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ABSTRACT: Supplementary oxygen during resuscitation of the asphyxiated newborn is associated with long-term detrimental effects including increased risk of childhood cancer. It is suspected that the resuscitation procedure results in accumulated DNA damage and mutagenesis. Base excision repair (BER) is the major pathway for repair of premutagenic oxidative DNA lesions. This study addresses DNA base damage and BER in brain, lung, and liver in neonatal mice (P7) after hyperoxic resuscitation. Mice were randomized to 8% oxygen or room air for 60 min in a closed chamber and subsequent reoxygenation with 100% oxygen for 0 to 90 min. During this treatment, 8-oxoguanine accumulated in liver but not in lung or cerebellum. We observed a linear relation between 8-oxoguanine and reoxygenation time in liver DNA from hypoxic animals (n = 28; B =0.011 [0.001, 0.020]; p = 0.037). BER activity was not significantly changed during resuscitation. Our data suggest that after hypoxia, the capacity for immediate repair in liver tissue is inadequate to meet increasing amounts of DNA damage. The duration of supplementary oxygen use during resuscitation should be kept as short as justifiable to minimize the risk of genetic instability. (Pediatr Res 66: 533-538, 2009)

In the transition from intra- to extrauterine life, the fetus is challenged by a sudden surge in oxygen partial pressure. Rapid alteration of physiological processes is required, but $\sim 10\%$ of newborns need assistance to begin breathing and 1% demand extensive resuscitation (1). Administration of hyperoxic gas has been widely used to reestablish normoxia and presumably ensure a vigorous resuscitation. Meta-analyses indicate that this practice may increase mortality (2,3).

Exposure to hypoxia produces higher levels of reactive oxygen species (ROS) in the newborn, and a further increase is seen if resuscitation is performed with hyperoxic gas (4). We have previously provided evidence that markers of DNA oxidation in urine show a dose-dependent increase with graded inspiratory fractions of oxygen (5). Direct damage to DNA by ROS formed in vivo is probably a key event to development of malignant disease, and the base excision repair (BER) pathway plays an important role in disease protection (6,7). Notably, neonatal exposure to supplementary oxygen during resuscitation has been associated with an increased risk of childhood leukemias and cancer (8.9).

Exposure of DNA to ROS results in oxidized bases, apurinic/apyrimidinic (AP) sites, and single-strand breaks, all lesions that are handled mainly by a multistep process in the BER pathway (6). In the first step of the repair pathway, lesions are recognized and removed by specific DNA glycosylases, leaving an abasic site, which is acted upon by an intrinsic AP lyase activity in the DNA glycosylase or by a separate AP endonuclease. The resulting strand break is processed by lyases and/or nucleotidases removing the sugarphosphate backbone, and the gap is filled and resealed by DNA polymerases and ligases, respectively.

8-oxoguanine (8-oxoG) is one of the major DNA lesions formed from ROS. Guanine has a low oxidation potential and is readily oxidized to 8-oxoG. The base modification is highly mutagenic as it may cause GC to TA transversions if not recognized and repaired (7). Removal of 8-oxoG is mainly exerted by the enzyme 8-oxoguanine-DNA glycosylase (OGG1) (10), and a deficiency in this enzyme leads to an age-dependent accumulation of 8-oxoG in mice (11). DNA glycosylases display partially overlapping substrate specificities and may act as backup for each other. The MutY homolog (MUTYH) excises adenine (A) from 8-oxoG:A mismatches when OGG1 has failed, and deficiencies in both these DNA glycosylases result in a strong predisposition to lung and ovarian tumors and lymphomas (12). Oxidized pyrimidines such as 5-hydroxycytosine (5-ohC) may also arise in DNA from exposure to ROS and are acted upon by endonuclease III homolog (NTH1) (13). Deamination of cytosine forms uracil (U), which can be removed by the four DNA glycosylases uracil-DNA glycosylase (UNG), single-strand selective monofunctional uracil-DNA glycosylase (SMUG1), thymine-DNA glycosylase (TDG), and methyl-binding domain protein 4 (MBD4) (14). Repair of AP sites, which are highly mutagenic and cytotoxic because DNA replication can result in either double strand breaks or misincorporations, starts with the actions of an AP endonuclease (6). DNA glycosylases show widespread but differential expression, presence, and incision

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Abbreviations: 5-ohC, 5-hydroxycytosine; 8-oxodG, 8-oxo-2'-deoxyguanosine; 8-oxoG, 8-oxoguanine; AP, apurinic/apyrimidinic; BER, base excision repair; dG 2'-deoxyguanosine; OGG1, 8-oxoguanine-DNA glycosylase; ROS, reactive oxygen species; THF, tetrahydrofuran; U, uracil

activity, suggesting important age dependent roles both in brain (15) and liver tissue (16).

To better understand the pathogenic outcome after treatment of the newborn, we explored the effect of hyperoxic resuscitation on oxidative DNA damage and BER in an *in vivo* model of the acidotic neonate. Newborn mice were exposed to hypoxia to produce metabolic acidosis. We provide evidence that while capacity for the initial steps of BER seems unchanged, 8-oxoG increases in liver tissue during resuscitation with 100% oxygen. This accumulation of premutagenic lesions indicates a vulnerability to hyperoxic gas in the asphyxiated newborn.

MATERIALS AND METHODS

Animals. Experiments were approved by the National Animal Research Authority (NARA) in Norway. Mice were housed and handled in accordance with the European Council Directive 86/609/EEC. C57BL/6J mice were stabled and bred at 24°C on a 12:12-h light/dark cycle with access to a diet of pellets and water *ad libitum*.

Experimental protocol. At 7 d of age, pups were block randomized within litters to either hypoxemic hypoxia for 60 min in a lucite chamber containing premixed 8% oxygen balance nitrogen (Yara, Oslo, Norway) or normoxemia in a chamber containing ambient air. The groups were reunited and exposed to hyperoxic conditions by flushing the chamber with 100% oxygen. Mice were unanesthesized and unrestrained throughout the experiment. To study the effect of time in 100% oxygen, mice were exposed for random duration, ranging from 0 to 90 min. Conditions were monitored by sampling of the air and maintained constant by continuous flow of 0.5 L gas/min through the chamber with an open outlet to avoid alterations in pressure. Carbon dioxide levels were kept at <0.3%. Pups were separated from their dams during the procedure and temperature in the chamber was maintained at 35°C by floor heating to avoid confounding from possible effects of hypothermia. As body temperature decreases during hypoxia (17), the temperature was set slightly above thermoneutral range in newborn rodents (18). Experiments commenced between 0900 and 1000 h.

Immediately following experiments, mice were killed by decapitation and rapidly dissected on ice. Organs were removed and split. The right brain hemisphere, the right lung, and two of the four liver lobes were suspended in cold PBS for nuclear extracts, and the other halves were snap-frozen in liquid nitrogen and stored at -70° C for later DNA extraction.

Tissue preparation. All steps were carried out at 4°C unless otherwise stated. Preparation of total DNA was performed as follows: frozen tissue was homogenized with OmniTip plastic probes (OMNI International, Marietta, GA). DNA was isolated using the Wizard Genomic DNA Purification Kit (Promega, Madison, WI). Isolated DNA was denatured at 90°C for 10 min and single-strand DNA was hydrolyzed with Nuclease P1 (EC 3.1.30.1, Sigma Chemical Co.-Aldrich, St. Louis, MO) in buffer (500 mM NaAc, 2 mM ZnCl₂, pH 5.3) at 50°C for 60 min. Nucleotides were incubated at 37°C for 90 min with FastAP Thermosensitive Alkaline Phosphatase (Fermentas International, Burlington, Canada) in buffer (10 mM MgCl₂, 50 mM Tris pH 8), and the resulting nucleosides were stored at -20° C for later composition analysis.

Nuclear extracts were prepared as follows: fresh tissues were resuspended in cold MSHE (0.21 M mannitol, 0.07 M sucrose, 10 mM HEPES pH 7.4, 1 mM EDTA, 1 mM EGTA) and homogenized with a glass pestle and mortar. Homogenized tissue was separated by centrifugation at 2000 \times g for 10 min. Pellets were frozen and thawed. Nuclei were extracted with 2 volumes of hypertonic buffer (20 mM HEPES KOH pH 7.7, 0.5 mM MgCl₂, 0.42 M NaCl, 0.2 mM EDTA, 25% glycerol, 1 M DTT, and 100 mM PMSF) and separated at 14,000 \times g for 10 min. Supernatants were recovered and protein concentrations measured on a spectrophotometer [BioPhotometer (Eppendorf, Hamburg, Germany)]. Extracts were stored at -70° C for later incision activity assays.

Analysis of oxidized DNA. Isolated genomic DNA from cerebellum, liver, and lung was hydrolyzed and dephosphorylated as described earlier. The nucleosides were separated by HPLC [two serial coupled columns CromSep Inertsil 5 ODS-2, 2 × (150 mm × 4.6 mm), Crompack] in buffer [10% vol/vol methanol, water of Milli-Q grade (Millipore), 50 mM sodium acetate pH 5.0] set at 0.5 mL/min. 8-Oxo-2' deoxyguanosine (8-oxodG) was detected with an electrochemical detector (ECD) (Coulochem II, ESA) with a graphite filter protected 5010 analytical cell (ESA, screen electrode + 150 mV; analytical

electrode + 300 mV), and 2'-deoxyguanosine (dG) was measured with a UV detector (UV2000, Thermo Separation Products) set at 254 nm (19). Data are expressed as the number of 8-oxodG per 50,000 dG in the same sample.

DNA substrates. Oligonucleotides containing a lesion on a defined position were radioactive phosphor isotope 32 (³²P)-endlabeled with T4 polynucleotide kinase (PNK) (New England Biolabs, Ipswich, MA) and [γ^{-32} P]ATP (Perkin Elmer, Waltham, MA) as previously reported (20). Substrate oligonucleotides contained the 8-oxoG, 5-ohC or U lesions, or alternatively tetrahydrofuran (THF) (an AP site analog), at a distinct position. DNA substrates were prepared by hybridizing substrate oligonucleotides with complimentary oligonucleotides. The sequence of the THF substrate was (5'-GCTCATGCGCAG[THF]CAGCCGTACTCG-3') with a complementary G opposite the abasic site at position 13. Substrates with 5-ohC and U contained the lesion at position 15 (5'-GCATGCCTGCACGG[5ohC/U]CATGGCCA-GATCCCCGGGTACCGAG-3') opposite a G. The sequence of the 8-oxoG containing substrate was (5'-GGCGGCATGACCC[8-oxoG]GAGGC-CCATC-3') with a complementary C opposite 8-oxoG at position 14.

In brief, DNA was radiolabeled at the 5'end by incubation for 30 min at 37°C, and PNK was deactivated at 80°C. Complementary oligonucleotides were annealed by heating the samples to 90°C for 2 min followed by slow cooling to room temperature. The double-stranded oligonucleotides were purified by 20% native PAGE and visualized by storage phosphor autoradiography [Typhoon 9410 (Molecular Dynamics, Sunnyvale, CA)]. Radiolabeled substrates were isolated from the gel, eluted in dH₂O, and stored at 4°C.

Assays for enzyme cleavage of DNA. Incision assays were performed as previously reported (20) to estimate the ability of nuclear proteins to recognize and remove small, non-helix distorting DNA lesions. To determine the linear range of basal incision activity in control animals, dose-dependent titration reactions were performed (Fig. 1). Nuclear extracts were then diluted in buffer until all samples reached an equal protein concentration ($\pm 10\%$), and assays were performed with appropriate protein amount for the linear range.

DNA substrates and nuclear extracts were mixed in reaction buffer (50 mM MOPS pH 7.5, 1 mM EDTA, 5% glycerol, 1 mM DTT, and ³²P-labeled DNA duplex, additionally 5 mM MgCl₂ for THF containing substrate as AP endonuclease is magnesium dependent) in a total volume of 10 μ L and incubated for 30 min at 37°C. To promote complete strand cleavage at the abasic sites in 8-oxoG and U substrates, 2.5 μ L 0.5M NaOH was added. The reaction mixture was heated to 70°C for another 20 min and neutralized with an equal amount of HCl. A 1:1 amount of formamide loading dye was added, and the samples were denatured at 95°C for 2 min. The reaction products were resolved in 20% denaturing polyacrylamide gels and visualized by autoradiography as for the substrates. Incision activity was calculated as for the titration reactions.

Statistical analyses. Statistical analyses were performed using SPSS 15.0. Linear trends were explored using regression analysis; *B* describes the increase in 8-oxodG per minute resuscitation. Comparisons between the groups were made using *t* test. Significance was accepted as p < 0.05. Values are presented as mean \pm SEM unless otherwise stated.

RESULTS

Characteristics of the model. Six litters comprising 6 to 10 pups each were included, totaling 48 animals. Of these, five died during experiments and one was excluded due to hydronephrosis. There was no significant difference in weight between the groups (ambient air n = 14, 4.0 ± 0.3 g vs hypoxia n = 28, 4.1 ± 0.2 g; p = 0.76). A separate experiment was conducted to determine pCO₂, pH, and base excess (BE) in mixed blood after exposure to hypoxia (Table 1). Mean BE was decreased by 13.7 mmol/L. During the 60-min course of hypoxemia, pups partially compensated a metabolic acidosis by hyperventilating, resulting in hypocapnia. Respiration rate was counted manually before and 10 min after start of hypoxemia and showed a near significant increase (n = 6, 194 ± 3 vs 207 ± 4 ; p = 0.058 by paired-samples t test).

Dose-dependent increase in oxidative DNA damage from resuscitation with 100% oxygen. We measured 8-oxodG in total DNA. Resuscitation with 100% oxygen produced $\sim 25\%$ increase in 8-oxodG levels in liver DNA after 90 min resuscitation. The increase was time dependent and displayed a



Figure 1. Determination of linear reaction range for incision activity. (*A*) Typical titration experiment exemplified by incision activity in nuclear extract from lung on 5-ohC substrate. Negative and positive controls indicated. S, substrate; P, cleavage product. (*B*–*E*) Titration curves for nuclear extracts from brain (- -), liver ($\cdot \cdot \cdot$), and lung (—) incising (*B*) 8-oxoG substrate, (*C*) 5-ohC substrate, (*D*) Uracil substrate, and (*E*) AP-site substrate. Incision activity was calculated as the amount of radioactivity in the band corresponding to the damage specific cleavage product over the total radioactivity in the lane using the ImageQuant TL Version 2003.02 (Amersham Biosciences) software. The lower detection limit for incision is ~1%. Curves will flatten due to interference with repair from proteins with nonrepair capacities in nuclear extracts.

Table 1. Whole blood analysis after exposure to 8% oxygen for60 min

	Mean \pm SEM	
	Ambient air, n = 5	Hypoxia, n = 6
pH*	7.42 ± 0.01	7.36 ± 0.02
BE (mmol/L)†	4.2 ± 1.1	-9.4 ± 1.6
Pco ₂ (kPa)†	6.0 ± 0.2	3.5 ± 0.2

 $p^* < 0.05$ by *t* test.

 $\dagger p < 0.001$ by *t* test.

significant, linear progression (Fig. 2). Lung and cerebellum samples did not show a trend to change. In control animals (n = 14), 8-oxodG tended to increase less than in hypoxic animals in all tissues during resuscitation, displaying no significant change (cerebellum, *B* [95% CI] = -0.014 [-0.043, 0.016]; liver, *B* [95% CI] = 0.004 [-0.012, 0.019]; lung, *B* [95% CI] = 0.000 [-0.013, 0.012]). Adjustment for confounding factors (weight, sex, litter size, dams) did not change the significance of the results.

Liver tissue contained lower baseline levels of 8-oxodG per 50,000 dG (3.88 ± 0.25) compared with lung (5.86 ± 0.27) and cerebellum (5.41 ± 0.78) in control animals exposed to ambient air and no resuscitation (n = 8, p < 0.001). In animals exposed to hypoxia and no resuscitation, 8-oxodG levels were not significantly different from baseline.

BER activities are not altered by resuscitation procedure. DNA glycosylase and AP endonuclease activities were measured in nuclear extracts (Fig. 3). Removal of the base lesions 8-oxoG, 5-ohC, and U did not display significant changes after resuscitation. A synthetic THF substrate, inert to AP lyase incision, was used to distinguish AP lyase cleavage of abasic sites from AP endonuclease cleavage, which corresponds to APE1 activity. As illustrated in Figure 1, basal AP endonuclease activity was more potent than the glycosylase activity but showed no apparent change after resuscitation with 100% oxygen, indicating that the observed increase in oxidative damage to liver DNA does not trigger immediate upregulation of repair.

DISCUSSION

The results presented indicate that newborn mice exposed to hypoxemic hypoxia accumulate 8-oxoG in a timedependent manner in liver when resuscitated with 100% oxygen. In rapidly proliferating fetal and neonatal tissue, increased levels of damaged DNA are of particular concern because of the risk of fixating mutations during DNA replication, exacerbating the pathophysiological effects (21). Oxidized purines such as 8-oxoG are strongly mutagenic because of their preferred base pairing with adenine, and increased levels of this base modification are associated with mutations (10) and early neoplastic changes in the liver (22). We note with interest the previous reports of an association between duration of oxygen treatment and hepatoblastoma incidence in extremely low birth weight infants (23,24). The results reported herein provide a possible explanation for this association.

In support of previous *in vitro* studies, we found that incision activity for single nucleotide base modifications or abasic sites is not inducible by oxidative damage (20,25). In accordance with data in Figure 1, C57BL/6 mice are reported to have approximately equal OGG1 activity in brain and liver extracts (26). This is somewhat intriguing considering baseline 8-oxoG values in Figure 2, which are higher in the cerebellum, and may be attributed to differential tissue proliferative status with replication- or transcription-associated repair of oxidized lesions in single-stranded DNA (27). Rapid changes in BER activity have been reported after focal cerebral ischemia in adult models and are designated to posttranslational modifications and intracellular redistribution of repair enzymes (28). We would thus expect to detect a change in BER activity if such had occurred in our experiment and



Figure 2. Relation between resuscitation time with 100% oxygen and 8-oxodG/50 000 dG ratio in total DNA (n = 28). Regression lines are indicated. (*B*) Liver DNA displayed a dose-dependent increase in 8-oxodG (*B* [95% CI] = 0.011 [0.001, 0.020]; p = 0.037), whereas no significant effect of resuscitation was detected in (*A*) cerebellum (*B* [95% CI] = -0.006 [-0.021, 0.009]) or (*C*) lung (*B* [95% CI] = 0.005 [-0.007, 0.018]).



Figure 3. Effect of resuscitation with 100% oxygen on incision activity in diluted nuclear extracts from brain (\Box), liver (\triangle), and lung (\bigcirc) from hypoxic animals and controls (\blacksquare , \blacktriangle , \bullet) not exposed to hypoxia, incising (*A*) 8-oxoG substrate, (*B*) 5-ohC substrate, (*C*) Uracil substrate, and (*D*) AP-site substrate. Dot and whiskers represent the mean ± SEM of 3 to 5 independent samples. Values are normalized, the baseline incision activity in control animals not resuscitated is defined as 0%.

speculate that a more severe asphyxiation would provide DNA damage even without subsequent hyperoxia.

In addition to repair of 8-oxoG, we examined the repair activity for pyrimidine modifications and AP sites. The oxidized pyrimidine derivate 5-ohC can cause C-to-T transitions (29), whereas deamination of cytosine creates U:G mispairs, which may lead to GC to TA tranversions if not repaired (14). AP sites arising spontaneously or during BER are highly cytotoxic and mutagenic as incorporation of inappropriate bases may occur.

Mechanistically, an increased endogenous production of ROS may be responsible for the observed increase in 8-oxodG in liver. Oxidative stress occurs when the capacity for antioxidant detoxification is exceeded by increasing amounts of ROS. In mammalian cells, a major source of ROS is from energy production in mitochondria, releasing mainly superoxide anion $(\bullet O_2^-)$, hydrogen peroxide (H_2O_2) , and the highly reactive hydroxyl radical $(\bullet OH^-)$, and formation of these radicals increases both during hyp-

oxia and during hyperoxia (4,30). This paradox has been explained by inhibition of cytochorome oxidase in the electron transport chain at low concentrations of oxygen, favoring production of $\cdot O_2^-$ (31), although the mechanisms are not fully elucidated. We speculate that the proposed alterations in mitochondrial function during hypoxia further enhance production of ROS upon reoxygenation, which might explain the current findings in hypoxic animals as opposed to controls. Furthermore, the sudden surge in partial oxygen pressure at birth has been reported to increase the amount of 8-oxoG in rat liver DNA (32), indicating a finely tuned equilibrium under physiological conditions. Mitochondrial dysfunction and defective translocation of OGG1, as seen with increasing oxidative stress in older mice (33), may cause disruption and a progressive accumulation of oxidative DNA damage.

To avoid *in vitro* oxidation during DNA extraction and analysis, we followed a protocol extracting DNA and converting to deoxynucleosides by nuclease P1 and phosphatase treatment avoiding the use of oxidants that may generate oxidative lesions. Chromatographic methods for measurement of 8-oxodG are associated with artifactual oxidation, and consensus over the background levels remain to be established, but HPLC-ECD is appreciated as a highly accurate technique for measurement in tissue and cells (34). The modified base can additionally be measured in urine and is used as a biomarker of whole body oxidative DNA damage. Urinary measurements of 8-oxodG are attributed entirely to the repair of DNA damage (35). We have previously shown a dose-dependent relation between graded inspiratory fractions of oxygen during resuscitation of the newborn pig and urinary 8-oxodG (5) but were not able to collect representative samples of urine for comparison with total DNA analysis in this study as the pups frequently had spontaneous miction during experiments.

The pathophysiological responses to hyperoxia, including antioxidant enzyme activity, vary significantly among mouse strains (36). The C57BL/6 is commonly used in studies of DNA damage and repair (10,26), and adult mice display base damage in the lungs when exposed to prolonged hyperoxia (37). Because of the small size of the P7 mouse, there are technical challenges in mechanical ventilation and intensive care, raising a predicament on animal welfare contra the importance of a robust and reproducible model of asphyxia. We performed titration experiments increasing the exposure time to hypoxic gas up to 120 min, resulting in a further, although not significant, decrease of pH and BE but obviously increasing mortality. With 60 min of hypoxia, we experienced a 10% mortality rate and a decrease in BE satisfying the definition of metabolic acidosis. Because of hypocapnia produced by hyperventilation, systemic pH will display merely small changes in this model (38).

To our knowledge, there has been no study measuring the effect of supplementary oxygen on oxidative damage and repair in an in vivo neonatal model. Our data suggest that oxidative DNA damage arising after hypoxia and resuscitation with hyperoxic gas is effectively prevented or removed by the normal BER capacity in brain and lung, whereas in liver, an increased oxidative stress load is not met by up-regulation of BER activity. We conclude that while BER activity seems unchanged, there is a dose-dependent association between liver damage and the duration of exposure to hyperoxic gas. Further studies should be conducted to determine whether this vulnerability is differential and age dependent. We are currently cross-breeding specific DNA glycosylase knockout mice aiming for a model deprived of all repair of oxidized DNA lesions to further address the impact of asphyxia and resuscitation on long-term outcome. Although this study has a short duration and conclusions for long-term effects should be drawn with caution, we recommend the use of supplementary oxygen should be limited to justifiable proportions to minimize the risk of detrimental effects.

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