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## Human and rodent red blood cells do not demonstrate xanthine oxidase activity or XO-catalyzed nitrite reduction to NO

Sara E. Lewis<sup>a,1</sup>, Courtney B. Rosencrance<sup>a,1</sup>, Evan De Vallance<sup>a</sup>, Andrew Giromini<sup>a</sup>, Xena M. Williams<sup>a</sup>, Joo-Yeun Oh<sup>d</sup>, Heidi Schmidt<sup>c</sup>, Adam C. Straub<sup>c</sup>, Paul D. Chantler<sup>b</sup>, Rakesh P. Patel<sup>d</sup>, Eric E. Kelley<sup>a,\*</sup>

<sup>a</sup>West Virginia University Departments of Physiology and Pharmacology, USA

<sup>b</sup>Exercise Physiology, USA

<sup>c</sup>University of Pittsburgh Vascular Medicine Institute, USA

<sup>d</sup>University of Alabama at Birmingham Center for Free Radical Biology, USA

### Abstract

A number of molybdopterin enzymes, including xanthine oxidoreductase (XOR), aldehyde oxidase (AO), sulfite oxidase (SO), and mitochondrial amidoxime reducing component (mARC), have been identified as nitrate and nitrite reductases. Of these enzymes, XOR has been the most extensively studied and reported to be a substantive source of nitric oxide (NO) under inflammatory/hypoxic conditions that limit the catalytic activity of the canonical NOS pathway. It has also been postulated that XOR nitrite reductase activity extends to red blood cell (RBCs) membranes where it has been immunohistochemically identified. These findings, when combined with countervailing reports of XOR activity in RBCs, incentivized our current study to critically evaluate XOR protein abundance/enzymatic activity in/on RBCs from human, mouse, and rat sources. Using various protein concentrations of RBC homogenates for both human and rodent samples, neither XOR protein nor enzymatic activity (xanthine → uric acid) was detectable. In addition, potential loading of RBC-associated glycosaminoglycans (GAGs) by exposing RBC preparations to purified XO before washing did not solicit detectable enzymatic activity (xanthine → uric acid) or alter NO generation profiles. To ensure these observations extended to absence of XOR-mediated contributions to overall RBC-associated nitrite reduction, we examined the nitrite reductase activity of washed and lysed RBC preparations via enhanced chemiluminescence in the presence or absence of the XOR-specific inhibitor febuxostat (Uloric®). Neither addition of inhibitor nor the presence of the XOR substrate xanthine significantly altered the rates of nitrite reduction to NO by RBC preparations from either human or rodent sources confirming the absence of XO enzymatic activity. Furthermore, examination of the influence of the age (young cells vs. old cells) of human RBCs on XO activity also failed to demonstrate detectable XO protein. Combined, these data suggest: 1) that XO does not contribute to nitrite reduction in/on human and

\*Corresponding author. West Virginia University, School of Medicine, Department of Physiology and Pharmacology, 3072B Health Sciences Center, PO Box 9229, Morgantown, WV, 26506-9229, USA. eric.kelley@hsc.wvu.edu (E.E. Kelley).

<sup>1</sup>These authors contributed equally.

Appendix A. Supplementary data

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rodent erythrocytes, 2) care should be taken to validate immuno-detectable XO by demonstrating enzymatic activity, and 3) XO does not associate with human erythrocytic glycosaminoglycans or participate in nonspecific binding.

## Keywords

Nitrite; Xanthine oxidoreductase; Nitric oxide; Red blood cells; Glycosaminoglycans

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## 1. Introduction

Xanthine oxidoreductase (XOR) catalyzes the final two steps in purine catabolism (hypoxanthine  $\rightarrow$  xanthine  $\rightarrow$  uric acid). XOR is a homodimer with each monomer consisting of four redox centers: a molybdenum cofactor (Mo-co), one FAD site, and two Fe/S clusters. The Mo-co is the site of purine oxidation while NAD<sup>+</sup> and O<sub>2</sub> reduction occur at the FAD [1–3]. The two Fe/S clusters operate to transfer electrons between the Mo-co and the FAD. The enzyme is translated as xanthine dehydrogenase (XDH) where purine-derived electrons reduce NAD<sup>+</sup> to NADH. Under inflammatory conditions post-translational modification of XDH produces xanthine oxidase (XO) where affinity for NAD<sup>+</sup> is diminished while affinity for O<sub>2</sub> is enhanced resulting in oxidant generation. This capacity to reduce O<sub>2</sub> has positioned XO as a critical source of oxidants in a variety of inflammatory processes [4–10].

While elevation in XO levels has long been associated with oxidant production, recent reports have described a salutary function for both XDH and XO whereby their catalytic activity can be utilized to reduce nitrate (NO<sub>3</sub><sup>-</sup>) to nitrite (NO<sub>2</sub><sup>-</sup>) and NO<sub>2</sub><sup>-</sup> to nitric oxide (NO) [11–18]. Several *in vitro* studies clearly demonstrate the capacity for XO to generate NO from NO<sub>2</sub><sup>-</sup> under inflammatory conditions [19–24]. Importantly, these hypoxic/ischemic conditions limit NO generation from endothelial nitric oxide synthase and have incentivized the supposition that XO- and XDH-mediated NO generation provides compensation [14]. Numerous *in vivo* reports validate the *in vitro* findings and demonstrate a clear XOR-dependence on beneficial outcomes associated with nitrite treatment in various inflammatory processes including those associated with the vasculature where vascular-associated XOR has been contributory [11,12,25–29]. However, the details regarding micro-environmental conditions requisite for XOR-mediated NO generation are not completely understood and may involve cooperation with alternative agents such as sulfide to afford maximal benefit [30].

While a detailed accounting of factors describing the biological relevance of XOR-mediated NO formation is ongoing, XOR's mobility may play a key role in its overall impact as a NO source. For example, XDH can be released from cells (*e.g.* hepatocytes) to the circulation, rapidly converted to XO by plasma proteases, and bind avidly to the glycocalyx of endothelial cells affording XO capacity to impact vessel homeostasis via alteration of NO levels [31–33]. Recently, immunocytochemical analysis of human RBCs demonstrated the presence of XO on the membrane where it was suggested to participate in NO<sub>2</sub><sup>-</sup> reduction

to NO [34,35]. Incentivized by these findings, we examined the extent of XO protein abundance, activity, and nitrite reductase activity in human and rodent RBCs.

## 2. Materials and methods

### Animals, Blood collection and Processing-

Both male and female rodents (Sprague Dawley rats and C57Bl/6 J mice, ages 28–42 weeks) were used for experiments in compliance with WVU IACUC-approved protocols. Blood was collected at time of euthanasia and kept on ice until spun at  $1,100\times g$  for 10 min at 4 °C. Plasma was collected and placed in a separate tube while the RBC pellet was washed 3x in a 1:1 ratio with ice-cold potassium phosphate buffer (KPi, 50 mM and pH 7.4) by gentle inversion. RIPA buffer was added and pellets homogenized by needle aspiration. Human blood samples were collected according to the WVU IRB-approved protocol (1303023857). Blood from healthy human subjects (3 men and 3 women, ages 20–50) was collected by venipuncture in tubes coated in potassium EDTA and processed as above. For experiments testing young and old RBCs, blood was collected from healthy volunteers (2 males and 3 females) according to UAB IRB-approved protocol (090504006) by venipuncture and centrifuged ( $1500\times g$ , 4 °C, 10 min) to pellet RBCs. Blood was collected from 2 male and 3 female volunteers (age range 25–48 years old). Human RBCs were then separated into young and old fractions as previously described [36].

### XOR Activity-

Plasma and RBC homogenates were assessed for XO and/or XDH activity as previously described using reverse phase HPLC coupled to electrochemical (CoulArray) detection of uric acid [37]. One unit of activity (U) is defined as 1  $\mu$ mole urate formed per min at 37 °C and pH 7.4. The lower limit of detection is 300 nM.

### Nitric oxide detection-

Nitric oxide was detected using chemiluminescence (Sievers Nitric Oxide Analyzer). Reaction chamber contained the following at the indicated concentrations using 5 mL KPi, pH 7.0: xanthine (25  $\mu$ M) and RBC homogenate (5 mg). Reactions were initiated by adding  $\text{NaNO}_2^-$  (500  $\mu$ M). Authentication of NO was accomplished by addition of cPTIO (200  $\mu$ M). Initial rates of NO generation were used for quantification.

### Western Blotting-

Electrophoresis used 4–12% Bis Tris gradient Criterion™ XT Precast Gels (Bio Rad, #3450123). Samples were resolved by SDS-PAGE and transferred to nitrocellulose. Membranes were blocked with Tris-Buffered Saline (TBS)-based LiCOR Intercept blocking buffer (LiCOR, 927–80001). The primary antibody was xanthine oxidase A-3 mouse monoclonal antibody (Santa Cruz, #398548), 1:500 dilution. Secondary antibody was LiCOR, 926–32210. Blots were imaged with the LiCOR Odyssey CLX.

### 3. Results

#### XO activity in plasma-

Plasma from human, mouse, and rat was collected as described in the methods. Data shown in Fig. 1A demonstrates robust XO activity in the plasma from rodents as well as XO activity values for human plasma. XDH activity was not detected in any of the plasma samples which was expected due to plasma protease-mediated partial proteolysis resulting in conversion to XO.

#### XO activity in RBC homogenates-

RBC homogenates (0.8 mg total protein) were assayed for XO activity as shown in Fig. 1B. Rodent samples contained oxonic acid (100  $\mu$ M) to eliminate potential contributions of urate oxidase. In the mouse samples, some positive XO activity was indicated whereas in rat and human samples activity was not detected. The presence of XO activity in the 0.2, 0.4 and 1.6 mg of murine RBC homogenate was not detectable suggesting contamination by existing UA to be responsible for the positive value in the 0.8 sample. In all cases, statistical analysis demonstrated no difference between the three groups. Analysis of XDH activity also revealed no detectable enzymatic activity in any of the groups.

#### XO binding to RBC GAGs-

In order to fully explore the potential for XO to bind RBC-associated GAGs, purified XO (50 mU) was added to the isolated RBCs (50 mM KPi, pH 7.4) and allowed to incubate with gentle inversion for 15 min. Following this incubation, the RBCs were subjected to gentle washing as described in the methods. In a control experiment, 25 mU of purified XO was added to a suspension of heparin-Sepharose 6B beads (HS6B-XO) and incubated and washed in the same manner as the RBCs. As seen in Fig. 2, RBCs with and without exposure to purified XO did not demonstrate XO activity whereas HS6B-XO showed abundant activity indicative of robust binding to the HS6B. Furthermore, addition of 100  $\mu$ M allopurinol only partially diminished HS6B-XO activity while febuxostat-mediated inhibition was not affected by XO-HS6B binding, a limitation of allo/oxypurinol-based inhibition of GAG-bound XO that we have previously reported [38,39].

#### XOR protein analysis of RBC homogenates and impact of cellular age-

To determine if XOR protein was present in/on RBC preparations from human, mouse, and rat samples, Western blot analysis was performed using a commercially-available mouse monoclonal antibody (Santa Cruz) as described in the methods. As seen in Fig. 3A, the dominant band, migrating at ~150 kD and representing a full monomer of the ~300 kD XOR homodimer, is detectable in the positive control (purified XO) lane whereas this band is absent in the lanes containing RBC homogenates. To further validate this monoclonal antibody we probed murine hepatocytes, rat liver homogenates and performed XDH cellular knockdown with siRNA (Suppl. Fig. 1).

Any preparation of human RBCs will comprise cells of different ages spanning the 120d lifespan. Moreover, older RBCs which are destined for clearance typically have altered cell surface characteristics that may allow more or less XO binding. To determine if

cellular, age-related alterations impact the content of XO in/on human RBCs, using the same monoclonal antibody, Western blot analysis was conducted on five human samples that were either of young or old cellular age. The positive control band (purified XO) is present at ~150 kD whereas it is not detectable in any other lane indicating the absence of immuno-detectable XO in the patient samples regardless of age, Fig. 3B.

#### **XOR-mediated nitrite reductase activity-**

Mouse, rat, and human samples were next analyzed for contributions of XOR to overall  $\text{NO}_2^-$  reductase activity, Fig. 4. Samples were prepared in the reaction chamber with 100  $\mu\text{L}$  of RBC homogenate (5 mg total protein) in KPi pH 7.0 with or without xanthine (20  $\mu\text{M}$ ) and with or without febuxostat (10  $\mu\text{M}$ ). Reactions were initiated by the addition of  $\text{NaNO}_2^-$  (500  $\mu\text{M}$ ) and authentication of NO was determined by addition of the NO spin trap carboxy-PTIO. As seen in an example of a typical experiment in Fig. 4A, NO formation was detected following addition of  $\text{NO}_2^-$  in human samples that was not altered by the presence of febuxostat. Quantitation revealed that neither the presence of xanthine nor febuxostat altered NO generation indicating the absence of XO-catalyzed  $\text{NO}_2^-$  reduction, Fig. 4B.

## **4. Discussion**

As sources of NO alternative to the canonical NOS-mediated pathway are beginning to be more clearly understood, a significant role for XOR is becoming apparent. It has been proposed that RBC-associated XOR is contributory to  $\text{NO}_2^-$  reduction to NO [34,35]. In these studies, XOR protein was reported to be present in/on the RBC membrane and a portion of the RBC-mediated  $\text{NO}_2^-$  reductase activity sensitive to the XOR inhibitor, allopurinol. While these studies did not assess enzymatic activity, they referenced previous publications that did report XOR activity in RBCs [40–42]. However, these reports are problematic in supporting RBC-associated XOR activity as they: 1) did not describe or reference their methods of XO activity assessment [41], 2) utilized approaches that did not account for existing uric acid (absence of controls with an XO inhibitor, [40,42], and 3) report levels of XO activity for human RBCs (mU/mg Hb or U/mL) that are greater than those seen in rat liver [40,42]. A potential explanation for the magnitude of the reported XO activity values in the studies described above is the absence of controls accounting for existing uric acid. Without this control, only variations in existing uric acid are realized unless there is pronounced XO activity. Herein, we accounted for existing uric acid by using sister tubes loaded with allopurinol and report no detectable XO activity in human and rodent RBCs. This is in agreement with a report showing the absence of XO activity in RBCs from humans using radio-labeled xanthine (8-  $^{14}\text{C}$ ) as a substrate and radiolabeled uric acid ( $^{14}\text{C}$ ) as the detected product [43]. In addition, we can also conclude that human RBCs do not bind and sequester XO via interaction with GAGs as seen in Fig. 2. While this experiment focused on binding to the artificial GAG, heparin-Sepharose, it also served to confirm the absence of observable non-specific binding to RBCs.

Whereas the absence of XO activity in RBCs seems conclusive, it does not dismiss the presence of XO protein as demonstrated previously [34]. This report clearly shows an abundance of immuno-detectable protein associated with the membrane of human RBCs

using a rabbit polyclonal anti-XOR antibody. We used a mouse monoclonal anti-XOR antibody (Santa Cruz) that has demonstrated, in our experience, the greatest specificity among those commercially available; yet, we did not detect XO protein, Fig. 3. Similar, yet inconclusive, results were obtained with an alternative rabbit polyclonal anti-XO antibody (Rockland), a notoriously nonspecific antibody (Suppl. Fig. 2). A plausible answer to why our findings oppose the immunocytochemistry findings reported previously lies in appreciating the literature describing why, with regard to XOR, immunocyto/histochemistry often countervails enzymatic activity results. In 1991, Clare and Lecce described the presence of antibodies (IgG, IgA, and IgM) in commercially-available purified XO preparations and concluded that antisera raised against these preparations would give rise to antibodies recognizing epitopes on these contaminant antibodies and thus could indicate the presence of XO where none existed [44]. This may also explain why we see faint hints of bands migrating near the 145 kD XO control band when using the above-described nonspecific rabbit polyclonal anti-XO antibody, Suppl. Fig. 2.

To further validate our findings, the  $\text{NO}_2^-$  reductase capacity of human and rodent RBCs was assessed for XO-mediated contributions. RBC homogenates were analyzed for  $\text{NO}_2^-$  reductase activity using xanthine as a reducing substrate for XO as seen in Fig. 4. Whereas the addition of  $\text{NO}_2^-$  to the reaction chamber induced some NO formation, the presence of xanthine and/or febuxostat did not alter the rate of generation. These results counter those reported previously that demonstrated a measurable component of RBC-associated, XOR-mediated  $\text{NO}_2^-$  reduction to NO [34]. This report differed slightly from our experiment as XOR-mediated contributions were apparent at pH 6.8 and not significant at pH 7.4. We also endeavored to favor acidic conditions similar to those encountered *in vivo* under hypoxia/inflammation and thus our study was conducted at pH 7.0. This is an important concept as pH plays a critical role in the capacity of XOR to catalyze  $\text{NO}_2^-$  reduction [14]. While the difference between 6.8 and 7.0 may be considered small, we must consider this discrepancy as a potential explanation for the difference between the previous report and our observations. However, when combined with the absence of detectable XO protein and enzymatic activity, we conclude that this slight difference in pH does not account for the absence of XO-mediated  $\text{NO}_2^-$  reductase activity. In fact, our results are in agreement with a previous report demonstrating the absence of XO-dependent  $\text{NO}_2^-$  reduction to NO in RBCs as evidenced by the absence of an effect of the XO inhibitor allopurinol on several readouts including: 1) nitrosyl-hemoglobin formation after addition of  $\text{NO}_2^-$  to red cells as measured by electron paramagnetic resonance, 2) NO release from a red cell/nitrite mixtures as measured by chemiluminescence and 3) platelet activation [45]. In addition, this report demonstrated that carbon monoxide (CO), which inhibits heme-mediated NO production from  $\text{NO}_2^-$  while not affecting XO activity, abrogated nitrosyl-hemoglobin formation measured by EPR.

In summary, data herein demonstrate the absence of XO protein, enzymatic activity, and  $\text{NO}_2^-$  reductase activity in human and rodent RBCs as well as the inability of these RBCs to bind purified XO. In addition, the cellular age of the RBCs does not affect XO content. While these findings affirm the concept that immuno-detectible protein does not equate to enzymatic activity, they are limited to the models tested herein and thus do not rule out all potential for a circumstance whereby XO may be present in or associated with RBCs.



## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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## Abbreviations:

<b>GAGs</b>	glycosaminoglycans
<b>HS6B</b>	heparin-Sepharose 6B
<b>H<sub>2</sub>O<sub>2</sub></b>	hydrogen peroxide
<b>NO</b>	nitric oxide
<b>NOS</b>	nitric oxide synthase
<b>O<sub>2</sub><sup>•-</sup></b>	superoxide
<b>XDH</b>	xanthine dehydrogenase
<b>XO</b>	xanthine oxidase
<b>XOR</b>	xanthine oxidoreductase

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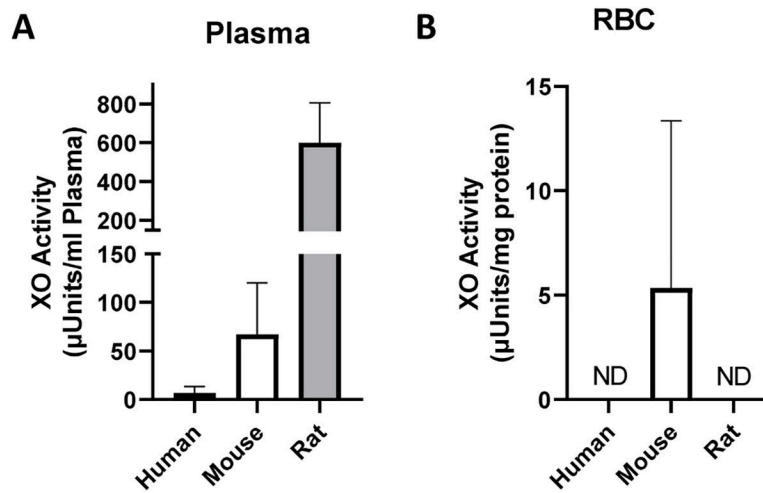
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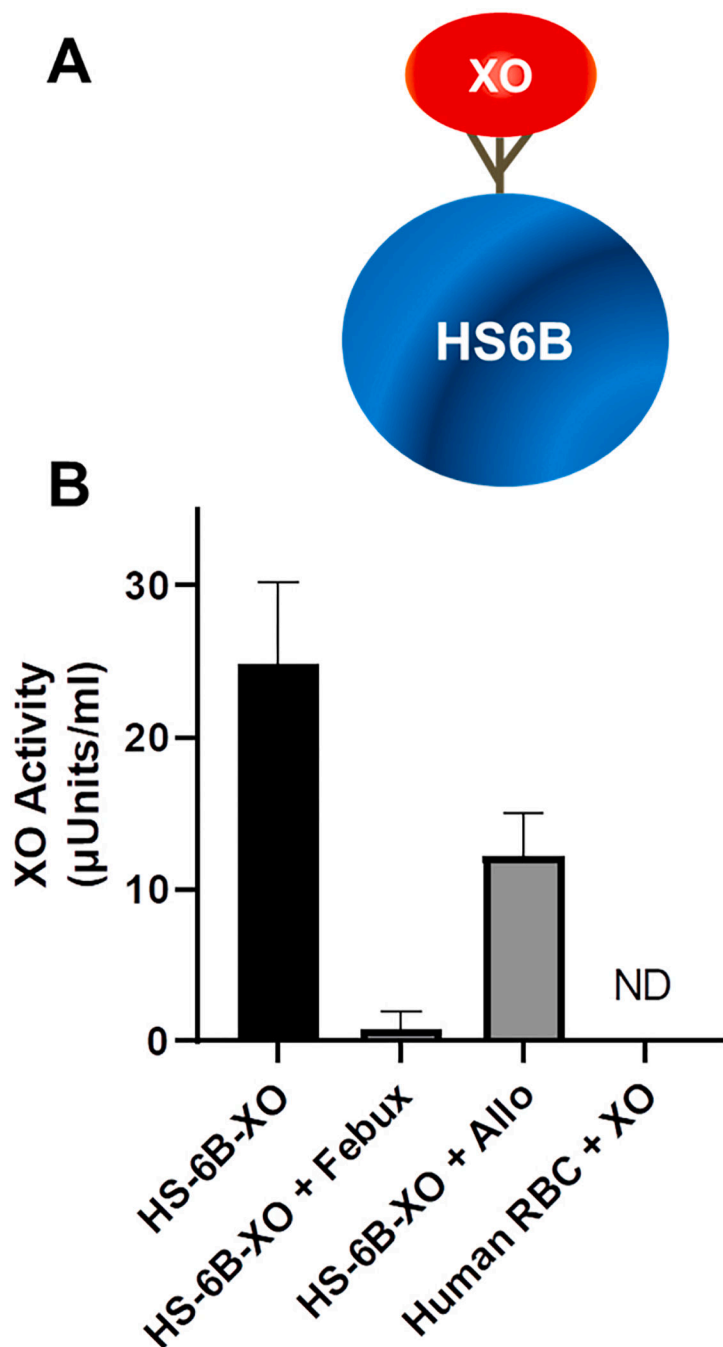
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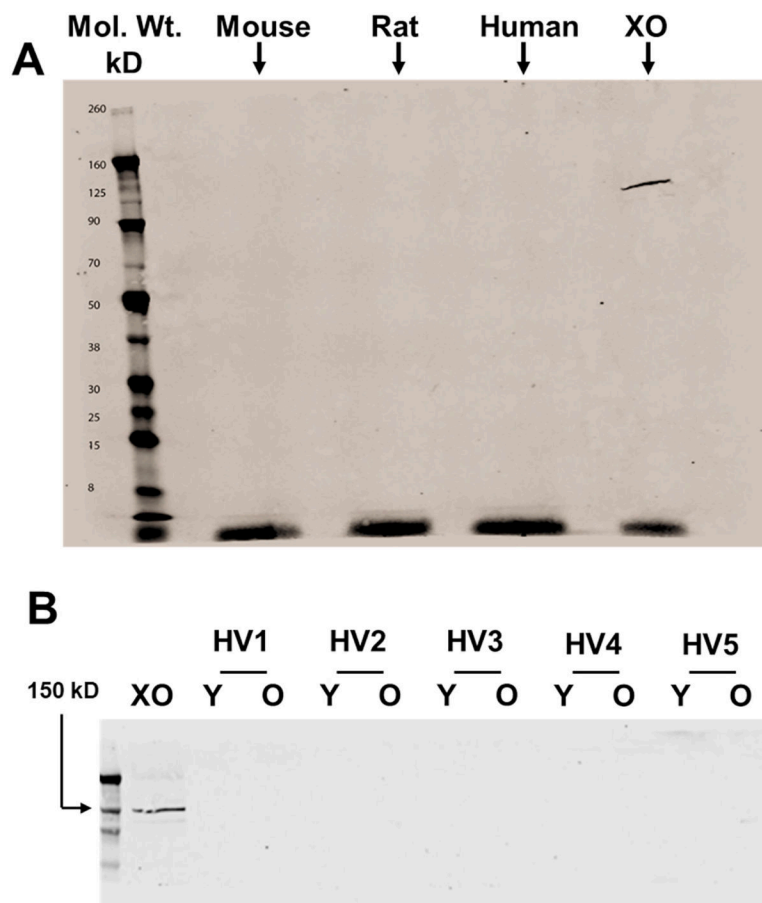


**Fig. 1.**

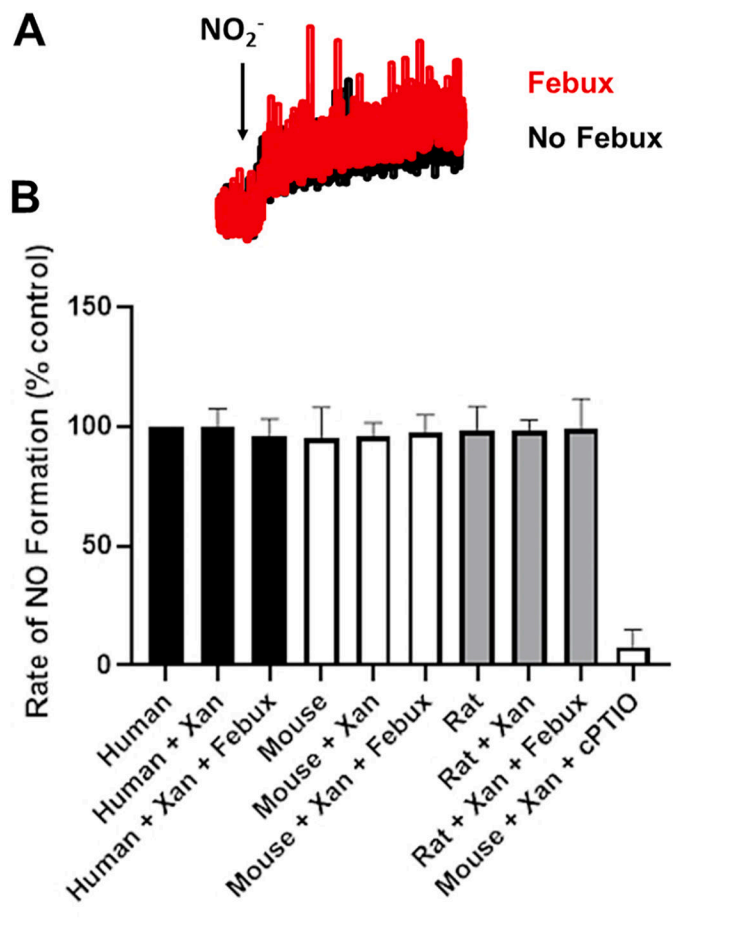
Human, mouse, and rat RBCs do not demonstrate XO activity. Human, mouse, and rat plasma (A) and washed RBC homogenates (B) were analyzed for XO activity as described in the methods. The volume of RBC homogenate used was 16  $\mu\text{L}$  which corresponds to 0.8 mg protein. Data represent the mean  $\pm$  standard deviation,  $n = 6$  for human,  $n = 3$  mice, and  $n = 4$  for rat samples.



**Fig. 2.** RBCs do not bind XO. A) Cartoon depiction of XO bound to heparin-Sepharose 6B. B) Purified XO (50 mU) was added to human blood for 15 min and then the samples washed under identical conditions as in Fig. 1 and in the methods. Samples were then assessed for XO enzymatic activity as in Fig. 1. Control samples were exposed to either the XO inhibitor allopurinol (Allo) or febuxostat (Febux) as indicated. Data represent the mean  $\pm$  standard deviation  $n = 4$ .



**Fig. 3.** Assessment of RBC-associated XOR protein abundance and impact of cellular age. A) Western blot analysis was performed on human, mouse, and rat RBC homogenates using a monoclonal anti-XOR antibody (Santa Cruz) as described in the methods. Purified XO was used as a positive control and electrophoresed using 0.5  $\mu$ g total protein. A total of 30  $\mu$ g of protein was electrophoresed for each homogenate (human, mouse, and rat). Shown is a representative of 3 independent blots representing a total of 3 mice, 3 rats and 3 humans. B) Western blot analysis was performed on five different human RBC homogenates, either young or old, using the same mouse monoclonal anti-XOR antibody as described above. Purified XO was used as a positive control and electrophoresed using 0.5  $\mu$ g total protein. (HV = healthy volunteer/Y = young/O = old).



**Fig. 4.** Human and Rodent RBCs do not demonstrate XO-catalyzed nitrite reductase activity. Human, mouse, and rat samples were analyzed for XOR-dependent contributions to overall  $\text{NO}_2^-$  reductase activity using the Sievers Nitric Oxide Analyzer (NOA) as detailed in the methods. Samples were prepared in the reaction chamber with 50  $\mu\text{L}$  of RBC homogenate (5 mg total protein) in KPi pH 7.0 with or without xanthine (25  $\mu\text{M}$ ) and with or without febxostat (10  $\mu\text{M}$ ). Reactions were initiated by the addition of  $\text{NaNO}_2^-$  (500  $\mu\text{M}$ ). A) Representative NO generation profiles from human RBC homogenates + xanthine with or without addition of febxostat (Febux). B) Human, mouse, and rat RBC homogenate values (alone or with xanthine or with xanthine and febxostat). Authentication of NO was determined by diminution of the signal by the presence of the electron spin resonance NO spin trap carboxy-PTIO. Data represent the mean  $\pm$  SD of  $n = 4$ .