A point mutation c.473A > G of ITGB3 is responsible for the formation of the Wo^a human platelet alloantigen

Sarah T. Holzwarth,¹ Julian Strobel,² Nina Cooper,^{1,3} Jörg Leyh,⁴ Behnaz Bayat,¹ Gregor Bein,^{1,3} Jürgen Zingsem,² and Ulrich J. Sachs ^{1,3}

CASE REPORT

36-year-old female (Wo) delivered a full-term girl in the 40th week of gestation with no signs of bleeding. Her neonatal platelet count was 29 G/L (dropping to 20 G/L), and fetal-neonatal alloimmune thrombocytopenia (FNAIT) was suspected. In the serological work-up, no anti-platelet antibodies were detectable using PAK Lx (Immucor Medizinische Diagnostik). Genetic testing did not reveal incompatibility for HPA-1, -2, -3, -4, -5, -6, -9 and -15. Two years later, Mrs. Wo gave birth to a white European full-term boy in the 39th week of gestation, again with no signs of bleeding. The neonatal platelet count was 31 G/L. Platelet antibody testing in the monoclonal antibody of platelet specific antigens (MAIPA) assay using random donor platelets was negative. In contrast, a MAIPA cross-match analysis between maternal serum and paternal platelets showed positive reactions with aIIbb3, indicating an alloimmunization against a lowfrequency antigen. On Day 1, the newborn was transfused with random donor platelets. The platelet count rose to 170 G/L, remained above 50 G/L without further transfusions, and was in normal range at Day 28.

METHODS

Full-length sequencing of *ITGA2B* and *ITGB3* was carried out as described previously.¹ Briefly, coding regions of paternal genomic DNA were PCR amplified with primers corresponding to intronic sequence surrounding all exons. PCR was carried out using a Fast-Start High Fidelity PCR system (Roche Diagnostic Corp.). Automated sequence analysis was performed in both directions on a genetic analyzer (ABI 3100, Applied Biosystems). Nucleotide sequences of PCR primers, sequencing, and reaction conditions are available upon request.

Full-length β 3 cDNA encoding for the Wo^a variant was produced by site-directed mutagenesis using the Quick Change Mutagenesis Kit (Stratagene) as previously described.² Allele specific constructs encoding wild-type β 3 (Gln₁₃₂) or mutant β 3 (Arg₁₃₂) were transfected into HEK 293F cells, together with an α IIb construct. Surface expression of α IIb β 3 was measured by flow cytometry (FACS Canto II, Becton Dickinson), as outlined previously.² To confirm that p.Gln132Arg is responsible for the formation of the Wo^a epitope, 3×10^5 transfected cells per well were used in a modified MAIPA.³

RESULTS

Sequencing *ITGB3* from paternal DNA showed the presence of a nucleotide substitution c.473A > G located in exon 4 of ITGB3 (Fig. 1A). This mutation predicts Gln (CAG) at position 132 in Wo^a-negative and Arg (CGG) in Wo^a-positive individuals (p.Gln132Arg). The mutation was deposited in GenBank (accession number MN624129). No sequence variation was detected in *ITGA2B*. The *ITGB3* c.473A > G mutation was also detectable in heterozygous state when sequencing DNA from the newborn. In a glycoprotein specific assay (Fig. 1B), HEK 293F cells expressing α IIb in complex with either wildtype (Gln₁₃₂) or mutant (Arg₁₃₂) β 3 were incubated with maternal serum, healthy control donor serum, or serum

From the ¹Institute for Clinical Immunology and Transfusion MedicineJustus Liebig University; the ³German Center for Feto-Maternal Incompatibility (DZFI), University Hospital Giessen and Marburg, Giessen; ²Department of Transfusion Medicine and Haemostaseology and ⁴Children's Hospital, Friedrich-Alexander-University, Erlangen, Germany.

Address reprint requests to: Ulrich J. Sachs, MD, Institute for Clinical Immunology and Transfusion Medicine, Justus Liebig University, Langhansstr. 7, 35392 Giessen, Germany; e-mail: ulrich. sachs@med.uni-giessen.de

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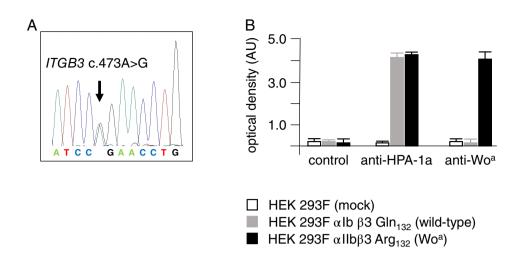


Fig. 1. (A) Nucleotide substitution c.473A > G located in exon 4 of ITGB3 as detected in paternal DNA (GenBank accession number MN624129, SNP database rs_748901429). (B) Serological confirmation of the Wo^a epitope. HEK 293F cells were used after mock transfection (white), transfection with wild-type α IIb β 3 (gray) and transfection with α IIb β 3 after introduction of the c.473A > G mutation in ITGB3 (black) in a modified MAIPA assay. Transfected cells were incubated with control serum from a healthy blood donor, with serum containing anti-HPA-1a antibodies, or with the maternal serum containing anti-Wo^a, as indicated. After cell lysis, α IIb β 3 was immobilized to the microtiter plate using complex-specific monoclonal antibody Gi5. Presence of human IgG in the immobilized complex was detected using horseradish peroxidase-labeled goat-anti-human-IgG and substrate buffer. Optical density (OD) was read at 492 nm. Wild-type HEK293F α IIb β 3 Gln₁₃₂ (gray) and mutant HEK293F α IIb β 3 Arg₁₃₂ (black) were reactive with anti-HPA-1a (OD 4.19 ± 0.19 and 4.32 ± 0.16, respectively). In contrast, the wild-type form (gray) did not react with anti-Wo^a (OD, 0.14 ± 0,19), whereas the mutant form (black) gave strong reactions (OD, 4.11 ± 0.47). All data given as mean ± standard deviation from n = 3 independent experiments.

containing anti-HPA-1a antibodies, demonstrating the specific recognition of p.Gln132Arg by anti-Wo^a.

ALLELE FREQUENCY

Summary data from the Exome Aggregation Consortium indicate allele frequencies of 0.99999 for ITGB3 c.473A and 0.00001 for ITGB3 c.473G. 4

SUMMARY

We observed a case of FNAIT in a newborn with a platelet count of 31 G/L. Serological analysis of the maternal serum revealed an immunization against α IIb β 3 on paternal platelets only, indicating the presence of an antibody against a rare alloantigen. Sequencing analysis and studies with mutant transfected cells showed that ITGB3 c.473A > G (p. Gln132Arg) is responsible for the formation of a new antigenic determinant on β 3, termed Wo^a. The antibody was overlooked in the initial work-up of the elder sibling where no cross-match study was performed. This demonstrates once more that a serological cross-match between paternal platelets and maternal serum should always be performed whenever FNAIT is suspected.

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REFERENCES

- Peterson JA, Gitter ML, Kanack A, et al. New low-frequency platelet glycoprotein polymorphisms associated with neonatal alloimmune thrombocytopenia. Transfusion 2010;50:324-33.
- Wihadmadyatami H, Heidinger K, Röder L, et al. Alloantibody against new platelet alloantigen (Lap(a)) on glycoprotein IIb is responsible for a case of fetal and neonatal alloimmune thrombocytopenia. Transfusion 2015;55:2920-9.
- Kiefel V, Santoso S, Weisheit M, et al. Monoclonal antibody-specific immobilization of platelet antigens (MAIPA): a new tool for the identification of platelet-reactive antibodies. Blood 1987;70:1722-6.
- 4. Reference SNP (rs) report for rs748901429. [cited September 26, 2019]. https://www.ncbi.nlm.nih.gov/snp/rs748901429#frequency_tab).