EXCEPTIONAL CASE REPORT

Thrombotic thrombocytopenic purpura and defective apoptosis due to CASP8/10 mutations: the role of mycophenolate mofetil

F. Fioredda,1 E. Cappelli,1 A. Mariani,1 A. Beccaria,1 E. Palmisani,1 A. Grossi,2 I. Ceccherini,2 R. Vené,3 C. Micalizzi,1 M. Calvillo,1 F. Pierr,1 I. Mancini,4 F. Peyvandi,5,6 F. Corsolini,7 C. Dufour,1 and M. Miano1

1Unit of Haematology and 2Medical Genetics Unit, IRCCS Istituto Giannina Gaslini, Genoa, Italy; 3Molecular Oncology and Angiogenesis Unit, IRCCS AOU San Martino–IST, Genoa, Italy; 4Fondazione IRCCS Ca’ Granda Ospedale Maggiore Policlinico, Angelo Bianchi Bonomi Hemophilia and Thrombosis Center, Università degli Studi di Milano, Department of Pathophysiology and Transplantation and Fondazione Luigi Villa, Milan, Italy; 5Department of Pathophysiology and Transplantation, Università degli Studi di Milano and Fondazione Luigi Villa, Milan, Italy; 6Fondazione IRCCS Ca’ Granda Ospedale Maggiore Policlinico, Angelo Bianchi Bonomi Hemophilia and Thrombosis Center, Milan, Italy; and 7Laboratory of Molecular Genetics and Biobanks, IRCCS Istituto Giannina Gaslini, Genoa, Italy

Key Points

• Immunological dysregulation may underlie unusual autoimmune diseases, which also deserve to be investigated from a genetic point of view.

Introduction

Thrombotic thrombocytopenic purpura (TTP) is a rare, life-threatening disorder characterized by microvascular platelet clumping and microangiopathic hemolytic anemia, often accompanied by renal dysfunction and neurological symptoms.1 The underlying pathophysiological mechanism is due to a defective ADAMTS13 protein, either congenital or secondary to the anti-ADAMTS antibodies that switch off its function. Acquired TTP occurs mainly in women, rarely in children, and has also been described in association with known autoimmune disorders (ie, systemic lupus erythematosus).1,2

The gold standard therapy for TTP is plasma exchange together with high-dose steroids or immunoglobulins. Drugs such as rituximab are indicated in cases of failure/relapse. Vincristine, cyclophosphamide, azathioprine, and cyclosporine have also been used with a certain efficacy3-5 and splenectomy has been performed with a positive outcome in patients with refractory or relapsing disease.6 To the best of our knowledge, correlation between acquired TTP and immune dysregulation due to a defect in components of the apoptosis pathway have never previously been described.

Case description

An Ecuador-born female patient came to us for the first time for observation at 18 years of age due to easy bruising and hematuria. Her platelet count was 4 3 10^9/L, and acute non-autoimmune hemolytic anemia was indicated (hemoglobin, 6 g/dL; lactate dehydrogenase, 2964 IU/L; reticulocyte count, 200 3 10^9/L; haptoglobin, <2 mg/dL; schistocytes on the blood smear; and negative results on direct and indirect antiglobulin tests). The patient carried antithyroglobulin antibodies; in addition, antinuclear antibodies, antibodies against double-stranded DNA, extractable nuclear antigens, antiphospholipid antibodies, and complement components (C3 and C4) were negative. Previously in Ecuador, the patient had had 2 similar episodes at the age of 10 and 15 years, successfully treated with transfusions and immunoglobulins. A long remission occurred between the 2 events without any therapy.

The differential diagnosis initially considered between relapsing Evans syndrome treated with high-dose steroids and immunoglobulins and atypical hemolytic uremic syndrome/ TTP. The patient was empirically treated with plasma infusions, with substantial improvement in clinical status.

Following steroid suspension, the patient’s platelet count dropped again below 10 3 10^9/L, and hemolysis relapsed. This oriented treatment toward a plasma exchange procedure. Unfortunately, the patient could not tolerate the procedure because of a severe reaction characterized by malaise, hypotension, and hives. Rescue therapy with a second cycle with immunoglobulin was administered, and mycophenolate mofetil (MMF) at a dose of 600 mg/mq twice a day was started as “maintenance” treatment.

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For original data, please contact the corresponding author, Francesca Fioredda (francescafioredda@gaslini.org).

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The choice of MMF was based on our recent encouraging results in refractory cytophenias.7 Given the stability of the hematological values, after 2 weeks of MMF, the therapeutic strategy was not shifted to rituximab. We preferred MMF to rituximab, even after confirmation of the TTP diagnosis, because the good initial response reported in autoimmune cytophenias is outbalanced by rarely maintained long-term remission and by the risk of iatrogenic, deep and prolonged hypogammaglobulinemia.8,9 Indeed, rituximab therapy, mainly in the pre-“low-dose” era, had a stronger impact on B cells and immunoglobulin G concentrations, which remain below the threshold for many months (requiring, in some cases, immunoglobulin replacement therapy).10

**Methods and results**

Despite normal platelet values and resolution of hemolysis, low levels of ADAMTS13 (<3% activity as measured by using the FRETs-VWF73 assay; normal range, 45%-138%) and positivity for antibodies against ADAMTS13 (62 U/mL as measured by using the Technozym ADAMTS-13 Inhibitor assay [Technoclone GmbH]) were shown.11 The patient continued MMF, and remission lasted >12 months. After that time span, another relapse occurred that was attributed to poor MMF compliance. A new plasma exchange cycle (plus steroids) was performed, and this time it was well tolerated. After 2 months, stable remission was again achieved; the patient has been motivated to adequately comply with the MMF regimen. To date, 19 months later, the patient is again in remission (steroid free), and the trough MMF plasma levels are in the therapeutic range (1-4 mg/mL).12

ADAMTS13 activity remained below the threshold, and antibodies against ADAMTS13 persisted as positive both in the “active” periods and during the remission phases of the disease. An expanded immunological evaluation was normal (even the CD3⁺ T-cell receptor αβ⁺ CD4⁺ CD8⁺ double-negative T cells), lymphoproliferation was absent, and the in vitro FAS-mediated apoptosis test result was normal. Molecular studies were performed by using an enlarged next-generation sequencing panel, which surprisingly showed a homozygous variant in CASP8 and a heterozygous variant in CASP10. The caspase-8 gene showed a homozygous mutation, c.2T>C (RefSeq NM_001080125.1), which led to a p.Met1Thr change, affecting the start codon. This variant, although rare in different populations, is relatively common in South America, with an allelic frequency of 12% and with ~1% to 2% of the population being homozygote-CC (ExAC Browser, http://exac.broadinstitute.org/).

In fact, western blotting of the extract of lymphoblast cells obtained from the patient showed normal expression of procaspase-8 (Figure 1A), probably a “different” protein with uncertain functionality. (Informed consent had been obtained from the patient, and all experiments were performed in accordance with the approved guidelines.) Treatment with FAS-L aimed to induce apoptosis generated a cleavage activation of caspase-8 only in the wild-type cell line and not in the patient’s lymphoblasts, suggesting that the mutation altered protein function and maintained an impaired apoptotic pathway. The heterozygous CASP10 variant (NM_032977.3: c.1564T>A) leading to a Leu522Ile missense change did not substantially alter the expression of the protein (Figure 1A-B). This variant is generally classified as a high-frequency polymorphism (EXAC DATABASE, http://exac.broadinstitute.org/) even if it was reported with a probably damaging role in most prediction tools; it has also been shown to be associated with an impaired apoptotic capacity.13,14

The additional weak cleavage inactivation of the PARP protein, along with FAS-L treatment that did not induce any apoptotic death (Figure 1A), suggests an overall reduction in apoptotic activity in the patient’s lymphoid cells.

The same experimental setting adopted for FAS-L was repeated with TRAIL, the other apoptosis inducer, thus confirming the impairment of the CASP pathway in the patient’s lymphoid cells (Figure 1B).15

**Discussion**

The current report contains 2 main peculiarities: the association between acquired TTP and mutations in the CASP8/10 genes, usually responsible for different disorders, and its management with a “nonconventional” immunosuppressive drug. Caspases are cysteine proteases that play a central role in apoptosis. Caspase-8 and caspase-10 may be the first enzymes in the
proteolytic cascade that are activated by FAS ligand and tumor necrosis factor. The absence of an immunomodulatory function of CASP8, enhanced by its lack of negative regulation by CASP10, interrupts the cleavage of CASP3, increases PARP pathway signaling, and causes hyperactivation of Akt–mammalian target of rapamycin. According to the literature, mutations in either CASP8 or CASP10 (being implicated in programming life and death of immune cells) influence immune surveillance of malignancies—inducing lymphoproliferation (autoimmune lymphoproliferative syndrome type IIA and B). In addition, the interplay between caspase-8 and caspase-10 proteins, which share substrates and functions, is crucial for the maintenance of regular apoptosis. The hypoplasia, recently addressed in an article from our group, is that the association of 2 variants on genes involved in the same pathway, even if polymorphic, might contribute to the development of immune dysregulation. Indeed, we speculate that the mutations found in our patient do not affect lymphocyte accumulation but generate a negative synergy expressed as autoimmunity against the ADAMTS13 protein. Even if direct causality between the hyperactivation of Akt–mammalian target of rapamycin and autoimmune TTP cannot be proved, an immune dysregulation was at the basis of the phenomenon affecting the patient. It is worth noting that our patient was also studied for 315 adjunctive genes involved in hematological disorders, immune deficiencies, and immune dysregulations, with negative results. Nevertheless, we cannot exclude that other genetic lesions in genes not covered by our panel might have contributed to the clinical phenotype of the study patient, but the knowledge of the impaired apoptosis pathway represents a suggestion for a tailored treatment.

Regarding the response to treatment, MMF, on account of the inhibition of B, T, and natural killer cells, has been shown to be effective in autoimmune cytopenia and in a number of other autoimmune diseases. In this patient, MMF seemed to have had a certain efficacy in the maintenance of autoimmunity suppression against ADAMTS13 with no undesirable effects. Moreover, according to a recent article, persistent low ADAMTS13 activity, as observed in our patient, seems to be a predictor of relapse

and thus indicates a possible role for a preemptive immunosuppression regimen.

The very last “lesson” of this experience is that patients presenting with refractory autoimmune cytopenia and/or unusual autoimmune targets should be studied by using an extended genetic panel directed to identify “disimmunity.” The aim is to better understand the mechanism and identify possible targeted drugs.

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