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Cooperative interaction of benzo[a]pyrene and ethanol on plasma membrane remodeling is responsible for enhanced oxidative stress and cell death in primary rat hepatocytes

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Abstract
Several epidemiologic studies have shown an interactive effect of heavy smoking and alcohol heavy drinking on the development of hepatocellular carcinoma. It has also been recently described that chronic hepatocyte death can trigger excessive compensatory proliferation resulting later in the formation of tumors in mice liver. As we previously demonstrated that both benzo[a]pyrene (B[a]P), environmental agent found in cigarette smoke, and ethanol alone possessed similar targets, especially oxidative stress, to trigger death of liver cells, we presently decided to study the cellular and molecular mechanisms of the effect of B[a]P/ethanol co-exposure on cell death. After an 18-hours incubation time with 100 nM B[a]P, primary rat hepatocytes were supplemented with 50 mM ethanol for 5 or 8 hours. B[a]P/ethanol co-exposure led to a greater apoptotic cell death that could be linked to an increase in lipid peroxidation. Plasma membrane remodeling, as depicted by a membrane fluidity elevation and physicochemical alterations of lipid rafts, appeared to play a key role, since both toxicants acted with specific complementary effects. Membrane remodeling was shown to induce an accumulation of lysosomes leading to an important increase in low molecular weight iron cellular content. Finally, ethanol metabolism, but not B[a]P, by providing ROS, induced the ultimate toxic process. Indeed, in lysosomes, ethanol promoted Fenton reaction, lipid peroxidation and membrane permeabilization thereby triggering cell death. To conclude, B[a]P exposure, by depleting cholesterol membrane content of hepatocytes, would constitute a favorable ground for a later toxic aggression such as ethanol intoxication. Membrane stabilization of both plasma membrane and lysosomes might be a potential target for further investigation considering cytoprotective strategies.

Key words:
Benzo[a]pyrene, ethanol, liver, primary rat hepatocytes, oxidative stress, iron, reactive oxygen species, lysosome, membrane fluidity, lipid rafts.
Introduction

Polycyclic aromatic hydrocarbons (PAHs) are widespread environmental agents formed during the incomplete combustion or pyrolysis of organic matter. Thus, they are found in ambient air, cigarette smoke, grilled and smoked food, so that humans are continuously exposed to these compounds by the inhalation and oral route. Among PAHs, benzo[a]pyrene (B[a]P) is the most thoroughly studied because of its ubiquitous presence in PAH mixtures in the environment and in food (1-3). Because of the strong evidence of its carcinogenicity in animals and of its genotoxicity in exposed humans, B[a]P is now recognized as a human carcinogen from group 1 by the International Agency for Research on Cancer (1). Focusing more on liver cancer, meta-analysis of epidemiologic studies indicated that tobacco smoking could be associated with hepatocellular carcinoma (4). In addition, subjects with elevated PAH metabolites-DNA adduct levels in the liver appear to exhibit a higher risk for hepatocellular carcinoma (5). Although the etiology of B[a]P-induced tumor initiation is very complex, three bulk mechanisms for this initiation can be put forward. First, although chemically inert, B[a]P undergoes metabolic activation to diol-epoxides within mammalian cells, which then bind covalently to cellular macromolecules, such as DNA (6,7). Second, during the metabolic process, B[a]P produces reactive oxygen species that can cause oxidative DNA damage (8). Both events can then start the mutagenic chain of events responsible for tumor initiation. Third, B[a]P can induce apoptotic death of liver cells (9,10). Considering this latter point, it has been recently described that recurrent aggression of the liver leading to chronic hepatocyte death can trigger excessive compensatory proliferation resulting later in the formation of tumors in mice liver (11-13). Consequently, thorough analyses of B[a]P-induced cell death will provide new insights in the pathogenic process for the tumor initiation in the liver. In this context, during the last decade, our team has been focusing its research on deciphering the mechanisms underlying the B[a]P-induced death of
liver cells, and has already demonstrated the involvement of both oxidative stress (14,15) and plasma membrane remodeling, especially changes in membrane fluidity (16) and in physicochemical characteristics of lipid rafts (17,18). Besides, we also found that membrane fluidity (19) and lipid rafts (20,21) were involved in ethanol-induced oxidative stress and apoptosis of primary rat hepatocytes. Interestingly, several epidemiologic studies have shown an interactive effect of heavy smoking and heavy drinking on the development of hepatocellular carcinoma (22,23); but the underlying molecular mechanisms still remain unknown, even if similar roles in hepatocellular carcinoma development have been pointed out for each exposure type (24). As we previously found that both B[a]P and ethanol alone possessed similar targets in liver cells, we presently decided to study the cellular and molecular mechanisms of the effect of the B[a]P/ethanol co-exposure on cell death by focusing on oxidative stress and membrane remodeling. In primary rat hepatocytes, B[a]P/ethanol co-exposure was found, for the first time in hepatocytes, to increase cell death when compared to toxicants alone. This could be related to oxidative stress and membrane remodeling, via a mechanism that involved lysosome permeabilization.
Materials and methods

Chemicals

Eagle minimum medium and medium 199 with Hanks’ salts were purchased from Gibco, Invitrogen (Cergy-Pontoise, France). Bovine serum albumin and fetal calf serum were purchased from Biowest (Nuaille, France). Liberase was from Roche Diagnostics (Meylan, France). Ethanol was obtained from Prolabo (Paris, France). Acridine orange, α-naphtoflavone (αNF), bafilomycin, benzo[a]pyrene, cholesterol oxidase (CHOX), cholesterol water soluble, N-Acetyl-Asp-Glu-Val-Asp-7-amido-4-methylcoumarin (Ac-DEVD AMC), diallylsulfide (DAS), 12-doxylstearic acid (12-DSA), Hoechst 33342, 4-methylpyrazole (4-MP), 2-nitro-4-carboxyphenyl-N,N-diphenylcarbamate (NCDC), pepstatin A, thiazolyl tetrazolium blue bromide (MTT), thiourea and α-tocopherol (vitamin E) were provided from Sigma-Aldrich. Deferiprone was from Acros (Fischer, Illkirch, France). Leucine-leucyl-O-methyl-HBr (LLoMe) was purchased from Bachem (Bubendorf, Switzerland). Ursodeoxycholate sodium salt (UDCA) and n-benzyloxy carbonyl-Val-Ala-Asp(O-Me) fluoromethyl ketone (Z-VAD) were purchased from Calbiochem (Millipore, Saint-Quentin Les Yvelines, France). 4,4-difluoro-5-(4-phenyl-1,3-butadienyl)-4-bora-3a,4a-diaza-s-indacene-3-undecanoic acid (BODIPY® 581/591 C11), dihydrofluorescein diacetate (H$_2$FDA) and cholera toxin subunit B (Recombinant), Alexa Fluor® 488 Conjugate were provided by Molecular Probes (Invitrogen, Cergy-Pontoise, France). CellTiter-Glo® Luminescent Cell Viability Assay was from Promega (Charbonnières, France). LDH-cytotoxicity assay kit II was provided by Abcam (Paris, France). InfinityTM cholesterol reagent was purchased from Thermo Scientific (Thermo Fisher Scientific, Courtaboeuf, France). Anti-mouse immunoglobulin G (IgG) was from DAKO (Trappes, France). Anti-
phospholipase C\(\gamma\)1 antibody was purchased from Upstate (Millipore, Saint-Quentin–Les Yvelines, France). Flotillin-1 was obtained from BD Biosciences (Le Pont de Claix, France).

**Cell isolation and culture**

Adult rat hepatocytes were isolated from 2-month-old Sprague-Dawley animals by perfusion of the liver as previously described, except that a 17 \(\mu\)g/ml liberase solution was used for dissociation (25). Routinely, cell viability was above 80%.

Cells were seeded at densities of \(30 \times 10^3\) cells/well in 96-well Falcon microplates (MTT assay, ATP content, LDH release, ROS and lipid peroxidation evaluation), \(1 \times 10^6\) cells in 35-mm Petri dish (Hoechst 33342 staining), \(3 \times 10^6\) cells in 60-mm Petri dish (caspase 3/7 activity, LDH release), \(10 \times 10^6\) cells in 100 mm Petri dish (membrane fluidity determination), and \(20 \times 10^6\) cells in 150-mm Petri dish (low molecular weight iron measurement and lipid raft isolation); they were then cultured in a medium composed of 75% minimum Eagle’s medium and 25% medium 199 with Hanks’ salts, supplemented with 10% fetal calf serum and containing 50 \(\mu\)g streptomycin, 5 \(\mu\)g penicillin, 5 \(\mu\)g bovine insulin, 1 mg bovine serum albumin, and 2.2 mg NaHCO\(_3\) per milliliter. The cells were kept at 37°C in an atmosphere of 5% CO\(_2\) and 95% air. The medium was changed 3h after seeding and substituted with the same medium as above but deprived of serum. Benzo[a]pyrene was added to the medium at a final concentration of 100 nM to rat hepatocytes. After an 18-hours incubation time at 37°C, some cultures were then supplemented with 50 mM ethanol for 5 or 8 hours, and untreated cultures (supplemented with 0.005 % DMSO) were used as controls. These incubation times were chosen because we previously demonstrated a maximal cytotoxicity at 24 hours for B[a]P (26), and 5 hours for ethanol (19).

**Cytotoxicity determination**
**Cell viability**

Metabolically active viable hepatocytes were evaluated both by the mitochondrial succinate dehydrogenase activity using a MTT assay and by the ATP content. Succinate dehydrogenase activity and ATP content were considered to be directly proportional to the number of cells in the culture.

**MTT assay.** MTT is a tetrazolium salt that is reduced to purple formazan crystals mainly by mitochondrial succinate dehydrogenase. Briefly, after discarding culture medium, hepatocytes were incubated for two hours with 0.5 mg/mL MTT. After MTT removal, formazan crystals were then solubilized in DMSO. Using a microplate reader (SPECTROSTAR Omega, BMG Labtech), absorbance was directly measured at 560 nm and 630 nm reference. For each sample, values from the 560 nm absorbance was subtracted from 630 nm to negate the background effect. Cell treatment with drug vehicle was considered 100% of viability.

**ATP content.** ATP content was evaluated using a luminescent assay kit based upon the reaction of luciferin with ATP in presence of luciferase (CellTiter-Glo® luminescent Cell Viability Assay, Promega), according to manufacturer instructions. The amount of ATP was proportional to the luminescent signal measured with a spectrophotometer (SpectraMax Gemini; Molecular Devices, France).

**Cell death**

Cell death was tested for both apoptotic and necrotic cell death.

**Visualization of chromatin condensation and fragmentation by nuclear staining.** Chromatin condensation and morphological changes in the nucleus were observed using the chromatin dye Hoechst 33342. After treatments, cells were stained with 50 μg/ml Hoechst 33342 in the dark for 30 min at 37 °C. Fluorescence was stabilized using FluorSave™ Reagent (Merck
Millipore). Cells were then examined under fluorescence microscopy (Olympus BX60, France). Total population was always more than 400 cells.

**Measurement of caspase-3/7 activity.** To determine caspases-3/7 activity, hepatocytes were lysed in the caspase activity buffer (27). Eighty micrograms of crude cell lysate were incubated with 80 μM DEVD-AMC for 2 h at 37 °C. Caspase-mediated cleavage of DEVD-AMC was measured by spectrofluorimetry (Spectramax Gemini; Molecular Devices) using 380 nm excitation and 440 nm emission wavelengths.

**LDH leakage.** Necrotic cell death is characterized by plasma membrane damage, thereby leading to the release of intracellular enzymes to the extracellular medium. Lactate dehydrogenase (LDH) leakage was thus measured by using the LDH-Cytotoxicity Assay Kit II (Abcam) based upon an enzymatic coupling reaction: LDH by oxidizing lactate produces NADH, which then reacts with WST to generate yellow color. Briefly, after adding manufacturer reactant, yellow color was quantified at 450 nm for 45 minutes at 37 °C by spectrophotometer (SPECTROSTAR Omega, BMG Labtech), both in culture medium and in cells. The background value at 650 nm was subtracted from the values at 450 nm for each sample, in order to negate the effect of cell debris and precipitated proteins. LDH leakage was then quantified by calculating the ratio of extracellular LDH/total LDH.

**Ultrastructural changes**

*Transmission electron microscopy (TEM).* Following drug exposure, the cells were rinsed with 0.15 M Na cacodylate buffer and fixed by drop wise addition of glutaraldehyde (2.5%) for 1 hour. After fixation, the specimens were rinsed several times with 0.15 M Na cacodylate buffer and post fixed with 1.5% osmium tetroxide for 1 hour. After further rinsing with cacodylate buffer, the samples were dehydrated through a series of graded ethanol from 70 to 100%. The specimens were infiltrated in a mixture of acetone-Eponate (50/50) for 3 h, then in pure Eponate for 16 h. Finally, the specimens were embedded in DMP30-Eponate for 24 h
at 60 °C. Sections (0.5 μm) were cut on a LEICA UC7 microtome and stained with toluidine blue. Ultra-thin sections (90 nm) were obtained, collected onto copper grids and counterstained with 4% uranyl acetate then with lead citrate. Examination was performed with JEOL 1400 transmission electron microscope operated at 120 kV.

Lysosomal membrane permeabilization. Lysosome disruption was estimated from acridine orange-stained hepatocytes knowing that an increase in green acridine orange fluorescence reflected such a disruption. Briefly, after 18 hours of B[a]P treatment, supernatant was removed apart and cultures were incubated with acridine orange (5 μM) for 15 minutes at 37°C. After a short wash, cells were treated with ethanol for 1 or 5 hours, and then analyzed by flow cytometry on a FACS calibur flow cytometer (Becton Dickinson, San Jose, USA). Fluorescence emission of acridine orange was detected at FL1 (Green acridine orange fluorescence) following excitation at 488 nm. Each measurement was conducted on 50 000 events and analyzed on Cell Quest software (Becton Dickinson).

Oxidative stress evaluation

Determination of reactive oxygen species production

Intracellular levels of reactive oxygen species (ROS) were measured using the nonfluorescent probe, dihydrofluorescein diacetate (H₂FDA), as previously described (19). Oxidation by ROS causes a conversion of this probe into fluorescent fluorescein. Fluorescence was directly recorded by a SpectraMax Gemini spectrofluorimeter (Molecular Devices, France) using 485 nm excitation and 530 nm emission wavelengths. Even though several limitations have been described for this probe such as a lack of selectivity for ROS detection, a possible oxidation by reactive nitrogen species or proteins without ROS production and a light dependent-autooxidation (28), we decided to use it because of its simplicity and the low cell quantity needed.

Evaluation of lipid peroxidation
As previously described (19), lipid peroxidation was measured using the fluoroprobe C11-BODIPY\textsuperscript{581/591} that incorporates into cell membranes and is sensitive to oxidation. C11-BODIPY\textsuperscript{581/591} shifts from red to green fluorescence upon oxidation. Briefly, fluorescence was recorded by a SpectraMax Gemini spectrofluorimeter (Molecular Devices) using two pairs of wavelengths to measure the amount of reduced (\(\lambda\text{ex } 590 \text{ nm}, \lambda\text{em } 635 \text{ nm}\)) and oxidized probe (\(\lambda\text{ex } 485 \text{ nm}, \lambda\text{em } 535 \text{ nm}\)). Lipid peroxidation was then quantified by calculating the ratio of oxidized probe/total probe.

**Measurement of low molecular weight iron**
Measurement of intracellular low molecular weight (LMW) iron was based upon the capacity of deferiprone to chelate only LMW iron and to give a paramagnetic chelate, which can be directly detectable by EPR in whole hepatocytes (29).

**Membrane remodeling characterization**
Membrane remodeling was estimated by the evaluation of bulk membrane fluidity and both physical and chemical alterations of lipid rafts.

**Membrane fluidity**
Membrane fluidity was measured, as previously described (19), in bulk membranes (i.e., plasma membrane and possibly endosome and lysosome membranes) of hepatocytes. Membranes were spin labeled by incubating hepatocyte suspensions at 37°C for 15 minutes with 12-doxyl stearic acid (50 \(\mu\)g/ml). The EPR spectra of labeled samples were acquired at ambient temperature on a Brucker Elexsys EPR spectrometer operating at 3509 G center field, 20 mW microwave power, 9.86 GHz microwave frequency, 1.77G modulation amplitude and 100 kHz modulation frequency. The fluidity of the labeled membrane was quantified by calculating the order parameter S, which is inversely related to membrane fluidity.
**Lipid raft fractionation by flotation method**

Lipid rafts are defined as being resistant to solubilization by nonionic detergents at low temperature and able to float and concentrate in low-density fractions of a sucrose gradient after ultracentrifugation, as previously described (20). Briefly, lysates were centrifuged at 39,000 rpm for 20 hours at 4 °C in a SW41 rotor, allowing eleven 1-ml fractions to be collected from the top of the gradient for protein and lipid analysis. The 1 to 6 fractions contained lipid rafts, as previously characterized (20). In some experiments, the whole raft fractions (fractions 1 to 6) were gathered, diluted in PBS and pelleted by another ultracentrifugation at 41000 rpm for 16-20 hours at 4°C in a SW41 rotor. This last procedure allowed us to pool lipid rafts in a single pellet.

**Protein assays in lipid raft fractions**

Flotillin-1 and phospholipase Cγ1 (PLCγ1) were determined using SDS polyacrylamide gel electrophoresis and western blot analysis. For each gradient fraction, 2 to 5 μg, for flotillin and PLCγ1 respectively, were separated by a 10 % SDS-polyacrylamide gel electrophoresis. Gel proteins were transferred overnight onto nitrocellulose membranes (Hyblond ECL, Amersham Biosciences Europe, Orsay, France). The membranes were then blocked for 2 hours with Tris-buffered saline (TBS) containing 2 % BSA and 0.1 % Tween 20, at room temperature. For flotillin-1 and PLC-γ1 detection, anti-flotillin-1 and anti-PLC-γ1 at a 1:8000 dilution and 1:1000, respectively, were added first, and the mixture was incubated 2 hours at room temperature. The membrane was then washed three times with 0.1% Tween 20 in PBS for 30 minutes and incubated in blocking buffer containing conjugated anti-mouse immunoglobulin G secondary antibody at a 1:2000 for 45 minutes. It was then washed three times with 0.1% Tween 20 in PBS for 15 minutes. Blots were examined using a solution of Tris-HCl 0.1 M containing coumaric acid (0.22 mM) and luminol (1.25 mM).
Alkaline phosphatase activity was determined by a colorimetric method using p-nitrophenylphosphate as a substrate, as previously described (20). Briefly, 10 microliters of each fraction were incubated, for 20 minutes at 37°C, with 90 μl PBS and 100 μl p-nitrophenylphosphate ready-to-use liquid reagent (Sigma-Aldrich, Saint Quentin Fallavier, France). The absorbance was measured at 405 nm using a microplate spectrophotometer (SPECTROSTAR Omega, BMG Labtech).

**Fluorescent staining of lipid rafts**

Lipid rafts were stained consecutively to the binding of the cholera toxin subunit B coupled to green-fluorescent Alexa Fluor 488 conjugate to the pentasaccharide chain of the lipid raft marker monosialotetrahexosyl ganglioside (GM1), using the Cholera Toxin Subunit B, Alexa Fluor® 488 Conjugate (Molecular Probes, Invitrogen) as previously described (20). Staining was analyzed by conventional fluorescence microscopy using Metavue software. When dual staining was performed, hepatocytes were incubated with both cholera toxin conjugate and the test primary monoclonal anti-phospholipase Cγ1 antibody (10 μg/ml) for two hours, followed by incubation with 1:2000 TRITC-conjugated anti-mouse immunoglobulin G secondary antibody for an additional two hours. Co-staining was examined by confocal fluorescence microscopy using a LEICA DM 5500 microscope with the Leica LAS AF software.

**Statistical analyses**

Data presented were acquired from a minimum of three independent experiments. They are expressed as means ± S.D. Statistical analyses were performed with a One-Way ANOVA with Newmann-Keuls post-test. All statistical analyses were performed with GraphPad Prism5 software (GraphPad Software, San Diego, CA, USA). The significance is shown as follows: *p < 0.05, **p < 0.01 and ***p < 0.001, treated versus untreated cells.
Results

*B[a]P/ethanol co-exposure increases cell death of primary rat hepatocytes*

Estimated by the MTT test, B[a]P/ethanol co-exposure was found to lead to a greater decrease in cell viability compared to either treatment alone (Figure 1A). This was confirmed by the decrease in ATP, that was found to be more marked with co-exposure compared to ethanol or B[a]P alone (Figure 1B).

An apoptotic cell death could result from this hepatocyte toxicity of B[a]P/ethanol co-exposure. Indeed, the 5-hours-treatment led only to a slight increase in cell number with condensed or fragmented nuclei (Figure 1C), but with neither significant effect of the coexposure compared to toxicant alone nor elevation of caspase 3/7 activity for any treatment (Figure 1D). By contrast, an 8-hours-co-treatment led to a further elevation of cell number with condensed or fragmented nuclei not only for either toxicant alone (ethanol : + 81 % at 8 hours instead of + 62 % at 5 hours; B[a]P : + 56 % at 8 hours instead of + 36 % at 5 hours), associated with a significant increase in caspase activity (Figure 1C), but also for the co-exposure compared to either treatment alone, with a significant increase in apoptotic cell number and caspase 3/7 activity (Figure 1C and 1D).

Note that there is a strong increase in apoptotic cell death between 5 hours and 8 hours in control cultures. This is due to the primary culture senescence. Indeed, it is well known that many antioxidant enzymes, such as Cu,Zn superoxide dismutase, glutathione peroxidase, or catalase, are less expressed over incubation time after isolation of rat hepatocytes (30), thereby increasing ROS production over incubation time (19). Although the increase in caspase activity remained moderate (+ 36 %) at 8 hours, the pan caspase inhibitor Z-VAD totally inhibited the increase in the apoptotic cell number (figure 1E), thus pointing to a caspase-dependent apoptotic cell death. In addition, data in Figure 1F clearly demonstrated the absence of necrosis whatever the
condition tested, since no significant increase in the extracellular LDH/total LDH ratio was detected compared to control conditions.

The possible involvement of B[a]P and ethanol metabolism in the observed decrease in cell viability was tested using α-naphthoflavone (to inhibit B[a]P metabolism via cytochromes P450 1), 4-methylpyrazole (to inhibit ethanol metabolism via alcohol dehydrogenase and cytochrome P4502E1) and diallylsulfide to inhibit cytochrome P4502E1. These compounds were found to prevent the co-exposure-induced decrease in cell viability (estimated by the MTT test) (Figure 1S). Altogether, these results therefore indicate an increase in cell death level when co-exposing hepatocytes to both B[a]P and ethanol, which involves metabolism of both toxicants.

**The toxicity of B[a]P/ethanol co-exposure involves oxidative stress**

As oxidative stress plays an important role in B[a]P or ethanol toxicity alone (15,20), we then decided to evaluate its involvement in the effects of co-exposure. Both vitamin E (a free radical chain-breaking antioxidant) and thiourea (a ROS scavenger), prevented the decrease in cell viability and in ATP content induced by co-exposure (Figures 2A and 2B). In addition, co-exposing hepatocytes to B[a]P and ethanol significantly enhanced lipid peroxidation compared to the toxicants alone (Figure 2C). However, no further increase in production of ROS, prooxidant factors, was observed upon co-exposure (Figure 2D). Whereas, as expected, ethanol was found to induce a significant elevation of ROS production, no such a rise was detected with B[a]P alone under our experimental conditions.

**LMW iron participates to the toxicity of B[a]P/ethanol co-exposure**

The lack of enhancement of ROS production by the co-exposure, despite the increased oxidative damage, led us to study another pro-oxidant factor, namely Low Molecular Weight
iron (LMW) iron. In addition, alterations in iron homeostasis have been already involved in the toxicity of both B[a]P and ethanol in liver cells (15,19). Thus, LMW iron (that is, iron species not contained in high-molecular-weight molecules, such as ferritin or mitochondrial ferroproteins), can trigger oxidative stress not only by catalyzing the formation of highly reactive hydroxyl radicals but also more directly by leading to the formation of the oxidant ferryl and perferryl species (31). Lipid peroxidation and the drop in cell viability were related to LMW iron since deferiprone (a known specific LMW iron chelator) inhibited these effects due to co-exposure as well as those due to toxicants alone (Figure 3A-C). Finally, whereas ethanol and B[a]P alone increased LMW iron pool, as measured by EPR, co-exposing hepatocytes to both molecules led to a further significant increase (Figure 3D), thereby corroborating the important role of iron in the co-exposure-induced oxidative stress and cell death.

**Role for lysosomal membrane permeabilization**

Our previous data regarding the toxicity of B[a]P or ethanol alone have clearly evidenced a role for lysosomes, notably through their important contribution to intracellular LMW iron content (21,32). Indeed, LMW iron could be delivered to these organelles by autophagic degradation of ferroproteins, such as ferritin or mitochondrial electron transport complexes. Using electron microscopy, co-exposure appeared to induce an increase in number of vesicles with degraded mitochondria, when compared to toxicants alone (See Supplementary Material, Figure 2S). An increase in lysosomes was also observed. Bafilomycin, an inhibitor of the lysosomal pathway, blocked the cytotoxicity (Figures 4A and 4B), caspase activity (Figure 4C) and lipid peroxidation (Figure 4D) induced by B[a]P/ethanol co-exposure. In addition, by using flow cytometry, an increase in permeabilization of lysosome membranes was evidenced upon co-exposure compared to toxicants alone, notably following a 5h co-
treatment (Figure 4E). Pepstatin A, an inhibitor of cathepsin D, protease likely to be released from permeabilized lysosomes, prevented the co-exposure toxicity as well as that of toxicants alone (Figures 4A and 4B). Interestingly, it was also able to inhibit caspase activity (Figure 4C). Once again, relationship between LMW iron, oxidative stress and lysosomes could be established, since deferiprone and vitamin E inhibited the lysosomal membrane permeabilization (LMP) induced by the co-exposure (See Supplementary Material, Figure 3S).

**Key role of membrane remodeling**

Membrane remodeling, characterized by both an increase in fluidity and lipid raft alterations, is known to play a key role in the cell death induced by B[a]P (15-17) and ethanol alone (19-21). A membrane stabilizer (UDCA) or a lipid raft disrupter (CHOX) inhibited the effect of co-exposure on cell viability (Figures 5A and 5B), lipid peroxidation (Figure 5C), LMW iron content (Figures 5D), and LMP (See Supplementary Material, Figure 2S). As we had demonstrated the involvement of LMW iron, oxidative stress and LMP in the toxicity of the co-exposure, these data therefore pointed to the key role of membrane remodeling in the toxic effects of co-exposure. The next set of experiments was carried out in order to gain further insight into the possible membrane remodeling upon co-exposure. Thus, co-exposure induced a further increase in bulk membrane fluidity compared to toxicants alone (Figure 6A), and lipid raft chemical alterations (Figures 6B and 6C). In addition, co-exposure decreased the raft cholesterol content compared to controls (Figure 6B); however no significant difference was observed relatively to B[a]P alone, suggesting the main role for B[a]P in this co-exposure effect. Nonetheless, a significant further decrease in the activity of alkaline phosphatase was observed as compared to B[a]P or ethanol alone (Figure 6C). Note that all treatments did not markedly change the flotillin-1 (a protein raft marker) distribution in
membranes (Figure 6D). In addition, raft staining of the raft-associated GM1 glycosphingolipid showed that, as expected, ethanol alone led to raft aggregation (as evidenced by punctuated staining), whereas a raft destabilization was observed upon B[a]P treatment (as visualized by diffuse staining). Regarding B[a]P/ethanol co-exposure, both types of alterations were observed, but at a lesser extent for raft aggregation (Figure 6E).

**Involvement of PLCγ1 relocation to lipid rafts**

As we previously showed for ethanol alone that lipid raft clustering promotes PLCγ1 relocation to lipid rafts, triggering its own activation and enhancing oxidative stress through lysosome alteration and increase in LMW iron (21), we decided to study the PLCγ1 relocation in the context of the co-exposure. It is worth stressing that the activity of alkaline phosphatase, a glycosylphosphatidylinositol (GPI)-anchored enzyme known to be a target of PLCγ1, was demonstrated to be decreased by the co-exposure when compared to toxicants alone (Figure 6E). As expected from our previous data concerning ethanol alone, merging of fluorescence staining led to the appearance of yellow patches indicating that GM1 and PLCγ1 co-localized (Figure 7A). More interestingly, the co-exposure maintained this translocation even though lipid rafts were partially destabilized. Note that B[a]P alone was not able to induce this translocation. By western blotting, PLCγ1 expression was found to slightly increase in lipid rafts when compared to ethanol alone, despite less raft aggregation (Figure 7B). In addition, NCDC, a specific PLCγ1 inhibitor, protected from both toxicity (Figures 7C and 7D) and LMP (See Supplementary Material, Figure 3S). It is noteworthy that this inhibitor did not protect from B[a]P toxicity (Figures 7C and 7D), unlike ethanol toxicity.

As membrane remodeling has been shown to be a major actor in the co-exposure toxicity, we decided to study its involvement, more specifically the raft cholesterol depletion, in the co-
exposure-increased PLCγ1 relocation into lipid rafts. Thus, in B[a]P/ethanol-treated hepatocytes, cholesterol supply protected from lipid raft destabilization and PLCγ1 relocation (Figure 7E). Note that, in this set of experiments, confocal fluorescence microscopy was used explaining that raft destabilization appeared as weaker fluorescence of membranes instead of diffuse fluorescence as observed with conventional fluorescence microscopy (Figure 6E). In addition, in cells treated with ethanol alone, neither lipid raft clustering nor PLCγ1 translocation into lipid rafts were altered whereas B[a]P alone-induced lipid raft destabilization was inhibited (Figure 7E). Therefore, relocation of PLCγ1 to lipid rafts was dependent on cholesterol depletion only for the co-exposure. It is worth remembering that B[a]P plays the main role in the co-exposure induced cholesterol depletion. Finally, cholesterol protected from this toxicity and also from that of B[a]P alone, but was ineffective on ethanol toxicity (Figures 7C and 7D); such differential effects for the cholesterol supply were also observed when considering LMP (See Supplementary Material, Figure 3S).
Discussion

In this paper, for the first time, ethanol was shown to act reciprocally with B[a]P in promoting hepatocyte oxidative stress and cell death through a novel mechanism of interaction between both toxicants, never described before, that involved plasma membrane remodeling and lysosomal membrane permeabilization. Indeed, both toxicants differentially acted on plasma membrane to ultimately lead to an increase in membrane fluidity and lipid raft alterations. By using a membrane stabilizer or a lipid raft disrupter, these effects were shown to be implicated in oxidative stress and cell death. First, as previously shown (16,19), both toxicants alone increased membrane fluidity, very likely through different mechanisms, which resulted in a larger membrane fluidization for the co-exposure. Second, they can also alter lipid rafts of the plasma membrane; these microdomains are detergent-resistant and rich in sphingolipids and cholesterol, which confer them a highly ordered spatial structure. The two toxicants opposingly altered lipid rafts, as previously reported (17,20). Thus, whereas B[a]P modified raft organization by decreasing their cholesterol content, ethanol induced raft clustering; as a result, co-exposure triggered raft clustering but at a lesser extent. Previously, we (20,21) and others (33) clearly demonstrated the involvement of lipid raft clustering in ROS production by activated various redox signaling pathways. Regarding ethanol, a phospholipase C-dependent pathway was described (20). Indeed, ethanol-induced lipid raft clustering was found to promote PLCγ1 translocation and subsequent activation leading to oxidative stress in primary rat hepatocytes (20). Upon co-exposure, to ethanol-induced raft clustering was associated the B[a]P effect i.e. the drop in raft cholesterol content; compared to toxicants alone, this latter effect would ultimately enhance PLCγ1 translocation into lipid rafts, thus explaining the increase in oxidative stress. Indeed, the decrease in raft cholesterol content, by fluidizing lipid rafts, may promote PLCγ1 raft insertion; regarding this point, it is noteworthy that PLCγ1 activity has been shown to increase with membrane fluidization (34).
Thus, cholesterol could appear as an attractive target since cholesterol supplementation was able to partially protect from cell death induced by B[a]P alone and B[a]P/ethanol co-exposure. Note that cholesterol increased ethanol alone toxicity likely by increasing raft clustering.

Previously, we showed that PLC\(_{\gamma}1\) participated to ethanol-induced oxidative stress and cell death by promoting lysosome accumulation, a major source of a known pro-oxidant, the LMW iron (21). Indeed, lysosomes can accumulate large amounts of LMW iron by degradation of macromolecules containing iron such as ferritin and mitochondrial electron transport complexes (35). Compared to toxicants alone, B[a]P/ethanol co-exposition was able to increase lysosome accumulation and LMW iron content, that could be linked to the elevation of PLC\(_{\gamma}1\) activity. Once again, a new mechanism involving a toxic cooperation between both compounds leading to lysosomal membrane permeabilization could explain the enhancement of lipid peroxidation and cell death due to the co-exposure. Indeed, in primary rat hepatocytes, ethanol, but not B[a]P, induced ROS production whereas both toxicants were found to participate to the elevation of LMW iron. Considering these differential effects on ROS production, one might propose the following scenario: in lysosomes, H\(_2\)O\(_2\) provided by ethanol metabolism would allow the formation of hydroxyl radical by a reaction catalyzed by LMW iron, leading to lipid peroxidation of lysosome membranes (35). As both toxicants work together to increase the LMW iron pool, lipid peroxidation would then be enhanced, thereby further destabilizing lysosomal membranes (Figure 8). In support to this, we found an enhancement of LMP upon co-exposure compared to toxicants alone, and pretreatment with vitamin E, a lipid peroxidation inhibitor, was shown to protect from it. Such a similar oxidative process has been reported in hepatocytes stressed by cytokines (36), free fatty acids (37) or hyperthermia (38); however our study is the first one to show that such an interaction between both toxicants can induce LMP through plasma membrane remodeling. Considering
this point, the depletion in membrane cholesterol due to B[a]P (previously reported to be due to inhibition of HMG CoA reductase, the first enzyme in cholesterol synthesis (17)), would also participate to lysosomal disruption. Indeed, recent studies described that cholesterol accumulation in lysosomes could protect from oxidant-induced LMP (39), whereas cholesterol depletion decreased lysosome stability (40). In our model, repletion in cholesterol protected from the co-exposure-induced LMP and cell death. In addition, a role for aryl hydrocarbon receptor (AhR), a B[a]P-activated transcription factor, in lysosomal disruption has also been established in hepatic epithelial cells (41). Although speculative, this could be linked to the cholesterol depletion since we (17) and others (42) demonstrated that AhR activation led to a decreased expression of several enzymes involved in cholesterol synthesis. Besides, it should also be noted that ethanol was reported to be able to activate AhR in mouse hepatic stellate cells (43). Taken altogether, our results showed that ethanol and B[a]P through mechanisms specific to each toxicant, i.e. cholesterol depletion for B[a]P and ROS production for ethanol, led to an important LMP. Accumulated evidence suggested that this process can lead to apoptotic or necrotic cell death by the translocation of lysosomal contents to the cytoplasm depending on the intensity of lysosome disruption (35,44). In our model, after 5 hours of ethanol incubation, lysosome disruption remains moderate (+ 29 %) explaining that no necrosis could be detected.

Caspase activity appeared to be crucial in the apoptotic process since pan caspase inhibitor totally inhibited the increase in apoptotic cell number. Interestingly, the caspase activation appeared to rely upon the release of cathepsins by LMP. Many papers reported the ability of cathepsins to directly activate caspases, or indirectly by promoting mitochondrial outer membrane permeabilization (44-46). This latter possibility seems to be the most likely in the present study since the delay for the co-exposure to induce apoptosis increase (8 hours). Indeed, even though these tests are not the best, mitochondrial alteration analyzed by MTT
test and ATP depletion was detected at 5 hours, that is before the occurrence of apoptosis; furthermore, pepstatin A, a cathepsin D inhibitor, protected from this mitochondrial damage suggesting the harmful effect of released cathepsin D on mitochondria.

Taken altogether, for the first time, it was demonstrated in this paper, that both toxicant-induced opposite effects on plasma membrane especially lipid rafts can finally interact to increase a toxic raft-dependent cell signaling pathway (phospholipase C-dependent), that, in addition, was not activated by one of the toxicants alone (namely B[a]P). Consequences of this plasma membrane interaction are important since several cytoprotective compounds (NCDC or cholesterol) identified for each toxicant alone could be used to target the coexposure toxicity, with a complete afforded protection, whereas they were ineffective for one or the other toxicants when applied alone. In conclusion, exposure to B[a]P, by depleting cholesterol membrane content of hepatocytes, would constitute a favorable ground for a later toxic aggression such as ethanol intoxication. By a cooperative remodeling of plasma membrane leading to lysosome accumulation (Figure 8), B[a]P and ethanol would act together to trigger lysosomal membrane permeabilization, cathepsin release and caspase activation. In this context, membrane stabilization of both plasma membrane and lysosomes might be a potential target for further investigation considering cytoprotective strategies.
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List of Abbreviations:
B[a]P, benzo[a]pyrene ; CHOX, cholesterol oxidase ; 12-DSA, 12-doxyloystearic acid ; EPR, electron paramagnetic resonance ; GM1, monosialotetrahexosyl ganglioside ; HMG CoA reductase, hydroxymethylglutaryl-CoA reductase; LDH, lactate dehydrogenase ; LMP, lysosomal membrane permeabilization ; LMW iron, low-molecular-weight iron ; MTT, thiazolyl blue tetrazolium blue; NCDC, 2-nitro-4-carboxyphenyl-N,N-diphenylcarbamate ; PAH, polycyclic aromatic hydrocarbon ; PI-PLC, phosphatidylinositol-specific phospholipase C ; ROS, reactive oxygen species ; TEM, transmission electron microscopy ; UDCA, ursodeoxycholate sodium salt ; Z-VAD, N-Benzoxycarbonyl-Val-Ala-Asp(O-Me) fluoromethyl ketone.
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Figure 1. Ethanol increased benzo[a]pyrene (B[a]P) cytotoxicity through an elevation of apoptotic cell death. Metabolically active viable hepatocytes were determined both by (A) the mitochondrial succinate dehydrogenase activity using a MTT assay and by (B) the ATP content using a luminescent assay. Apoptosis was evaluated by (C) staining nuclear chromatin condensation with Hoechst 33342 (D) analysing DEVDase activities of caspase 3/7 by spectrofluorimetry and (E) the effect of a pan caspase inhibitor (Z-VAD) on the...
number of cells with condensed and fragmented nuclei. (F) Necrosis was estimated by calculating the ratio of LDHe (extracellular lactate dehydrogenase) over LDHtot (total LDH = LDHe + intracellular LDH). Primary rat hepatocytes were treated or not for 18 hours with 100 nM B[a]P before 50 mM ethanol addition (or not) for a further 5- or 8-hours incubation time. Some cultures were pretreated, for one hour, with 10 µM ZVAD, a pan caspase inhibitor. In this case, ethanol was added for 8 hours. Values are given as means ± standard deviation of at least seven independent experiments. Toxicant treated versus untreated cultures: *p<0.05 ; ** p<0.01 ; ***p<0.001. Z-VAD pretreated versus untreated cultures : #p<0.05 ; ## p<0.01 ; ###p<0.001.

Figure 2. The increase by ethanol of benzo(a)pyrene (B[a]P) cytotoxicity could be related to oxidative stress. Metabolically active viable hepatocytes were evaluated both by (A) the mitochondrial succinate dehydrogenase activity using a MTT assay and by (B) the ATP content using a luminescent assay. (C) The increase in lipid peroxidation was measured by the relative fluorescence of the probe C11-Bodipy 581/591. (D) ROS production was
followed by the fluorescence of fluorescein. Primary rat hepatocytes were treated or not for 18 hours with 100 nM B[a]P before 50 mM ethanol addition (or not) for a further 5-hours incubation time. Some cultures were pretreated, for one hour, with 100 mM thiourea, a reactive oxygen species scavenger, or with 250 µM vitamin E, a free radical chain-breaking antioxidant. Values are given as means ± standard deviation of at least five independent experiments. Toxicant treated versus untreated cultures: ***p<0.001. Vitamin E or thiourea pretreated versus unpretreated cultures: #p<0.05; ##p<0.01; ###p<0.001.

Figure 3. Involvement of LMW iron in the B[a]P/ethanol co-exposure cytotoxicity and oxidative stress. (A) Lipid peroxidation was measured by the relative fluorescence of the probe C11-Bodipy 581/591. Metabolically active viable hepatocytes were evaluated both by (B) the mitochondrial succinate dehydrogenase activity using a MTT assay and by (C) the ATP content using a luminescent assay. (D) Intracellular low-molecular-weight (LMW) iron content was determined by electron paramagnetic resonance analysis after chelation by 3 mM
deferiprone. Primary rat hepatocytes were treated or not for 18 hours with 100 nM B[a]P before 50 mM ethanol addition (or not) for a further 5-hours incubation time. Some cultures were pretreated with 30 µM deferiprone, a LMW iron chelator, for one hour. Values are given as means ± SD of at least three independent experiments. Toxicant treated versus untreated cultures: ** p<0.01 ; *** p<0.001. Deferiprone pretreated versus unpretreated cultures : #p<0.05 ; ## p<0.01 ; ###p<0.001.

Figure 4. Involvement of lysosomes in the B[a]P/ethanol co-exposure cytotoxicity and oxidative stress. Metabolically active viable hepatocytes were evaluated both by (A) the
mitochondrial succinate dehydrogenase activity using a MTT assay and by (B) the ATP content using a luminescent assay. (C) DEVDase activities of caspase 3/7 were analyzed by spectrofluorimetry. (D) Lipid peroxidation was measured by the relative fluorescence of the probe C11-Bodipy \( ^{581/591} \). (E) Lysosomal membrane permeabilization was analyzed by flow cytometry using the green fluorescence of acridine orange. Primary rat hepatocytes were treated or not for 18 hours with 100 nM B[a]P before 50 mM ethanol addition (or not) for a further 5-hours incubation time, except for caspase activity measurement for which ethanol was added for 8 hours. Some cultures were pretreated with 1 nM bafilomycin or 15 µM pepstatin A for one hour. Values are given as means ± standard deviation of at least three independent experiments except for bafilomycin experiments for which \( n \geq 4 \). Toxicant treated versus untreated cultures: * \( p<0.05 \); ** \( p<0.01 \); *** \( p<0.001 \). Bafilomycin or pepstatin pretreated versus unpretreated cultures: #\( p<0.05 \); ## \( p<0.01 \); ###\( p<0.001 \).

Figure 5. Membrane remodeling, characterized as membrane fluidization or lipid raft alteration, was involved in the B[a]P/ethanol co-exposure cytotoxicity and oxidative
Stress. Metabolically active viable hepatocytes were evaluated both by (A) the mitochondrial succinate dehydrogenase activity using a MTT assay and by (B) the ATP content using a luminescent assay. (C) Lipid peroxidation was measured by the relative fluorescence of the probe C₁₁-Bodipy₅₈₁/₅₉₁. (D) Intracellular low-molecular-weight iron content was determined by electron paramagnetic resonance analysis after chelation by 3 mM deferiprone. Primary rat hepatocytes were treated or not for 18 hours with 100 nM B(a)P before 50 mM ethanol addition (or not) for a further 5-hours incubation time. Some cultures were pretreated for one hour with 100 µM ursodeoxycholic acid (UDCA), a membrane stabilizer, or with 0.05 U/ml cholesterol oxidase (CHOX), a lipid raft disruptor. Values are given as means ± standard deviation of at least four independent experiments. Toxicant treated versus untreated cultures: ** p<0.01 ; *** p<0.001. UDCA or CHOX pretreated versus unpretreated cultures : # p<0.05 ; ## p<0.01 ; ### p<0.001.
Figure 6. Ethanol modified the B[a]P-induced membrane remodeling, depicted by an enhancement of membrane fluidity and new physicochemical alterations of lipid rafts. (A) The membrane fluidity was monitored by electron paramagnetic analysis of 12-doxylstearic acid embedded in membranes. This analysis resulted in the calculation of the order parameter which is inversely related to membrane fluidity. (B) Total amounts of
cholesterol and (C) alkaline phosphatase activities were determined in pooled fractions 1-6 (detergent-resistant raft fractions) and 7-11 (soluble non raft fractions). Lipid rafts were obtained after lysis of hepatocytes in ice-cold 1% Triton X-100 and fractionation on a sucrose density gradient by ultracentrifugation. (D) Flotillin-1 (protein raft marker) expression. An equal protein amount of each fraction was submitted to western blotting for analysis of flotillin-1. One representative of three independent experiments is shown. (E) Visualization of lipid rafts by conventional fluorescence microscopy using choleratoxin subunit B conjugated with Alexa Fluor 488 which binds the raft-associated glycosphingolipid GM1. One representative of three independent experiments is shown. Primary rat hepatocytes were treated or not for 18 hours with 100 nM B[a]P before 50 mM ethanol addition (or not) for a further 5-hours incubation time. Values are given as means ± SD of at least three independent experiments. Toxicant treated versus untreated cultures: * p<0.05; ** p<0.01; *** p<0.001.
Figure 7. The phospholipase Cγ1 translocation to lipid rafts induced by B[a]P/ethanol co-exposure via a cholesterol depletion enhanced cell death. (A, E) Rat hepatocytes were co-stained by Alexafluor cholera toxin subunit B (CTX ; green fluorescence, left panels) and
tetramethyl rhodamine isothiocyanate (TRITC)-conjugated antibody to phospholipase Cγ1 (antiPLCγ1; red fluorescence, middle panels). The merge of the images obtained by confocal fluorescence microscopy shows yellow areas (right panels) that represent colocalization of monosialotetrahexosylganglioside and PLCγ1. One representative of three independent experiments is shown. (B) Lipid rafts were obtained after lysis of hepatocytes in ice-cold 1% Triton X-100 and fractionation on a sucrose density gradient by ultracentrifugation. An equal protein amount of each fraction was submitted to western blotting for analysis of phospholipase Cγ1 expression, followed by band densitometry (below the blots). One representative of three independent experiments is shown. Metabolically active viable hepatocytes were evaluated both by (C) the mitochondrial succinate dehydrogenase activity using a MTT assay and by (D) the ATP content using a luminescent assay. Primary rat hepatocytes were treated or not for 18 hours with 100 nM B[a]P before 50 mM ethanol addition (or not) for a further 5-hours incubation time. Some cultures were pretreated for 30 minutes with 50 µM 2-nitro-4-carboxyphenyl-N,N-diphenylcarbamate (NCDC), a phospholipase Cγ1 inhibitor, or, for 1 hour, with 50 µg/ml cholesterol. Values are given as means ± SD of at least four independent experiments. Toxicant treated versus untreated cultures: ** p<0.01 ; *** p<0.001. NCDC or cholesterol pretreated versus unpretreated cultures: #p<0.05 ; ## p<0.01 ; ###p<0.001.
Figure 8. Proposed mechanism for the role of membrane remodeling and lysosome membrane permeabilization in the enhancement by ethanol of benzo[a]pyrene-induced oxidative stress and cell death. Both toxicants could induce plasma membrane remodeling, but with different complementary effects, that finally would promote an elevation of phospholipase C translocation into lipid rafts. Then, activation of phospholipase C would induce the accumulation of lysosomes and lead to an important elevation of LMW iron cellular content. Ethanol metabolism but not B[a]P would provide hydrogen peroxide to lysosomes, which, in turn, would trigger Fenton reaction and excessive hydroxyl radical production, thereby enhancing lysosomal lipid peroxidation and membrane permeabilization leading to the release of hydrolase capable of activating caspases. Thus, both toxicants may cooperate by similar and various mechanisms to induce an increase in lysosomal membrane permeabilization that could lead to cell death.