

Effects of Aging and Oxidative Stress on Spermatozoa of Superoxide-Dismutase 1- and Catalase-Null Mice¹

Johanna S. Selvaratnam³ and Bernard Robaire^{2,3,4}

³Department of Pharmacology and Therapeutics, McGill University, Montréal, Québec, Canada

⁴Department of Obstetrics and Gynecology, McGill University, Montréal, Québec, Canada

ABSTRACT

Advanced paternal age is linked to complications in pregnancy and genetic diseases in offspring. Aging results in excess reactive oxygen species (ROS) and DNA damage in spermatozoa; this damage can be transmitted to progeny with detrimental consequences. Although there is a loss of antioxidants with aging, the impact on aging male germ cells of the complete absence of either catalase (CAT) or superoxide dismutase 1 (SOD1) has not been investigated. We used CAT-null (*Cat*^{-/-}) and SOD1-null (*Sod*^{-/-}) mice to determine whether loss of these antioxidants increases germ cell susceptibility to redox dysfunction with aging. Aging reduced fertility and the numbers of Sertoli and germ cells in all mice. Aged *Sod*^{-/-} mice displayed an increased loss of fertility compared to aged wild-type mice. Treatment with the pro-oxidant SIN-10 increased ROS in spermatocytes of aged wild-type and *Sod*^{-/-} mice, while aged *Cat*^{-/-} mice were able to neutralize this ROS. The antioxidant peroxiredoxin 1 (PRDX1) increased with age in wild-type and *Cat*^{-/-} mice but was consistently low in young and aged *Sod*^{-/-} mice. DNA damage and repair markers (γ -H2AX and 53BP1) were reduced with aging and lower in young *Sod*^{-/-} and *Cat*^{-/-} mice. Colocalization of γ -H2AX and 53BP1 suggested active repair in young wild-type mice but reduced in young *Cat*^{-/-} and in *Sod*^{-/-} mice and with age. Oxidative DNA damage (8-oxodG) increased in young *Sod*^{-/-} mice and with age in all mice. These studies show that aged *Sod*^{-/-} mice display severe redox dysfunction, while wild-type and *Cat*^{-/-} mice have compensatory mechanisms to partially alleviate oxidative stress and reduce age-related DNA damage in spermatozoa. Thus, SOD1 but not CAT is critical to the maintenance of germ cell quality with aging.

aging, oxidative stress, Sertoli cells, spermatogenesis

INTRODUCTION

Spermatogenesis is a continuous process that allows males to produce sperm from the onset of puberty and well into old age. Although spermatozoa are continuously produced with advanced paternal age (APA), there is a growing body of

evidence indicating that APA is associated with negative impact on the quality of male germ cells [1]. Epidemiological studies indicate that APA may be a factor that is as critical as maternal age in determining pregnancy outcomes and offspring health. The effects of APA on pregnancy include increased time to pregnancy [2], preimplantation loss [3], and miscarriage [4]. The children of older fathers show increased risk for congenital birth defects and diseases such as achondroplasia, schizophrenia [5], and autism [6, 7].

The free radical theory of aging [8, 9] suggests that oxidative stress due to excessive free radicals or reactive oxygen species (ROS) is responsible for aging [10, 11]. ROS are the natural by-products of cellular metabolism and are involved in various cellular functions [12, 13]. If not maintained within normal physiological levels, ROS can damage cellular macromolecules, inducing stress signaling and, at high levels, cell death [14]. Thus, cellular antioxidant defenses [15] that neutralize ROS are essential, particularly for the protection of rapidly dividing male germ cells. The antioxidant superoxide dismutases (SODs) break down superoxide (O_2^-) radicals into hydrogen peroxide (H_2O_2), which can then be further neutralized by either catalase (CAT) or peroxiredoxins (PRDXs) to water. While CAT is localized to the peroxisomes and responds to high levels of H_2O_2 , the PRDXs are widely distributed in various subcellular organelles [16, 17] and neutralize low levels of H_2O_2 , hydroxyl (OH^-) and peroxynitrite ($ONOO^-$) radicals [18, 19].

Aging results in the loss of antioxidant activity and elevated levels of ROS [20, 21]. That this imbalance between ROS and antioxidants causes oxidative stress is well documented in the male reproductive tract [22, 23] and in the spermatozoa of aging rodents [21]. Despite the presence of complex antioxidant pathways with redundant antioxidant enzymes, germ-line knockouts (KOs), such as thioredoxins 2 and 3, which are part of the thioredoxin-glutaredoxin system, display increased susceptibility in spermatozoa to age-related oxidative stress [24]. Gene expression studies using isolated germ cells from aging rodents have highlighted that the antioxidants SOD1 and CAT play a critical role in the response to oxidative stress during aging [25, 26]. Mice lacking SOD1 (*Sod*^{-/-}) show increased spermatogenic cell susceptibility to heat stress [27] and have spermatozoa with impaired fertilizing ability [28]; to the best of our knowledge, there are no studies examining the germ cells of CAT-null (*Cat*^{-/-}) mice. The effects of aging on the quality of male germ cells in *Sod*^{-/-} and *Cat*^{-/-} mice have yet to be investigated. In addition to SOD1 and CAT, reduced levels of antioxidant PRDX1 have been linked to pathologies caused by increased oxidative stress, such as testicular torsion and varicocele [29], as well as male infertility [30, 31]. Furthermore, deficiencies in PRDX1 are associated with accelerated aging [32, 33]. The role of PRDX1 in the aging testis remains unclear.

In this study, we investigated the effects of the loss of SOD1 and CAT in both young and aged *Sod*^{-/-} and *Cat*^{-/-} mice on

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²Correspondence: Bernard Robaire, Department of Pharmacology & Therapeutics, McGill University, 3655 Promenade Sir-William-Osler, Montréal, Québec, Canada H3G 1Y6.
E-mail: bernard.robair@mcgill.ca

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testicular germ cells. Furthermore, to assess response to oxidative stress, mice were exposed to an acute oxidative insult by the pro-oxidant molsidomine (SIN-10) followed by measurement of ROS levels in live spermatocytes. We hypothesize that germ cells from aged mice lacking SOD1 will display increased ROS and greater susceptibility to DNA damage, while aged mice lacking CAT compensatory antioxidant will show a milder phenotype.

MATERIALS AND METHODS

Animals

The mice used for this study were bred in-house. C57BL6 strain wild-type (WT) and *Sod*^{-/-} (B6.129S7-*Sod1*^{tm1Leb}/DnJ) breeders were purchased from the Jackson Laboratory. *Cat*^{-/-} global KO breeders [34] were acquired from Dr. Eugene Chen at the University of Michigan. *Sod*^{-/-} mice have exons 1 and 2 replaced with a PGK-hprt expression cassette [35], and *Cat*^{-/-} mice have part of intron 4 and exon 5 replaced by a neocassette [34]. All mice were on C57BL/6 background and aged 3 mo (young) and 18 mo (aged). Mice were housed at the McIntyre Animal Resources Centre at McGill University, maintained under controlled temperature (22°C) and lighting (12L:12D), and provided with food and water ad libitum. Genotypes were determined by polymerase chain reaction (PCR) analysis of DNA extracted from tail biopsies using Qiagen DNeasy Tissue Kits (Qiagen). *Sod*^{-/-} and *Cat*^{-/-} genotyping was done using PCR protocols provided by the Jackson Laboratory and Dr. Eugene Chen, respectively. All animal studies were conducted in accordance with the principles and procedures outlined in the Guide to the Care and Use of Experimental Animals prepared by the Canadian Council on Animal Care (McGill Animal Care Committee protocol #4687).

Fertility Studies

To assess the fertility of male mice, we bred young WT female mice with either 3-mo (young) or 12-mo (aged) WT, *Sod*^{-/-}, or *Cat*^{-/-} males. The numbers of pups per litter that were sired from these breeding pairs were counted for each mouse group. The results were presented as number of pups per litter sired by 3-mo and 12-mo males of each genotype. The number of litters examined per experimental group ranges from 11 to 14. While all female breeders were WT and aged between 3 and 6 mo, male breeders were chosen from each genotype (WT, *Sod*^{-/-}, or *Cat*^{-/-}) and aged 12 mo, which is typically the oldest age of retired breeders [36].

Animal Treatments

Oxidative stress was induced with administration of molsidomine (SIN-10) (Sigma Aldrich Canada). Mice aged 3 or 18 mo were given a single intraperitoneal injection of 30 mg/kg of body weight 2 h prior to being euthanized [37]. SIN-10 treatment time was chosen based on previous studies that indicated that serum levels of the SIN-10 metabolite SIN-1 peaked at 2 h [38], and experimental trials in which the treatment time of 2 h was found to induce a peak increase in ROS in pachytene spermatocytes from young WT animals (Supplemental Figure S1A; Supplemental Data are available online at www.biolreprod.org). The SIN-10 dose was based on information in the literature [37, 38] and preliminary dose-response trials using aged animals to measure ROS response to the oxidative insult (Supplemental Figure S1B). In control animals, SIN-10 was replaced with saline injections. After each treatment, mice were euthanized using carbon dioxide followed by cervical dislocation. Tissues were removed, weighed, flash frozen, and stored at -80°C for later analyses. Five to seven animals were used for each of the four groups of animals (young and old serving as either control or SIN-10 treated) for both the *Sod*^{-/-} and the *Cat*^{-/-} studies.

Spermatocyte ROS Measurements

Mice were euthanized 2 h after SIN-10 or saline (control) injections, and their testes were removed. Preparation of testicular germ cell suspensions was adapted from a previously described method [39]. Briefly, the testes were decapsulated, and the parenchyma was minced using a scalpel until in a semiliquid state and mixed with an equal volume of phenol red-free media (Sigma Aldrich Canada); the cell suspension was strained through a 56- μ m nylon mesh. The cells were then diluted and counted using a hemocytometer, and 150,000 cells were aliquoted per tube from the mixed testicular cells.

The mixed testicular cells were then incubated for 30 min at 32°C with Hoechst (2,5'-bi-1H-benzimidazole, 2'-[4-ethoxyphenyl]-5-[4-methyl-1-piper-

azinyl]) (ThermoFisher) and probes to measure ROS that included CellROX DeepRed Reagent and peroxynitrite sensor aminophenyl fluorescein (Invitrogen). After incubation, the cells were washed in media and transferred to a 96-well Cell Carrier (PerkinElmer) plate with an optically clear bottom. The plate was immediately scanned using the Operetta (PerkinElmer) high-content imaging system, and images were then analyzed using Columbus software (version 2.5.0; PerkinElmer) with a previously validated algorithm designed to select for spermatocytes [40]. The overall ROS and peroxynitrite intensities in spermatocytes were measured per well per cell using this software. Information for each probe used is listed in Supplemental Table S1.

Sperm Counts

Testicular and epididymal sperm counts were done as previously described [41]. Briefly, a frozen testis or epididymis from each mouse was weighed and homogenized (Polytron PTA7, setting 5; Brinkman Instruments) for two 15-sec periods separated by a 30-sec interval in 2 ml of 0.9% NaCl (Millipore), 0.1% thimerosal (Sigma), and 0.5% Triton X-100 (BDH). Sperm heads were counted in aliquots of diluted homogenate using a hemocytometer. Sperm counts from the testis and epididymis are shown as the number of sperm $\times 10^6$ per tissue.

Testis Fixation

Young and aged mice were euthanized from each group (WT, *Sod*^{-/-}, and *Cat*^{-/-}); their testes were removed and immersion fixed in modified Davidson fixative [42] for 2 h, then the testes were sliced in half and further fixed for 6–8 h. After fixation, the testes were transferred to 70% ethanol, further dehydrated, and then embedded in paraffin.

Histology

Several samples from different testes were sectioned (4 μ m) and placed on a single slide; each had sections from different mice of the same group, with the addition of one WT control animal on each slide. The slides were deparaffinized (HistoClear; Diamed Inc.) and loaded in 30-slide batches into the Tecan GenePaint (Tecan) solvent delivery robotic system, programmed to rehydrate the tissues by passing them through a graded ethanol series, and then stained with Periodic Acid Schiff (PAS) (Sigma) according to the manufacturer's protocol. The slides were mounted with Permount (Fisher Scientific) and coverslips and left to dry before observation with a light microscope. Tubule diameters and epithelial heights were measured using ImageJ (NIH) for 200 tubules per animal, with five to six animals per experimental group. Using similar methods, a fixed epididymis from each mouse was sectioned (4 μ m) and stained using Toluidine Blue (Sigma) following the manufacturer's protocol.

Immunofluorescence

The Tecan GenePaint robotic system was used to do automated immunofluorescence labeling of batches of slides. Before loading the slides, they were deparaffinized with HistoClear, rehydrated through a graded ethanol series, washed in PBS for 5 min, and finally boiled in sodium citrate buffer (10 mM sodium citrate, 0.05% Tween 20, pH 6.0) for 30 min. The slides were left to cool to room temperature for approximately 20 min and then rinsed in PBS and loaded into the Tecan GenePaint robotic system, which was programmed to add blocking reagents (10% normal goat serum) for 1 h, primary antibodies for 1 h, secondary antibodies for 45 min, and 4',6'-diamidino-2-phenylindole (DAPI) combined with lectin PNA from *Arachis hypogaea* (peanut) Alexa Fluor 647 conjugate (ThermoFisher) (used as a staging tool) for 10 min. Finally, slides were mounted with ProLong Gold antifade mountant (ThermoFisher), left to dry at room temperature for 24 h, and then stored at 4°C before imaging. Information for each antibody used is listed in Supplemental Table S1.

Oxidative damage-induced DNA lesions, assessed using 8-oxo-2'-deoxyguanosine (8-oxodG) as a marker, was detected in sperm heads by immunofluorescence, as previously described [25]. Briefly, frozen slides with sperm smears were placed in PBS for 5 min and blocked in goat serum for 30 min. Slides were then incubated in primary antibody specific for 8-oxodG (1:10) overnight at 48°C (controls incubated in serum alone). Slides were washed in PBS and incubated in goat-anti-mouse conjugated to Alexafluor 488 (ThermoFisher) for 30 min, washed in PBS, and mounted with DAPI. Using a Zeiss LSM 510 Axiovert confocal microscope to visualize sperm, a minimum of 100 spermatozoa per slide per sample were counted as either positive or negative for 8-oxodG. Positive control samples were pretreated for 30 min with 5% H₂O₂ (Sigma).

Measurement of Immunofluorescence Intensity

Sequential images were captured (resolution 2488×2488) using a multiphoton Leica TCS SP8 MP microscope with $20\times$ and $63\times$ objectives. Quantification of positive cells per tubule was assessed using Imaris version 3.8 software (Bitplane). Briefly, tubules were manually selected and cells identified by Imaris using DAPI nuclear stain; intensity data for each cell for each channel were exported. Data from control slides with no primary antibody and slides with blocking peptides were used to set baseline thresholds and the intensity range for each marker.

Statistical Analyses

All statistical analyses were done using two-way ANOVA followed by Bonferroni tests or one-way ANOVA with Newman-Keuls tests with GraphPad Prism 4.0; significance was set at $P < 0.05$.

RESULTS

The Effects of Aging on Body and Tissue Weights, Sperm Counts, and Fertility

WT mice at 3 mo had significantly higher body weights than KO mice of the same age. Mice were littermates, indicating that the body weight was reduced in response to the antioxidant loss in *Sod*^{-/-} and *Cat*^{-/-} mice (Fig. 1A). Aging resulted in increased body weight for all animals. WT mice had the greatest increase in weight, while *Cat*^{-/-} and *Sod*^{-/-} mice had significantly lower body weights compared to WT (Fig. 1A). Fertility was unaffected at 3 mo among WT, *Cat*^{-/-}, or *Sod*^{-/-} mice. By 12 mo, the number of pups per litter was significantly reduced in all mice; however, the number of pups per litter in *Sod*^{-/-} mice at this age was significantly lower than that in aged controls (Fig. 1B).

Testis weight remained the same with age; *Cat*^{-/-} and *Sod*^{-/-} mice maintained lower testis weights compared to WT mice at both 3 and 18 mo (Fig. 1C). Testes sperm counts were significantly reduced in young *Sod*^{-/-} mice; by 18 mo, all mice had significantly reduced testis sperm counts (Fig. 1D). There were no significant changes observed in epididymal weights with either aging or KO mice (Fig. 1E). Yet, similar to the testis sperm counts, the epididymal sperm counts were significantly reduced in young *Sod*^{-/-} mice compared to WT; epididymal sperm counts were significantly reduced in all mice with aging (Fig. 1F). Since tissue weights were only minimally affected, expressing sperm counts per milligram of tissue, as opposed to per tissue, resulted in the same significant differences (data not shown).

Epididymal histology revealed lumina filled with sperm in the caput and cauda of young WT mice; in aged WT mice, some sloughed-off cells and sperm were visible in the epididymal lumina. Similar observations were made in the caput and cauda epididymides of young and aged *Cat*^{-/-} mice (Supplemental Figure S2). In contrast, at 3 mo, *Sod*^{-/-} mice displayed sloughed-off cells in all segments of the epididymis, with collections of cells in the lumina; by 18 mo, the epididymal epithelia of the different segments were visibly altered with accumulation of a few sloughed-off cells and fluids in the epididymal lumina (Supplemental Figure S2).

Accelerated Loss of Germ Cells and Sertoli Cells with Aging in Antioxidant KO Mice

Testes histology (Fig. 2, A–F) indicated loss of germ cells in the seminiferous tubules of all mice with aging (Fig. 2, B, D, and F). Seminiferous tubule diameters and epithelium heights were significantly reduced with aging in WT mice (Fig. 2, G and H). In the testes of young *Cat*^{-/-} mice, some seminiferous tubule lumina were filled with sloughed-off cells (Fig. 2C).

Compared to WT mice, young *Cat*^{-/-} mice had significantly smaller seminiferous tubule diameters and epithelial heights; diameters were maintained in aged *Cat*^{-/-} animals, while epithelial heights were significantly reduced at 18 mo (Fig. 2H). Young *Sod*^{-/-} mice had significantly reduced tubule diameters compared to WT; these were further reduced with aging (Fig. 2G). The seminiferous tubule epithelial heights were significantly smaller in young *Sod*^{-/-} mice when compared to WT and were unaltered with aging (Fig. 2H).

Using ZBTB16 as a marker for spermatogonia and vimentin as a Sertoli cell marker (Fig. 3A), we found that the mean germ cell numbers per seminiferous tubule were significantly lower in *Cat*^{-/-} and *Sod*^{-/-} mice compared to WT at 3 mo; all mice displayed a loss in germ cell numbers with aging (Fig. 3B). This loss of germ cells in aged animals was significant at all stage groupings. Sertoli cell numbers were reduced only in young *Sod*^{-/-} compared to WT mice; this reduction was significant in the later stages of spermatogenesis in stage groupings VIII–IX and X–XII. Aging resulted in the loss of Sertoli cells in all mice; interestingly, aged *Cat*^{-/-} mice displayed significantly higher numbers of Sertoli cells compared to aged *Sod*^{-/-} mice (Fig. 3C), indicative of possible compensatory protective mechanisms in *Cat*^{-/-} mice. The numbers of spermatogonia among all groups were unaltered (Supplemental Figure S3). Serum levels of gonadotropins LH and FSH and serum testosterone were not significantly changed between WT, *Cat*^{-/-}, and *Sod*^{-/-} mice or with aging (Supplemental Figure S4).

Spermatocyte ROS in SIN-10-Treated Mice

Mice were treated with the pro-oxidant SIN-10 (Fig. 4A), their germ cells were isolated, and the mean ROS intensities were quantified following spermatocyte selection. Young WT mice responded to the oxidative insult with no change in spermatocyte ROS versus saline-treated controls, suggesting effective removal of cellular ROS (Fig. 4B). However, spermatocytes from aged WT mice were not able to remove ROS following oxidative insult and displayed significantly increased ROS with SIN-10 treatment (Fig. 4B). In contrast, spermatocytes from both young and aged *Cat*^{-/-} mice were able to respond to SIN-10-induced oxidative insult by neutralizing ROS efficiently (Fig. 4C). In *Sod*^{-/-} mice, the ROS levels of spermatocytes showed similar responses to SIN-10 as WT mice; spermatocytes from young mice were able to neutralize SIN-10-induced ROS, but in the aged mice they were not able to neutralize this ROS (Fig. 4D). Based on MitoSOX measurements, spermatocyte O₂⁻ levels were not significantly changed among the groups (data not shown), while ONOO⁻ measurements, detected using aminophenyl fluorescein, displayed identical results to those seen with overall cellular ROS (Supplemental Figure S5).

PRDX1 Localized to Germ Cell Cytoplasm and the Acrosome

Live cell ROS measurements indicated that ONOO⁻ radicals were generated in the spermatocytes following SIN-10 treatment in aged WT and *Sod*^{-/-} (Supplemental Figure S5), so we decided to further investigate the antioxidant involved in peroxynitrite neutralization, PRDX1 [33]. In addition, PRDX1 has been associated with aging and oxidative stress-linked male infertility [29]. Interestingly, previous studies localized PRDX1 specifically to the Leydig cells in the mouse testis [43]. Similar to these studies, we localized PRDX1 to the Leydig cells (Supplemental Figure S6); however, unlike these previous

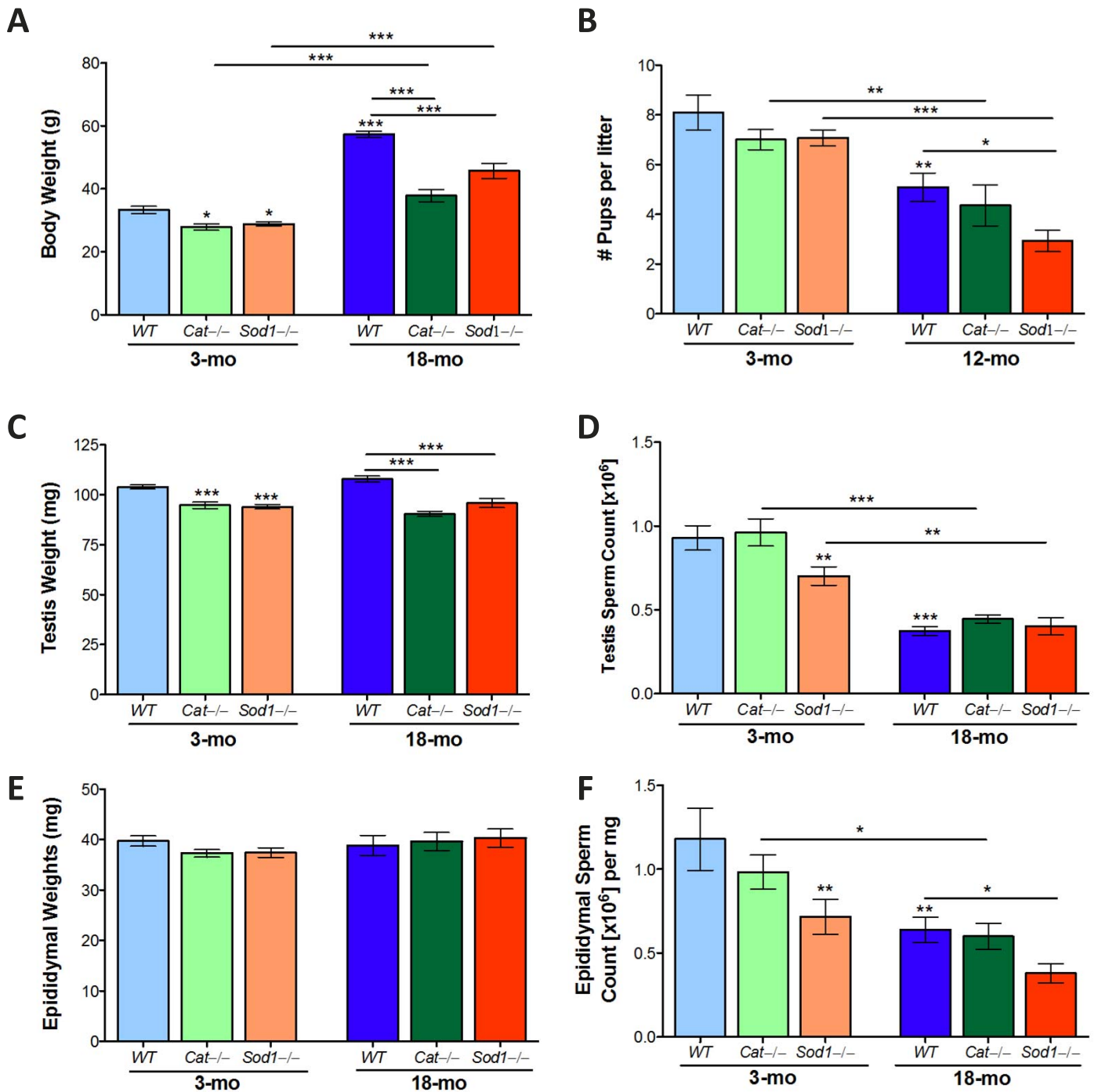


FIG. 1. Reproductive organ weights and testis and epididymal sperm counts in aged wild-type (WT), *Cat*^{-/-} and *Sod1*^{-/-} mice. Body weights increased with age in WT, *Cat*^{-/-}, and *Sod1*^{-/-} mice (A), while *Cat*^{-/-}, and *Sod1*^{-/-} mice remained consistently lower in weight than WT mice. The numbers of pups per litter decreased with aging (B) and were significantly lower in aged *Sod1*^{-/-} mice. Testis weights (C) were reduced in *Cat*^{-/-} and *Sod1*^{-/-} compared to WT mice. Testis sperm counts (D) were lower in young *Sod1*^{-/-} mice and all aged mice. Epididymal weights were not significantly changed (E); sperm count in the epididymis (F) reflected the changes observed in the testis sperm counts. Mean values with SEM error bars; two-way ANOVA with Bonferroni test; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$, $n = 10-14$ (A, B, D, F), $n = 5-8$ (E, G), $n = 4-5$ (C, F).

studies, PRDX1 was also detected in the cytosol of spermatocytes and developing spermatids, specifically within the developing acrosome (Fig. 5A, orange arrow heads indicate PRDX1 in the acrosome). Furthermore, the localization of PRDX1 revealed that there was a progressive increase in PRDX1 immunoreactivity in germ cells as they developed through the different stages of spermatogenesis (Fig. 5A).

Up-Regulation of PRDX1 with Aging Occurs in All Mice, Except *Sod1*^{-/-} Mice

The percentage of PRDX1-positive cells per seminiferous tubule section was quantified in the testes of each experimental group. The results showed that PRDX1-positive cells increased with age in WT and *Cat*^{-/-} mice, with the most significant

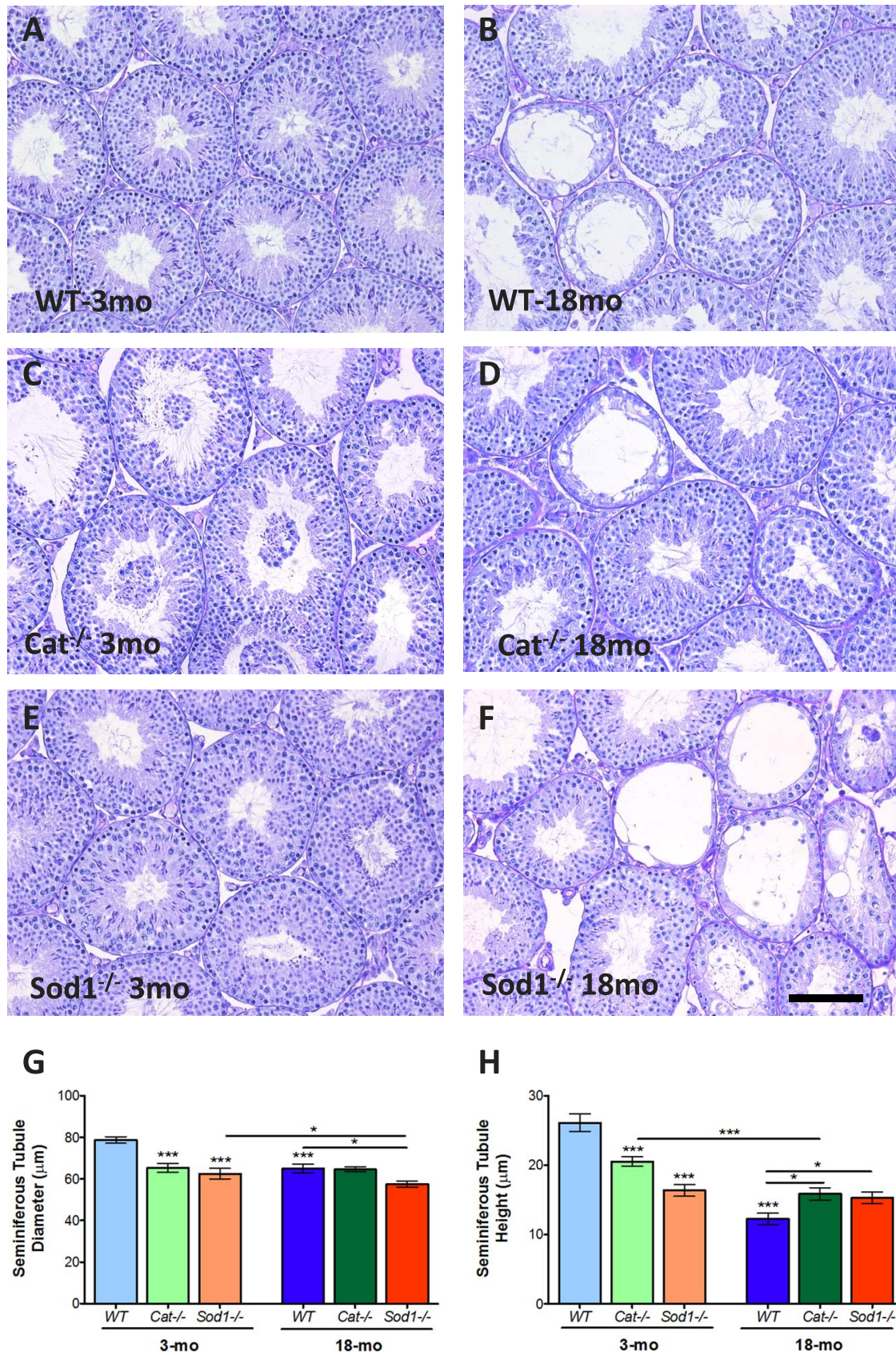
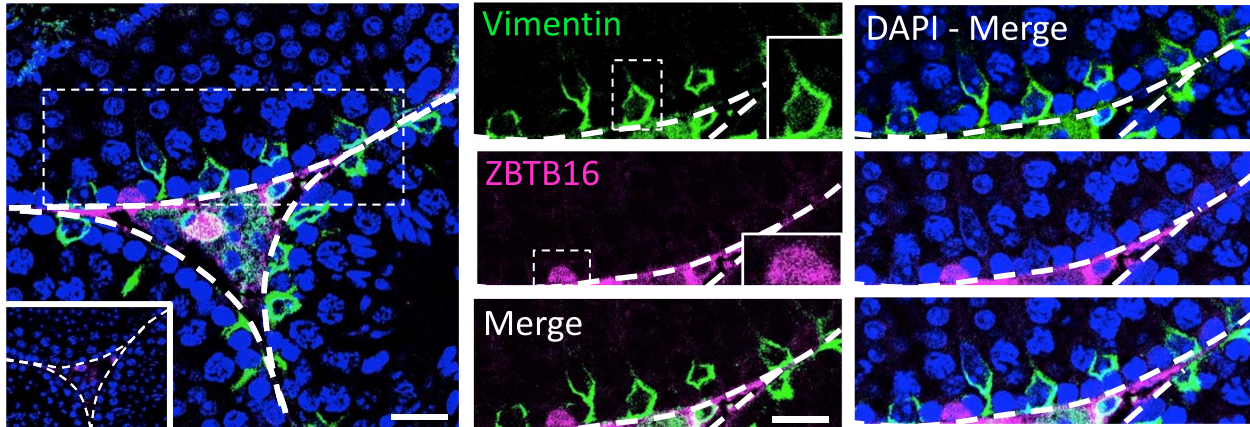
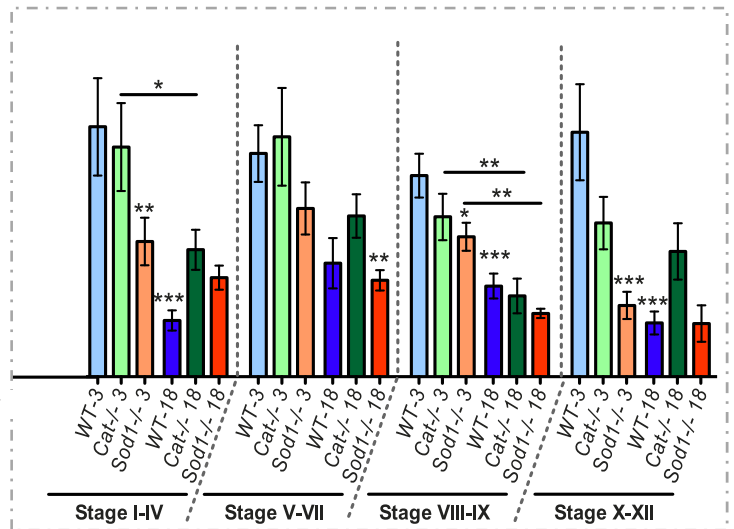
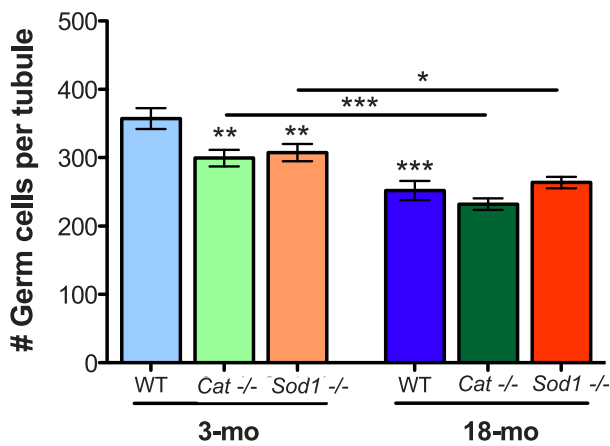


FIG. 2. Histological analyses of young and aged *Cat*^{-/-} and *Sod1*^{-/-} mouse testes. Testis histology in young and aged mice stained with PAS-hematoxylin (A–F). Seminiferous tubule diameters (G) were reduced in knockout (KO) mice and in all mice with aging. The seminiferous tubule epithelium height (H) was reduced in KO mice, with the most extreme loss in height in *Sod1*^{-/-} mice. Loss of tubule height was observed with aging in all mice. Mean values with SEM error bars; two-way ANOVA with Bonferroni test; **P* < 0.05; ***P* < 0.01; ****P* < 0.001, *n* = 6 (A–F); *n* = 8 (G, H). Bar = 100 μm .

A



B



C

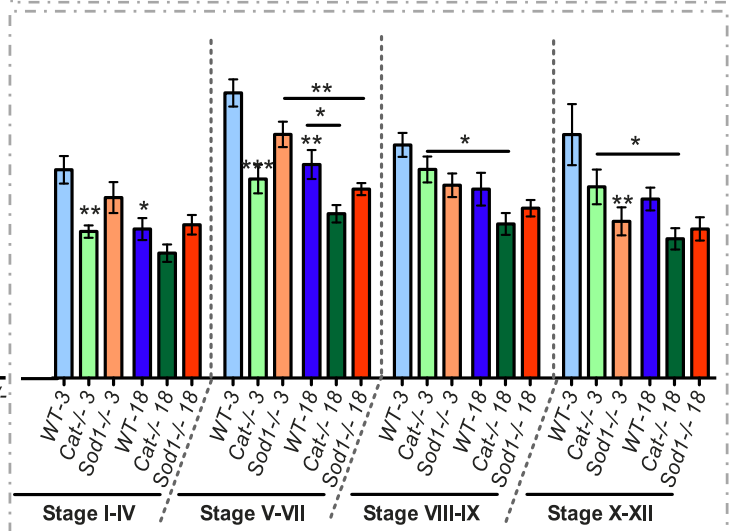
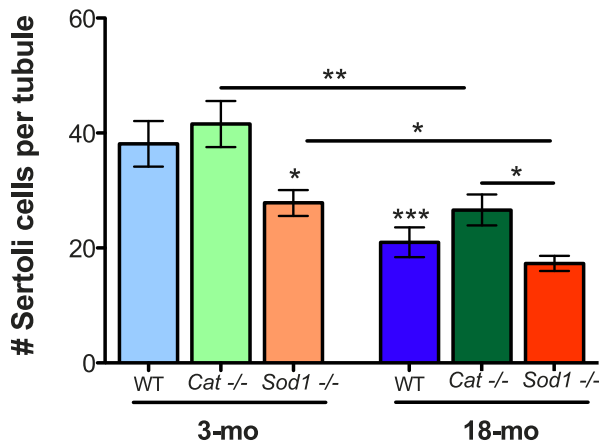


FIG. 3. Increased Sertoli cell loss in *Sod1*^{-/-} mice and with aging in all mice. Representative image with DAPI (nuclear stain), ZBTB16 (spermatogonial marker), and vimentin (Sertoli marker in the testis epithelium) (A). Imaris quantification of the number of germ cells per tubule (B) and the number of Sertoli cells per tubule (C) are also shown sorted by stage groupings. Mean values with error bars (\pm SEM); two-way ANOVA with Bonferroni test; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; $n = 4$. Bar = 20 μ m.

increase seen during stages VIII–IX (Fig. 5B). There was no significant difference between WT and *Cat*^{-/-} mice at 3 mo,

but by 18 mo, WT mice had significantly more PRDX1-positive cells than *Cat*^{-/-} mice (Fig. 5B), particularly during

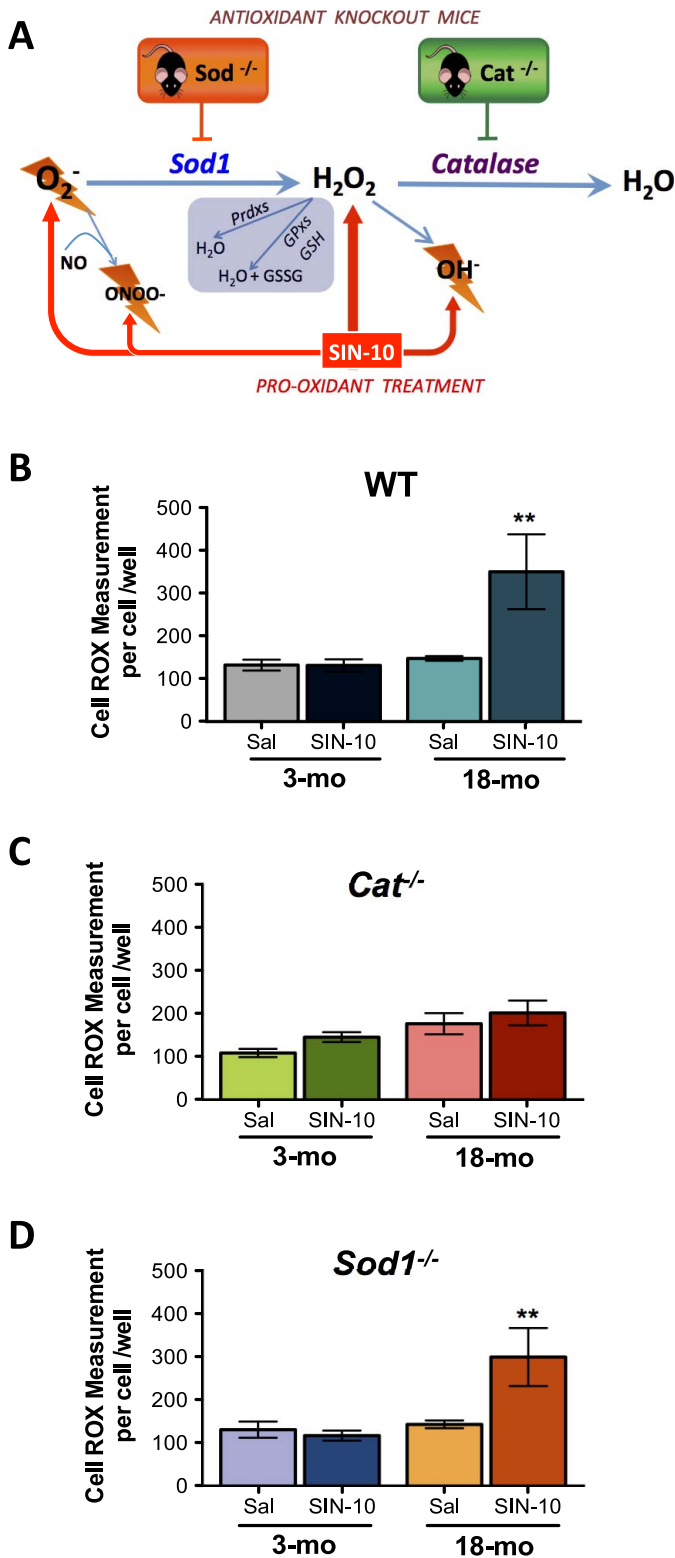


FIG. 4. Redox dysfunction with aging in WT and *Sod1*^{-/-} spermatocytes and efficient management of oxidative stress in spermatocytes from *Cat1*^{-/-} mice following SIN-10 induced oxidative insult. Schematic of SIN-10 induced ROS generation (A). Overall ROS, measured using CellROX, showed efficient removal of ROS in all young animals, WT (B), *Cat1*^{-/-} (C), and *Sod1*^{-/-} (D). While aged *Cat1*^{-/-} mice responded to SIN-10 with removal of ROS, aged WT and *Sod1*^{-/-} did not efficiently remove ROS following SIN-10 treatment (D). Mean values shown with error bars (\pm SEM); one-way ANOVA with Newman-Keuls; ** $P < 0.01$, $n = 4-5$.

spermatogenic stages VIII–IX. Young *Sod1*^{-/-} mice displayed significantly fewer PRDX1-positive cells per tubule compared to young WT mice; this trend was most evident during the later stage groupings 8–9 and 10–12. Unlike the increases observed in WT and *Cat1*^{-/-} mice, in *Sod1*^{-/-} mice there were no significant changes in the percentage of PRDX1-positive cells per seminiferous tubule with aging (Fig. 5B). No significant changes were observed between young and aged *Sod1*^{-/-} in PRDX1 at any stage groupings. Overall, compared to both aged WT and *Cat1*^{-/-} mice, the percentages of PRDX1-positive cells per tubule were significantly lower in the aged *Sod1*^{-/-} mice (Fig. 5B).

Increased DNA Double-Strand Breaks and Reduced Repair in Aged Mice

DNA double-strand breaks (DSBs) lead to the modification of histone H2AX; phosphorylation of this histone forms γ -H2AX, which then recruits DNA damage repair proteins, such as p53 binding protein 1 (53BP1). γ -H2AX is normally present within germ cells going through meiosis in the testis [44]. The percentage of cells positive for γ -H2AX per tubule in young WT mice is indicated in Figure 6A. No significant differences were observed in γ -H2AX-positive cells in young WT, *Cat1*^{-/-}, or *Sod1*^{-/-} mice. However, during stages I–IV, young *Cat1*^{-/-} displayed a significantly higher percentage of γ -H2AX-positive cells (Fig. 6A). Aging caused the percentage of γ -H2AX-positive cells to decrease in WT mice at all stages. In contrast, aging resulted in an increased percentage of γ -H2AX-positive cells in both *Cat1*^{-/-} or *Sod1*^{-/-} mice compared to WT control mice; this was most pronounced during stages V–VII, suggesting persistent DNA damage in *Cat1*^{-/-} and *Sod1*^{-/-} mice with advanced age (Fig. 6A). The DNA damage repair response occurs with the accumulation of 53BP1 at the site of γ -H2AX foci. The percentage of 53BP1-positive cells was highest in young WT mice; this was evident throughout all stages of spermatogenesis (Fig. 6B). Overall 53BP1 was reduced significantly in both young *Cat1*^{-/-} and *Sod1*^{-/-} mice; this was significant in stages I–IV, V–VII, and VIII–IX. In addition, aging resulted in a decrease in 53BP1-positive cells in all mice compared to young WT mice (Fig. 6B). Nevertheless, in aged *Cat1*^{-/-} and *Sod1*^{-/-} mice, significantly higher numbers of 53BP1-positive cells were evident in the later stages VIII–IX and X–XII.

The percentage of cells with colocalization of γ -H2AX and 53BP1, an indication of DNA damage repair, showed that repair was decreased in young *Cat1*^{-/-} and *Sod1*^{-/-} mice, which had fewer γ -H2AX- and 53BP1-positive cells (Fig. 6C). In addition, stage-specific identification of DNA damage repair was clear, with stages VIII–IX being the highest for colocalization, while stages X–XII displayed the lowest colocalization. Aging led to fewer cells with colocalized γ -H2AX and 53BP1 in all mice, across all spermatogenic stages, suggesting that WT, *Cat1*^{-/-}, and *Sod1*^{-/-} may have less DSB repair occurring in aged control and KO mice.

Increased Oxidative Stress-Induced DNA damage in the Spermatozoa of Sod1-/- Mice Regardless of Age

Immunofluorescence was used to determine the extent of 8-oxodG, an ROS-induced DNA damage lesion, in spermatozoa (Fig. 7A). Positive control samples pretreated with H₂O₂ had 100% 8-oxodG-positive spermatozoa. The lowest numbers of 8-oxodG-positive spermatozoa were found in young WT mice. The percentage of 8-oxodG-positive spermatozoa increased in WT mice as a result of aging (Fig. 7B). 8-OxodG-labeled

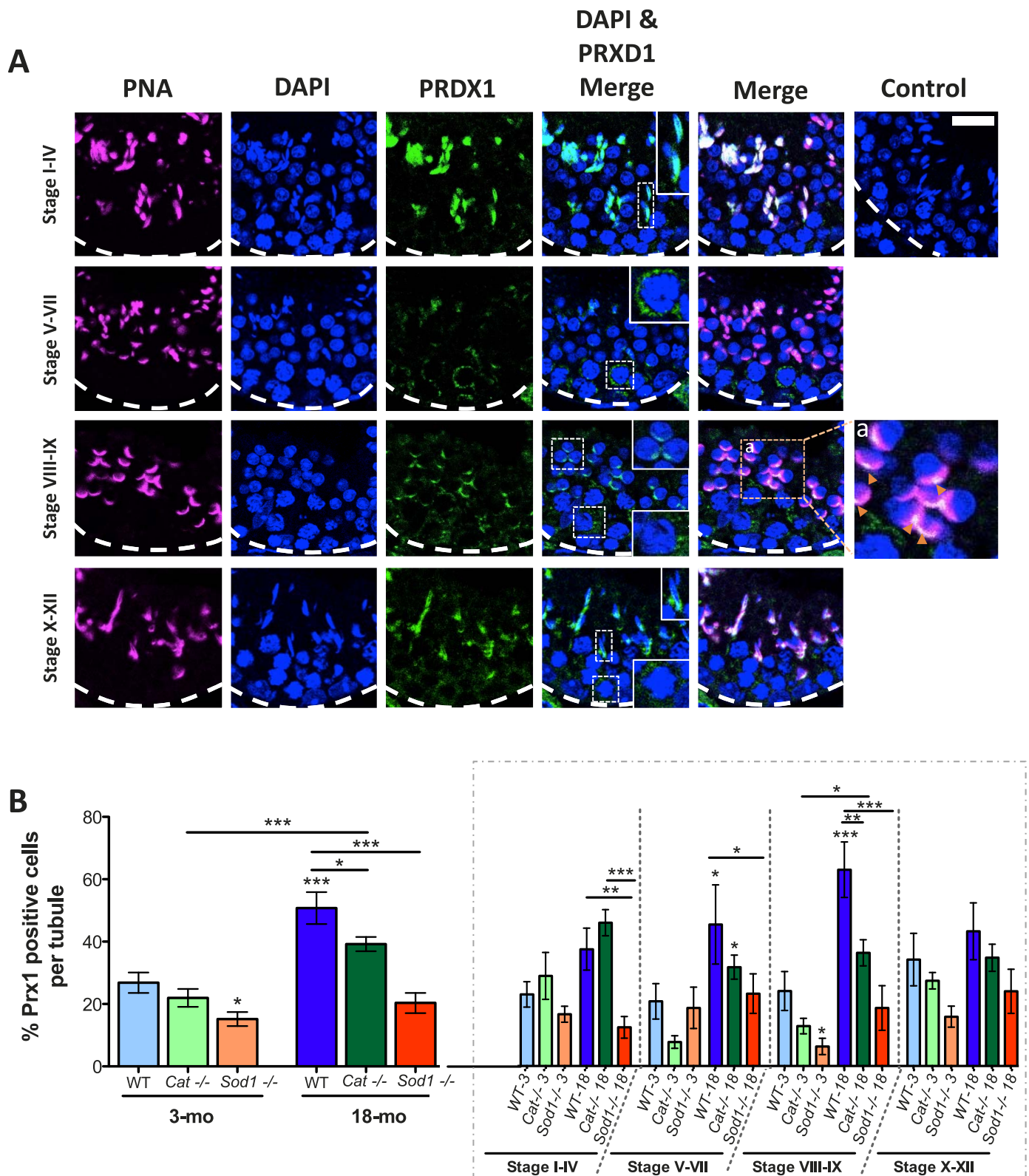


FIG. 5. Immunolocalization of PRDX1 to the developing acrosome and inhibition of the increase in PRDX1 with aging in *Sod*^{-/-} mice. PRDX1 was localized in the seminiferous tubules to the cytoplasm of spermatocytes (portions of the images showing this are magnified in insets) and within the acrosome of spermatids (A) (see insets). Imaris quantified percentage of PRDX1-positive cells per tubule showed increased PRDX1 with age in WT and *Cat*^{-/-} mice; however, *Sod*^{-/-} mice did not display this increase in PRDX1 with age (B). Mean values shown with error bars (\pm SEM); two-way ANOVA with Bonferroni test; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; $n = 5$. Bar = 20 μ m.

spermatozoa were significantly elevated in young *Cat*^{-/-} compared to young WT mice; however, the proportion of

these 8-oxodG lesions in *Cat*^{-/-} mice remained unaltered with aging. The *Sod*^{-/-} mice displayed the highest percentage of 8-

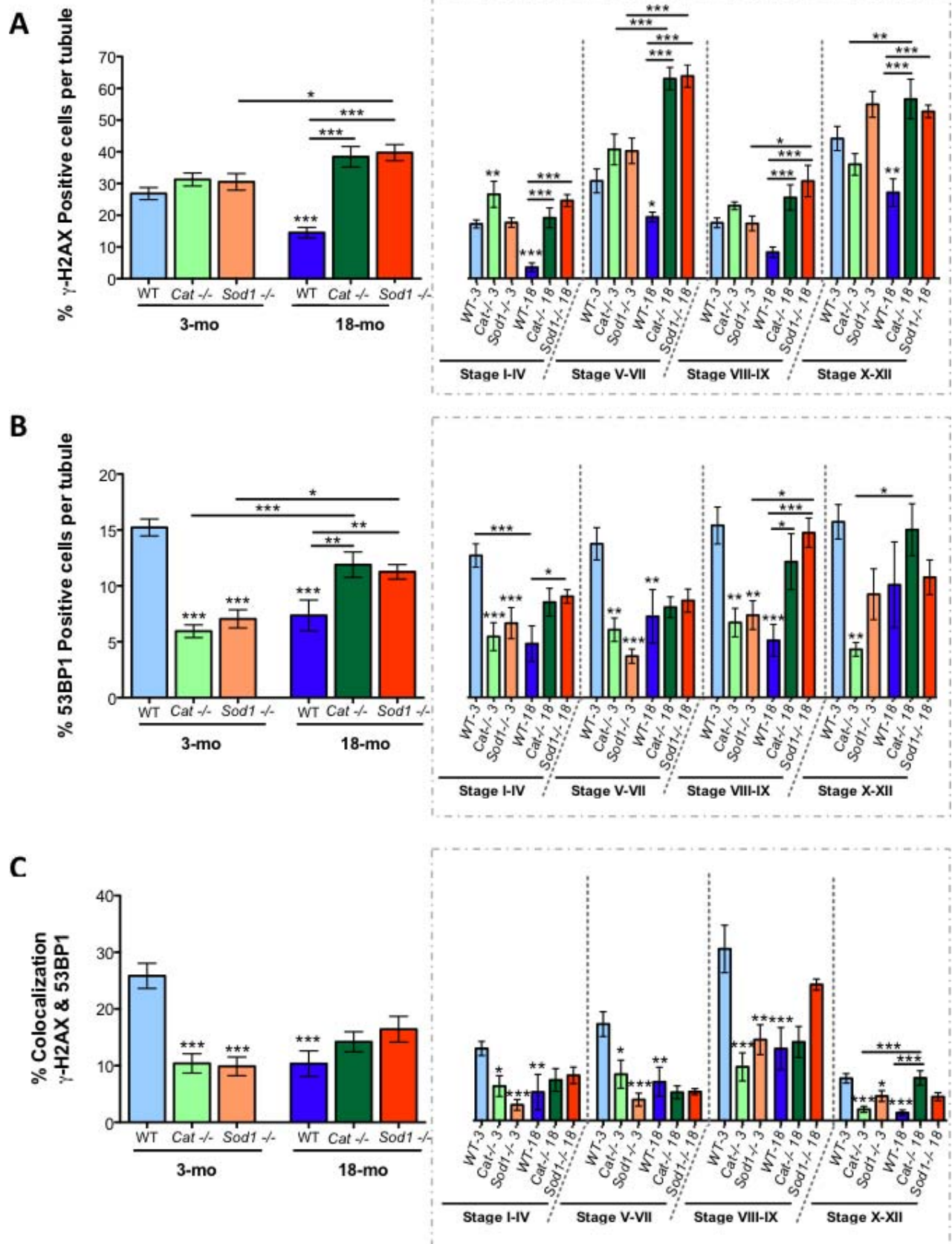


FIG. 6. Reduced DNA damage repair in seminiferous tubules of young *Cat*^{-/-} and *Sod1*^{-/-} mice and in all aged mice. The percentage of γ -H2AX-positive cells per seminiferous tubule increased with age in knockout (KO) mice (A). The percentage of 53BP1-positive cells per tubule was reduced in young KO mice and increased in aged KO mice, while the opposite was seen in aged WT mice (B). Colocalization of γ -H2AX- and 53BP1-positive cells showed decreased DNA damage repair in young KO mice and in all aged mice (C). Mean values shown with error bars (\pm SEM); two-way ANOVA with Bonferroni test; **P* < 0.05; ***P* < 0.01; ****P* < 0.001; *n* = 5.

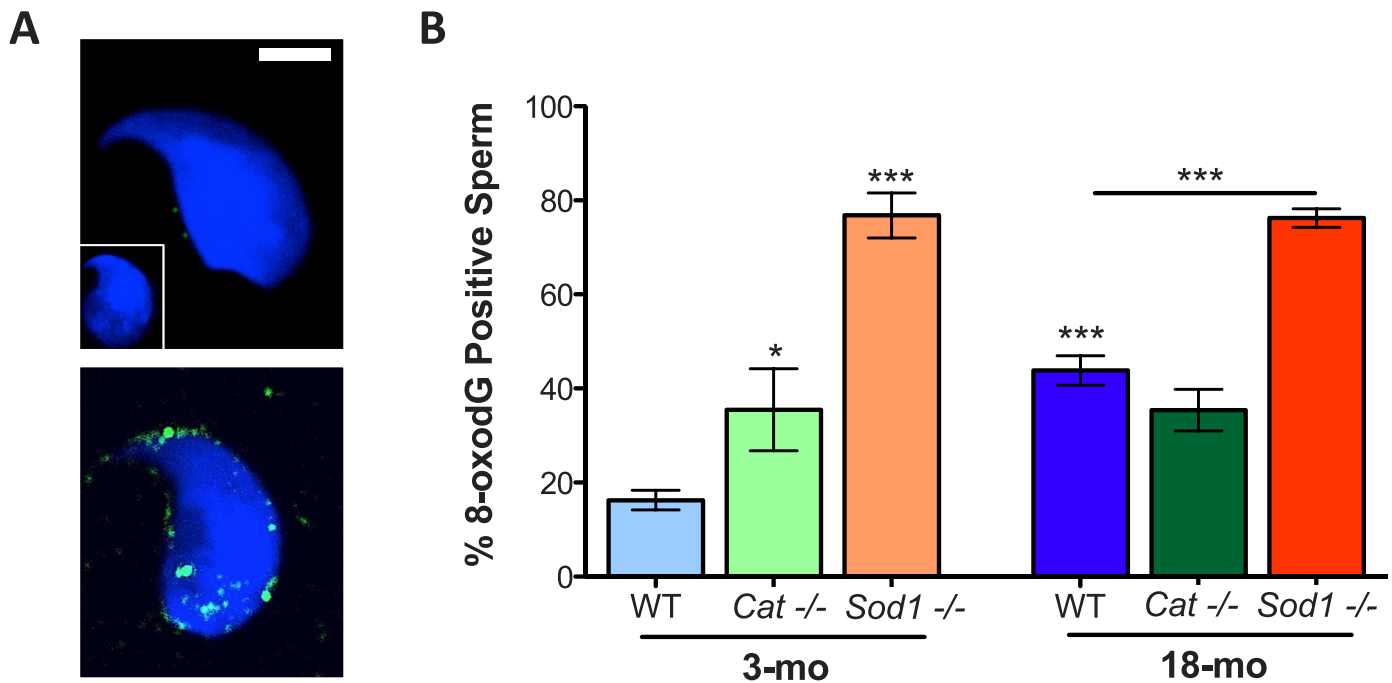


FIG. 7. Elevated oxidative stress-induced DNA damage in young and aged *Sod1*^{-/-} mice. Sample immunofluorescence images of sperm heads, with 8-oxodG (green) counterstained with DAPI (blue); inset shows negative control (A). The quantified percentage of 8-oxodG-positive sperm heads showed an increase in 8-oxodG with aging in WT and an increased 8-oxodG in *Cat*^{-/-} mice that remained at the same level with aging. The highest amounts of 8-oxodG were observed in *Sod1*^{-/-} mice in both the young and the aged groups. Mean values shown with error bars (\pm SEM); two-way ANOVA with Bonferroni multiple comparison test; * $P < 0.05$; *** $P < 0.001$; $n = 5$. Bar = 2.5 μ m.

oxodG-positive spermatozoa (approximately 80%) in both young and aged groups, showing the highest amount of ROS-induced damage regardless of age (Fig. 7B).

DISCUSSION

Aging is associated with the loss of antioxidants and increased ROS; previous studies have indicated that germ cells are sensitive to aging-associated oxidative stress [21, 25, 26]. To determine the consequences of losing key antioxidants, either CAT or SOD1, on maturing male germ cells, we bred and aged *Cat*^{-/-} and *Sod1*^{-/-} mice and examined the effects of aging on their developing testicular germ cells.

The original studies that had generated these null mutant (*Cat*^{-/-} and *Sod1*^{-/-}) mice found them to be viable and phenotypically normal despite the lack of these major antioxidant enzymes [34, 35]. Later studies examining the lack of SOD1 revealed increased vulnerability to ROS and also displayed a 30% reduction in lifespan [45–47]; these are highly significant findings in the context of aging studies. Similar to recent studies [48, 49], our examination of the overall health of these animals revealed that the loss of either antioxidant, CAT or SOD1, resulted in mice weighing less than their WT littermates at both 3 and 18 mo of age. These results were not surprising considering that CAT imparts various metabolic health benefits and SOD1 is a major cytoplasmic superoxide scavenger with minimal compensatory mechanisms [50–52]; thus, the global and chronic loss of these antioxidants could contribute to the overall decrease in the weights of these mice compared to WT mice throughout their lives.

Although it is natural to hypothesize that the loss of an antioxidant will result in accelerated aging, the results of systemic oxidative stress are difficult to predict. The effects of compensatory mechanisms and chronic stress, which are observed with aging, make predicting the effects of antioxidant

null mutations particularly difficult. Previous studies found both *Cat*^{-/-} and *Sod1*^{-/-} mice to be fertile [34, 35]. However, many null mutation mouse studies have reported male mice to be fertile and overlooked the possible effects on sperm counts and testicular histology. Mice can have very large reductions in their sperm counts [53, 54] and reduced Sertoli cell numbers [54, 55] and yet be fertile. Our data on breeding indicate that despite a clear reduction in the number of sperm in young *Sod1*^{-/-} mice, these males show no difference in fertility (number of pups per litter). Sperm counts in all mice declined with aging. In addition, aging reduced fertility in all groups of mice, with significantly fewer pups per litter sired by 12-month-old males. The 12-month-old *Sod1*^{-/-} mice displayed the most severe reduction in pup numbers.

Testis weights were consistently lower in KO mice compared to WT mice, while epididymal weights were unaltered. The histology of aged testes displayed the typical presentations associated with aging, including loss of germ cells and shrunken seminiferous tubules [56]. Further analyses revealed that, in addition to the loss of germ cells, Sertoli cell numbers were reduced in young *Sod1*^{-/-} mice and all aged mice.

Epididymal histology revealed that the most severely affected epididymides were in aged *Sod1*^{-/-} mice; all segments of the epididymides showed degenerated epithelia and cellular debris in the lumina. The epididymides from aged *Sod1*^{-/-} mice also had few or no visible spermatozoa (Supplemental Figure S2), providing corroborating evidence of the significant decline in spermatozoa counts and supporting the reduced fertility observed in *Sod1*^{-/-} mice. SOD1 is expressed in the epithelium of the epididymis from the caput to the cauda [57], accompanied by an extracellular secreted SOD (SOD3) [58] as well as several other antioxidants, including GPX1, 3, 4, and 5 and GSH [59]. Aging decreases antioxidants GPX4 and 5 in the epididymis [57]; the lack of SOD1 in *Sod1*^{-/-} mice could further exacerbate the redox dysfunction occurring with aging.

Future investigations will address changes in other antioxidants and delve into the mechanisms contributing to the advanced *Sod*^{-/-} epididymal degeneration that was observed with aging.

The spermatocytes of aged WT and *Sod*^{-/-} mice were least able to cope with elevated ROS after mice were administered SIN-10 as an oxidative insult. As seen with previous studies on the *Cat*^{-/-} mice [34, 60], the lifelong absence of CAT could lead to the up-regulation of compensatory antioxidant systems. The protective effects observed in the *Cat*^{-/-} Sertoli cell number with aging could be attributed to compensatory antioxidants, such as PRDX1. These compensatory mechanisms are demonstrated in the spermatocytes of aged *Cat*^{-/-} mice, which are able to detoxify SIN-10-induced ROS, suggesting that similar mechanisms may be protecting Sertoli cell numbers with aging in these animals.

Currently, there are no data to suggest that compensation could be occurring in *Sod*^{-/-} mice. One particular antioxidant that is affected by both aging and oxidative stress and is linked to male infertility is PRDX1 [29]. As previously reported, PRDX1 was localized to Leydig cells [42]; however, we also found it localized to the cytoplasm of spermatocytes and developing acrosome of spermatids. We found that PRDX1 was altered with aging and reduced in the germ cells of null mutant mice. More specifically, aging resulted in an increase in PRDX1 in the germ cells of WT and *Cat*^{-/-} mice, confirming the hypothesized genetic robustness in the *Cat*^{-/-} mice model [61, 62]. In *Sod*^{-/-} mice, PRDX1 was consistently at lower levels than WT and was not increased with aging. This lower PRDX1 in *Sod*^{-/-} mice points toward redox dysfunction as the primary cause of increased vulnerability in the *Sod*^{-/-} mice. More specifically, the low levels of PRDX1 in the germ cells of aging *Sod*^{-/-} mice suggest that compensatory removal of radicals was disrupted and that the normal expression of antioxidant was altered. Furthermore, the ability of high levels of ROS to hyperoxidize antioxidants, in particular PRDXs [29, 63], contributes to the hypothesis that antioxidants may become catalytically inactive during aging [29, 63, 64]. While antioxidant enzymatic activity needs to be examined in the aging testis of these mice, the present findings indicate that the loss of SOD1 may result in increased redox dysfunction in the germ cells of aging mice.

Although previous studies have confirmed glutathione peroxidase 1 (GPX1) activity to be unaltered between WT and *Cat*^{-/-} mice [33] and in *Sod*^{-/-} mice [26], it remains possible that the compensatory mechanisms observed between *Cat*^{-/-} may be due to the thioredoxin-glutaredoxin system [24]. Further studies are required to clarify this point and to determine the effects of aging on the thioredoxin-glutaredoxin antioxidants in these KO mice.

Erythrocytes are constantly under oxidative stress; however, the absence of CAT and GPX1 does not lead to hemolytic anemia due to damage and premature death of erythrocytes [65, 66]. The only antioxidant KO that did display this vulnerability was the PRDX1 KO [67], leading our investigation toward PRDX1. The altered PRDX1 levels with aging and in the KO animals suggests that although PRDXs are not considered to be primary enzymatic scavengers, they can act in compensatory ROS removal with aging.

Next, we decided to examine DNA damage and repair markers in the seminiferous tubules of these animals. The DNA DSB marker γ -H2AX increased with aging in animals with null mutations, indicating the presence of elevated levels of DSBs in the germ cells of both aged *Cat*^{-/-} and *Sod*^{-/-} mice. Normally, in response to this damage, elevated DNA damage-repair protein 53BP1 accumulates at the site of DSBs. 53BP1 accumulation at the site of γ -H2AX leads to downstream

signaling for DNA damage repair, with NHEJ repair promoted by 53BP1. We speculate that DNA damage repair mechanisms could be compromised in the young KO animals, which display fewer 53BP1-positive cells. Our results indicate that with aging, additional oxidative stress induces increased γ -H2AX in these animals and results in increased 53BP1 accumulation to promote signaling for DNA damage repair. Moreover, the colocalization of γ -H2AX and 53BP1 revealed active repair in young WT mice but reduced DNA repair in all other mice. In support of these findings, recent studies with models of controlled DNA DSBs suggest that DNA damage itself can contribute to deregulation of genes during aging [68, 69].

Finally, our examination of 8-oxodG, a DNA lesion that is directly induced by oxidative stress, corroborated previous observations that 8-oxodG significantly increased in spermatozoa with aging [60]. In addition, these results suggested that the greatest oxidative damage occurred in the spermatozoa of both young and aged *Sod*^{-/-} mice. In addition, this damage was not repaired in *Sod*^{-/-} mice during spermatogenesis. While damage was detected in young *Cat*^{-/-} mice, this damage was not further increased by aging, suggesting that while loss of CAT increased DNA damage in spermatozoa of young animals, aging did not further contribute to this oxidative stress-induced damage. These findings reinforce the importance of maintaining SOD1 to prevent irreversible oxidative stress induced damage [11, 45], leading us to postulate that in the absence of SOD1, ONOO⁻ could be the species of radicals inflicting excessive damage in *Sod*^{-/-} spermatozoa.

In conclusion, the present study displays the complexity of the antioxidant systems and their differential effects on the quality of aging male germ cells. The loss of CAT resulted in loss in germ cells but no loss in Sertoli cells in young mice. Furthermore, spermatocytes from aged *Cat*^{-/-} mice displayed an enhanced ability to detoxify ROS following SIN-10 treatment. Aged *Cat*^{-/-} mice had significantly increased levels of PRDX1, indicating that the up-regulation of this antioxidant may have increased ROS removal capabilities. However, the levels of DNA damage in the spermatozoa from young animals were still high in *Cat*^{-/-} mice. The loss of SOD1 was markedly detrimental when compared to CAT loss. The negative consequences of SOD1 loss, in young and aged mice, were evident through reduced sperm counts; abnormal epididymal histology, significant germ cell and Sertoli cell loss, and increased spermatozoa DNA damage were observed. However, the effects on fertility were manifested only with aging. Finally, the inability of spermatocytes from *Sod*^{-/-} mice to remove ROS following SIN-10 demonstrated the loss of redox capabilities, while the failure of *Sod*^{-/-} mice to up-regulate PRDX1 with aging identified redox dysfunction.

In conclusion, the results of these studies may have clinical implications in relation to sperm selection for artificial reproductive technologies. The present study suggests that SOD1-mimicking compounds may represent a critical component of potential antioxidant supplements for infertile/subfertile men and aspiring aging fathers who have spermatozoa with high levels of oxidative damage.

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