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Oxidative stress resulting from exposure of a human salivary gland cells to paraoxon: an *in vitro* model for organophosphate oral exposure

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Abstract

Organophosphate (OP) compounds are used as insecticides, acaricides, and chemical agents and share a common neurotoxic mechanism of action. The biochemical alterations leading to many of the deleterious effects have been studied in neuronal cell lines, however, non-neuronal toxic effects of OPs are far less well characterized *in vitro*, and specifically in cell lines representing oral routes of exposure. To address this void, the human salivary gland (HSG) cell line, representing likely interactions in the oral cavity, was exposed to the representative OP paraoxon (PX; O,O-diethyl-p-nitrophenoxy phosphate) over a range of concentrations (0.01 μM to 100 μM) and analyzed for cytotoxicity. PX induced cytotoxicity in HSG cells at most of the exposure concentrations as revealed by MTT assay, however, the release of LDH only occurred at the highest concentration of PX tested (100 μM) at 48 h. Slight increases in cellular ATP levels were measured in PX-exposed (10 μM) HSG cells at 24 h. Exposing HSG cells to 10 μM PX also led to an increase in DNA fragmentation prior to loss of cellular membrane integrity implicating reactive oxygen species (ROS) as a trigger of toxicity. The ROS genes *gss*, *gstm2*, *gstt2* and *sod2* were upregulated, and the presence of superoxide following 10 μM PX exposure was determined via dihydroethidium fluorescence studies further implicating PX-induced oxidative stress in HSG cells.

Keywords

paraoxon; HSG cells; oxidative stress; organophosphate; MTT; DNA fragmentation

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Conflict of interest statement

No conflict of interest exists with any authors or this institution.

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1. Introduction

Organophosphate compounds (OPs) are widely used to control insect pests but are also known as a chemical class used as warfare agents. Exposure to OPs can result in neurotoxicity (Ballantyne and Marrs, 1992; Casida and Quistad, 2004; Fukuto, 1990) that is typically attributed to the inhibition of acetylcholinesterase (AChE) (Scheme 1). In contrast to the warfare agents, OP insecticides must be converted from the less reactive thionate (P=S) into the oxon (P=O) form to inhibit AChE. When AChE is inhibited, an accumulation of acetylcholine (ACh) occurs resulting in an overstimulation of acetylcholine receptors (AChR's) (Ballantyne and Marrs, 1992; Broomfield et al., 1995). OPs have also been shown to interact with a number of cholinergic and non-cholinergic protein targets (Bomser and Casida, 2000; Bomser and Casida, 2001; Ehrich et al., 1994; Proskocil et al., 2010; Quistad et al., 2001; Richards et al., 1999; Schuh et al., 2002).

The formation of oxons occurs during *in vivo* metabolic oxidation but also during the manufacture, storage, or environmental lifetime, for example in chlorinated water (Kamel et al., 2009). Therefore, direct exposure to OP oxons may occur prior to *in vivo* metabolism. Skin absorption is the most common pathway to OP exposure, yet more OP poisonings occur when pesticides enter the mouth. Food and water contaminated with OPs, airborne OPs (spraying or foggers), and hand to mouth transfer in infants/children (Chambers et al., 2007) each contribute to oral and inhalation exposures of parent and oxon forms of the insecticide.

A number of studies have examined the *in vitro* interactions between OPs and cell lines. Altered signaling pathways in neuronal and glial cells (Garcia et al., 2001; Hong et al., 2003; Qian et al., 2007; Schuh et al., 2002) adverse effects on mitochondrial integrity and ATP production (Hong et al., 2003; Knoth-Anderson et al., 1992; Massicotte et al., 2005), and an increase in stress response (Garcia et al., 2001; Sachana et al., 2001) have been reported. In PC-12 cells, chlorpyrifos exposure induced apoptosis via mitochondrial damage (Lee et al., 2012).

Paraoxon (PX) is the oxidative metabolite of parathion (Scheme 1), an inhibitor of AChE, and has been used as a model OP compound to investigate and identify toxicologically relevant targets in cell lines of neuronal origin (Bharate et al., 2010; Casida and Quistad, 2005; Pope, 1999; Richards et al., 1999) including the SH-SY5Y cell line (Bharate et al., 2010; Carlson and Ehrich, 1999; Ehrich et al., 1997; Prins et al., 2010; Saleh et al., 2003), and non-neuronal HepG2 cells (Hreljac et al., 2008). PX inhibits AChE to form a diethoxyphosphorylated serine residue (Scheme 1) that is identical to the OP-AChE conjugates formed from the oxidative metabolites of diazinon, phorate, chlorpyrifos, etc. PX inhibits the AChE present in SH-SY5Y cells within minutes at sub-micromolar levels but the loss of cell viability occurs at millimolar levels (24–48 h) suggesting deleterious biochemical mechanisms that may be downstream effects of AChE inhibition or may result from modification of proteins other than AChE (Ehrich et al., 1997). Likewise, protein expression changes occur when SH-SY5Y cells were treated with micromolar PX (Prins et al., 2010). However, none of these studies address possible interactions of OP compounds in the oral cavity.

In order to elucidate possible non-neuronal effects resulting from OP oral exposure, the human salivary gland (HSG) cell line was selected. The HSG cell line was derived from isolated epithelial cells from the irradiated submandibular salivary gland of a squamous cell carcinoma. This cell line is characterized as an epithelial duct cell type and expresses fibrinolytic activity (Shirasuna et al., 1981). HSG cells have been used in cellular response studies of toll-like receptor-mediated immune responses (Kawakami et al., 2007), growth factor stimulation of signal transduction pathways (Crema et al., 2006), and ATP-dependent activation of potassium channels (Liu et al., 1999). In the event of a human exposure to foodstuffs or aerosols of OP insecticides, the salivary glands would be a likely target in the oral cavity. Therefore, the HSG cell line was selected an appropriate model to study non-neuronal OP toxicity responses. In this study, HSG cells were exposed to paraoxon (Scheme 1), and select cytotoxic assays were conducted and validated by gene expression changes to assess toxicity resulting from low-level exposure.

2. Material and methods

2.1. Reagents and chemicals

Ethyl paraoxon was purchased (catalog number PS-610; Chem Service, Inc. West Chester, PA).

2.2. Culture of human salivary gland cells

The human salivary gland (HSG) cell line was established from an irradiated human salivary gland (Shirasuna et al., 1981), and was generously provided by Dr. Bruce Baum (National Institute of Health/National Institute of Dental and Cranial Facial Research (NIH/NIDCR)). HSG cells were cultured using Dulbecco's Modified Eagle's medium and Ham's F-12 nutrient mixture (DMEM/F12) (GIBCO BRL, Grand Island, NY) supplemented with 5% fetal bovine serum (FBS) (Hyclone; Thermo Scientific, Rockford, IL), 100 U/ml penicillin, 100 µg/ml streptomycin, and 2 mM L-glutamine in a CO₂ incubator maintained at 5% CO₂ and 37 °C. The medium was changed every two days, and cells were allowed to reach 80% confluence before exposure to PX.

2.3 Acetylcholinesterase activity in HSG cells

Total cellular proteins were harvested from HSG cells by lysis (150 mM NaCl, 1% Triton X-100, 50 mM Tris pH 8.0) and used immediately for analysis. Ellman assays (Ellman et al., 1961) were based on published reports, and were performed by incubation of crude protein lysates with 5,5-dithiobis(2-nitrobenzoic acid) (final concentration, 0.32 mM in 0.1 M potassium phosphate buffer, pH 7.6) and initiated with acetylthiocholine iodide (ATCh-I) (final concentration, 0.75 mM) in a final volume of 200 µl. (George et al., 2003). All kinetic assays performed under these conditions were linear with respect to time and corrected for protein concentration (Redinbaugh and Turley, 1986; Zhang and Halling, 1990). Negative controls were lysates plus DTNB (no ATCh-I). Changes in absorbance were measured at OD 412 nm on a VersaMax microplate reader with Softmax Pro V. 3.0 software. The optimum lengths of time for incubations were determined empirically to allow maximum changes in absorbance for samples versus blanks. Recombinant mouse AChE (rMoAChE)

was used as a positive control for AChE activity. Ellman assays were performed in triplicate.

2.3. HSG cell exposure to PX

A 10 mM stock solution of PX was prepared in 100% ethanol and concentrations from 0.01 μ M to 100 μ M were prepared by dilution of the stock into DMEM/F12 medium (50:50), 1% FBS, 100 U/ml penicillin, 100 μ g/ml streptomycin, and 2 mM L-glutamine. Prior to PX treatment (24 h), the FBS in the HSG cell culture media was reduced from 5% to 1%. Culture media was replaced with PX culture media and allowed to incubate at 37°C for the various time points described below.

2.4. MTT and LDH cell viability assays

Effects of PX exposure on HSG cell viability was estimated using MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) and LDH (lactate dehydrogenase) release assays (Cell Proliferation Kit (MTT) and Cytotoxicity Detection Kit (LDH), Roche Applied Science, Indianapolis, IN). Cells were grown in 96-well culture plates to 80% confluence before exposure to PX. For both MTT and LDH assays, cells were exposed to PX concentrations (or ethanol alone as the vehicle control) ranging from 0.01 μ M to 100 μ M for 24, or 48 h with a 2% Triton X-100 solution in assay medium used as a positive control for cell death (all additions were at final v/v of 0.1%).

For the MTT assay, cells were rinsed several times with culture medium prior to PX exposure, the culture medium was removed and 100 μ l of fresh medium containing the various PX concentrations or Triton X-100 was added to each well (n = 6–8 for each PX concentration and Triton X-100). At various incubation time points, 10 μ l of the supplied MTT labeling reagent (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) was added to each well, incubated for 4 h, after which 100 μ l of solubilizing solution was added to each well. Plates were incubated overnight and then measured at 575 nm (formazan) using a VersaMax microplate reader with Softmax Pro V. 3.0 software. Cytotoxicity was determined by comparing the absorbance readings of the wells containing the PX-treated cells with those of the vehicle- (0.1% ethanol) treated cells.

For the LDH assay, cells were rinsed several times with culture medium prior to PX exposure. Culture medium was removed and 100 μ l of fresh medium containing the various PX concentrations was added to each well (n = 6–8 for each PX concentration and Triton X-100). At the appropriate time points, 100 μ l of supernatant was removed from each well and transferred into the corresponding wells of a new 96-well plate. LDH activity was determined by adding 100 μ l of reaction mixture (included with kit) to each well and incubating for 20 min in the dark at room temperature. Absorbance measurements were taken at 490 nm on a microplate reader with a reference wavelength at 650 nm. Cytotoxicity was determined by comparing the absorbance readings of the wells containing the PX-treated cells with those of the vehicle- (0.1% ethanol) treated cells.

2.5. Assay for cellular ATP levels

Cellular levels of ATP were measured using an ATP Bioluminescence Assay Kit (HS II, Roche Applied Science, Indianapolis, IN). Briefly, HSG cells were cultured in a 96-well plate to 80% confluence, exposed to 10 μ M PX or ethanol control for 8–24 h, and harvested and lysed. Total protein was quantitated by BCA assay. The level of cellular ATP was calculated relative to a standard wherein the bioluminescence was quantified on a Luminex 100. Samples were analyzed as $n = 4-6$.

2.6. DNA fragmentation assay

Cellular DNA fragmentation was measured using a Cellular DNA Fragmentation ELISA kit (Roche Applied Science, Indianapolis, IN). In this assay, 5'-bromo-2'-deoxy-uridine (BrdU)-labeled DNA fragments in cell lysates or cell supernatants indicate the cell death type (e.g., apoptosis or necrosis). HSG cells were cultured in 96-well culture plates to a density of 80% confluence in culture media containing 10 μ M BrdU-labeling solution. Cells were incubated overnight (~16 h), and BrdU media was replaced with culture media containing 10 μ M PX (or 0.1% ethanol) for 2–24 h ($n = 6-8$). BrdU-labeled DNA fragments released from the HSG cells were measured from 100 μ L samples of cell culture media supernatant at various time points. Samples for each time point were stored at -20°C prior to analysis. After removal of all the remaining supernatant, 200 μ L of cell lysis buffer was added to each 100 μ L sample well followed by a 30 min incubation to lyse the remaining cells. Lysate (100 μ L) was removed and placed in a new 96-well plate to measure the amount of BrdU-labeled DNA fragment in the cytosol. The amounts of BrdU-labeled DNA released were measured by sandwich ELISA using antibodies against DNA and BrdU. Absorbance was measured at 450 nm using a microplate reader with a reference wavelength at 690 nm.

2.7. Reactive oxygen species analysis

The presence of reactive oxygen species, superoxide and peroxide, was determined using dihydroethidium (DHE) (Sigma-Aldrich, St. Louis, MO) and 2',7'-dichlorodihydrofluorescein diacetate ($\text{H}_2\text{DCF-DA}$) (C400, Molecular Probes, Grand Island, NY). For superoxide analysis, HSG cells were cultured in plates containing glass cover slips until 75% confluence was reached. At this point, culture medium was removed; cells were rinsed with phosphate-buffered saline (PBS), and media containing 10 μ M PX was added to cells for 4 h. After 4 h, the PX-treated media was removed, cells rinsed with PBS, and media plus 5 μ M DHE was added for 30 min. Cells were rinsed with PBS and imaged for fluorescence at 525/610 nm (excitation/emission) using a NIKON epifluorescence microscope. Image Pro Analyzer 6.3 was used for DHE quantitation using the Integrated Optical Density (IOD) tool. For peroxide analysis, HSG cells were seeded in black 96 well plates to a density of 80% confluence and allowed to acclimate for 24 h. Cells were incubated with 10 μ M PX for 4 h. With 30 min remaining in the incubation, $\text{H}_2\text{DCF-DA}$ was added at a final concentration of 10 μ M. Cells were washed with PBS and fluorescence intensity was measured at 488/525 nm excitation/emission using a Molecular Devices fluorescent microplate reader.

2.8. Quantitative real-time PCR

HSG cells were exposed to 10 μM PX or ethanol (control) as described for 4 and 24 h. Total RNA was extracted from control and 10 μM PX-exposed cells with TRIzol/chloroform, DNaseI treated, and column purified (Omega Bio-Tek, Inc., Norcross, GA). PCR primers were designed using the Roche Universal Probe Library (UPL) Assay Design Center (Supplemental Table 1). Primers were synthesized by Integrated DNA Technologies (IDT, Coralville, IA). All primer sets spanned an exon–exon junction to reduce errors due to contaminating genomic DNA. Probes were selected from the Roche UPL. Amplification and detection of the fluorescence were measured using a Stratagene Mx3005p (Stratagene, La Jolla, CA). All signals were normalized to *glyceraldehyde 3-phosphate dehydrogenase* (*gapdh*). The fold change and p-values were determined by comparing mRNA expression in PX-treated to control cells. Once data had been normalized ($-\text{Ct}$), the average of 4 replicates in each group was used to calculate the $-\text{Ct}$ and fold change (fold change = $2^{(-\text{Ct})}$). p-values were calculated using $-\text{Ct}$ values, $n = 4$.

2.9. Statistical analyses

For the MTT and LDH cytotoxicity assays, statistical significances were determined by one-way ANOVA followed by a Tukey-Kramer *post hoc* test. For the cellular ATP, DNA fragmentation, and the q-rt-PCR experiments, statistical significances were determined using Student's *t*-test. For Mean IOD quantification of DHE fluorescence, statistical significance was determined by a two-tailed, unpaired *t*-test.

3. Results

3.1. AChE activity in HSG cells

The activity of AChE present in HSG cells was measured using a colorimetric assay (Ellman et al., 1961) at 5 min and 1 h, and afforded an acetylthiocholine hydrolysis rate of about 0.15 mU/ μg (nanomole/min/ μg). The low hydrolysis rate is just above background, non-enzymatic hydrolysis and slightly above the limit of detection (Supplemental Data Fig. 1). Enzyme activity was measured at 5 min and 1 h to determine any time-dependent variability in the observed AChE activity due to the lysis conditions, and to correlate the activity with prior studies (Strömbland, 1959) that measured the AChE activity in salivary gland cells over a 30 min time course. Because AChE activity was not significantly above baseline in HSG cells, the inhibition by PX could not be accurately measured and no further experiments with AChE were conducted.

3.2. HSG cell viability – MTT and LDH assays

Results from the MTT assay showed that there was no cytotoxicity at 24 h exposure to 0.01 to 0.1 μM PX concentrations (Fig. 1; Panel A). However, significant levels of cytotoxicity were observed at 24 h in HSG cells when treated with 1 μM , 10 μM and 100 μM PX (17%, 25%, and 35%, respectively) (Fig. 1; Panel A). Likewise, HSG cells at 48 h post-exposure to PX also showed significant cytotoxicity at doses of 1–100 μM (17–32%) as compared to control. In contrast, when LDH release was measured as an indicator of cytotoxicity, only 100 μM PX at 24 h showed deleterious effects (Fig. 1; Panel B). After 48 h of exposure, the cytotoxicity measured at 100 μM PX treatment increased to 13%. From this data, 10 μM PX

was selected for exposure experiments at 48 h since this concentration caused low, yet significant cytotoxicity in the MTT assay.

3.3. Cellular ATP levels

The results from the MTT assay indicate that mitochondria function may be impaired by PX exposure. If mitochondria function is affected, a corresponding decrease in cellular ATP could result. In order to further explore this possibility, cellular ATP levels were measured following 10 μ M PX exposure (Fig. 2). However, no change in cellular ATP levels was found up to 12 h of exposure. By 24 h, a slight, yet significant increase in cellular ATP levels was observed.

3.4. DNA fragmentation assay

In order to further characterize the cytotoxicity as apoptotic or necrotic, cellular DNA fragmentation resulting from PX exposure was determined using an ELISA to detect BrdU-labeled DNA fragments in HSG cell lysates or in cell culture supernatants. DNA fragments detected in cell lysates indicate apoptosis, whereas DNA fragments detected in cell culture supernatants is indicative of loss of membrane integrity and necrosis. HSG cells were exposed to 10 μ M PX for 2–24 h and DNA fragmentation was measured as BrdU incorporation. In PX-treated cell lysates, a significant amount of BrdU was observed within 12–24 h (Fig. 3A). There was a slight, yet significant increase in DNA fragments early (6–8 h post-exposure) in the supernatant fractions of the control cells (Fig. 3B), which decreased after 8 h.

3.5. Reactive oxygen species analysis

To determine if reactive oxygen species (ROS) contributed to the cytotoxicity of PX-treated HSG cells, analyses for the presence of superoxide and peroxide were conducted. Superoxide oxidizes dihydroethidium (DHE) into ethidium that subsequently intercalates into DNA and stains the nucleus a fluorescent red (Bindokas et al., 1996). At 4 h following 10 μ M PX exposure, nuclear fluorescence increased 2.6-fold in the PX-treated HSG cells compared to control cells indicating the formation of superoxide anion (Fig. 4). Conversely, an assay for peroxide using 2',7'-dichlorofluorescein diacetate (H₂DCF-DA), which reacts with intracellular H₂O₂ to form the fluorescent compound dichlorofluorescein (Saulsbury et al., 2009) indicated no difference between control and PX-exposed cells (Supplemental Table 2).

3.6. q-RT-PCR analysis

To further substantiate the possibility that PX exposure induces oxidative stress, the change in transcription of several genes associated with oxidative stress was assessed by q-RT-PCR. The genes selected for the q-RT-PCR experiments were based on a previous study that identified transcriptional changes in oxidative stress genes in PC12 cells after chlorpyrifos exposure (Slotkin and Seidler, 2009). The results from the q-RT-PCR experiments indicated that the genes *gstt2* (*glutathione S-transferase theta 2*) and *sod2* (*superoxide dismutase 2*) were significantly upregulated following exposure to 10 μ M PX for 4 h. Exposure to 10 μ M

PX for 24 h resulted in significant upregulation of the genes *gss* (*glutathione synthetase*), *gstm2* (*glutathione S-transferase mu 2*), and *gstt2* (Fig. 5).

4. DISCUSSION

This study tested the cellular response of the HSG cell line to a sub-lethal dose of the model organophosphate anticholinesterase agent, paraoxon (PX) – a dose that reduces mitochondrial function in approximately 20% of cells but does not cause cell death. Acetylcholinesterase (AChE) activity was determined to be negligible in the HSG cell line although substrate hydrolysis occurred slightly above background, possibly indicative of non-specific esterases. Previous reports have identified AChE activity in salivary glands or saliva using different enzyme activity assays (Ng et al., 2009; Strömbland, 1959). When AChE activity was measured manometrically (activity was expressed in ~ml CO₂ evolved/30 min/g of tissue), contamination by blood may have played a significant role in the enzyme activity since the samples were minced whole salivary glands (Strömbland, 1959). A more recent study measured extremely low AChE activity levels in salivary glands using a sensitive fluorescence-based assay (Amplex Red ACh/AChE Assay from Molecular Probes), which supports our conclusion (Ng et al., 2009). Based on the AChE activity assays and data from previous reports, we hypothesize that the cellular toxicity measured in this report most likely occurs through mechanisms other than AChE inhibition.

In order to determine if PX affected HSG cell viability, MTT assays and LDH release assays were employed to assess cytotoxicity. Significant increases in PX-induced cytotoxicity occurred in the HSG cells at most of the exposure concentrations tested beginning at 1 μM PX concentration at 24 and 48 h time points, as revealed by MTT assay. In contrast, significant changes in LDH concentrations only occurred at the highest concentration of PX tested (100 μM) following 48 h of exposure. The discrepancy in the results obtained from the MTT and LDH assays may be explained, in part, by differences in how cytotoxicity is determined. The MTT assay assesses mitochondria function whereas the LDH assay measures loss of cell membrane integrity. Therefore, the disparate assay results possibly suggest that exposure to PX may lead to impaired mitochondrial function (MTT reduction) in HSG cells prior to reduction in membrane integrity (LDH release).

If exposure to PX results in mitochondrial dysfunction, a resultant decrease in cellular ATP levels would be expected. However, there was no significant decrease in cellular ATP levels at 24 h, clearly contrary to expectation. These results may suggest that mitochondria are sensitive to PX exposure through mechanisms that are unrelated to ATP levels. For comparison, exposure to 100 μM of PX reduced HSG viability to 65% of control at 24 h, as analyzed by MTT assay, which is much less than that observed for SH-SY5Y neuroblastoma cells (Bharate et al., 2010; Prins et al., 2010). After 24 h exposure to 100 μM PX, SH-SY5Y viability was reduced to 20% of control (Bharate et al., 2010). The difference in cytotoxicity susceptibility between HSG and SH-SY5Y cells may be due to the inhibition of AChE and related OP-susceptible proteins in SH-SY5Y cells that are not present in HSG cells.

The amount of DNA fragmentation was measured to differentiate apoptotic or necrotic mechanisms. The results of the BrdU-labeling study indicate that DNA fragmentation is

occurring prior to loss of cellular membrane integrity, and therefore, cell cytotoxicity is most likely occurring via an apoptotic pathway (Elmore, 2007). However, BrdU-labeled DNA fragments in the cell lysates appear after 12 h of exposure, suggesting a time-dependent cytotoxicity possibly through the generation of ROS.

The levels of ROS like peroxide and superoxide can be increased by environmental stressors resulting in damage to cellular components, including damage to DNA, RNA, and proteins (Roberts et al., 2010). Cellular damage induced by ROS is cumulatively referred to as oxidative stress. In order to determine if PX induced ROS formation and subsequent oxidative stress, the HSG cells were analyzed for the presence of superoxide and peroxide (Roberts et al., 2010). The results of the DHE and H₂DCF-DA assays indicated that superoxide but not peroxide formed in PX-treated HSG cells, suggesting that the cells underwent oxidative stress within 4 h of exposure to 10 μ M PX. This outcome raises the possibility that oxidative stress is a trigger of cytotoxicity (Lukaszewicz-Hussain, 2010).

Cells defend themselves against ROS damage by upregulating proteins such as superoxide dismutase (*sod2*) and glutathione metabolism enzymes (*gss*, *gstm2*, and *gstt2*) that play an important role as cellular antioxidants (Laskin et al., 2010). In order to further establish that oxidative stress may play a role in PX induced cytotoxicity, q-RT-PCR was used to monitor the expression levels for several genes associated with oxidative stress. The genes *gstt2* (glutathione S-transferase theta 2) and *sod2* (superoxide dismutase 2) were upregulated following exposure to 10 μ M PX for 4 h, and at 24 h PX exposure induced an upregulation of the genes *gss* (glutathione synthetase), *gstm2* (glutathione S-transferase mu 2), and *gstt2*. This further confirms that oxidative stress plays an important role in PX induced cytotoxicity in HSG cells.

Although these studies suggest that oxidative stress plays a role in the cytotoxic mechanism in HSG cells exposed to PX, additional studies must be undertaken to exclude other possible biochemical routes. PX is a potent phosphorylating agent and likely to covalently modify a number of proteins and biomolecules. Therefore, it may be difficult to elucidate whether PX-induced oxidative stress in HSG cells results from direct phosphorylation of biomolecules involved in regulating oxidative species or indirect modification of biomolecules resulting from upregulation of oxidative stress proteins. This report demonstrates the potential utility of the HSG cell line as an effective *in vitro* model system to examine the non-cholinergic effects of exposure to organophosphate chemicals, which will ultimately lead to the discovery of novel mechanisms of toxicity and lead to the discovery of new biomarkers of OP exposure.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Highlights

- Human salivary gland (HSG) cells show a cytotoxic response to paraoxon exposure when analyzed by MTT assays but not LDH assays;
- Paraoxon induces the formation of superoxide in HSG cells as demonstrated by dihydroethidium analysis;
- Genes associated with reactive oxygen species were upregulated in paraoxon-treated HSG cells.

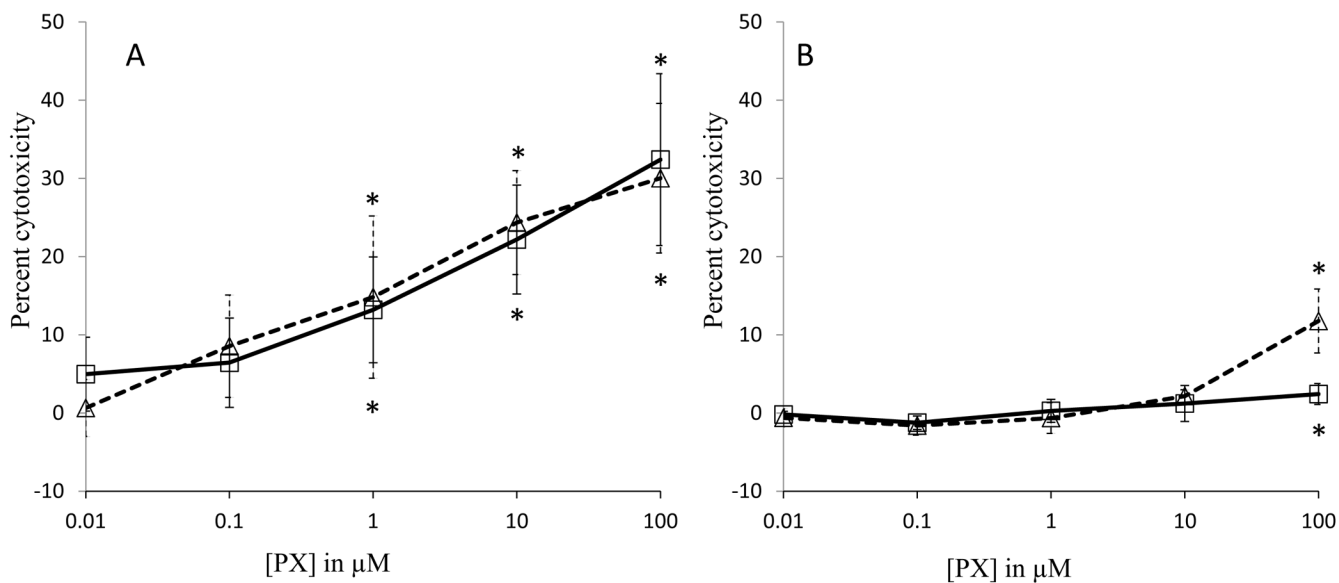


Fig. 1. Cytotoxicity of paraoxon (PX) on HSG cells. Cells were treated with PX and analyzed at 24 h (squares) and 48 h (triangles). Cytotoxicity was estimated by a MTT colorimetric assay (A), and an LDH release assay (B). Results are presented as average with standard deviation compared to cells treated with vehicle (0.1% ethanol). Triton X-100 was used as a positive control for cell death (100% cytotoxicity). *A significant difference between the treatment and negative control (no PX) was determined ($p < 0.05$) using one-way ANOVA followed by a Tukey-Kramer *post host* test ($n = 6-8$).

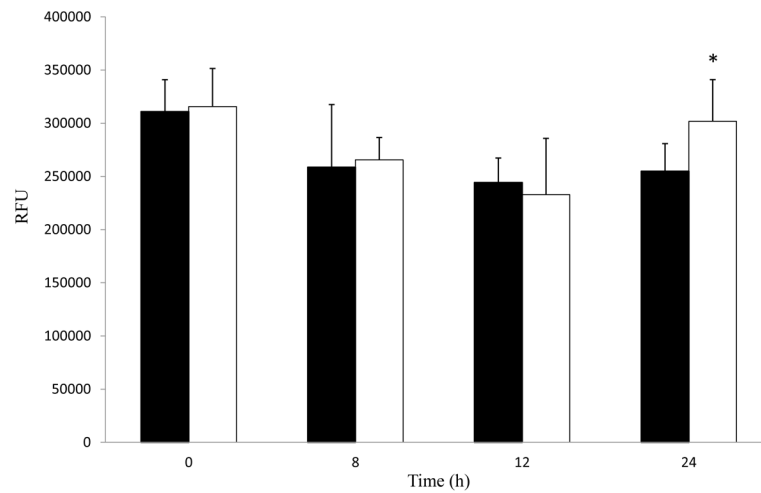
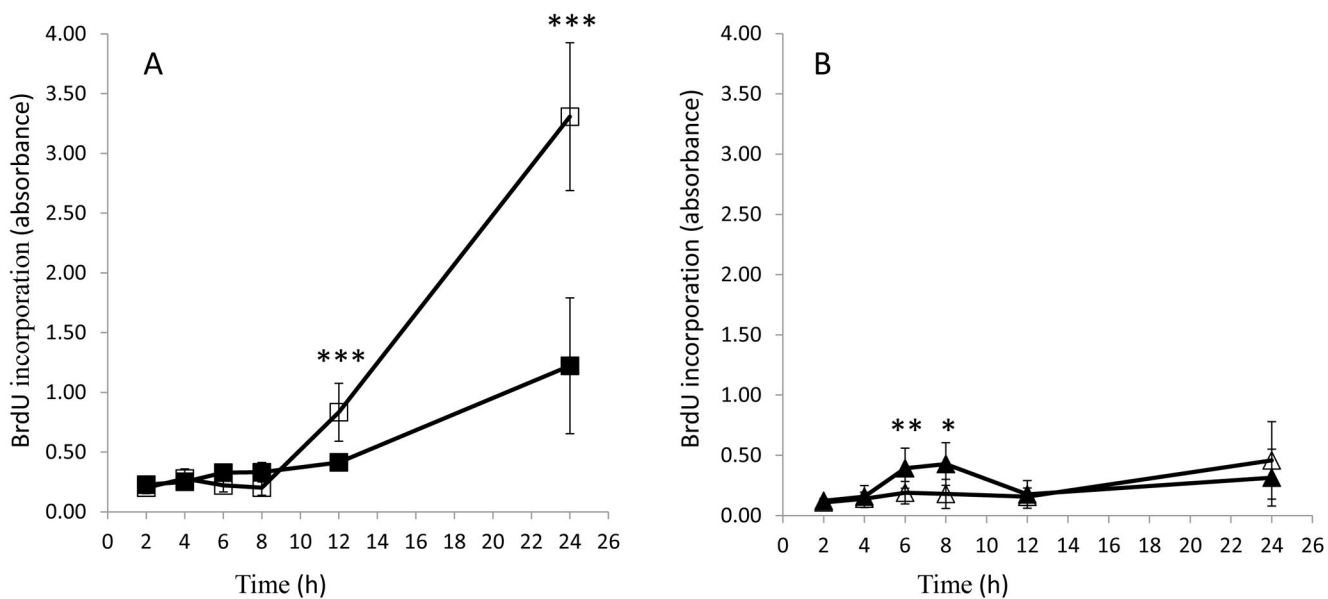


Fig. 2.

Paraoxon (PX) exposure slightly alters cellular ATP levels in HSG cells. Cells were treated with 0.1% ethanol as control (black bars) or 10 μ M PX (white bars). Cellular lysates were analyzed for cellular ATP levels (RFU, relative fluorescence units) from 8 to 24 h. Samples were analyzed and results shown are averages with standard deviations. Samples were analyzed for significance with Student's t-test, * represents p values < 0.05 (n = 4–6).

**Fig. 3.**

Paraoxon (PX) exposure induces DNA fragmentation in HSG cell lysates. Cell lysates (Panel A) or cell culture supernatants (Panel B) were analyzed for BrdU incorporation at 2 to 24 h following 10 μ M PX treatment (or ethanol control). Black squares, control lysates; black triangles, control supernatants; white squares, PX lysates; white triangles, PX supernatants. Results were analyzed using Student's *t*-test and data are represented as average with standard deviation. * represents *p* values < 0.05, ** represents *p* values < 0.01, and *** represents *p* values < 0.001 (*n* = 6–8).

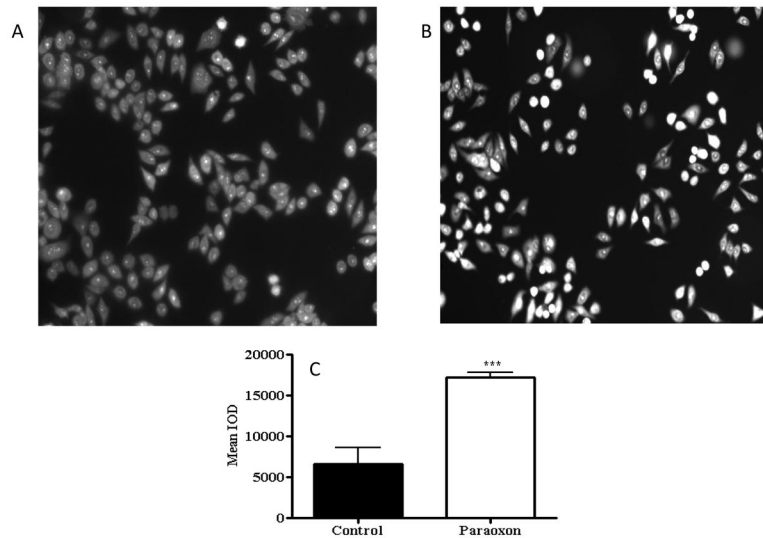


Fig. 4. Paraoxon (PX) exposure induces superoxide formation in HSG cells. Cells were treated with 0.1% ethanol as control (Panel A) or 10 μ M PX (Panel B) and treated with DHE at 4 h. Quantification of fluorescent staining (Panel C) shows that the mean IOD was significantly higher in PX-treated cells compared to controls (Student's *t*-test, *** $p < 0.001$).

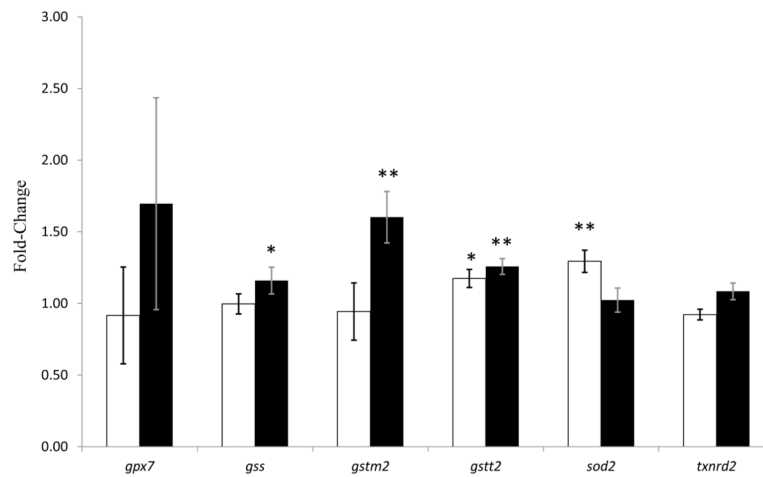
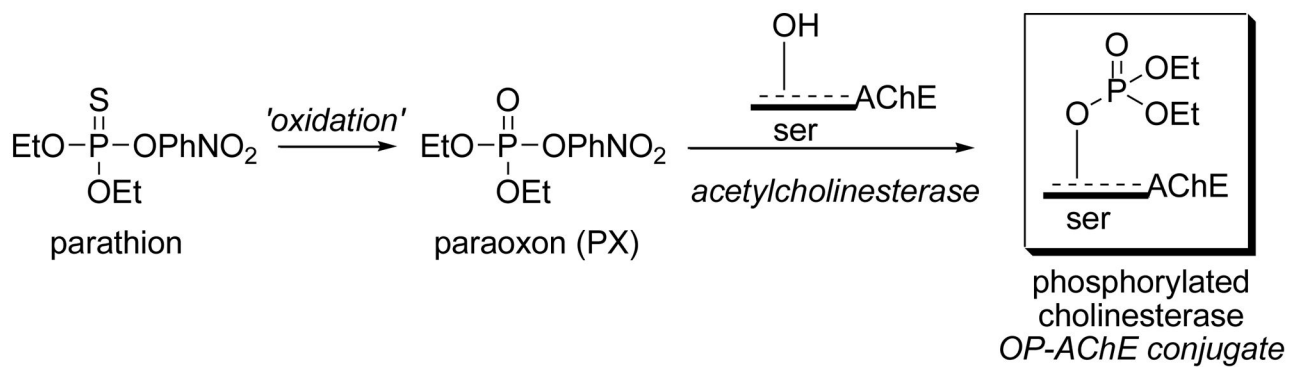


Fig 5.

Expression of oxidative stress genes in paraoxon-treated HSG cells. Cells were treated with either ethanol (0.1%) as a control or 10 μ M PX for 4h (white bars) or 24 h (black bars). Fold-change measured by q-RT-PCR of *gpx7*, *gss*, *gstm2*, *gstt2*, *sod2*, and *txnrd2* was measured. Expression was normalized to *gapdh* expression. Once data had been normalized ($-\Delta\Delta C_t$), the average of 4 replicates in each group was used to calculate the ΔC_t and fold change (fold-change = $2^{(-\Delta\Delta C_t)}$). p-Values were calculated using ΔC_t values, n=4, compared to the control using Student's *t*-test. * represents p values < 0.05, ** represents p values < 0.01.

**Scheme 1.**

Oxidation of parathion to paraoxon and the inhibition of acetylcholinesterase by paraoxon to form an OP-AChE adduct.