

# Humanized mouse model of glucose 6-phosphate dehydrogenase deficiency for in vivo assessment of hemolytic toxicity

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Edited by Thomas E. Wellems, National Institutes of Health, Bethesda, MD, and approved September 10, 2013 (received for review June 26, 2013)

**Individuals with glucose 6-phosphate dehydrogenase (G6PD) deficiency are at risk for the development of hemolytic anemia when given 8-aminoquinolines (8-AQs), an important class of antimalarial/anti-infective therapeutics. However, there is no suitable animal model that can predict the clinical hemolytic potential of drugs. We developed and validated a human (hu)RBC-SCID mouse model by giving nonobese diabetic/SCID mice daily transfusions of huRBCs from G6PD-deficient donors. Treatment of SCID mice engrafted with G6PD-deficient huRBCs with primaquine, an 8-AQ, resulted in a dose-dependent selective loss of huRBCs. To validate the specificity of this model, we tested known nonhemolytic antimalarial drugs: mefloquine, chloroquine, doxycycline, and pyrimethamine. No significant loss of G6PD-deficient huRBCs was observed. Treatment with drugs known to cause hemolytic toxicity (primaquine, sitamaquine, tafenoquine, and dapsone) resulted in loss of G6PD-deficient huRBCs comparable to primaquine. This mouse model provides an important tool to test drugs for their potential to cause hemolytic toxicity in G6PD-deficient populations.**

**G**lucose-6-phosphate dehydrogenase (G6PD) deficiency is the most common human enzyme deficiency, with an estimated 400 million people worldwide affected by this enzymopathy (1). G6PD-deficient RBCs are uniquely sensitive to oxidative stress. Several drugs induce oxidative stress in G6PD-deficient RBCs, including the antimalarial drug primaquine (PQ), an 8-aminoquinoline (8-AQ) (2). The damaged G6PD-deficient RBCs are subsequently cleared from the peripheral circulation, resulting in “hemolytic” anemia (2). 8-AQs are the only approved drug class able to eliminate the hypnozoite stages of the malaria parasite (3), as well as stage V *Plasmodium falciparum* gametocytes (4, 5). These characteristics make them an ideal drug class for malaria elimination campaigns (6, 7). However, because of the hemolytic toxicity associated with PQ, this drug has sharply limited utility in public health programs for the treatment of malaria. Although new 8-AQs have been developed (8), the lack of a relevant animal model to predict hemolytic toxicity in the context of G6PD deficiency has hindered further development of this class of antimalarial drugs (2).

SCID mice have routinely been used as models of human disease (9–12). Lacking in functional immune responses, and therefore able to accept xenogenic transplants, SCID mice have been widely used as hosts for the engraftment of both normal and malignant human cells including human RBCs (huRBCs) (13, 14). Subsequent studies on the engraftment of huRBCs improved the degree and persistence of engraftment by using sublethal irradiation, chemical treatment protocols, or inclusion of denatured human serum and repeated administration of huRBCs to enhance engraftment efficiency (15–18). Recent advances in the development of immunodeficient mouse models have yielded mice with higher engraftment capacities and with minimal manipulation before engraftment. SCID mice developed on the nonobese diabetic (NOD) background (NOD/SCID) are

able to support huRBCs for prolonged periods after repeated i.p. injections of huRBCs (19–21).

We report here the development of an NOD/SCID mouse model engrafted with G6PD-deficient huRBCs. Treatment with PQ and other drugs known to induce hemolytic anemia in humans produced hemolytic responses in mice engrafted with G6PD-deficient huRBCs, suggesting that this model can be used for testing drugs for the prediction of the hemolytic toxicity. Use of this model will expedite the development of safer drugs for malaria treatment and prophylaxis, as well as other antiparasitic diseases for which 8-AQs have demonstrated efficacy.

## Results

**Engraftment of G6PD-Deficient Human Erythrocytes into NOD/SCID Mice.** A number of genetic variants resulting in loss of G6PD activity have been described (22). Depending on the genetic variant, drug-induced hemolysis may be mild and self-limiting as observed in individuals with the African variant (A-) or severe and life threatening as described in individuals with the Mediterranean variant (Med-) (23). To determine whether huRBCs from G6PD-deficient donors could be successfully engrafted into NOD/SCID mice, RBCs from A- or Med- donors or donors with normal G6PD activity were injected i.p. daily for 14 d (herein referred to as A- mice, Med- mice, or normal huRBC SCID

## Significance

This article describes the development and validation of a novel mouse model that can be used to predict hemolytic toxicity of drugs that occurs in individuals with an enzyme deficiency known as glucose-6-phosphate dehydrogenase (G6PD) deficiency. G6PD deficiency affects more than 400 million people worldwide. In this model, nonobese diabetic/SCID mice are transfused with human RBCs from G6PD-deficient donors. Treatment with drugs known to cause hemolytic anemia in humans do not cause damage to the mouse RBCs nor to the transfused normal human RBCs; but a robust hemolytic response is observed in the mice transfused with G6PD-deficient human RBCs. The immediate impact of this model will be in advancing the development antimalarial drugs.

This work was presented in abstract form at the 59th Annual Meeting of the American Society of Tropical Medicine and Hygiene, Atlanta, GA, November 3–7, 2010.

Author contributions: R.R., C.O., P.C.B., B.C., A.S., A.J.M., B.L.T., and L.A.W. designed research; R.R. and P.C.B. performed research; R.R., C.O., P.C.B., B.C., A.S., A.J.M., B.L.T., and L.A.W. analyzed data; and R.R., C.O., P.C.B., B.C., A.S., A.J.M., B.L.T., and L.A.W. wrote the paper.

The authors declare no conflict of interest.

This article is a PNAS Direct Submission.

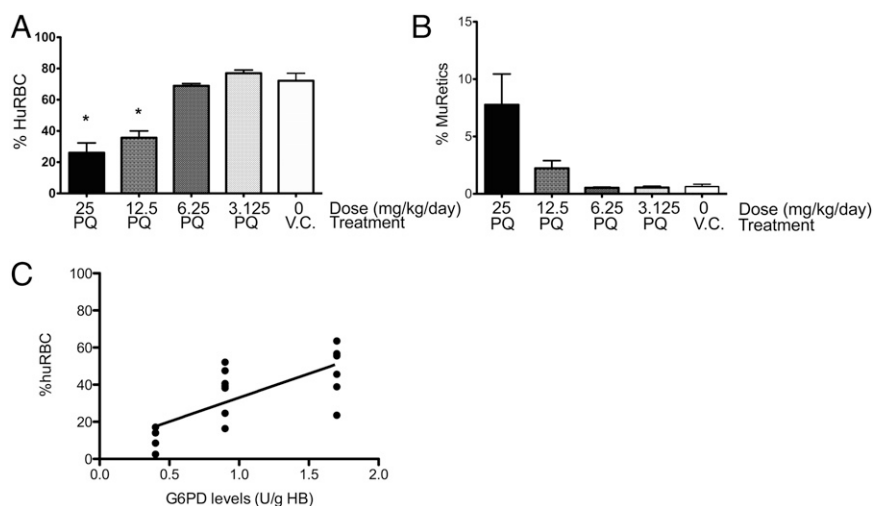
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**Fig. 3.** Evaluation of dose of PQ and G6PD levels on hemolytic response. A- huRBC SCID mice were treated with PQ p.o. daily for 7 d with indicated doses of PQ (25, 12.5, 6.25, or 3.125 mg/kg per day) or given PBS as a vehicle control and assessed for percentage huRBCs (A) and muRetics (B) at day 7 by FACS analysis. Each treatment group consisted of 4 mice. \*Statistically significant differences (one-way ANOVA with Bonferroni posttest,  $P < 0.001$ ). (C) G6PD levels were measured at start of engraftment and correlated with percentage of huRBCs at 7 d after treatment with PQ (25 mg/kg per day, p.o., 7 d) using Spearman rank correlation ( $P = 0.0005$ ,  $r = 0.7373$ ). Data pooled from three experiments with three donors. Each dot represents an individual mouse.

known to be free from hemolytic toxicity. Thus, A- huRBC SCID mice were treated with pamaquine, one of the first synthetic antimalarial drugs to be introduced clinically (2), with sitamaquine, another 8-AQ analog that has been under development for the treatment of leishmaniasis (26, 27), and with tafenoquine, currently under development for malaria prophylaxis (28). Like PQ, pamaquine, sitamaquine, and tafenoquine cause hemolysis in individuals with G6PD deficiency (26, 27). A- huRBC SCID mice were treated with pamaquine (50 mg/kg or 75 mg/kg) or with sitamaquine (40 mg/kg) p.o. once daily for 7 d (Fig. 4). The pamaquine doses of 75 and 50 mg/kg per day and sitamaquine dose of 40 mg/kg per day were selected on the basis of the clinical dose regimen relative to PQ. Tafenoquine (2.5 mg/kg p.o. once daily for 3 d) was also tested using a dose regimen shown to be effective for causal prophylaxis in mice. The percent loss of huRBCs was calculated at the end of 7 d of treatment (relative to percentage of huRBCs at start of treatment). Pamaquine, sitamaquine, and tafenoquine treatment resulted in loss of huRBC in A- G6PD mice at levels similar to PQ-treated mice but significantly different from vehicle control-treated mice (one-way ANOVA,  $P < 0.0001$ ).

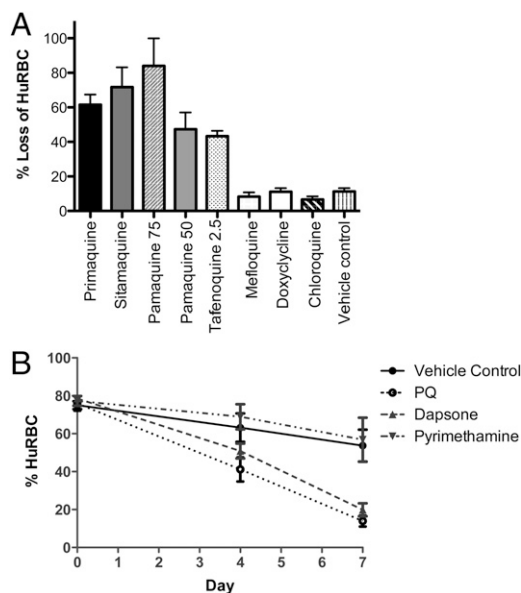
We next tested two other antimalarial drugs with the basic quinoline structure but that are known to be free of hemolytic potential clinically: chloroquine, a 4-aminoquinoline, and mefloquine, a quinoline methanol. An additional structurally unrelated drug, doxycycline, was also used. Doses were selected on the basis of doses known to produce efficacy in mouse blood stage malaria infection. HuRBC levels in chloroquine-, mefloquine-, and doxycycline-treated A- huRBC mice were similar to those seen in the vehicle controls (Fig. 4A). These results corroborate that the loss of huRBCs we observe in G6PD-deficient huRBC SCID mice was observed with 8-AQ drugs known to have hemolytic toxicity in G6PD individuals and was not a nonspecific response to quinoline antimalarial drugs. Finally, in a separate experiment we tested a known hemolytic drug, dapsone—which is not an 8-AQ but is known to cause hemolysis in both G6PD-deficient and normal individuals (29)—along with a known nonhemolytic drug used in malaria chemotherapy, pyrimethamine. As seen in Fig. 4B, dapsone induced loss of huRBCs similar to PQ, whereas pyrimethamine had no effect. These data suggest that this model reliably reflects known human red cell toxicity in G6PD-deficient subjects.

## Discussion

Primaquine, an 8-AQ, is an important antimalarial drug because treatment with PQ can eradicate the liver stage of *Plasmodium vivax* (6) and the stage V *P. falciparum* gametocytes (7). However, PQ treatment causes hemolytic anemia in G6PD-deficient individuals (4, 27, 30), a major drawback for its application in malaria elimination campaigns. Development of nonhemolytic 8-AQ antimalarials (or at least 8-AQ derivatives with an improved therapeutic window) is therefore critical so that this class of compounds can achieve more widespread use for the treatment and elimination of malaria. A major limitation in drug development has been the lack of a suitable animal model to test potential hemolytic toxicity of novel drugs. In this article we describe the development of a humanized NOD/SCID mouse model containing circulating G6PD-deficient human RBCs that reproduces drug-induced hemolytic toxicity and thus can be used for the testing and validation of the hemolytic capacity of new 8-AQ analogs and other drugs with potential hemolytic toxicity.

The optimum engraftment protocol for the human red cells into the NOD/SCID mice results in high relative huRBC levels—approximately 80% of total red cells—that are stably maintained for at least 7 d with cells from a G6PD normal donor. With an A- type G6PD-deficient donor, stability is a bit lower but still relatively comparable to normal huRBCs. However, with the Med- deficiency, stability is much reduced, likely reflecting the almost complete lack of G6PD activity and resultant marked depletion of cellular NADPH and reduced glutathione. These would compromise red cell oxidative defenses and reduce cellular metabolic competency.

PQ-induced hemolytic anemia in G6PD individuals is not immediate but occurs after several days of treatment (31). Clinical indicators of hemolytic anemia in humans include decreased hemoglobin concentration, low hematocrit levels, low haptoglobin, reticulocytosis, elevations in unconjugated bilirubin and lactate dehydrogenase, and Heinz body formation (2, 32, 33). In our model, drug-induced hemolysis was evaluated using two primary parameters: loss of human RBCs and induction of murine reticulocyte production. These were measured using sensitive and quantitative flow cytometry-based assays. We observed both loss of huRBCs and an increase in circulating mouse reticulocytes in our model after PQ treatment, regardless of



**Fig. 4.** Qualification of the G6PD-deficient huRBC SCID mouse model with other antimalarial drugs. (A) A- huRBC SCID mice were treated p.o. with PQ (25 mg/kg per day for 7 d), vehicle control (PBS for 7 d), pamaquine (75 or 50 mg/kg per day for 7 d), sitamaquine (40 mg/kg per day for 7 d), tafenoquine (2.5 mg/kg per day for 3 d), chloroquine (25 mg/kg per day for 7 d), mefloquine (40 mg/kg per day for 3 d), or doxycycline (60 mg/kg per day for 7 d). The percentage of huRBCs as measured by flow cytometry at day 7 of treatment was subtracted from the percentage of huRBCs at start of treatment and then divided by percentage of huRBCs at start of treatment to determine the percent loss of huRBCs.  $n = 15$  for PQ and vehicle control,  $n = 4$  for other groups. (B) A- huRBC SCID mice were treated p.o. with PQ (25 mg/kg per day for 3 d), dapson (20 mg/kg per day for 3 d), pyrimethamine (50 mg/kg per day for 3 d), or vehicle control. Percentages of huRBCs were assessed at posttreatment day 0, day 4, and day 7.

route of administration. The decline in huRBCs was not observed until after 3 d of PQ treatment and mirrors the kinetics in huRBC loss reported by Beutler et al. (31) in early studies of PQ drug trials in human volunteers that were G6PD-deficient. The delay is thought to be due to the accumulation of red cell damage over time with recycling of redox-active metabolites. The red cell loss is likely “extravascular,” in that it results from the removal of damaged RBCs from the circulation by tissue (e.g., splenic) macrophages via phagocytosis; this phenomenon was demonstrated in an *in vitro* erythrophagocytosis assay (34). In the present studies, posttreatment spleen and liver weights were also evaluated as indirect/secondary markers of hemolysis. The major organ of red cell sequestration and removal is the spleen (34). The degree of red cell damage and removal determines the size of spleen; high RBC clearance results in larger spleen size. Consistent with this model, we observed increases in spleen size and no effect on liver size in A- and Med- huRBC NOD/SCID mice given doses of PQ that induced loss of huRBCs.

We observed increases in murine reticulocytes in both A- and Med- huRBC NOD/SCID mice, also consistent with PQ-induced hemolysis. During severe and rapid hemolysis of RBCs, the bone marrow increases the rate of RBC production, resulting in significantly higher than normal reticulocyte numbers in the peripheral blood; reticulocytes normally account for less than 1% of total red blood cells but after hemolysis may comprise up to 15% (35). Interestingly, we observed a biphasic increase in murine reticulocytes, with an early lower response observed at 2 to 3 d when huRBCs first declined and then a second larger increase by day 7, greater than 10% murine reticulocytes in Med-huRBC mice. These observations would suggest a switch from

bone marrow to extramedullary erythropoiesis, which can occur under conditions of oxygen stress.

The effects of the PQ were specific to NOD/SCID mice engrafted with A- or Med- G6PD-deficient blood; NOD/SCID mice engrafted with G6PD-normal blood did not exhibit hemolysis. As expected, the more severe hemolysis after PQ treatment occurred in the Med- huRBC SCID mice compared with A-huRBC SCID mice, reflecting the differences in susceptibility to oxidative stress of RBC and greater fragility (as seen by poorer engraftment and reduced stability after engraftment) in Med-erythrocytes.

Med- G6PD-deficient humans given PQ also show an exaggerated hemolytic response (23). These findings suggest that the hemolytic effect of PQ is specific to G6PD-deficient huRBCs and that the more severe the G6PD deficiency (e.g., Med-), the more rapid the loss of huRBCs after PQ treatment. The correlation between G6PD levels and loss of huRBC observed in this study (Fig. 3C) is also consistent with this model.

Our model is therefore an appealing screening tool for the hemolytic potential of drugs at the preclinical stage. We further validated the model by demonstrating that two other 8-AQ derivatives (pamaquine and sitamaquine), as well as dapson, a non-8-AQ, all known to produce clinical hemolysis, also induced hemolysis in our model; further, drugs known not to have hemolytic potential (mefloquine, chloroquine, pyrimethamine, and doxycycline) showed no activity, thus suggesting that this model can be used to evaluate potential human hemolytic toxicity.

The clinical picture for hemolysis in malaria patients treated with PQ may be considerably more complicated, because the parasite-induced effects in red blood cells also result in oxidative stress, and this could conceivably render cells more susceptible to PQ-induced hemolysis. The G6PD-deficient erythrocytes are considered relatively resistant to the malaria infection, predictably owing to preexisting oxidative stress (36). The *P. falciparum* infection in antioxidant-blunted G6PD-deficient RBCs results in a remarkable increase in expression of enzymes associated with antioxidant defense (37).

Taken together, these findings demonstrate the development of a G6PD-deficient huRBC SCID mouse model that can be used to evaluate hemolytic toxicity of new drugs for the treatment of malaria in the G6PD-deficient populations. Use of this model to identify nonhemolytic 8-AQ analogs, 8-AQs that have less hemolytic effect, and coadministration of 8-AQ plus partner drugs that would protect huRBCs from the hemolytic effect of 8-AQ will also afford a more robust and comprehensive approach to reduce the hemolytic liability associated with 8-AQs in G6PD-deficient individuals and aid in the campaigns to eliminate malaria.

## Materials and Methods

**Mice.** Eight- to nine-week-old female NOD.CB17-Prkdc<sup>scid</sup>/J mice (herein referred to as NOD/SCID mice) were purchased from the Jackson Laboratories. Mice were maintained on sterile food and water in a specific pathogen-free animal facility at Upstate Medical University and were routinely acclimated for 1 wk in the facility before initiation of experimental procedures. Softened food was provided in sterile water in the cage during drug treatment as part of standard of care. All animals were treated according to Upstate Medical University’s Committee for the Humane Use of Animals and in compliance with the Animal Welfare Act and other federal statutes and regulations relating to animals and experiments involving animals and adhere to the principles stated in the *Guide for the Care and Use of Laboratory Animals* (38).

**Collection and Processing of Donor Blood.** G6PD-deficient and G6PD-normal volunteer blood donors were recruited through the Walter Reed Army Institute of Research (WRAIR), Silver Spring, MD, under approved institutional review board protocols. The donor blood samples were analyzed for G6PD levels using the Kinetic method (Quest Diagnostics) and confirmed by genomic sequencing for G6PD deficiency. Donor G6PD levels were 0–0.1 IU/g HgB for the Med- donors ( $n = 2$ ) and 0.4–1.9 IU/g HgB for the A- donors ( $n = 8$ ). Genomic DNA was isolated from whole blood using Blood Mini kits

(Qiagen) according to the manufacturer's protocol. Specific PCR reactions were performed to isolate exons 3–5 that correspond to the common A/A-allele found in patients of African descent, or exon 6 that corresponds to the Mediterranean mutant allele (39). PCR products were checked for size on agarose gels and purified using a Qiagen PCR Purification kit and sequenced on both forward and reverse strands across the regions of interest to verify presence of mutant allele. Up to 450 mL of leukocyte-depleted whole blood was obtained by venous puncture in citrate phosphate dextrose adenine anticoagulant. Blood was transported overnight in a controlled temperature-cooling chamber to Upstate Medical University. Whole blood was centrifuged at  $900 \times g$  for 40 min at room temperature (RT). Plasma was carefully aspirated and stored at 4 °C. RBCs were resuspended in RPMI 1640 (HyClone Laboratories) and washed twice at  $900 \times g$  for 20 min each at RT. Plasma-free RBCs were reconstituted in RPMI 1640 with 25% autologous plasma at a concentration of  $5 \times 10^9$  RBC/mL. RBC were aliquoted into 14 equal proportions and stored at 4 °C.

**Engraftment of RBCs into NOD/SCID Mice.** NOD/SCID mice were injected i.p. once daily with a 1-mL suspension of RBCs for 14 d. An aliquot of RBCs was prewarmed to 37 °C before injection. Engraftment levels were determined at the end of 14 d. Approximately 5  $\mu$ L of tail snip blood was collected into 100  $\mu$ L of sterile heparin-PBS (10 U sodium heparin per mL). Using V-bottomed 96-well plates,  $2 \times 10^5$  RBCs were stained for 30 min at 4 °C in the dark with phycoerythrin (PE)-conjugated anti-human CD235a antibody (Abcam) in staining buffer (0.5% BSA in PBS). Cells were washed twice and reconstituted in staining buffer for assessment by flow cytometry. Baseline levels of muRetics were also determined. RBCs ( $2 \times 10^5$ ) were stained with an FITC-labeled anti-mouse CD71 (eBioscience), a reticulocyte marker, and a PE-labeled anti-TER119 (Invitrogen) monoclonal antibody for mouse erythroid

cell detection. Cells were acquired on a Guava EasyCyte Plus flow cytometer (Millipore). Analysis of the flow data was done using FlowJo software (TreeStar).

**Drug Treatment.** Mice with peripheral huRBC levels greater than 60% were randomized for drug treatment, with 4 to 5 mice per group assigned. Drugs were provided by the Division of Experimental Therapeutics, WRAIR, or by Swiss Tropical and Public Health Initiative and were reconstituted in appropriate vehicle (PBS, hydroxyethyl cellulose-Tween, or Tween-EtOH). Tail-snip blood was collected on days 2, 4, 5, 6, and 7 of treatment for the determination of huRBC and muRetic levels. On the last day of treatment, mice were killed 1 hour after last dose and were assessed for hematocrit levels and body, spleen, and liver weights. Liver and spleen weights were calculated as a percent of total body weight for each mouse to account for variability in animal weights.

**Statistics.** Statistical analysis was performed using GraphPad Prism (GraphPad Software). Mann-Whitney or one-way ANOVA with Bonferroni correction at 95% confidence intervals was used to determine differences between groups as appropriate.

**ACKNOWLEDGMENTS.** We thank Caroline Othoro, Marino Mauro, Julie Ritchie, and Nancy Fiore for excellent technical assistance; and Jeff Friedman (The Scripps Research Institute) for genotyping the G6PD-deficient donors. This work was supported by grants from the US Army Medical Research and Materiel Command, via Awards W81XWH-07-2-0095 and WX81XWH-10-2-0059 to the University of Mississippi (to L.A.W.). Funding was also provided by the Medicines for Malaria Venture (to R.R.).

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