunoblotting analysis of PERK showed the shift pattern in its immunoreactivity upon stress (Tg and Aripiprazole), which is because of its autophosphorylation (Fig. 1A). This mobility shift was not observed by coincubation with the PERK inhibitor (Supplemental Fig. 1), and thus both arms of the UPR are activated by Aripiprazole. These data are supported by the analysis of newly synthesized proteins, which indicate a concentration-dependent reduction by Aripiprazole, a feature of eIF2α phosphorylation by PERK (Supplemental Fig. 2, A and B).

Ola and Aripiprazole affect schizophrenia in similar mechanisms by curtailing dopamine signaling. However, Ola did not induce the UPR at concentrations up to 50 μM (Fig. 1D). This suggests that the mechanisms by which Aripiprazole activates the UPR may not be related to dopamine signaling. To exclude a relationship of Aripiprazole-mediated UPR and the activation of dopamine receptors, we treated the HepG2 cells with a combination of Aripiprazole and dopamine. Dopamine alone neither promoted the UPR nor inhibited UPR induction by Aripiprazole (Supplemental Fig. 2C).
We treated a second HCC cell line, Hep3B, to explore whether the ER stress caused by Ari is a general effect of the drug. We observed an even stronger induction of ER stress in this line (Supplemental Fig. 3A). Ari, though to a lesser extent, also promoted ER stress in MEL526 and glioblastoma cells, indicating that this is a general pharmacological response to the drug. The fact that Ari induces ER stress in specialized drug-metabolizing cells such as liver cells and in nondrug metabolizing cells suggests that drug metabolism is not required for UPR induction. To address this, we treated Hep3B cells with the CYP3A4 inducer rifampicin and compared the induction of the UPR. Although CYP3A4 mRNA was induced by rifampicin as expected, we did not observe differences in UPR induction (Supplemental Fig. 3B), indicating that Ari-mediated induction of the UPR is independent of drug catabolism.

Aripiprazole Induces Calcium Mobilization from the ER. ER homeostasis is perturbed by several insults, such as depletion of calcium content, a metal ion essential for chaperone activity, and protein folding at millimolar concentrations. Once calcium leaks to the cytoplasm, protein folding is inhibited and UPR is induced. We performed live-cell calcium imaging in HepG2 cells after an overnight treatment to test whether ER calcium levels are affected by Ari. The experiment was designed in a manner that after the overnight treatment, the remaining calcium in the ER is released by Tg treatment in the course imaging (Fig. 2A). Thus, a positive signal indicates levels of ER calcium concentration. As expected, DMSO (negative control) provided the highest signal, whereas Tg overnight treatment (positive control) completely dwindled the calcium storage in the ER, resulting in no effect of Tg in the course of imaging. Similar to Tg, which depleted the storages of calcium in the ER when added overnight, Ari used at concentrations above 20 μM resulted in the diminishing of calcium levels in the ER (Fig. 2B). We challenged Hep3B cells with a 30-μM concentration to generalize the effect of Ari on calcium mobilization. We observed that calcium ER was mobilized in these cells as well (Supplemental Fig. 3, D and E). The dose-response correlation of ER calcium mobilization and the induction of the UPR reflect a causative relationship, indicating that Ari induces the UPR by causing calcium depletion from the ER.

Aripiprazole Cytotoxicity is Regulated by the UPR. The UPR is an adaptive response to ER stress that promotes survival, a response termed “adaptive UPR.” However, it also

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**Fig. 1.** Aripiprazole, but not Olanzapine, induces ER stress in HepG2 cells. (A) HepG2 cells were treated overnight with the indicated concentrations of Ari. Total cell extracts of the adhered cells were analyzed by immunoblotting for phosphorylated IRE1, total IRE1, total PERK, and p97 as loading control. The lower panel shows RT-PCR analysis of XBP1 splicing after Ari treatment. A representative experiment of three independent repetitions is shown. (B) Relative quantification of phosphorylated IRE1 (24 hours). Shown is an average of three independent experiments ± S.D. (C) Time course of IRE1 phosphorylation after Ari treatment. Tg (2.5 μg/ml) was used as a positive control. (D) Similar analysis as in (A) was conducted for Ola. Statistical significance is indicated by **P < 0.01; ***P < 0.001 by ANOVA.
participates in induction of apoptosis if the stress is not resolved, termed as the "terminal UPR" (Hetz et al., 2015). The transition between adaptive to terminal UPR is dependent on the stressor and the cell type. Cytotoxicity of Ari has been previously demonstrated, but the mechanism is not well understood and was postulated to involve inhibition of tyrosine kinases such as Src and Syk (Kim et al., 2017; Yoo et al., 2018). In dopaminergic neurons, Ari protected from glutamate-induced cytotoxicity. The dual activities of Ari may suggest that, like the UPR, it controls prosurvival and prodeath programs. We therefore investigated whether, at concentrations that UPR is induced, Ari exerts a cytotoxic effect. Figure 1 shows mild ER stress in HepG2 cells at 20 μM, and its intensity increases in a concentration-dependent manner up to 30 μM. However, at concentrations higher than 30 μM, very few cells survived after 24 hours of treatment (Supplemental Fig. 4). At first, we verified whether the mechanism of Ari-mediated cell death is caspase-dependent. Cell death was evaluated by flow cytometry, examining light-scattering properties, and positive staining of dead cells with propidium iodide (PI). Inclusion of the pan caspase inhibitor Z-VAD-FMK inhibited Ari toxicity when used at a 30-μM concentration (Fig. 3B). To examine whether the UPR plays a role in the cytotoxicity of Ari, we generated IRE1 knockout (KO) and PERK KO HepG2 cells by CRISPR/Cas9 gene editing (Supplemental Fig. 5A). Again, cells were treated overnight with 30 μM of Ari. As shown in Fig. 3A, deletion of PERK mildly reduced the number of dead cells, although with no statistical significance, whereas IRE1 deletion significantly protected the cells against death (Fig. 3A). This effect was also supported in MEL526 treated with an IRE1 inhibitor (Supplemental Fig. 3B). Altogether, these data indicate that although both arms of the UPR may participate in Ari cytotoxicity, IRE1 seems to play a dominant role. Unfortunately, we were unable to address ATF6 contribution because of the absence of specific inhibitors to this UPR arm as well as specific antibodies.

**Glutathione Protects Cellular Viability Downstream to the UPR.** ER stress and particularly calcium leakage from the ER induce ROS production (Cao and Kaufman, 2014). Conversely, oxidative stress promotes ER stress by affecting the process of protein folding. To obtain better insight into this interaction in the context of Ari toxicity, we treated HepG2 cells with ROS and calcium scavengers and assessed viability by flow cytometry. It should be noted that the addition of DMSO as the vehicle control resulted in 2%–7% of cell death, a variability probably associated with culture conditions. We thus normalized the cell death in each experiment to that of its own DMSO control. The addition of N-acetyl L-cysteine (NAC) and the calcium chelators BAPTA and EGTA did not affect the cytotoxicity of Ari in HepG2 cells (Fig. 4). In fact, BAPTA and EGTA were toxic by themselves (Supplemental Fig. 7B). However, when reduced glutathione (GSH) was added using GSH ethyl ester, we observed a significant improvement in survival of the cells relative to Ari alone (Fig. 5). Under these conditions, UPR activity was not affected (Supplemental Fig. 6), suggesting that GSH protects cell viability by mechanisms that are either downstream or independent of the UPR.

**Aripiprazole Induces the UPR, Lipid Accumulation, and Calcium Mobilization from the ER in Primary Mouse Hepatocytes.** We isolated primary hepatocytes from mouse livers and treated them with Ari at different concentrations for different time periods to examine if Ari elicits ER stress in untransformed hepatocytes. Tunicamycin (Tm) was used as a positive control for UPR induction. Figure 6A illustrates Ari-induced UPR as early as 4 hours upon addition to primary mouse hepatocytes as indicated by IRE1 and PERK phosphorylation and the splicing of XBP1 mRNA (Fig. 6, A–E). Analysis of ER calcium in the primary hepatocytes confirmed its depletion by Ari, as was observed in human HCC cells (Fig. 6, G and H). This also occurred at concentrations lower than 20 μM. In contrast to HCC cells, Ari-induced UPR in the primary cells was maintained until
12 hours of treatment, and then it was reduced (data not shown), suggesting that primary hepatocytes are able to adapt to this stress. However, the viability of the primary hepatocytes was compromised after 24 hours of treatment (Supplemental Fig. 8C). Interestingly, Oil Red-O positive lipid droplets were visualized in the surviving cells (Fig. 6F). These data suggest that in contrast to transformed cells, which die by apoptosis, primary hepatocytes adapt to Ari-induced ER stress, and the adaptation involves accumulation of intracellular lipids. To demonstrate a connection between Ari-induced UPR and the accumulation of intracellular lipids, we performed a dose-response analysis. To avoid artifacts of cell death, we limited the incubation to 12 hours and elevated the concentration of Ari up to 40 \( \mu M \). Lipids accumulated in association with the development of UPR, assayed by XBP1 mRNA splicing (Supplemental Fig. 8, A and B). These data are consistent with the induction of liver steatosis by Ari reported in rats (Soliman et al., 2013). Evidence of ER stress was not observed in vivo after a single administration of Ari at a dose of up to 50 mg/kg in mice (data not shown). It is noteworthy to mention that solubilization of the drug was not achieved in a mixture of Tween-80, DMSO, and water, and the drug was eventually administered as a suspension.

Fig. 3. IRE1 contributes more than PERK to Ari-induced cytotoxicity. (A) WT, IRE1 KO, and PERK KO HepG2 were treated with DMSO and 30 \( \mu M \) of Ari. Cells were stained with PI and analyzed by flow cytometry. Representative dot plots of forward vs. side scattering properties and percentage of PI-positive cells are shown. (B) Percentage of PI-positive cells, measured by flow cytometry, after overnight treatment with DMSO, Ari (30 \( \mu M \)), and Ari with Z-VAD-FMK (25 \( \mu M \)). Average of three independent experiments ± S.D. is shown. Statistical significance is indicated by *\( P < 0.05 \); **\( P < 0.01 \) by ANOVA for Z-VAD-FMK treatment and for the KO cells by paired Student’s t test.

Discussion

Stress responses evolved to respond to transient insults and protect the hosts until stress conditions dissipate. However, if stress lingers in a chronic manner, adaptation may involve irreversible plasticity that ultimately affects tissue function and leads to pathology. As such, chronic conditions of ER stress have been implicated in metabolic and inflammatory diseases in a manner that cannot be reversed even if the stress eventually subsides. Congruent with this hypothesis is the difficulty to identify the initial mechanisms associated with drug-induced liver injuries. In some cases, after exposure to the drug even for a short duration, pathologic conditions develop in a manner that cannot be reversed upon cessation of the drug treatment. Typical examples are amiodarone- and tamoxifen-induced liver steatosis. For amiodarone, we have previously showed that ER stress can be the initial trigger for the steatosis after a single administration of the drug (Erez et al., 2017).

The development of weight gain, insulin resistance, diabetes, and dyslipidemia in SGAs treated patients is a clinical problem worldwide that limits compliance. The molecular mechanisms underlying these metabolic changes may be subtle and short-lived. Thus, stress responses triggered by
SGAs are a plausible mechanism for the side effects developed in patients treated with these drugs. With this premise, we sought to test whether Ari and Ola induce ER stress and UPR signaling. Although Ola treatment is more prone to induce metabolic derangement and weight gain in patients with schizophrenia than Ari (Musil et al., 2015), we found that Ari, but not Ola, induces ER stress, as evident by the UPR signaling (Fig. 1). A second difference between Ari and Ola relates to cytotoxicity. Whereas Ari in vitro is toxic at low micromolar concentrations, Ola at this concentration range is not toxic. This suggests that the two drugs have fundamental pharmacodynamic properties that are not related to their effects in dopamine signaling.

Most likely, the induction of ER stress by Ari occurred by the parental drug in all cell lines tested in a dopamine receptor–independent manner (Supplemental Figs. 2 and 3). These results indicate a new off-target effect of Ari that may occur during treatment when its concentrations in the liver exceed the threshold for adaptive UPR. Our data clearly show that ARI-induced ER stress correlates with calcium mobilization from the ER (Fig. 2), but it is unclear which mechanisms mediate this response. Calcium homeostasis in the ER is controlled by its active transport to the ER by SERCA, its leakage through the Sec61 translocon (Lang et al., 2017), or by triggering ionotropic calcium channels such as the inositol trisphosphate receptor (IP3R) (Taylor and Tovey, 2010). Thus, further investigation is needed to delineate which of these mediators of the ER machinery are responsible for the depletion of calcium induced by Ari.

The cytotoxicity of Ari has been documented for several cancer cell lines (Baek et al., 2015; Kim et al., 2017; Yoo et al., 2018). Some studies suggested that Ari inhibits key tyrosine kinases of the Src family as the mechanism responsible of cell death (Kim et al., 2017). Although this mechanism has not been addressed in our study, assessment of the contribution of the different molecular pathways to Ari-induced cell death is rather complex. Our data confirm the cytotoxic properties of Ari (Fig. 3; Supplemental Fig. 4). In particular, in HepG2 cells, death was associated with ER stress, suggesting a correlation between both phenomena. Apoptosis induced by ER stress is a complex process that is modulated by the UPR itself. Each branch of the UPR has been shown to be implicated in promoting both viability and cell death. This is largely dependent on the magnitude of the stressor and the cell type.

**Fig. 4.** N-acetyl cysteine (NAC), BAPTA-AM, and EGTA do not protect from Arizapridole-induced cell death. (A) HepG2 cells were treated overnight with DMSO and 30 μM of Ari in the absence or presence of NAC, BAPTA-AM, or EGTA. Percentage of PI-positive cells of a representative experiment, as measured by flow cytometry after treatment, is shown. (B) Average of three independent experiments ± S.D. is shown. No statistical significance was measured.
In fact, IRE1, primarily by XBP1-independent mechanisms, promotes cell death (Maurel et al., 2014). Curiously, it is also implicated in promoting survival of several tumor types (Tang et al., 2014). Likewise, PERK also promotes cell death by enhancing C/EBP homologous protein (CHOP) levels and elevating ROS production (Zeeshan et al., 2016). However, PERK is also a potent oncogene in certain tumors and confers protection against certain chemotherapies (Luo and Lee, 2013; Ju et al., 2017). At least for HepG2 cells, removal of IRE1 circumvents Ari-induced cell death (Fig. 3A). We have not investigated further downstream mediators responsible for this effect, and in this regard, further research is needed. Removal of IRE1 results in hyperactivation of PERK and ATF6 pathways; therefore, the mechanism of protection seen in IRE1 KO may be PERK-dependent. Preliminary data show that the protective effect of IRE1 deletion is partially lost in IRE1/P ERK double-KO HepG2 cells (results not shown), supporting a role for PERK in promoting survival rather than death of HepG2 cells in the absence of IRE1. Because both IRE1 and PERK are druggable, we investigated whether pharmacological inhibition of IRE1 can protect from Ari-induced cell death. Inhibitors for IRE1 are being developed for cancer and diabetes (McGrath et al., 2018), which include inhibitors to both kinase and RNAase activities of the protein. Whereas the effect of IRE1 RNAse inhibitor in HepG2 cells was minor, in melanoma cells, it protected from Ari-induced cell death. These data may provide a novel indication for IRE1 inhibitors and should be explored in animal models.

Because ER and oxidative stress are intimately connected, to further investigate the mechanisms of cell death induced by Ari, we analyzed whether calcium chelators or elevating the reductive potential of the cells may offer protection. Both EGTA and BAPTA-AM were toxic to HepG2 cells; thus, we could not establish calcium leakage as the reason for cell death. However, addition of NAC at low millimolar concentrations did not affect HepG2 viability. A possible explanation

**Fig. 5.** GSH ethyl ester protects from Aripiprazole-induced cell death. HepG2 cells were treated overnight with DMSO and 30 μM of Ari in the absence or presence of GSH ethyl ester. Percentage of PI-positive cells of a representative experiment, as measured by flow cytometry after treatment, is shown. Average of three independent experiments ± S.D. is shown. Statistical significance is indicated by *P < 0.05 by ANOVA.
for such a lack of effect could be the inability to change the redox balance at this timeframe. We did observe a mild protective effect for reduced glutathione when added directly; however, cells were still susceptible to Ari-induced cell death (Fig. 5). UPR induction by Ari was not affected by GSH addition, suggesting that depletion of GSH is not an upstream trigger for ER stress, and its restoration improves the cellular fitness to curtail Ari cytotoxicity in an ER stress–independent manner.

Whereas cancerous cell lines responded to Ari by triggering ER stress resulting in cell death, primary hepatocytes responded to Ari with much faster kinetics of stress induction and accumulation of intracellular lipids. The prolipogenic effect of Ari in primary hepatocytes is consistent with the in vivo effect of Ari in humans and rodents. Although there is little evidence of liver damage after Ari treatment in the clinics, increased incidence of liver steatosis has been reported for Ari- and other SGA-treated patients (Morlán-Coarasa et al., 2016). Becuase ER stress induced genetically or by overexpression of misfolded proteins is also involved in liver steatosis (Kim et al., 2018), Ari-mediated ER stress should be further investigated primarily in individuals who already suffer from indolent nonalcoholic liver diseases with no apparent symptoms. Our work suggests that these patients should be carefully monitored when treated with SGAs.

Authorship Contributions
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References