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Lessons in gene hunting: a RAG1 mutation presenting with agammaglobulinemia and absence of B cells

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To the Editor

Recombination activating genes (*RAG*) 1 and 2 are critical for V(D)J recombination and lymphocyte development.¹ Mutations in *RAG1/2* associated with <1% of wild-type recombination activity typically result in severe combined immunodeficiency (SCID) lacking T and B cells.² Hypomorphic mutations with residual RAG activity can result in Omenn syndrome, characterized by erythroderma, lymphadenopathy, and the presence of oligoclonal T cells. More recently, studies have broadened the phenotypic variability associated with *RAG1/2* mutations, which now includes granulomatous disease, $\gamma\delta$ T-cell expansion, CD4⁺ lymphopenia with intact T cell function, and hyper-IgM syndrome.^{1, 3} We report a patient with a homozygous missense mutation in *RAG1*, who presented initially with agammaglobulinemia, absent B cells, and normal numbers of T cells, followed by evolution into a T⁻B⁻NK⁺ SCID phenotype with opportunistic infections and granulomas.

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The patient was born to Kuwaiti first-degree cousin parents and presented at 6 months of age with pneumonia and failure to thrive. The physical examination was notable for the absence of tonsils. Computerized tomography of the chest revealed a prominent thymus. He had undetectable IgG and IgA. IgM was 30 mg/dL. The absolute lymphocyte count was 5010 cells/ μ L with normal numbers of total CD3⁺ cells, CD4⁺ and CD8⁺ T cells, CD16⁺/CD56⁺ NK cells, but virtually no CD19⁺ B cells (Table 1). Genetic testing did not reveal any mutations in the genes encoding BTK, μ heavy chain, Ig α , Ig β , λ 5, and VpreB. He was given a diagnosis of agammaglobulinemia with absent B cells and was treated with IVIG.

The patient subsequently developed recurrent otitis media and cytomegalovirus-associated retinitis at 4 years of age. Immunologic evaluation performed at 5 years of age revealed virtually absent B cells, CD3⁺ and CD4⁺ T cell lymphopenia, and normal numbers of NK cells (Table I). Lymphocyte proliferation to PHA was severely decreased (Table I). NK cell cytotoxic function was normal (data not shown). At 8 years of age, he developed fever and diarrhea due to Epstein-Barr virus viremia and *Salmonella enteritidis*. At 10 years of age, the patient developed persistent colitis; endoscopy and colonoscopy revealed esophageal candidiasis and colonic granulomas. At that time, he had progressive T cell lymphopenia with a predominantly CD45RO⁺ activated phenotype, and nearly absent CD4⁺CD45RA⁺CD31⁺ recent thymic emigrants (Table I). HLA typing of the patient and his mother revealed no evidence of maternal engraftment. T cell proliferation to PHA and anti-CD3 stimulation was minimal. Proliferation to anti-CD3 stimulation increased with the addition of anti-CD28, suggesting residual signaling through the CD3 pathway. The combination of absent B cells, agammaglobulinemia, and progressive T cell lymphopenia with poor T cell function led to a diagnosis of combined immunodeficiency. The patient died at the age of 12 years from sepsis.

Whole genome sequencing was performed on DNA obtained from the patient prior to his death, his father, and his healthy sibling. Thirteen candidate genes had novel mutations that were homozygous in the patient, heterozygous in the father, and heterozygous or absent in the sibling (Table E1). Only the c.1211G>A mutation in *RAG1* correlated with the patient's phenotype of combined immunodeficiency. This mutation resulted in the substitution of arginine to glutamine at position 404 (p.R404Q) in the nonamer-binding region (NBR) of the protein. The NBR of RAG1 binds to recombination signal sequences that flank the Variable (V), Diversity (D) and Joining (J) elements of the immunoglobulin and T cell receptor genes, and is critical for V(D)J recombination activity.⁴ Using a flow cytometry-based assay that permits analysis of RAG1 recombination activity on an intrachromosomal substrate,⁵ we have demonstrated that the R404Q RAG1 mutant protein is normally expressed, but has markedly reduced recombination activity ($1.18 \pm 0.14\%$ of wild-type) (Fig. 1). This mutation has been previously reported in patients with T⁻B⁻ SCID or Omenn syndrome.⁶ In contrast to previously published patients with this mutation, our patient presented with clinical and laboratory features of agammaglobulinemia with absent B cells, a prominent thymus, normal numbers of circulating T cells, and no features of Omenn syndrome such as erythroderma or eosinophilia. Normal numbers of T cells with absence of B cells have been associated with another *RAG1* mutation (c.T2686C, p.W896A) in a patient who presented with isolated culture-negative pneumonitis at 6 months of age.⁷ This patient,

like ours, had a normal thymic shadow and normal T cell numbers. The W896A mutation resulted in impaired proliferation to mitogens and a highly restricted TCR β repertoire that was not of maternal origin. We have previously demonstrated that the VDJ recombination activity of the W896A mutant is associated with minimally preserved RAG1 function ($0.9 \pm 0.1\%$ of wild-type)⁵, similar to the R404Q mutation in our patient,

This case broadens the spectrum of disease associated with a *RAG1* mutation that nearly abolishes recombination activity. This phenomenon has been seen in families where the same mutation gives rise to different clinical and immunological phenotypes, suggesting that the manifestations of *RAG1/2* mutations may reflect interactions with other uncharacterized genetic modifiers, epigenetic factors, or environmental exposures. The progression of the patient's disease also underscores how a single mutation can have variable time-dependent effects in the same individual. Although he had normal numbers of T cells at 6 months of age, T cell function may have been abnormal at that time as suggested by his history of pneumonia and failure to thrive. He developed progressively worsening T cell numbers and function, opportunistic infections, granulomas, and ultimately died of sepsis, which, in retrospect, are consistent with an underlying *RAG1/2* mutation. The diagnostic delay in this case underscores the importance of SCID newborn screening, particularly in areas of the world where lymphocyte proliferation or TCR repertoire studies are not available. Quantification of T cell receptor excision circles (TRECs) can identify poor thymic output, even in the case of normal T cell numbers due to maternal engraftment or oligoclonality.⁸ Additionally, cases of SCID which present with absent B cell numbers, as in our patient and others⁷, support the inclusion of quantification of κ -deleting recombination excision circles, which can detect absent B cells, as part of newborn screening for SCID for improved early detection.⁹

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Abbreviations

| | |
|-------------|----------------------------------|
| RAG | Recombination activating genes |
| SCID | severe combined immunodeficiency |

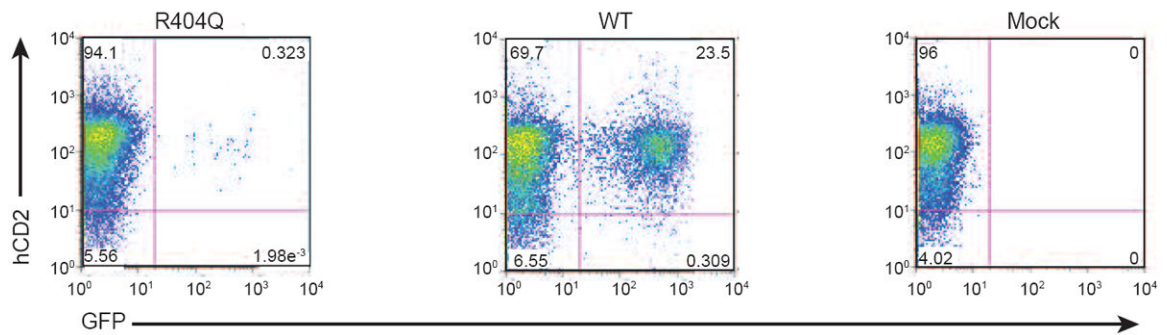


Figure 1.

Recombination activity of the RAG1 R404Q mutant protein. Abelson (Abl) virus-transformed *Rag1*^{-/-} mouse pro-B cells engineered to contain a single integrant of an inverted green fluorescent protein (GFP) cassette flanked by recombination signal sequences (RSS) were transduced with retroviral vectors allowing expression of wild-type (WT) human RAG1 or the mutant R404Q RAG1 protein, and human CD2 (hCD2) as a reporter. Mock-transduced cells served as a negative control. Following MACS-purification of hCD2-expressing cells and treatment *in vitro* with imatinib, recombination activity of the R404Q mutant protein was measured by normalizing GFP expression to levels observed in the presence of wild-type RAG1, as previously described.⁵ One representative experiment of three is shown.

Table 1

Immune profiles

| | 6 months | Proband 5 years | 11 years |
|----------------------------------------------------------|-------------------------------|-------------------------|-----------------------------------|
| Lymphocytes, cells/ μ L (normal range) ⁱ | 5010 (3400-9000) | 1550 (2300-5400) | 1620 (1900-3700) |
| CD3 ⁺ | 4520 (1900-5900) | 1150 (1400-3700) | 541 (1200-2600) |
| CD3 ⁺ CD4 ⁺ | 3300 (1400-4300) | 388 (700-2200) | 260 (650-1500) |
| CD3 ⁺ CD8 ⁺ | 1220 (500-1700) | 679 (490-1300) | 279 (370-1100) |
| CD4 ⁺ /CD8 ⁺ ratio | 2.7 (1.6-3.8) | 0.51 (0.9-2.6) | 0.93 (0.9 – 3.4) |
| CD4 ⁺ CD45RA ⁺ | ND | ND | 3.7 (46-77%) |
| CD4 ⁺ CD45RO ⁺ | ND | ND | 95 (13-30%) |
| CD4 ⁺ CD45RA ⁺ CD31 ⁺ | ND | ND | 0.96 (45.7%) ^{iv} |
| CD19 ⁺ | 2 (610-2600) | 4 (390-1400) | 61 (270-860) |
| CD16 ⁺ /CD56 ⁺ | 340 (160-950) | 349 (130-720) | 1004 (100-480) |
| Eosinophils, cells/ μ L (normal range) ⁱⁱ | 184 (200-300) | 95 (0-600) | 210 (0-600) |
| Immunoglobulins, mg/dL (normal range) ⁱⁱⁱ | | | |
| IgG | Undetectable (215-704) | ND | ND |
| IgA | Undetectable (8.1-68) | ND | ND |
| IgM | 30 (35-102) | ND | ND |
| Proliferation, cpm (normal control) ^{iv} | | | |
| Phytohemagglutinin | ND | 1578 (16,485) | 541 (65,369) |
| Anti-CD3 | ND | ND | 232 (3,814) |
| Anti-CD3+CD28 | ND | ND | 6,797 (19,170) |
| Background | ND | 32 (192) | 131 (221.5) |

ND, Not Determined

ⁱ Shearer WT, Rosenblatt HM, Gelman RS, Oyomopito R, Plaeger S, Stiehm ER, Wara DW, Douglas SD, Luzuriaga K, McFarland EJ, Yogev R, Rathore MH, Levy W, Graham BL, Spector SA; Pediatric AIDS Clinical Trial Group. Lymphocyte subsets in healthy children from birth through 18 years of age: the Pediatric AIDS Clinical Trials Group P1009 study. *J Allergy Clin Immunol* 2003; 112(5):973-80.

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^{iv} Value in parenthesis for lymphocyte proliferation are derived from studies done on a healthy control the same day.