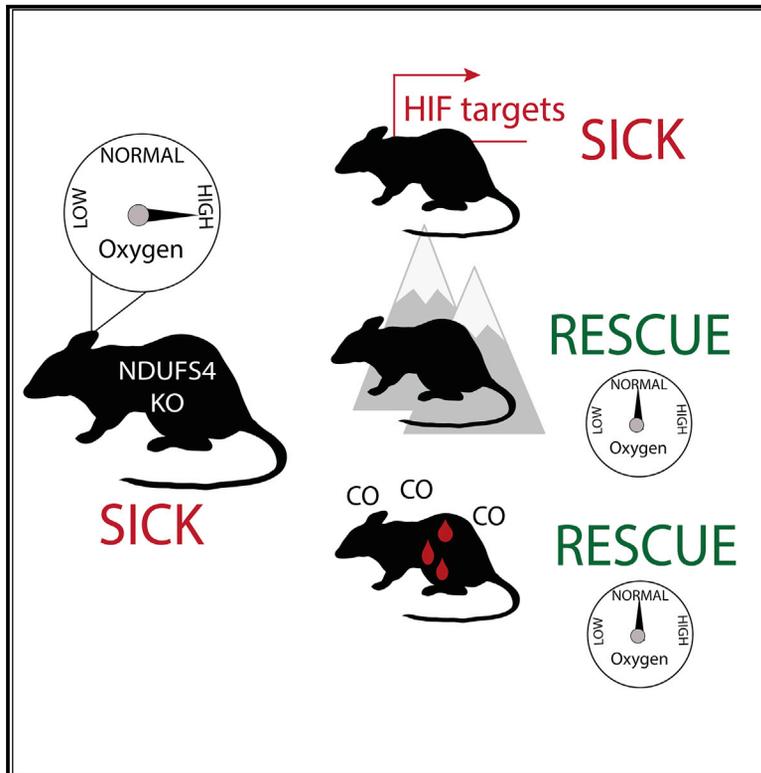


# Cell Metabolism

## Leigh Syndrome Mouse Model Can Be Rescued by Interventions that Normalize Brain Hyperoxia, but Not HIF Activation

### Graphical Abstract



### Authors

Isha H. Jain, Luca Zazzeron, Olga Goldberger, ..., Fumito Ichinose, Warren M. Zapol, Vamsi K. Mootha

### Correspondence

wzapol@mgh.harvard.edu (W.M.Z.), vamsi@hms.harvard.edu (V.K.M.)

### In Brief

Leigh syndrome is a severe mitochondrial disorder. Here, Jain et al. show, in a mouse model of the disease, that excess oxygen in the brain is a likely cause of tissue damage and that distinct interventions that reduce oxygen delivery to the brain can prevent and even reverse the neurological disease, likely by normalizing tissue hyperoxia.

### Highlights

- The *Ndufs4* KO mouse model of mitochondrial Leigh syndrome exhibits brain hyperoxia
- Genetically activating the hypoxia transcriptional response is not beneficial
- CO treatment and anemia reverse disease by decreasing oxygen delivery



# Leigh Syndrome Mouse Model Can Be Rescued by Interventions that Normalize Brain Hyperoxia, but Not HIF Activation

Isha H. Jain,<sup>1,2,3,9,10</sup> Luca Zazzeron,<sup>4,10</sup> Olga Goldberger,<sup>1,2,3</sup> Eizo Marutani,<sup>4</sup> Gregory R. Wojtkiewicz,<sup>5</sup> Tsilil Ast,<sup>1,2,3</sup> Hong Wang,<sup>1,2,3</sup> Grigorij Schleifer,<sup>4</sup> Anna Stepanova,<sup>6</sup> Kathleen Brepoels,<sup>7,8</sup> Luc Schoonjans,<sup>7,8</sup> Peter Carmeliet,<sup>7,8</sup> Alexander Galkin,<sup>6</sup> Fumito Ichinose,<sup>4</sup> Warren M. Zapol,<sup>4,\*</sup> and Vamsi K. Mootha<sup>1,2,3,11,\*</sup>

<sup>1</sup>Department of Molecular Biology and Howard Hughes Medical Institute, Massachusetts General Hospital, Boston, MA, USA

<sup>2</sup>Department of Systems Biology, Harvard Medical School, Boston, MA, USA

<sup>3</sup>Broad Institute of Harvard and MIT, Cambridge, MA, USA

<sup>4</sup>Department of Anesthesia, Critical Care, and Pain Medicine, Massachusetts General Hospital, Boston, MA, USA

<sup>5</sup>Center for Systems Biology, Massachusetts General Hospital and Harvard Medical School, Boston, MA, USA

<sup>6</sup>Department of Pediatrics, Division of Neonatology, Columbia University, New York, NY, USA

<sup>7</sup>Laboratory of Angiogenesis and Vascular Metabolism, VIB-KU Leuven, Center for Cancer Biology, Leuven, Belgium

<sup>8</sup>Laboratory of Angiogenesis and Vascular Metabolism, Department of Oncology and Leuven Cancer Institute (LKI), KU Leuven, Leuven, Belgium

<sup>9</sup>Present address: Department of Physiology, University of California, San Francisco, San Francisco, CA, USA

<sup>10</sup>These authors contributed equally

<sup>11</sup>Lead Contact

\*Correspondence: [wzapol@mgh.harvard.edu](mailto:wzapol@mgh.harvard.edu) (W.M.Z.), [vamsi@hms.harvard.edu](mailto:vamsi@hms.harvard.edu) (V.K.M.)

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## SUMMARY

Leigh syndrome is a devastating mitochondrial disease for which there are no proven therapies. We previously showed that breathing chronic, continuous hypoxia can prevent and even reverse neurological disease in the *Ndufs4* knockout (KO) mouse model of complex I (CI) deficiency and Leigh syndrome. Here, we show that genetic activation of the hypoxia-inducible factor transcriptional program via any of four different strategies is insufficient to rescue disease. Rather, we observe an age-dependent decline in whole-body oxygen consumption. These mice exhibit brain tissue hyperoxia, which is normalized by hypoxic breathing. Alternative experimental strategies to reduce oxygen delivery, including breathing carbon monoxide (600 ppm in air) or severe anemia, can reverse neurological disease. Therefore, unused oxygen is the most likely culprit in the pathology of this disease. While pharmacologic activation

of the hypoxia response is unlikely to alleviate disease *in vivo*, interventions that safely normalize brain tissue hyperoxia may hold therapeutic potential.

## INTRODUCTION

Mitochondrial disorders comprise a large collection of individually rare, inborn errors of metabolism, with an estimated incidence of at least 1 in 4,300 births (Gorman et al., 2015). The most severe forms of disease occur because of mutations in structural components or assembly factors of the electron transport chain (ETC) and ATP synthase (CV). Of these disorders, Leigh syndrome is the most common pediatric manifestation of mitochondrial diseases (Lake et al., 2015). Mutations in over 75 distinct genes can underlie this syndrome, with patients first showing symptoms at any stage from birth to childhood (Lake et al., 2015). Patients often develop seizures, failure-to-thrive, ataxia, developmental delay, and clinical regression, ultimately leading to death, typically within the first few years of life. Diagnosis of Leigh syndrome is based on the clinical presentation

### Context and Significance

Leigh syndrome is a devastating, inherited mitochondrial disease characterized by marked neurological and metabolic deficits and early death. At present, there are no effective, approved therapies. Here, the authors show, in a mouse model, that excess, unused oxygen plays a key role in the pathology of Leigh disease. Multiple interventions that decrease oxygen delivery to the brain can normalize the elevated oxygen concentration and reverse the disease. While many pathologies are associated with reduced tissue oxygen (e.g., stroke and heart attack), this appears to be one of the first associated with excessive oxygen. These findings suggest a new mechanism of action for mitochondrial disease and may aid in the design of future therapies based on the control of oxygen delivery.



and radiographic findings of bilaterally symmetric lesions on T2-weighted brain MRI. While significant advances have been made in defining their genetic basis, there are no generic treatments with proven efficacy in these patients. Recently, we demonstrated that chronic, continuous normobaric exposure to hypoxia (11% FiO<sub>2</sub>) can prevent and even reverse the neurodegenerative brain disease in the *Ndufs4* mouse model of Leigh syndrome, not only extending lifespan, but also normalizing body weight, body temperature, disease biomarkers, and improving behavioral performance (Ferrari et al., 2017; Jain et al., 2016).

The initial motivation for testing hypoxia as a therapy arose from our cell culture experiments, where we found that artificially activating the vHL-PHD-HIF hypoxia transcriptional program was sufficient to rescue cell proliferation defects caused by ETC inhibition (Jain et al., 2016). However, the *in vivo* therapeutic benefit was achieved by exposing *Ndufs4* KO mice with a partial complex I deficiency to breathing chronic, normobaric 11% O<sub>2</sub>. In subsequent work, we showed that neither intermittent hypoxia nor more moderate hypoxia (17% O<sub>2</sub>) could rescue the neurodegenerative disease (Ferrari et al., 2017). Moreover, breathing 55% O<sub>2</sub>, which is harmless in wild-type mice, was extremely toxic for *Ndufs4* KO mice, leading to death within days, suggesting the possibility of HIF-dependent and independent mechanisms.

The aim of this study was to explore the sufficiency of the hypoxia transcriptional response in the hypoxic rescue of disease, as well as the role of brain oxygen levels in pathogenesis. We report that genetic activation of the canonical vHL-PHD-HIF hypoxia response is insufficient to prevent disease in the *Ndufs4* KO mouse. Rather, we find that oxygen consumption is impaired and brain oxygen tissue levels increase in KO mice, likely resulting in increased oxygen toxicity. We show that chronic hypoxic breathing normalizes brain PO<sub>2</sub> in *Ndufs4* KO mice, and that other diverse interventions that reduce brain oxygen levels are effective at preventing neurological disease in this mouse model.

## RESULTS

### Activation of the Hypoxia Response Pathway Is Not Sufficient to Rescue Disease

We had originally performed a genome-wide CRISPR screen and identified activation of the hypoxia transcriptional response as being protective in cell culture models of mitochondrial disease. However, we never tested whether activation of this pathway is sufficient to rescue the *Ndufs4* KO mouse model. A major component of the cellular and organismal hypoxia response is mediated by the hypoxia-inducible transcription factors (HIFs). In well-oxygenated environments, the prolyl hydroxylase enzymes (PHDs) use molecular oxygen as a substrate and hydroxylate the HIF transcription factors (Wang et al., 1995; Majmundar et al., 2010; Ivan et al., 2001). It is this hydroxylated form that is recognized by the E3-ligase, von Hippel Lindau (vHL) factor and targeted for proteasomal degradation. There are three known PHD enzymes involved in oxygen sensing, PHD1-3. PHD2 inhibition is typically associated with HIF activation. However, PHD1 and PHD3 have also been reported to have redundant roles in signaling to the HIF transcriptional program. There are also two isoforms of HIF—HIF1 $\alpha$  and HIF2 $\alpha$ . While these transcription fac-

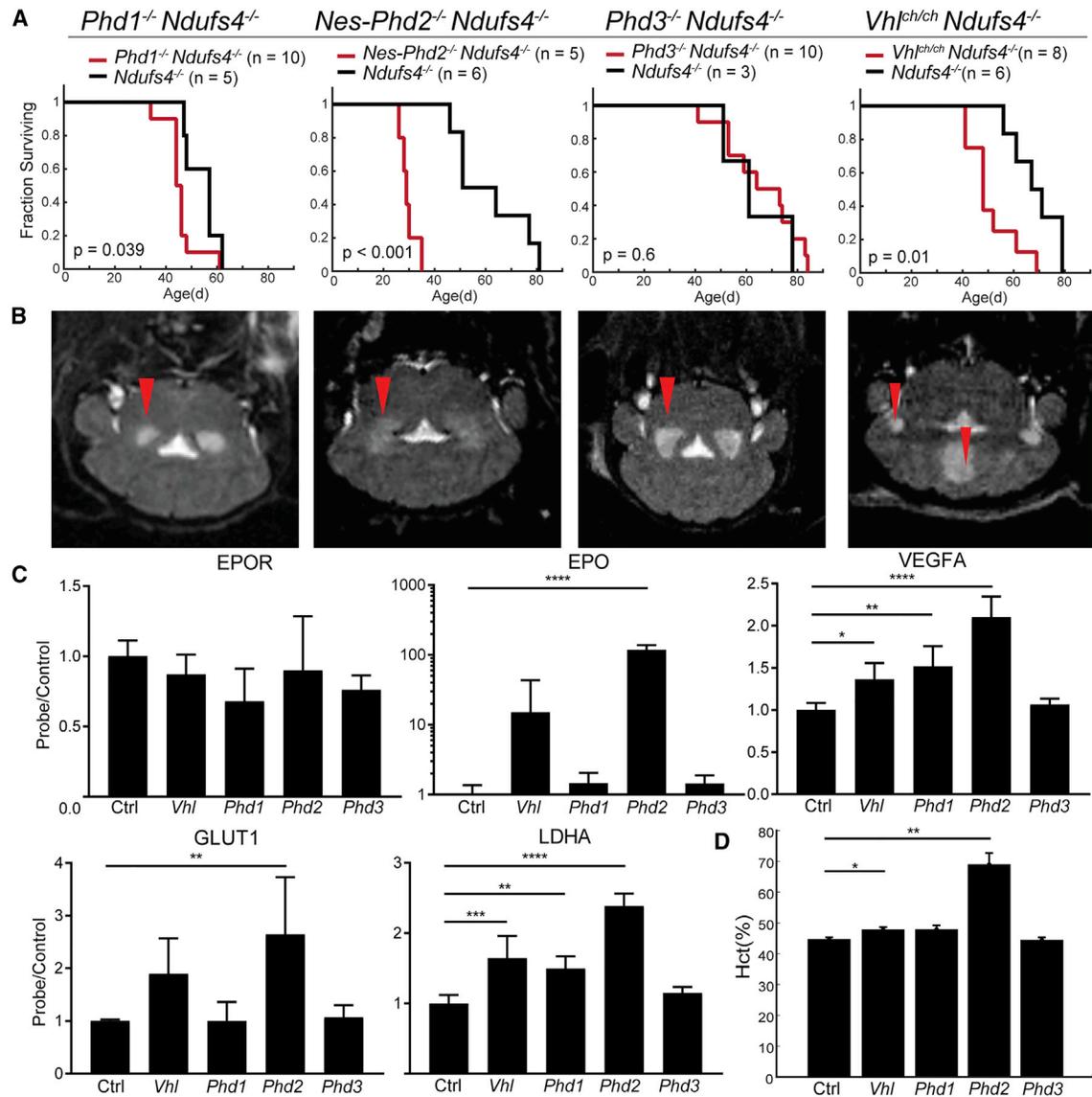
tors have partially overlapping roles, HIF1 $\alpha$  is thought to primarily regulate cellular hypoxia adaptations (e.g., increasing glycolysis, modifications of the tricarboxylic acid cycle [TCA]) (Majmundar et al., 2010). HIF2 $\alpha$  is more relevant for whole-body physiological adaptations such as erythropoiesis and angiogenesis. In mouse models, genetic disruption of the PHD enzymes or vHL activate the hypoxia transcriptional program and have been previously utilized to explore the therapeutic potential of targeting this pathway in a range of conditions, including stroke and hindlimb ischemia (Quaegebeur et al., 2016). Here, we used the same genetic models to assess whether activating the HIF transcriptional program is sufficient to rescue neurological disease in the *Ndufs4* KO mouse model.

We crossed the *Ndufs4* KO strain with mice lacking *Phd1*, *Nestin-Phd2*, *Phd3*, or the Chuvash mouse harboring a point mutation in *Vhl* (*Vhl*<sup>ch/ch</sup>) (Quaegebeur et al., 2016; Hickey et al., 2007). We asked whether the hypoxia response was indeed activated in the different genetic models of the hypoxia transcriptional program. The *Phd2* strain consistently showed elevation of canonical HIF targets in the cerebellum, where disease pathology eventually appears (Figure 1C). For example, Epo expression was increased 100x in this strain. Moreover, *Ldha* and *Vegfa* transcripts were similarly elevated in *Phd1*, *Phd2*, and *Vhl* mice. *Phd3* mice did not show a canonical HIF response in the cerebellum. These findings are consistent with previous reports that *Phd2* and vHL are more direct mediators of the HIF response. Furthermore, HIF2 $\alpha$  activation results in elevated hematocrit levels. In our hands, only *Nestin-Phd2* mice had an elevated hematocrit (>60%, Figure 1D).

Notably, none of these genetic crosses were sufficient to prevent disease. Overall survival of double KOs was equivalent or even worse than *Ndufs4*<sup>-/-</sup> mice (Het or WT for *Phd/Vhl* genes) (Figure 1A). Specifically, *Ndufs4* KO mice that were lacking *Phd1* had a median survival of 45 days versus control *Ndufs4* KO mice with a median survival of 57 days (matched for genetic background). Even more strikingly, *Ndufs4*<sup>-/-</sup>; *Nestin-Phd2*<sup>-/-</sup> had a median survival of 29 days versus matched *Ndufs4* KO controls which had a median survival of 64 days. *Ndufs4*<sup>-/-</sup>; *Phd3*<sup>-/-</sup> survived for a median of 69 days versus matched *Ndufs4*<sup>-/-</sup> with median survival of 61 days. Finally, *Ndufs4*<sup>-/-</sup>; *Vhl*<sup>ch/ch</sup> lived to 48 days versus 69 days for matched *Ndufs4*<sup>-/-</sup> mice. Notably, loss of *Phd2* or *Vhl* robustly led to HIF activation, and these same strains were particularly sensitive to complex I deficiency. Moreover, all four genetic crosses resulted in double KOs that developed lesions in the vestibular nucleus (and additional cerebellar lesions in the *Vhl* cross) as imaged by T2-weighted MRI (Figure 1B). These genetic crosses indicate that artificially activating the HIF response is insufficient to rescue disease. Thus, small molecule inhibitors of the PHD enzymes or vHL enzyme are predicted to be ineffective *in vivo*, and in fact, could be toxic, in the context of a whole organism with complex I deficiency and Leigh syndrome.

### Impaired Whole-Body Oxygen Consumption and Brain Hyperoxia in *Ndufs4* KO Mice

Mitochondria consume 90% of the oxygen in the body (Rich, 2003). Thus, it is possible that genetic defects in the electron transport chain will have a secondary impact on tissue oxygenation. In fact, venous hyperoxia has been documented in



### Figure 1. Genetic Activation of the Hypoxia Response Is Insufficient to Prevent Leigh Disease

(A) Kaplan-Meier curves for double KO mice relative to single *Ndufs4*<sup>-/-</sup> KO mice. Log-rank p values and sample size within insert. Single KO mice are homozygous-null for *Ndufs4* locus and WT or Het for other genetic locus of given cross.

(B) MRI of vestibular nucleus (red arrow for *Phd1*, *Phd2* and *Phd3* crosses) of late-stage disease mice for each genotype. In *Vhl* cross, red arrow corresponds to lateral lesions (dentate nucleus) and midline lesion (central lobule, cerebellum).

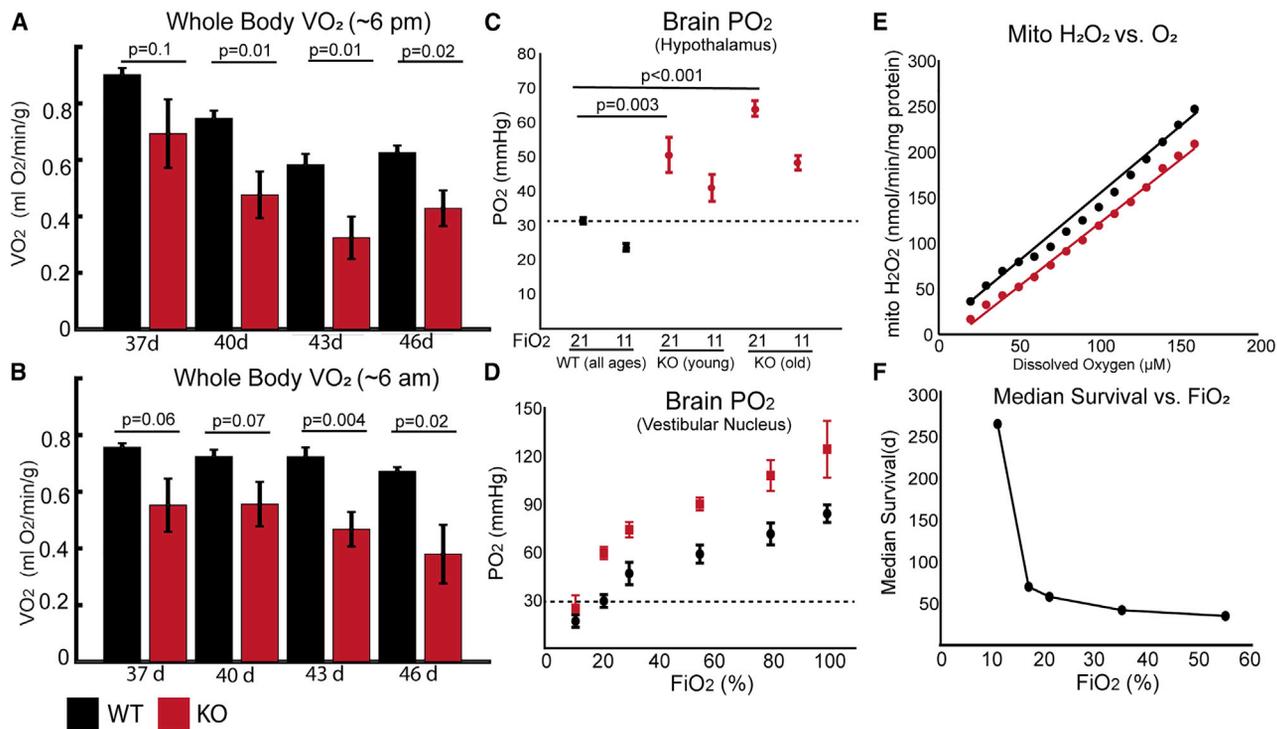
(C) qPCR for canonical HIF targets (mean  $\pm$  SD, n = 4 per group). Mice are homozygous for given genotype and WT or Het for *Ndufs4* genotype.

(D) Hematocrit of control or *Phd* and *Vhl* genetic models (WT or Het for *Ndufs4*) (mean  $\pm$  SEM, Ctrl [n=3], *Vhl* [n=6], *Phd1* [n=3], *Phd2* [n=4], *Phd3* [n=5]). One-tailed unpaired t test p value shown for genetic crosses relative to control mice. (\*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001, \*\*\*\*p < 0.0001 by one-way ANOVA, with Dunnett's test for multiple comparison).

patients with mitochondrial disease and has also been proposed to be a diagnostic test in patients with mitochondrial myopathy (Taivassalo et al., 2002). We monitored whole body and cerebral oxygenation in WT and *Ndufs4* KO animals. We continuously measured whole body oxygen consumption in WT and KO animals, as a function of age. At ~37 days of age, whole body oxygen consumption (normalized for total body weight) was not significantly different between KO and WT animals. However, in awake mice this parameter decreased in KO animals as a function of age, reaching nearly half the value of WT animals at

~45 days of age (Figure 2A). Thus, it appears that the decline in whole body oxygen consumption correlates with the overt disease phenotype until death at ~60 days of age. This difference is observed during both active and inactive periods of the murine day (Figure 2B).

We reasoned that such overt defects in oxygen consumption might affect brain tissue PO<sub>2</sub>. We used two different types of probes for measuring the partial pressure of oxygen in the brain—an amperometric Clark electrode, and an optical probe whose phosphorescence is sensitive to oxygen levels (see



**Figure 2. Whole-Body and Brain Oxygenation of KO and WT Mice**

(A and B) Whole-body oxygen consumption (mean SEM), normalized for body weight, during (A) active hours or (B) inactive hours. Data smoothed over 70 collection points (~5.5 h). Values normalized to body weight at each collection point ( $n = 6$  per group). Unpaired t test p value shown for each genotype comparison at each day.

(C) Tissue  $PO_2$  (mean  $\pm$  SEM) measured stereotaxically in hypothalamus of WT and KO mice as a function of age using a Clark electrode (young = 36–41 days, old 50–54 days).  $n = 6$  for WT,  $n = 5$  for young KO and  $n = 5$  for old KO. Dotted line represents  $PO_2$  of WT mice breathing room air. Unpaired t test p value shown for comparison of KOs to WT breathing room air.

(D) Tissue  $PO_2$  (mean  $\pm$  SD) measurement using an optical  $PO_2$  probe stereotaxically placed near vestibular nuclei of WT and KO mice breathing varying oxygen tensions ( $n = 4$ –6 per group). Dotted line represents  $PO_2$  of WT mice breathing room air.

(E) Hydrogen peroxide released during complex I-mediated respiration by isolated mitochondria from WT and KO brains as a function of dissolved oxygen.

(F) Median survival of KO mice as a function of inhaled oxygen, using data from this paper and from previously published studies (Ferrari et al., 2017; Jain et al., 2016).

**STAR Methods**). Using a Clarke electrode, we stereotaxically monitored cerebral  $PO_2$  in the hypothalamus (as a representative brain region that was easily accessible) of anesthetized KO and WT mice as a function of age (Figure 2C). In 30-day-old KO mice, the partial pressure of oxygen ( $PO_2$ ) in brain tissue was significantly higher than in WT mice (51 versus 31 mmHg,  $p = 0.003$ ). Remarkably, KO mice showed an age-dependent increase in brain tissue  $PO_2$ , reaching 64 mmHg at 40–50 days of age ( $p < 0.001$  versus WT mice and  $p = 0.04$  versus 30-day-old KO mice). Once again, age-dependent changes in brain tissue  $PO_2$  appear to correlate with the disease severity in KO mice. We then asked how breathing various oxygen concentrations translated into changes in brain  $PO_2$ . We monitored brain  $PO_2$  in the vestibular nuclei of anesthetized KO and WT mice while  $FiO_2$  was increased from 11% to 100%. These measurements were made using an optical probe. For any given  $FiO_2$ , brain  $PO_2$  was significantly higher in late-stage KO mice versus age-matched WT mice. Indeed, 11%  $FiO_2$  in KO mice resulted in an equivalent brain  $PO_2$  as 21%  $FiO_2$  in WT mice (Figure 2D).

We determined the dose-response relationship between  $FiO_2$  and survival of *Ndufs4* KO mice. Combining previously published data (Ferrari et al., 2017) with newly generated sur-

vival curves, we show a remarkable drop-off in survival at 11%–13%  $FiO_2$  (Figure 2F). Survival between 17% and 55%  $FiO_2$  also shows a clear worsening in the disease state. Therefore, it is likely that localized and age-dependent changes in brain  $PO_2$  result in tissue hyperoxia, which directly causes brain tissue damage.

Previous studies have demonstrated a strong dependence of isolated mitochondrial superoxide production on local  $O_2$  concentrations (Kussmaul and Hirst, 2006; Stepanova et al., 2019). We asked whether this relationship varied between KO and WT tissue. We collected forebrain and midbrain tissue from ~60-day-old mice from each genotype, isolated mitochondria, and assessed hydrogen peroxide production as a function of local oxygen tension. We modulated oxygen concentration in the medium with respiring mitochondria while simultaneously measuring the resulting  $H_2O_2$  production. Using malate and pyruvate as a substrate combination for complex I, we find that this relationship is comparable in KO and WT mice (Figure 2E). Therefore, for a given oxygen tension, mitochondria from both genotypes produce an equivalent amount of  $H_2O_2$ . In principle, the increased brain  $PO_2$  observed in KO mice is predicted to provide a higher rate of ROS production at a fixed  $FiO_2$ .

Collectively, these findings suggest that impaired complex I activity in aged KO mice results in relative tissue hyperoxia, presumably due to a mismatch between oxygen delivery and its utilization. Future work is needed to understand how this elevated oxygen pressure results in the observed neuroinflammatory signature and neuronal cell death.

### Chronic Hypoxia Normalizes Brain $PO_2$ in *Ndufs4* KO Mice

In the context of acute hypoxia exposure, we observed a decrease in brain  $PO_2$  in both WT and KO mice (Figure 2C). We asked whether a similar effect would be observed in mice chronically exposed to 11% oxygen. During chronic hypoxia a compensatory increase in Hb concentration and ventilation may preserve oxygen delivery to the brain with no change in brain  $PO_2$ . We exposed both WT and KO mice to 11%  $O_2$  and we measured brain  $PO_2$  after three weeks. All hypoxic mice had a lower brain  $PO_2$  as compared to mice breathing 21%  $O_2$  with the same genotype (17 mmHg versus 30 mmHg,  $p = 0.0007$  in WT and 29 versus 64 mmHg,  $p < 0.0001$  in KO, Figure 2D). As we observed in the acute setting, chronic 11%  $O_2$  exposure in KO mice resulted in a brain  $PO_2$  similar to WT mice breathing air. These results show that chronic hypoxic breathing normalizes the brain  $PO_2$  of *Ndufs4* KO mice despite an increase in the circulating Hb level (Jain et al., 2016).

### CO Decreases Oxygen Content, Normalizes Brain Tissue $PO_2$ , and Rescues Brain Disease

If brain  $PO_2$  is a key parameter dictating neurological disease progression in the mouse model of Leigh syndrome, other interventions that lower brain  $PO_2$  should similarly rescue neurodegenerative disease. Hypoxia breathing reduces arterial oxygen content by lowering both arterial oxygen saturation and the partial pressure of arterial oxygen ( $PaO_2$ ). A well-known mode of altering tissue oxygenation is exposure to carbon monoxide (Blumenthal, 2001). Carbon monoxide binds to hemoglobin with a 200-fold increased affinity relative to oxygen, outcompeting the binding pocket for oxygen. In doing so, CO displaces oxygen from hemoglobin, forming carboxyhemoglobin (COHb) and lowering arterial oxygen saturation ( $O_2Hb$ ). CO also increases the affinity of hemoglobin for oxygen, reducing the off-loading of bound oxygen to the tissues. We decided to test whether breathing a high but sublethal concentration of CO in 21%  $O_2$  could rescue neurological disease in *Ndufs4* KO mice.

Exposure to chronic, continuous 600 ppm CO in air resulted in ~40% arterial COHb and ~60% arterial  $O_2Hb$  (Figure 3C). Of note, breathing 11%  $O_2$  results in a similar arterial  $O_2Hb$  saturation of 55%–60%. After three weeks of CO treatment hematocrit showed a marked increase from 48% to 79% in WT and from 53% to 76% in KO mice, suggesting this dose was sufficient to cause renal hypoxia and erythropoietin production (Figures 3A and 3B). CO treatment was started in KO mice at a late-stage of brain disease, after body weight loss had already occurred and MRI lesions were apparent. Upon commencing 600 ppm CO treatment, body weight gradually regained, similar to the trend observed during chronic inhaled hypoxia treatment in previous experiments (Figure 3D) (Ferrari et al., 2017). Moreover, survival was substantially prolonged in KO mice with CO treatment, with a median survival of ~150 days (Figure 3E). We

additionally performed MRI in late-stage mice before and after treatment with CO. Remarkably, CO treatment for 2 to 3 weeks completely reversed lesions in the vestibular nucleus (Figure 3F).

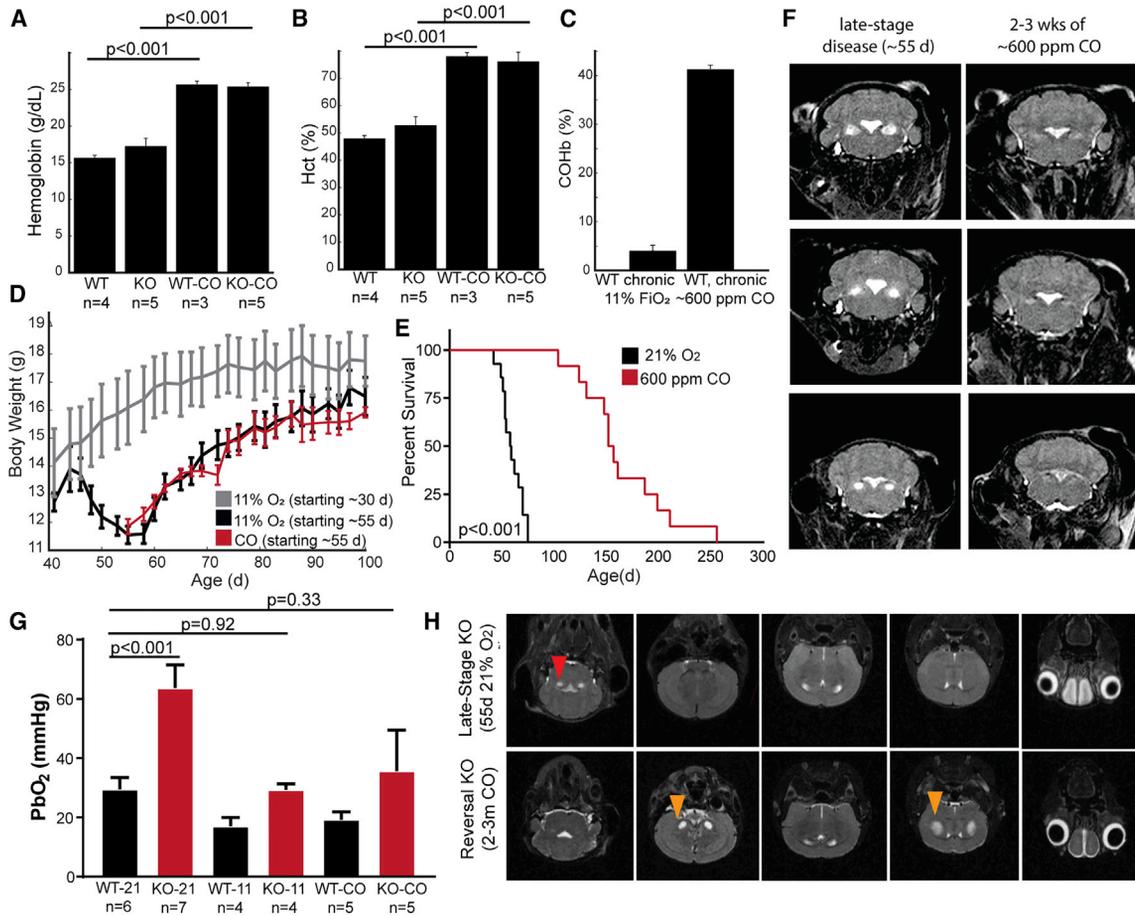
Since chronic 11%  $O_2$  breathing normalized the relative brain hyperoxia observed in KO mice breathing room air, we asked whether CO treatment similarly acted by reducing brain tissue  $PO_2$ . Both WT and KO mice chronically breathing CO at 600 ppm in air had a lower brain  $PO_2$  as compared to WT and KO mice breathing air (19 versus 30 mmHg,  $p = 0.0008$  in WT and 36 versus 64 mmHg,  $p = 0.001$  in KO, Figure 3G), based on measurements using an optical probe. The results show that chronic administration of CO at 600 ppm reduces brain  $PO_2$ , improves survival, and reverses brain disease in *Ndufs4* KO mice.

Of course, CO presents a significant health hazard and high doses can result in death or permanent neurological dysfunction (Blumenthal, 2001). We asked whether KO mice that were exposed to chronic low-dose CO were able to chronically maintain healthy brain tissue, both in the regions affected by CI deficiency and more generally. We scanned KO mice that had been exposed to CO for 2 to 3 months. While the Leigh's disease lesions were not present, these mice did develop hyperintense lesions apparent in the caudoputamen region of the brain, presumably due to chronic CO treatment (Figure 3H). Lesions that are present in humans that have suffered from CO poisoning can be found in different anatomical structures, such as the globus pallidus. However, this pattern is not completely stereotyped and can vary between individuals or states of poisoning (Jeon et al., 2018; O'Donnell et al., 2000). Although inhaled CO is unlikely to be utilized clinically as a therapeutic strategy given its remarkably narrow therapeutic index, these data (Figure 3) provide valuable proof of concept that reducing brain oxygen delivery alleviates the progression of Leigh disease in this mouse model.

### Severe Anemia Can Reverse Disease Progression in *Ndufs4* KO Mice

The ability of CO treatment to rescue disease serves as a proof of concept that additional therapeutic regimens which affect oxygen delivery might also be protective. In theory, affecting several physiological parameters (arterial Hb saturation, Hb content, cardiac output, oxygen consumption, etc.) should also lower brain  $PO_2$  and rescue disease (Figure 4A). We explored an additional mode of decreasing oxygen delivery—extreme anemia achieved by phlebotomy in combination with dietary iron deficiency.

One simple strategy to decrease oxygen delivery is to limit the number of circulating red blood cells. In humans, chronic anemia causes weakness, lethargy, and reversible hair loss (Lopez et al., 2016). However, phlebotomy remains the first line therapy for a small number of diseases, including certain forms of polycythemia and hemochromatosis. We therefore phlebotomized our KO mice. Mice were bled ~200–400 microliters by tail vein, 5 to 6 times with 2 to 3 days in between bleeding sessions to allow blood volume to recover. Mice were also placed on an Fe-deficient diet to prevent the regeneration of RBCs. Using this therapeutic regimen, we were able to cause extreme anemia in KO mice, reaching an average Hb of 2 to 3 g/dL (Figure 4B). While this is an extreme level of anemia, mice appeared relatively normal and active even at these low levels of Hb. To determine



**Figure 3. Breathing Carbon Monoxide Reduces Brain Hyperoxia and Reverses Disease in *Ndufs4* KO Mice**

(A) Hemoglobin and (B) Hematocrit (mean  $\pm$  SEM) of WT and KO mice exposed to air or  $\sim$ 600 ppm carbon monoxide in air for  $\sim$ 3 weeks. Unpaired t test p value shown within inserts.

(C) Percentage of carboxyhemoglobin (mean  $\pm$  SEM) of WT mice breathing 11%  $F_iO_2$  versus 600 ppm CO.

(D) Body weight trajectory of mice exposed to preventative (gray) hypoxia therapy, late-stage hypoxia therapy (black), or late-stage carbon monoxide therapy (red). (n = 8 for KO prevention in gray, n = 8 for KO disease reversal with hypoxia breathing in black, and n = 6 for KO disease reversal with CO breathing). Note: hypoxia body weight curves taken from historical data previously reported in (Ferrari et al., 2017) and adapted for this figure.

(E) Survival of mice breathing room air (historical data from (Ferrari et al., 2017)) versus 600 ppm CO in room air. Log-rank p value shown within insert. n = 12 for CO survival in red, n = 14 for mice breathing room air in black.

(F) Reversal of T2-weighted MRI lesions in mice exposed to 600 ppm CO starting a late stage of disease. Pre-treatment scans on left, post-treatment of same mice after 2 to 3 weeks of CO on right. Three rows are three individual mice.

(G) Brain (vestibular nucleus)  $PO_2$  of WT and KO (mean  $\pm$  SD) mice breathing 21%  $F_iO_2$ , 11%  $F_iO_2$ , or 600 ppm CO in 21%  $O_2$ . (data for WT taken from same data as Figure 2).

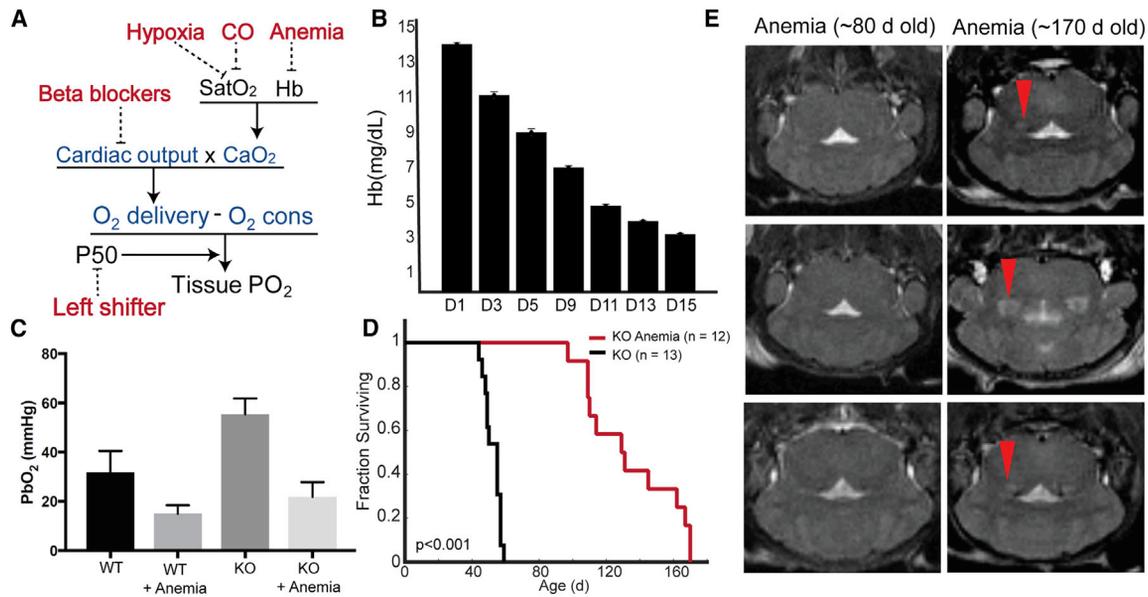
(H) T2-weighted MRI scan sections through late-stage KO mouse breathing room air versus lesions observed in KO mouse chronically exposed to 600 ppm CO. From left to right in 3H, lesions correspond to top 1 (vestibular nucleus), bottom 2 (red nucleus), bottom 4 (caudoputamen) and top 5 (olfactory bulb). Typical disease lesions in vestibular nucleus shown with red arrow. New lesions in CO-treated KO mice shown with orange arrow.

the effects of anemia on brain  $PO_2$ , we once again used the optical probe to measure cerebral oxygenation in WT and KO mice that were untreated or subjected to our therapeutic anemia protocol. Similar to CO treatment and inhaled hypoxia, anemia reduced the brain hyperoxia observed in *Ndufs4* KO mice, down to WT levels (Figure 4C). This anemia protocol extended the life of KO mice from a median of 55 days [4] to a median of  $\sim$ 130 days (Figure 4D). Moreover, MRI lesions were not apparent at 50–70 days of age when they normally appeared in untreated mice. However, brain lesions did become apparent closer to death at  $\sim$ 150 days (Figure 4E). The ability of this relatively

simple procedure—severe anemia—to rescue disease, further establishes proof of concept that brain tissue oxygenation is a key parameter in the pathogenesis of neurological disease in Leigh syndrome.

## DISCUSSION

Chronic, continuous 11%  $O_2$  breathing results in a striking rescue of brain disease in the *Ndufs4* KO mouse model of Leigh syndrome (Jain et al., 2016; Ferrari et al., 2017). Our initial studies were motivated by a CRISPR/Cas9 screen that spotlighted the



**Figure 4. Severe Anemia Decreases Brain Oxygenation and Rescues Neurological Disease in the *Ndufs4* KO Mouse**

(A) Schematic of variables affecting tissue oxygenation. Theoretical therapeutic interventions shown in red, physiological variables represented in blue or black. (B) Gradual decrease in hemoglobin following serial phlebotomy every 2 to 3 days for ~20 days, in combination with an Fe-deficient diet (mean ± SEM, n = 7). (C) Brain PO<sub>2</sub> (vestibular nuclei) of WT and KO mice that are untreated or made anemic using phlebotomy, in combination with an Fe-deficient diet (WT data same as Figure 2).

(D) Survival of untreated (Ferrari et al., 2017) or anemic *Ndufs4* KO mice.

(E) MRI of anemic *Ndufs4* KO mice at 80 days or 170 days of age. Note: Mice scanned in left and right panels are distinct. Rows correspond to different mice to show reproducibility. Red arrow depicts lesions in vestibular nuclei.

hypoxia response and suggested that shifting metabolism from oxidative phosphorylation toward glycolysis can alleviate growth defects in cultured cells. In cell culture, forced activation of this pathway is sufficient to buffer the growth defects in cells with impaired electron transport chains (Jain et al., 2016; Li et al., 2018). However, subsequent clues suggested that this pathway may be insufficient for the hypoxic rescue *in vivo* of the *Ndufs4* KO mouse. For example, intermittent hypoxia and more moderate hypoxia were able to trigger HIF-dependent hematocrit elevation but did not result in even an intermediate rescue (Ferrari et al., 2017). In our current work, we show that genetically activating the hypoxia transcriptional program by decreasing or eliminating activity of PHD1/2/3 or vHL is insufficient to rescue brain disease, even though canonical HIF targets are activated. While our work strongly argues against sufficiency of HIF activation for the rescue of Leigh syndrome, we have not tested the necessity of this pathway for rescue by hypoxia.

Not only were the genetic crosses we performed unable to rescue brain disease, but in some cases disease progression was hastened. At a systemic level, hypoxia adaptations mediated by HIF include several homeostatic mechanisms to increase oxygen delivery and dampen oxygen consumption. Thus, if these pathways are activated in normoxic conditions, they may paradoxically increase the amount of local brain tissue PO<sub>2</sub> (the key culprit identified in the current study) thereby exacerbating disease. One caveat is that the PHD enzymes are also involved in oxygen sensing more broadly. For example, PHD1 inhibition is protective during ischemia stroke (Quaegebeur et al., 2016). In this case, the proposed mechanism is shunting of

glucose toward the pentose phosphate pathway and increased NADPH production for ROS-scavenging enzymes. Several other HIF-independent, PHD-dependent adaptations have been proposed including an increase in flux through gluconeogenesis. If PHD enzymes have redundant roles in HIF-independent hypoxia adaptations, it is possible that distinct hypoxia adaptations contribute to disease rescue. While our studies demonstrate that genetic activation of the hypoxia response is not sufficient to rescue Leigh syndrome, we cannot exclude the possibility that the vHL-PHD-HIF program is necessary for some aspects of hypoxia's *in vivo* rescue of disease trajectory.

Mitochondria are widely regarded as the energy-producing organelles of the cell and the textbook explanation for mitochondrial disease is centred on defective ATP production. However, in most tissues mitochondria are also the primary oxygen-consumers, allowing intracellular oxygen tensions to remain low and preventing the toxic oxidation of intracellular biomolecules. In fact, humans with mitochondrial disease are known to exhibit venous hyperoxia and impaired oxygen extraction which is most apparent during exercise (Taivassalo et al., 2002), and this venous hyperoxia has even been proposed as a biomarker for the disease. We show that with advanced disease, CI deficiency results in a decline in whole body oxygen consumption. This is associated with elevated brain tissue PO<sub>2</sub>, which is normalized by chronic hypoxic exposure. A recent study showed that the defect in CI-mediated electron transport occurs in a regional and temporal manner in the *Ndufs4* KO mouse. Kayser et al. showed that CI activity is relatively well-preserved in young mice but declines with age (Kayser et al., 2016). It is tempting

to speculate that a vicious cycle could lead to the end pathology: an initial defect in mitochondrial CI could lead to local hyperoxia that exacerbates mitochondrial dysfunction, creating a feed-forward loop of ETC damage. Since whole body oxygen consumption declines with age, it is likely that other tissues are also experiencing relative hyperoxia in KO mice.

The current study proposes that in the face of mitochondrial dysfunction, unused oxygen can accumulate and this can be directly toxic. Oxygen toxicity is traditionally associated with the damaging effects of superoxide radicals or hydrogen peroxide. In principle, the generation of such reactive oxygen species should be higher in the brains of *Ndufs4* KO mice, as both NADH and O<sub>2</sub> can contribute to the generation of toxic reactive oxygen species (Kussmaul and Hirst, 2006; Hirst et al., 2008). Complex I deficiency leads to a direct increase in NADH/NAD<sup>+</sup> ratio and as we have shown in the current paper, will lead to increased brain O<sub>2</sub> tensions *in vivo*. Moreover, an impaired ETC with structural defects may cause increased one-electron leak to molecular oxygen. However, we have shown that isolated mitochondria from KO and WT mice produced similar amounts of H<sub>2</sub>O<sub>2</sub> at a given oxygen tension, when respiring on malate and pyruvate. If ROS are the culprit species, KO mice may be producing greater amounts of ROS simply because more oxygen is available as a substrate. An alternative hypothesis for oxygen toxicity is that the increased oxygen levels may be directly oxidizing and damaging biomolecules, e.g., those containing Fe-S clusters or iron centers, without the need to invoke the formation of toxic reactive oxygen species, as we have recently shown for the Fe-S cluster biosynthetic machinery itself (Ast et al., 2019). Consistent with this perhaps unconventional hypothesis for oxygen toxicity, various antioxidant strategies have been tested in mitochondrial diseases with little or no proven benefit. It is possible that hypoxia is more effective than antioxidants, since it prevents oxygen toxicity at its root base. Further work is needed to determine the nature of dioxygen toxicity in the *in vivo* setting.

A key finding from the current work is that very different strategies that reduce brain oxygenation are protective in the *Ndufs4* KO mouse model of Leigh syndrome. Our original work (Ferrari et al., 2017; Jain et al., 2016) utilized hypoxic gas mixtures to create normobaric inhaled hypoxia. Here, we have reported the use of two different strategies to reduce oxygen delivery to also alleviate disease in this mouse model. First, we have demonstrated that lowering brain tissue PO<sub>2</sub> by using CO to decrease arterial oxygen content (without altering arterial PO<sub>2</sub>), is sufficient to rescue disease (Figure 3). Second, we have used severe anemia (through a mix of an iron deficient diet and phlebotomy) to reduce oxygen delivery to the brain (Figure 4). Both of these interventions are severe, but serve as important proof of concept that decreasing arterial oxygen content (rather than arterial PaO<sub>2</sub>) or overall oxygen delivery should rescue disease by lowering tissue PO<sub>2</sub>. Although low-dose CO (250 ppm) is known to have anti-inflammatory effects and is in phase 1 clinical trials for idiopathic pulmonary fibrosis (Rosas et al., 2018) in humans, its therapeutic index is too narrow and hence unlikely to serve as a candidate therapeutic in humans. Similarly, although phlebotomy is used as a therapy for other disorders, including hemochromatosis, its extension to mitochondrial disorders will be challenging, as anemia is known to be a risk factor for worse

prognosis in certain inherited mitochondrial diseases, including *POLG* deficiency (Hikmat et al., 2017). While hypoxia breathing, inhaled CO, and severe anemia are each individually highly pleiotropic and could be having a beneficial effect on the *Ndufs4* KO mice by distinct mechanisms, the most parsimonious explanation is that they are all alleviating Leigh syndrome by directly reducing brain oxygen tensions.

We anticipate that other approaches, such as small molecule HIF inhibitors, now in trials for renal cell cancer (Martínez-Sáez et al., 2017), may be more practical to limit an adaptive increase in RBC production, or orally available oxy-Hb left-shifters, which are now in clinical trials for sickle cell anemia (Oksenberg et al., 2016), are worthy of future exploration. Further understanding of the precise mechanisms by which elevated tissue oxygenation contributes to end-organ pathology in mitochondrial disease may lead to additional novel and practical combination therapies that are needed to treat Leigh syndrome.

### Limitations of Study

(1) Our study demonstrates that HIF activation is not sufficient to rescue complex I deficiency in mouse models. However, it is possible that HIF is still necessary. Future studies will be required to determine whether HIF is necessary for the inhaled hypoxia rescue mechanism *in vivo*. (2) We show increased oxygen levels in the brains of the *Ndufs4* KO mouse model. While this suggests oxygen toxicity, we have not directly measured toxic reactive oxygen species *in vivo*. We have also not determined the exact nature of the damage (e.g., which enzymes and biomolecules are failing). (3) Our current work suggests that therapeutic strategies which decrease oxygen delivery might be protective in this mouse model. However, future studies are needed to optimize such a therapeutic strategy and investigate any long-term negative side effects.

### STAR★METHODS

Detailed methods are provided in the online version of this paper and include the following:

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#### AUTHOR CONTRIBUTIONS

I.H.J., L.Z., W.M.Z., and V.K.M. designed study. I.H.J., L.Z., O.G., E.M., G.R.W., T.A., H.W., G.S., A.S., A.G., K.B., L.S., P.C., F.I., W.M.Z., and V.K.M. performed, interpreted, and analyzed experiments. I.H.J., L.Z., W.M.Z., and V.K.M. compiled data and wrote the manuscript.

#### DECLARATION OF INTERESTS

V.K.M., W.M.Z., I.H.J., and L.Z. are listed as inventors on a patent application submitted by the Massachusetts General Hospital on technology described in this paper. V.K.M. is a paid advisor to Janssen Pharmaceuticals, Raze Therapeutics, and 5AM Ventures.

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## STAR★METHODS

### KEY RESOURCES TABLE

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Chemicals, Peptides, and Recombinant Proteins		
Mannitol	Sigma	Cat # 63559
Bovine Serum Albumin	Sigma	Cat # A6003
Digitonin	Sigma	Cat # D141
Horseradish Peroxidase	Thermo Fisher Scientific	Cat #012001
EPOR TaqMan Assay	Thermo Fisher Scientific	Mm00833882_m1
EPO TaqMan Assay	Thermo Fisher Scientific	Mm01202755_m1
VEGF TaqMan Assay	Thermo Fisher Scientific	Mm00437306_m1
LDHA TaqMan Assay	Thermo Fisher Scientific	Mm01612132_g1
SLC2A1 TaqMan Assay	Thermo Fisher Scientific	Mm00441480_m1
Critical Commercial Assays		
TaqMan Gene Expression Master Mix	Thermo Fisher Scientific	Cat # 4369016
M-MLV Reverse Transcriptase	Promega	Cat # M1701
Amplex UltraRed	Thermo Fisher Scientific	Cat # A36006
Qiagen RNeasy Lipid Tissue Mini Kit	Qiagen	Cat # 74804
Experimental Models: Organisms/Strains		
Mouse: <i>Ndufs4</i> <sup>-/-</sup>	Richard Palmiter	N/A
Mouse: <i>Phd1</i> <sup>-/-</sup>	Peter Carmeliet	N/A
Mouse: Nestin- <i>Phd2</i> <sup>-/-</sup>	Peter Carmeliet	N/A
Mouse: <i>Phd3</i> <sup>-/-</sup>	Peter Carmeliet	N/A
Mouse <i>Vhl</i> <sup>ch/ch</sup>	Josef Prchal	N/A

### LEAD CONTACT AND MATERIALS AVAILABILITY

Further information and requests for resources and reagents should be directed to and will be fulfilled by the Lead Contact, Vamsi K. Mootha ([vamsi@hms.harvard.edu](mailto:vamsi@hms.harvard.edu)). This study did not generate new unique reagents.

### EXPERIMENTAL MODEL AND SUBJECT DETAILS

#### Mice

All animal work in manuscript was approved by the Subcommittee on Research Animal Care and the Institutional Animal Care and Use Committee of Massachusetts General Hospital. *Ndufs4*<sup>-/-</sup> were obtained from Palmiter laboratory (University of Washington). Mice lacking *Phd1* (Aragonés et al., 2008), nestin-*Phd2* (Mazzone et al., 2009) and *Phd3* (Bishop et al., 2008) were provided by the Carmeliet laboratory (KU Leuven). *Vhl*<sup>ch/ch</sup> mice were provided by Josef Prchal (University of Utah). Genetic crosses were performed by crossing *Ndufs4*<sup>+/-</sup> mice with remaining strains. Throughout this study, *Ndufs4*<sup>+/+</sup> and *Ndufs4*<sup>+/-</sup> mice were indiscriminately used as healthy controls, since there is no apparent phenotypical difference between these genotypes. When possible, controls from the same genetic cross were used, in order to make comparisons between reasonably similar genetic backgrounds. Pups were genotyped and weaned from mothers at 25-30 d of age. Cages and water were changed weekly. Humane euthanasia criteria were used throughout the study. Natural death or 20% body weight loss were used to generate survival curves. Animals were housed in either conventional or barrier housing. Males and females were used throughout the study. The *NDUFS4*, *PHD1* and *PHD3* KO strain are C57/BL6. The nestin-*PHD2* strain is mixed background (C57/BL6 and 129S6).

### METHOD DETAILS

Experiments were conducted in experimental and technical replicates (sample size described throughout figures). Mice were randomized within each genotype. Careful consideration was used to ensure groups were balanced for gender, age and starting weight. When possible, samples were blinded. For example, survival curves, body weight measurements, brain PO<sub>2</sub> measurements, etc.

were all made blinded. Sample sizes and power calculations were based on data from our previous studies using the *Ndufs4* KO mouse model (Ferrari et al., 2017; Jain et al., 2016).

### Respirometry

Whole body oxygen consumption was monitored using the Sable system (Promethion Cages) continuously from ~30 d to ~45 d of age. During the experiment, mice were continuously provided food/water and exposed to a day/night cycle of 12h/12h. A smoothing window of 3h was used for each timepoint during the analysis. Oxygen consumption values were normalized to body weight.

### H<sub>2</sub>O<sub>2</sub> Assay

A high-resolution respirometer (O2k, Oroboros Instruments GmbH) equipped with home-built two-channel fluorescence optical module was used to monitor simultaneously oxygen concentration and H<sub>2</sub>O<sub>2</sub> production as described in detail in [14]. The module was connected to the amperometric port of the respirometer and the fluorescence was recorded using the Datlab software. Mitochondrial H<sub>2</sub>O<sub>2</sub> release was measured as the fluorescence of resorufin formed from Amplex UltraRed in the presence of horseradish peroxidase (excitation 525nm, emission filter 580 nm cut off). The fluorescence signal was calibrated into an H<sub>2</sub>O<sub>2</sub> concentration by addition of 90nM of H<sub>2</sub>O<sub>2</sub> to the chamber at the end of each experiment. To assess H<sub>2</sub>O<sub>2</sub> release at different oxygen level, humidified gaseous pre-purified argon was purged into the 1ml gas headspace of respirometer chamber at a flow rate of 10–60ml/min to rapidly decrease oxygen concentration in the measuring assay.

Intact brain mitochondria (0.1–0.5 mg/ml) were added to assay media containing 125 mM KCl, 14 mM NaCl, 0.2 mM EGTA, 4 mM KH<sub>2</sub>PO<sub>4</sub>, 2 mM MgCl<sub>2</sub>, 20 mM HEPES-Tris (pH 7.4), supplemented with 2 mg/ml BSA, 10 μM Amplex UltraRed, 4 U/ml horseradish peroxidase, and 40 U/ml Cu/Zn superoxide dismutase at 37°C. Combination of 2 mM malate and 5 mM pyruvate was used to reduce matrix NADH for respiration mediated by CI. Pyruvate is oxidized by pyruvate dehydrogenase generating NADH and acetyl-CoA which reacts with matrix oxaloacetate to shift the malate dehydrogenase reaction towards oxidation of malate and generation of another NADH molecule.

### Brain Tissue PO<sub>2</sub> Measurement

*Ndufs4*<sup>-/-</sup> and WT mice were anesthetized with Isoflurane (induction at 2–4%, maintenance at 1–1.5%), intubated, and mechanically ventilated with a tidal volume of 8 ml/kg, a respiratory rate of 110 breaths per minute and an inspired fraction of oxygen (FiO<sub>2</sub>) of 21%. Mice were placed in a prone position and the head was stabilized using a stereotaxic frame (ASI Instruments, MI). After incision and dissection of the skin, an opening in the skull was performed using a micro-drill (MD-1200, Braintree Scientific, MA) and a PO<sub>2</sub> probe was inserted at the desired location. The coordinates for the thalamic measurements were mediolateral [ML] = –1.00 mm, antero-posterior [AP] = –2.00 mm, and dorsoventral [DV] = –4.00 mm from the bregma, while the coordinates for the vestibular nuclei were ML = –1.25 mm, AP = –6.00 mm, and DV = –3.90 mm from the bregma. For the thalamic measurements a flexible polarographic Clark-type oxygen microprobe was used (LICOX; GMS, Kiel-Mielkendorf, Germany). For the vestibular nuclei measurements an optical PO<sub>2</sub> probe was employed (OxyLab, Oxford Optronix, Abingdon, UK). During the brain PO<sub>2</sub> measurement the depth of anesthesia was reduced by lowering the Isoflurane concentration to 0.5–1% to minimize the impact of anesthesia on the brain PO<sub>2</sub>. During measurements mice were ventilated with various gas mixtures according to the experimental group: 21% O<sub>2</sub>, 11% O<sub>2</sub>, 600 ppm CO balanced air. In some experiments the FiO<sub>2</sub> was modulated over the course of the test and the brain PO<sub>2</sub> was simultaneously measured.

### Hematocrit and COHb Measurements

Eighty microliters of blood were collected by tail snip into a heparinized capillary. Hemoglobin concentration, hematocrit and the saturation of hemoglobin with oxygen (O<sub>2</sub>Hb) and carbon monoxide (COHb) were measured using a blood gas analyzer (ABL800 FLEX, Radiometer, Copenhagen, Denmark).

### Chambers for Various Chronic Breathing Regimens

Mice were housed in 80-L transparent acrylic boxes. Control chambers were provided with a constant flow of medical air at 21% FiO<sub>2</sub>. A concentration of 11% or 17% oxygen was obtained by mixing medical air with nitrogen obtained from a nitrogen generator (MAG-20; Higher Peak). A concentration of 600 ppm CO was obtained by mixing medical air with CO from a cylinder containing 2% CO (20,000 ppm) (Airgas). The total gas flow through the chamber was adjusted to maintain chamber CO<sub>2</sub> concentrations below 0.4% (CO200; Exttech). Soda lime was added to the chambers as a CO<sub>2</sub> scavenger. The oxygen concentration was measured at the outlet port of the chamber with an O<sub>2</sub> sensor (MiniOx 1; Ohio Medical), which was calibrated using an 8.55% O<sub>2</sub> reference tank (Airgas). In a similar way the CO concentration was measured a CO sensor (MSA Altair PRO; MSA Safety, Pittsburgh, Pa), which was calibrated using a 1000-ppm CO reference cylinder (Airgas). Oxygen levels inside the chambers were tolerated within a 0.4% offset from the target concentration by adjusting nitrogen flow as needed. Temperature was maintained at 24–26°C, and humidity was maintained at 30–70%. A standard light-dark cycle of ~12h light exposure was used. Mice were housed in cages with standard bedding and given unlimited access to food and water.

### Anemia

To induce anemia, mice were placed on a low iron diet. Since the time to achieve adequate levels of anemia would have been too long and KO mice would have died of their mitochondrial disease, 150–200 μL of blood were collected every other day until hemoglobin

concentration reached the desired values. Within 2 weeks, hemoglobin concentrations reached approximately 3–4 g/dL and remained stable while the mice remained on a low iron diet.

### MRI

MRI scans of the brain were performed as previously described (Ferrari et al., 2017), using respiratory gated T2-weighted RARE (rapid acquisition of refocused echoes) MRI images acquired on a 4.7-T small animal scanner (Pharmascan; Bruker) with the following parameters: RARE factor: 10, echo time: 60 ms, repetition time: 6000 ms, Averages: 8, 192 x 192 x 24 image matrix with a voxel size of 0.130 x 0.130 x 0.7 mm. Mice were under anesthesia (isoflurane between 0.5–1.5% in room air) during the imaging procedure. Images were generated using a DICOM reader (OsiriX, University of Geneva).

### qPCR

Animals were sacrificed by CO<sub>2</sub> asphyxiation followed by cervical dislocation. Individual cerebella were immediately harvested and snap frozen in liquid N<sub>2</sub>. The tissue was then disrupted with two 5mm stainless steel beads (Qiagen) using a Qiagen TissueLyser for 2 mins at 25 Hz. RNA was extracted with the RNeasy Lipid Tissue Mini Kit (Qiagen), before murine leukemia virus (MLV) reverse transcription using random primers (Promega). qPCR was performed using the TaqMan technology (Life Technologies), using the following probes:

Mm00833882\_m1 (EPOR)  
Mm00437306\_m1 (VEGFA)  
Mm01612132\_g1 (LDHA)  
Mm00441480\_m1 (SLC2A1)  
Mm01202755\_m1 (EPO)  
Mm00437762\_m1 (B2M)

All data were normalized to B2M (Mm00437762\_m1).

### QUANTIFICATION AND STATISTICAL ANALYSIS

Analyses were performed using GraphPad Prism 7.0 software. The two-sample Student t-test was used for two-group comparisons. Unless otherwise indicated, one-way ANOVA with Bonferroni's correction was used for multiple comparisons. The log-rank test was performed, and HRs with 95% CIs were calculated to compare survival rates. A p value < 0.05 was considered to indicate statistical significance. Data are reported as mean ± SD.

### DATA AND CODE AVAILABILITY

There are no relevant datasets or code in this study.