CASE REPORT

Case Report: Paroxysmal nocturnal hemoglobinuria in a woman heterozygous for G6PD A- [version 1; peer review: 2 approved with reservations]

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Abstract
We describe a case of paroxysmal nocturnal hemoglobinuria (PNH) in a woman who is heterozygous for the glucose-6-phosphate dehydrogenase A- (G6PDA-) allele. PNH is associated with one or more clones of cells that lack complement inhibition due to loss of function somatic mutations in the PIGA gene. PIGA encodes the enzyme phosphatidylinositol glycan anchor biosynthesis, class A, which catalyses the first step of glycosylphosphatidylinisotol (GPI) anchor synthesis. Two GPI anchored red cell surface antigens regulate complement lysis. G6PD catalyses the first step of the pentose phosphate pathway and enzyme variants, frequent in some populations have been because they confer resistance to malaria, are associated with hemolysis in the presence of oxidizing agents including several drugs. The patient had suffered a hemolytic attack after taking Bactrim, a drug that precipitates hemolysis in G6PD deficient individuals. Since both G6PD and PIGA are X-linked we hypothesized that the PIGA mutation was on the X-chromosome carrying the G6PDA- allele. Investigations showed that in fact the PIGA mutation was on the X-chromosome carrying the normal G6PD B allele. We speculate that complement activation on G6PD A- red cells exposed to Bactrim might have triggered complement activation inducing the lysis of G6PD B PNH Type II red blood cells or that the patient may have had a PNH clone expressing G6PDA- at the time of the hemolytic episode.
Corresponding author: Philip Mason (masonp@email.chop.edu)

Competing interests: No competing interests were disclosed.

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**Materials and methods**

Peripheral blood from patient CHOP277.01 was obtained after obtaining written informed consent according to the declaration of Helsinki. The Internal Review Board of the Hospital of the University of Pennsylvania approved this study. DNA and RNA were extracted by using QIAamp DNA and RNA Blood mini Kits, respectively, according to manufacturers’ instructions. Blood samples for fluorescent cytometry and electrophoretic analyses were obtained from EDTA tubes and experiments were performed within 2 hours of blood withdrawal.

PCR primers to detect mutations confirming the G6PD A- genotype were designed with Primer3 v4.0. (primers for c.202 G->A mutation: forward 5’- agagaagatctaccctccacatct-3’ and reverse 5’-cttgtacagggcagagccagc-3’; primers for c.376A->G: forward 5’-catctgtctgtgctgtctgtc-3’ and reverse 5’-ctcagagttcttgagggac-3’). Sanger sequencing was done by the Nucleic Acids core facility at CHOP.

The HUMARA assay was performed as previously described. Briefly, Hhad digested and non digested DNA was subjected to PCR amplification of the first exon of the HUMARA locus (containing a CAG repeat) using fluorochrome-coupled primers. Amplification products were then migrated on an ABI Prism 3100 Automated Genetic Analyzer (Applied Biosystems). Allele calling and the area under the curve (AUC) were determined using GeneMapper v.4.0 software (Applied Biosystems). The AUC was used to calculate the skewing from X chromosome inactivation (XCI). The XCI ratio of the digested fraction was corrected with that of the undigested fraction to allow for preferential amplification of the smallest allele (i.e., the allele containing less CAG repeats). Skewing is present when the percentage of the predominant allele exceeds 74%. A percentage of predominant allele between 90% and 100% is considered extreme skewing.

Measurements of oxidative stress ROS assay was performed as previously described. Briefly, red blood cells were incubated with 20-70-dichlorofluorescein diacetate (DCF; Sigma) dissolved in methanol. After incubation at 37°C for 15 minutes in a humidified atmosphere of 5% CO2 in air, the cells were washed, resuspended in PBS and analyzed by flow cytometry (FACSCalibur; Becton-Dickinson, Immunofluorometry Systems, Mountain View, CA, USA). The mean fluorescence channel (MFC) was calculated by FACSDiv software. The identity of the red cell population was verified by staining with an antibody anti to glycophorin-A. To determine the presence of GPI proteins, cells were labeled with a phycoerythrin-conjugated anti-CD55 antibody. For our experiment, cells from a non PNH- non G6PD individual served as control. The MFC of cells stained with DCF, is proportional to generation of ROS.

The electrophoretic mobility of the protein was performed in cellogel strips as previously described. Hemolysates treated with and without acidified serum were run in order to assess differences in mobility of the G6PD enzyme.

**Case report**

The patient, a 23 year old African American woman, presented with left upper abdominal pain, vomiting and blurry vision. She had a four year history of episodes of hemolysis, abdominal pain and dark urine.
She was diagnosed with acute hemolytic exacerbation of PNH and was admitted. The patient was found to have anemia, leukocytosis and mild transaminitis. A Magnetic Resonance Venogram (MRV) of the abdomen and pelvis revealed several ill-defined low attenuation lesions of the posterior segment of the right hepatic lobe consistent with liver thrombosis so she was started on anticoagulation. A concomitant urinary tract infection was treated with Trimethoprim-Sulfamethoxazole (Bactrim) and she was discharged.

After discharge, she followed up with a hematologist. On review of systems, the patient complained of a hemolytic attack with dark urine after taking Bactrim. On physical exam, she was found to have mildly icteric sclera. On further questioning of family history of other hematologic disorders, the patient said that one of her nephews has Glucose-6-Phosphate-Dehydrogenase deficiency. Because of her diagnosis of PNH and abdominal thrombosis, she was started on Eculizumab. The laboratory workup revealed a white blood cell count 4.6 k/uL, RBC count of 3.8 k/uL, hemoglobin of 11.8 g/dL, hematocrit 35%, MCV 91 fl, platelets 327 k/uL, reticulocytes 24.5%, PTT 27.8, INR 1.7, D dimer 2.5 ug/mL, bilirubin 1.3 mg/dL, bilirubin direct 0.2 mg/dL, bilirubin indirect 1.1 mg/dL, ANC 2.7 k/uL, ALC 1.3 k/uL, G6PD screen normal, LDH 720 U/L. The flow cytometry at this point revealed: 69% of the red blood cells were partially deficient for CD59 and 9.1% of the red blood cells were completely deficient for CD59; 86.3% of the granulocytes were partially deficient for CD59 and 6.4% were completely deficient for CD59.

**Results**

The family history and the case history suggested that the patient may have both G6PD deficiency and PNH, since the G6PD deficiency might explain the hemolysis precipitated by Bactrim, a drug that is reported to cause hemolysis in G6PD patients. Sequencing of DNA from her granulocytes confirmed that she was heterozygous for G6PD A- having a *G6PD B* allele on one X-chromosome and a *G6PD A* allele on the other. This finding raised the question as to whether the somatic *PIGA* mutation causing her PNH took place on the X-chromosome carrying the *B* or the *A* *G6PD* gene. The flow cytometry data showed that the patient most likely had 2 PNH clones, a class I clone (partial deficiency) of about 86% and a class II clone (complete deficiency) of about 6%. The HUMARA assay showed a single clone of about 90% (Figure 1), suggesting that in both clones the mutation had taken place on the same X-chromosome. We hypothesized that the mutations in *PIGA* would have taken place on the chromosome carrying the *G6PD A*- allele since this would help explain the patient’s reaction to Bactrim. Another factor was that treatment with eculizumab, by inhibiting lysis of PNH red cells may have led to a higher level of PNH (and concomitantly G6PD deficient) red cells than would be present in untreated patients. To determine which *G6PD* allele was expressed in the PNH clone we sequenced cDNA from the patient’s granulocytes. The sequencing trace showed that the vast majority of expressed G6PD cDNA contained the wild type (*G6PD B*) sequence at both nucleotides where it differs from *G6PD A* (Figure 2), leaving us to conclude that the PIGA mutations had taken place on the X-chromosome containing the *G6PD B* allele. This was confirmed at the protein level since the red blood cells lysed by acidified serum (the PNH cells) contained most of the G6PD activity while the residual cells did not contain detectable G6PD activity. While developing our hypothesis, which turned out to be incorrect, we also considered whether PNH/G6PD A- cells might have high levels of oxidative stress since both G6PD deficiency and PNH have been shown to be associated with elevated levels of reactive oxygen species. We found that the patient’s red blood cells contained ROS levels that were significantly higher than those from healthy controls, though surprisingly we did not detect any difference in ROS between PNH (CD55-) and normal (CD55+) cells (Figure 3).

**Figure 1.** Clonal hematopoiesis in patient CHOP 277.01. The panel at the bottom shows the migration of the 2 microsatellite alleles in patient CHOP277.01 revealed by PCR analysis of a region of the Androgen receptor gene on the X-chromosome. The amplified region contains both a polymorphic repeat and a site for the methylation sensitive restriction enzyme HhaI. This site is methylated on the inactive X-chromosome. The top panel shows the same sample digested with HhaI before the PCR so only the fragment on the inactive X-chromosome is amplified. An imbalance in the allelic ratio reflects an imbalance in X inactivation and therefore indicates clonality.
**Figure 2.** Patient CHOP 277.01 is heterozygous for G6PD B/A- and the B allele is predominantly expressed. A1 and A2 shows sequencing of genomic DNA around the 202 G→A and the 376 A→G mutation that lead to Val68Met and Asn126Asp changes in G6PD, respectively. Figures B1 and B2 show the cDNA sequence of the exact same mutations in the peripheral blood of the patient. The WT (B) allele is expressed in the majority of cells.

**Discussion**

PNH is a rare condition, having an incidence of about 1 in a million, so the co-incidental finding of a female with PNH and heterozygous for G6PD deficiency was an opportunity to observe the interaction between these 2 conditions which both involve red blood cell hemolysis mediated by X-linked genes. Notably the first demonstration that PNH was a clonal disease took advantage of a female PNH patient who was heterozygous for the electrophoretic variant 

\[ \text{G6PD A} \]

In this patient both isozymes were present in a lysate of total red blood cells, but only one was present after acidified serum lysis, demonstrating clonality of the PNH cells.

In the case discussed here a female African American patient with PNH suffered episodes of hemolysis, often following treatment with Trimethoprim-Sulfamethoxazole (Bactrim), one of the drugs that is known to cause hemolysis in G6PD patients\(^1\). When it emerged that she was heterozygous for 

\[ \text{G6PD A} \]

we hypothesized that her expanded PNH clones may be expressing only the 

\[ \text{G6PD A} \]

protein. The hypothesis proved incorrect and the clone expressed the wild type 

\[ \text{G6PD} \]

allele. A possible explanation is that complement activation on G6PD A- red cells exposed to Bactrim might have triggered complement activation inducing the lysis of G6PD B PNH Type II red blood cells\(^1\). Alternatively since PNH patients often have several clones\(^1\) which change in prevalence it is possible that earlier she did have a G6PD A- clone but this was replaced by wild type clones. Finally we can also speculate that the combination of G6PD and PIGA deficiency confers a serious growth disadvantage and PNH clones in this situation are more likely to be G6PD wild type. There is no clear mechanism for this however as G6PD A- nucleated cells have similar enzyme activity to WT cells\(^1\) – the deficiency becoming apparent in red blood cells, which do not synthesize new protein\(^1\).

**Patient consent**

Written informed consent for publication of their clinical details was obtained from the patient.

**Author contributions**

PJM and MB conceived the study, NP designed the experiments and carried out the research. MM collected and collated clinical data. All authors contributed to preparing a draft of the manuscript and have agreed to the final content.

**Competing interests**

No competing interests were disclosed.

**Grant information**

The work has been supported by the Buck Family Endowed Chair in Hematology, and by NCI NIH grants 2R01CA106995 to PJ Mason, and 2R01 CA105312 to M Bessler.

The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

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London, UK

Perdigones and colleagues describe the rare co-occurrence in the same patient of two X-linked
disorders, one acquired the other inherited, both causing intravascular haemolysis.

However how the interaction of the two disorders led to the clinical episode described in the case
report is not clear because the temporal analysis of the clinical events and the associated
laboratory tests are not presented in sufficient detail.

Charting clinical events, labs and therapeutic interventions might make association of the
haemolytic episodes with co-trimoxazole or intercurrent infection clearer.

It appears that the majority of the PNH clone (in both granulocytes and red cells) is type II, i.e.,
only partially deficient of GPI. One assumes that this picture was obtained after the haemolytic
attack. If so, it could be that type III, i.e., severely deficient RBC were indeed G6PD-deficient. Was
flow-cytometry performed after Eculizumab treatment? Eculizumab would protect type III RBC
from lysis and thus would allow re-assessment of G6PD activity.

Minor points
- Was anti-CD55 or –CD59 was used for flow analysis? The authors state anti-CD55 in methods
but describe anti-CD59 in results.

- Normal ranges of lab tests need to be provided

- ‘Another factor was that treatment with eculizumab, by inhibiting lysis of PNH red cells may
have led to a higher level of PNH (and concomitantly G6PD deficient) red cells than would be
present in untreated patients.’ This statement is rather irrelevant because eculizumab was
started after treatment with co-trimoxazole.

- Need to use Greek characters appropriately, i.e., ul should be ml
Competing Interests: No competing interests were disclosed.

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard, however I have significant reservations, as outlined above.

Author Response (F1000Research Advisory Board Member) 15 Oct 2014

Philip Mason, Abramson Pediatric Research Center, The Children’s Hospital of Philadelphia, USA

It appears that the majority of the PNH clone (in both granulocytes and red cells) is type II, i.e., only partially deficient of GPI. One assumes that this picture was obtained after the haemolytic attack. If so, it could be that type III, i.e., severely deficient RBC were indeed G6PD-deficient. Was flow-cytometry performed after Eculizumab treatment? Eculizumab would protect type III RBC from lysis and thus would allow re-assessment of G6PD activity.

We agree with this comment. Of course it is possible that at the time of the observation of association of co-trimoxazole associated hemolysis a different PNH clone was more prevalent than when analyzed for G6PD deficiency. We included this possibility in our discussion. The most recent flow cytometry was performed when the patient was on eculizimab.

Minor points

- Was anti-CD55 or –CD59 was used for flow analysis? The authors state anti-CD55 in methods but describe anti-CD59 in results. The diagnosis for PNH was performed in a CLIA approved clinical laboratory of the hospital.

  We present the results for CD59. CD55 was also tested however the discrimination between the three populations is more difficult.

- Normal ranges of lab tests need to be provided.

  We have included normal ranges for laboratory tests in parentheses after the patient's values.

- Another factor was that treatment with eculizumab, by inhibiting lysis of PNH red cells may have led to a higher level of PNH (and concomitantly G6PD deficient) red cells than would be present in untreated patients.’ This statement is rather irrelevant because eculizumab was started after treatment with co-trimoxazole.

  We think this statement is valid because treatment with eculizimab increases the level of PIGA deficient red cells. On our hypothesis that the G6PDA- allele was linked with PIGA- then PIGA-,G6PDA- red cells would increase. Our hypothesis however was not correct.

- Need to use Greek characters appropriately, i.e., ul should be ml
Rosario Notaro
Laboratory of Genetics and Gene Transfer, Core Research Laboratory, Istituto Toscano Tumori, Florence, Italy

Perdigones and collaborators report an interesting clinical case about the association of PNH and G6PD deficiency. However, the timing of the clinical history is not clear and the report is scanty of some relevant clinical/laboratory data.

**Major scientific points.**

1. The Authors should provide the exact timing of entire clinical history: diagnosis; infective episode resulting in the prescription of *trimethoprim-sulfamethoxazole* -*co-trimoxazole*- (which dose?); start and stop of *co-trimoxazole*; “dark urine episode” associated with *co-trimoxazole*; start of *eculizumab*; time of biological studies.

2. It is extremely important that the Authors provide more details about the timing and the features of the hemolytic attack apparently associated with *co-trimoxazole*: how long after the start of *co-trimoxazole* treatment the patient experienced the “dark urine episode”? There are any objective data at the time of this “dark urine episode” or the episode has been just self-reported? At the time of this “dark urine episode” there were any signs/symptoms of an ongoing infective condition?

3. The Authors provide clinical/laboratory details only at one time point, that seems be after the start of eculizumab (how long after?). They should provide such clinical/laboratory details at time of diagnosis, at the time of the co-trimoxazole associated “hemolytic attack”, at start of eculizumab treatment, etc.: blood count, absolute reticulocyte count, LDH levels (providing the normal range), flow citometry, etc.

4. In PNH patients the presence of red blood cells with partial deficiency of GPI-linked molecules is relatively common. However, the presence of granulocyte/monocyte with partial deficiency of GPI-linked molecules is uncommon: thus, it would be interesting to show the dot plot of the “CD59/lineage marker” analysis of granulocytes and monocytes of
this patient.

5. The Authors, at variance with their starting hypothesis, have clearly proven that in this patient PNH cells express the wild type G6PD B. They provide 2 possible explanations for the co-trimoxazole-associated “hemolytic crisis” observed in this patient. These hypotheses are interesting but their probability is very low. I suggest that the Authors should discuss a much more likely explanation: this “hemolytic crisis” was just due to the infective condition that led to the prescription of co-trimoxazole.

Minor points

1. In the Introduction the Authors report the classical classification of PNH cells as Type III (completely deficient in GPI anchored proteins), Type II (partially deficient) and Type I (normal display of GPI-linked proteins). However, in the Results they write about “class I clone (partial deficiency) ... and class II clone (complete deficiency) “: this is extremely confusing.

2. In the Results the 2 sentences (from “We hypothesized that ...” to “...red cells than would be present in untreated patients.”) seem to suggest that the relative expansion of the “PNH (and concomitantly G6PD deficient) red cells” following eculizumab treatment could have played a role in the co-trimoxazole associated hemolytic attack. The Authors should rephrase these sentences since eculizumab has been started after the co-trimoxazole associated hemolytic attack.

3. Trimethoprim-sulfamethoxazole (co-trimoxazole) is a generic drug name thus it should not be capitalized. In the Results, “eculizumab” should not be capitalized.

4. The brand name Bactrim should be replaced with the generic drug name.

5. The levels of ROS in the patient red cells should be compared with a group of healthy controls: the comparison with only one control does not allow any conclusion.

6. The Authors should report the concentration of DCF used for ROS detection.

**Competing Interests:** No competing interests were disclosed.

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard, however I have significant reservations, as outlined above.
some relevant clinical/laboratory data.

**Major scientific points.**

1. The Authors should provide the exact timing of entire clinical history: diagnosis; infective episode resulting in the prescription of trimethoprim-sulfamethoxazole -co-trimoxazole- (which dose?); start and stop of co-trimoxazole; “dark urine episode” associated with co-trimoxazole; start of eculizumab; time of biological studies.

In our revised manuscript we include a fuller, more detailed case history. Of note is the fact that the patient presented to the specialty clinic with a probably 6 year history of PNH and that most of the patients past history relies on records and notes. Here we demonstrate that the patient has indeed both conditions PNH and G6PD deficiency. With these results in hand we try to explain the patient’s history and clinical observations. It was not our intention to prove that indeed the hemolysis described by the emergency physician after co-trimoxazole was indeed triggered by the drug, but rather to evaluate and discuss the possibility as the patient carries on her chart the diagnosis of co-trimoxazole hypersensitivity due to this incident. We try to make this clearer in the case description. We hope this will answer the criticism of the reviewer.

2. It is extremely important that the Authors provide more details about the timing and the features of the hemolytic attack apparently associated with co-trimoxazole: how long after the start of co-trimoxazole treatment the patient experienced the “dark urine episode”? There are any objective data at the time of this “dark urine episode” or the episode has been just self-reported? At the time of this “dark urine episode” there were any signs/symptoms of an ongoing infective condition?

The discussion above and the revised version provides many more details.

3. The Authors provide clinical/laboratory details only at one time point, that seems be after the start of eculizumab (how long after?). They should provide such clinical/laboratory details at time of diagnosis, at the time of the co-trimoxazole associated “hemolytic attack”, at start of eculizumab treatment, etc.: blood count, absolute reticulocyte count, LDH levels (providing the normal range), flow citometry, etc.

We have provided all the history we can obtain in the revised version.

4. In PNH patients the presence of red blood cells with partial deficiency of GPI-linked molecules is relatively common. However, the presence of granulocyte/monocyte with partial deficiency of GPI-linked molecules is uncommon: thus, it would be interesting to show the dot plot of the “CD59/lineage marker” analysis of granulocytes and monocytes of this patient.

The diagnosis of PNH and the subtype analysis of PNH granulocytes was performed in a CLIA approved clinical laboratory, and their test results are reported here. The
individual dot blots are not available to us. However the senior author who was involved in the setup and quality assessment of flow-cytometric PNH testing at this institution fully trusts their analysis. We agree with the reviewer that partial GPI-anchor deficiency in granulocytes is less frequently observed and very much depends on the underlying mutation and the antibody chosen for the analysis.

5. The Authors, at variance with their starting hypothesis, have clearly proven that in this patient PNH cells express the wild type G6PD B. They provide 2 possible explanations for the co-trimoxazole-associated “hemolytic crisis” observed in this patient. These hypotheses are interesting but their probability is very low. I suggest that the Authors should discuss a much more likely explanation: this “hemolytic crisis” was just due to the infective condition that led to the prescription of co-trimoxazole.

We agree that this is a possible alternative explanation and have added this in the discussion however considering the rather bland urine sediment we actually favor the other two possible explanations.

Minor points

1. In the Introduction the Authors report the classical classification of PNH cells as Type III (completely deficient in GPI anchored proteins), Type II (partially deficient) and Type I (normal display of GPI-linked proteins). However, in the Results they write about “class I clone (partial deficiency) ... and class II clone (complete deficiency)” – this is extremely confusing.

This has been corrected

2. In the Results the 2 sentences (from “We hypothesized that ...” to “...red cells than would be present in untreated patients.”) seem to suggest that the relative expansion of the “PNH (and concomitantly G6PD deficient) red cells” following eculizumab treatment could have played a role in the co-trimoxazole associated hemolytic attack. The Authors should rephrase these sentences since eculizumab has been started after the co-trimoxazole associated hemolytic attack.

The timing of the eculizimab treatment and the hemolysis are given in the detailed case report that we now provide.

3. Trimethoprim-sulfamethoxazole (co-trimoxazole) is a generic drug name thus it should not be capitalized. In the Results, “eculizumab” should not be capitalized.

We have corrected this.

4. The brand name Bactrim should be replaced with the generic drug name.

We have corrected this.

5. The levels of ROS in the patient red cells should be compared with a group of healthy
controls: the comparison with only one control does not allow any conclusion.

The experiment was carried out with several controls and it confirms the results of Amer et al., cited as our ref 14 that ROS are elevated in PNH. Our point is a small one, that we might expect a further elevation if we compare PNH and normal cells from this patient, if indeed PIGA- is linked with G6PDA-. We didn't find this. We include the sentence “The figure shows a representative example of 4 normal controls that gave similar results” to clarify this.

6. The Authors should report the concentration of DCF used for ROS detection.

We have done this.

**Competing Interests:** No competing interests were disclosed.