

## Case Report

# Concomitant a novel *ALAS2* mutation and *GATA1* mutation in a newborn: a case report and review of the literature

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**Abstract:** *GATA-1*, an X-linked gene, encodes a transcription factor that plays a role in erythropoiesis and megakaryopoiesis. *GATA-1* mutations have been associated with various diseases, such as X-linked thrombocytopenia. *ALAS2* is an X-linked erythroid-specific isoenzyme expressed during erythropoiesis. Mutations of *ALAS2* were associated with X-linked sideroblastic anemia. We report a case of newborn twin boy with anemia and thrombocytopenia at birth. A bone marrow biopsy at 4 months of age showed marked dyserythropoiesis, dysmegakaryopoiesis, and rare ringed sideroblasts. Gene sequencing study showed a previously reported mutation in *GATA-1* at c.622G>A location (G208R) and a novel *ALAS2* mutation at c.1436G>A location (R479Q).

**Keywords:** *GATA1* mutation, *ALAS2* mutation, macrothrombocytopenia, dysmegakaryopoiesis, dyserythropoiesis, ringed sideroblasts

### Introduction

*GATA1* is an X-linked gene that encodes a transcription factor that regulates the expression of a very large number of genes in many cell types. It contains a DNA binding domain and a trans-activator domain. The N-finger in the DNA binding domain plays an important role in *GATA1*'s activity, both by binding DNA to increase the stability, and by recruiting cofactors, such as Friend of *GATA1* (FOG-1), which is important for erythropoiesis and megakaryopoiesis [1-3]. Cases associated with transient myeloproliferative disease and acute megakaryocytic leukemia in children with Down syndrome have somatic mutations in exon 2 of the *GATA-1* gene, resulting in the short isoform, *GATA-1s* [4]. Germline missense mutations in exon 4 of *GATA-1* that lead to alterations in various amino acids of the N-terminal zinc-finger domain have been linked to various diseases, such as X-linked thrombocytopenia and anemia and congenital erythropoietic porphyria [5-9].

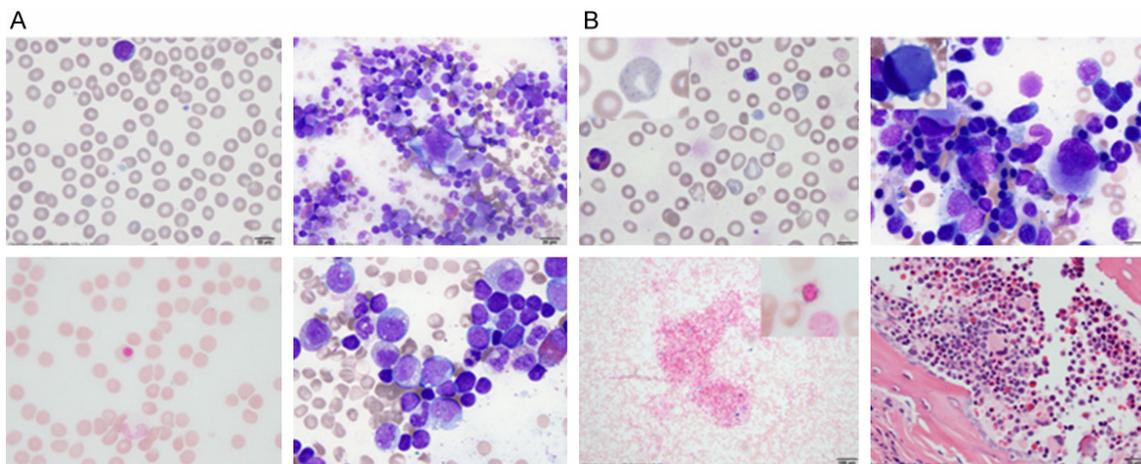
*ALAS*, 5-aminolevulinatase synthase, catalyzes the formation of 5-aminolevulinatase from succinyl CoA and glycine. It is a rate-limiting mitochondrial enzyme. The *ALAS2* is an erythroid-specific isoenzyme that is expressed during erythropoiesis [10]. The *ALAS2* mutations known are heterogeneous and clustered in the exons that encode the catalytic domain.

A case of a newborn male with anemia and thrombocytopenia with concomitant *GATA1* and a novel *ALAS2* mutation is reported.

### Case presentation

The patient was a male born at 36 weeks gestation as the product of a dichorionic diamniotic twin pregnancy resulting from non-consanguineous parents. At birth, CBC showed anemia and thrombocytopenia with hemoglobin of 10.2 g/dL, reticulocyte count of 7.9%, and platelet count of 22,000/ $\mu$ L. Physical examination was significant for hepatosplenomegaly and "blueberry muffin" rashes. There were no dysmor-

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**Figure 1.** Comparison of peripheral blood and bone marrow morphology at 2 months (A) and 4 months of age (B). In panel A: Top left: Peripheral blood showed normocytic RBCs without anisopoikilocytosis. A very rare agranular large platelet is seen (MPV 10.3). Top right: Bone marrow aspirate showed trilineage hematopoiesis with marked erythroid hypoplasia. The megakaryocytes appeared unremarkable. Bottom left: Iron stain on the bone marrow aspirate showed a rare ringed sideroblast. Bottom right: Occasional early stage myeloid cells contained small cytoplasmic vacuoles. In panel B: Top left: Peripheral blood showed normocytic anemia with marked anisopoikilocytosis. The insert showed the presence of one of the many RBCs with coarse basophilic stippling. There were many giant agranular platelets (MPV 12.9). Top right: The bone marrow aspirate showed moderate dyserythropoiesis with increased monolobated/hypolobated megakaryocytes with eccentrically located nucleus (insert). Bottom left: The bone marrow showed increased iron storage for age with rare ringed sideroblasts (insert). Bottom right: The bone marrow core biopsy showed the presence of the small megakaryocytes with crescent shaped eccentrically located nuclei.

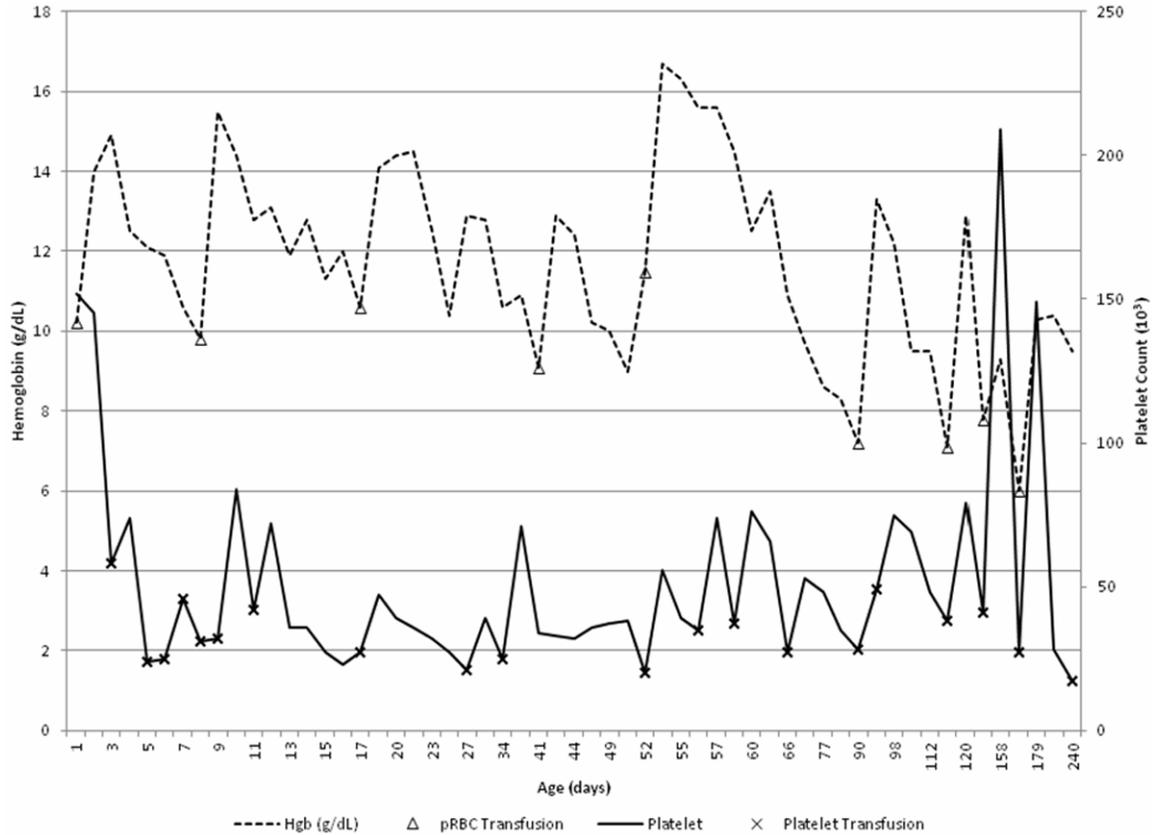
phic features or skeletal abnormalities. Work-up for infections and genetic diseases was negative. Review of the placenta showed immature placenta without acute or chronic villitis and viral inclusions. There were many nucleated RBCs within the villi. The co-twin placenta was mature and unremarkable.

The patient required several packed red cell and platelet transfusions within the first month of life. The first bone marrow aspirate was done at 2 months of age and showed pauci-spicular bone marrow with marked erythroid hypoplasia and mild dyserythropoiesis. The megakaryopoiesis was unremarkable without obvious dysmegakaryopoiesis. Few early myeloid precursors contained cytoplasmic vacuoles. An iron stain showed rare ringed sideroblasts (**Figure 1A**). At the time, sideroblastic anemia, especially Pearson syndrome, was considered but ruled out when mitochondrial DNA deletion was not detected. A repeat bone marrow biopsy was performed at 4 months of age due to persistent anemia and thrombocytopenia requiring multiple packed red cell and platelet transfusions (**Figure 2**). CBC at this time showed anemia and macrothrombocytopenia (**Table 1**). The RBCs

on the peripheral smear showed marked anisopoikilocytosis with coarse basophilic stippling and many large agranular platelets (**Figure 1B**). The bone marrow aspirate showed marked dyserythropoiesis. The megakaryocytes were dysplastic with small, eccentric crescent shaped nuclei and clear cytoplasm. An X-linked *GATA1* gene sequencing study was performed (Prevention Genetics, Marshfield, WI) and showed a missense mutation at c.622 G>A location resulting in a glycine to arginine amino acid change (G208R). In addition, a comprehensive mitochondrial nuclear gene panel was performed (GeneDX, Gaithersburg, MD) and showed a novel *ALAS2* mutation at c.1438 G>A resulting in an arginine to glutamine amino acid change (R479Q). This variant has not been published as a mutation, nor has it been reported as a benign polymorphism.

At the age of one, the patient continues to have persistent thrombocytopenia with platelet counts averaging 30,000 and mild anemia (**Figure 2**). He continues to have significant bleeding episodes requiring regular platelet transfusions. Of note, this patient's twin lacks the *GATA-1* mutation as well as any anemia or

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**Figure 2.** Hemoglobin, platelet count and transfusion requirements over time. The patient's platelet count (solid line) and hemoglobin concentration (dashed line) over time until the age of 240 days (8 months) are shown. Transfusion of packed red cells ( $\Delta$ ) and platelets (x) over time are also displayed on these curves.

**Table 1.** CBC and bone marrow findings at initial and subsequent bone marrow biopsies

		2 month old (normal range)	4 month old (normal range)
CBC	WBC ( $\times 10^3/\mu\text{l}$ )	16.31 (5.0-19.5)	23.46 (5.0-19.5)
	Hgb (g/dl)	10.9 (9.4-13.0)	7.1 (9.4-13.0)
	MCV (fl)	91.7 (84-106)	98.5 (84-106)
	Platelet ( $\times 10^3/\mu\text{l}$ )	27 (150-400)	38 (150-400)
	MPV (fl)	10.3 (7.0-11.3)	12.9 (7.0-11.3)
Bone marrow	Erythropoiesis	Decreased; Mild dysplasia	Predominant; Marked Dysplasia
	Granulopoiesis	Normal	Normal
	Megakaryopoiesis	Mostly normal	Dysplasia

thrombocytopenia. The twin's *ALAS2* mutation status is unknown at this time.

### Discussion

*GATA1*, an X-linked gene, is the founding member of the GATA transcription factor family. It is primarily expressed in hematopoietic tissues and plays an important role in erythropoiesis and megakaryopoiesis [12-14]. There seems to

be a dose-related effect: erythroid precursors can mature at low levels of *GATA1* whereas megakaryocytes need higher levels. In murine studies, deletion of *GATA1* has shown a compensatory increase in *GATA2*, which may explain why these proerythroblasts express some genes at wild type levels such as beta globin H1 and alpha globin [3]. *GATA1* deficient megakaryocytes are more numerous, smaller, have larger and segmented nuclei, have a paucity of

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alpha granules, do not express GPIb and stain weaker with acetylcholinesterase [3, 15]. In essence, these megakaryocytes are developmentally arrested and proliferate excessively [16].

Our case demonstrated the presence of previously described missense mutation that led to G208R substitution within the N-finger of GATA-1 [5]. It was thought that this mutation reduces the hydrophilic interaction by substitution of a positively charged amino acid, which caused more profound destabilization of the GATA1 and FOG-1 interaction, than other amino acid substitutions at the same location. Thus it was hypothesized that a patient carrying this mutation would have a more severe clinical presentation. This mutation was initially identified in a 17 year old boy, the third child born from the family [5]. Subsequently, there have been two reported cases of the same mutation, one who on bone marrow analysis at age 12 had 5% ringed sideroblasts, but without ringed sideroblasts in the other [17, 18]. Some patients also carried a diagnosis of idiopathic thrombocytopenia purpura prior and some had splenectomy.

The finding of co-existing previously unknown hemizygous R479Q mutation in the *ALAS2* gene is interesting. Mutations in the *ALAS2* gene are linked to X-linked hereditary sideroblastic anemia [19]. The patient with known mutations mainly presented with microcytic anemia, iron overload and ringed sideroblasts. However, the R479Q mutation identified in our patient is a novel variant at a conserved region. Although in-silico analysis indicates that this mutation is unlikely to be damaging to the *ALAS2* protein (per GeneDx report), a functional study might be helpful in further understanding this mutation. Continued clinical follow up might be helpful as some patients might present in late childhood or even in late adulthood [20]. Clinically, our patient has normocytic anemia initially, and later macrocytic anemia without evidence of iron overload.

The relationship of GATA1 mutation and *ALAS2* mutation in our patient is unknown. Studies have shown that GATA1 binds to *ALAS2* at a GATA consensus sequence in intron 8 of the gene [21], presumably to regulate heme synthesis. The mutation in this patient is located in the coding region in *ALAS2* gene, not the GATA1 binding sequence. While the patient's mutation

in *ALAS2* alone may not affect protein function, it remains to be determined what kind of interaction there may be between the dysfunctional GATA1 protein and the *ALAS2* protein. It is possible that the hemizygous GATA1 mutation plays an insignificant role in the *ALAS2* regulated heme synthesis, as long as the *ALAS2* mutation does not affect the protein function.

Another interesting aspect of this case is that the initial bone marrow biopsy at 2 months of age did not reveal typical morphologic features that are suggestive of a typical GATA1 mutation. The reasons for this are not clear. Transfusions and the underlying medical conditions in the perinatal period might mask the peripheral blood and bone marrow findings. In addition, the role of GATA1 and *ALAS2* in fetal and postnatal hematopoiesis is not well understood. This case presents the challenge in diagnosing certain genetic disorders in hematopoiesis in a newborn.

### Disclosure of conflict of interest

The authors have no conflicts of interest to disclose.

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