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## Gain-of-function *STAT1* mutations are associated with PD-L1 overexpression and a defect in B-cell survival

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### Keywords

STAT1; CMC; T<sub>H</sub>17 cells; PD-L1; antibody deficiency; B cell lymphopenia

To the Editor:

Heterozygous gain-of-function mutations in the coiled-coil domain of *STAT1* were recently identified as a cause of chronic mucocutaneous candidiasis (CMC) (1, 2). As with *STAT3* mutations in hyper-IgE syndrome, the candidal susceptibility associated with gain-of-function *STAT1* mutations appears secondary to T<sub>H</sub>17 cell deficiency (3). The mechanistic link between constitutive *STAT1* activity and diminished T<sub>H</sub>17 cells has yet to be clearly defined. Here we present a kindred with a novel gain-of-function *STAT1* mutation associated with a complex clinical phenotype including candidiasis, humoral immunodeficiency, overexpression of programmed cell death protein ligand 1 (PD-L1) and increased B-cell apoptosis.

We have identified four related individuals each heterozygous for a novel E235A missense mutation in a highly conserved segment of the coiled-coil domain of *STAT1* (Fig 1, A). To our knowledge this is the first mutation described in exon 9 of the *STAT1a* gene locus to be associated with CMC. The mutation is not present in unaffected adult family members. The index patient (II.6) is a 60-year-old woman with CMC and progressive antibody deficiency.

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All patient samples were collected in accordance with IRB-approved protocols.

Beginning in infancy, the patient experienced candidal infections at sites including the oral cavity, esophagus, vagina, skin and nails. Later, as a young woman, the patient was diagnosed with IgG2 subclass deficiency that progressed to frank hypogammaglobulinemia (IgG 514, IgA 7, IgM<10 mg/dl) and B-cell lymphopenia (2 CD20<sup>+</sup> cells/ $\mu$ l, nml range 97–440) requiring IVIG. Over her lifetime, the patient suffered recurrent pulmonary infections from *P. aeruginosa*, *S. pneumonia*, *Serratia* species, *M. avium* and respiratory syncytial virus that resulted in severe bronchiectasis requiring lobectomy. Consistent with published reports of patients with gain-of-function *STAT1* mutations, our index patient (II.6) has experienced HPV<sup>+</sup> squamous cell carcinoma of palate, basal cell carcinoma, shingles and fibromuscular dysplasia with carotid and celiac/splenic artery dissection.

The index patient's daughter (III.2) is a 30-year-old female with CMC, B-cell lymphopenia (6 CD20<sup>+</sup> cells/ $\mu$ l) and IgG2 subclass deficiency manifesting in adolescence. Despite antifungal therapy, the patient experienced candidal infections at sites including the vagina, skin and nails. She has also experienced non-candidal infections including pneumonia, otitis media, sinusitis and chronic bronchitis. The third and fourth persons carrying the E235A allele are children of patient III.2, two males ages 6 weeks and 24 months (IV.1 and IV.2). They have not yet manifested symptoms of immunodeficiency.

Stimulation of T cells from patients II.6 and III.2 with IL-21 significantly increased phosphorylation of STAT1 compared to a healthy control (Fig 1, B). Furthermore, stimulation of patient PBMCs with PMA/ionomycin demonstrated diminished IL-17 secreting CD4<sup>+</sup> cells compared to a related healthy control (Fig 1, C). Hence the E235A mutation confers to STAT1 a gain-of-function and is associated with T<sub>H</sub>17-cell deficiency.

A remarkable feature of our kindred is overexpression of PD-L1 on the surface of naïve CD4<sup>+</sup> T cells. All four family members carrying the E235A allele had higher PD-L1 staining compared to members without it (Fig 1, D). To investigate if overexpression of PD-L1 was a common feature to gain-of-function *STAT1* mutations, we obtained PBMCs from two additional patients carrying either the I156T or the E353K missense *STAT1* mutation. Both patients, unrelated to our kindred, revealed a similar increase in PD-L1 expression on their naïve T cells (Fig 1, D). Recent data from mice demonstrate that the expression of PD-L1 on undifferentiated naïve T cells prevents commitment to the T<sub>H</sub>17 lineage through a PD-1/PD-L1 interaction. In this context, PD-L1 expression is dependent on IL27/IL27R binding and STAT1 (4). Accordingly, constitutively active STAT1 molecules in subjects carrying gain-of-function *STAT1* mutations may be responsible for increased PD-L1 expression on naïve T cells thereby discouraging differentiation into T<sub>H</sub>17 cells.

As previously described, the clinical phenotype of patients with gain-of-function *STAT1* mutations is quite broad and can include candidiasis, anti-thyroid autoimmunity, squamous cell carcinoma and vascular anomalies. In this issue two reports show broader phenotypes to gain-of-function *STAT1* mutations with an IPEX-like autoimmune syndrome in one report, and disseminated coccidioidomycosis and histoplasmosis in the other. Here, we report a not yet appreciated feature associated with gain-of-function *STAT1* mutations: humoral immunodeficiency.

STAT1's function as a key modulator of cell death is thoroughly described. Perhaps the best illustration of this is growth arrest of the STAT1-negative U3A fibroblast line upon transfection with wild type STAT1 $\alpha$  and treatment with interferon- $\gamma$  (5). The interferon- $\gamma$  receptor requires STAT1 for intracellular signaling. Moreover, transfection of the same cell line with constitutively activated STAT1 initiates caspase mediated apoptosis (6). Related experiments implicate STAT1 activation during apoptosis in B-cell lymphoma cells (7).

Interestingly, progressive B-cell lymphopenia is a remarkable feature of patients (II.6 and III.2) in our kindred suggesting a defect in cell survival. Indeed, CD19<sup>+</sup> B cells from subjects carrying the E235A allele appear apoptotic with increased Annexin V staining (Fig 2, A top row) and elevated caspase activity (Fig 2, B). In culture for 24 hours, B cells from patient III.2 demonstrated even greater Annexin V and considerable 7-AAD staining, evidence of accelerated cell death (Fig 2, A middle row). The patient's B cells were only partially rescued by stimulation of their B-cell receptors (Fig 2, A bottom row). We also found enhanced caspase activity in B cells from the two additional patients that were heterozygous for either the E353K or the I156T missense *STAT1* mutation (Fig 2, B). B cell lymphopenia was a significant finding in the former patient (39 cells/ $\mu$ l) but not the latter (335 cells/ $\mu$ l). Altogether, our data reveal that gain-of-function *STAT1* mutations increase B-cell apoptosis. Over time this may result in B-cell lymphopenia and antibody deficiency.

In summary, we identified individuals heterozygous for gain-of-function *STAT1* mutation with two unappreciated features. The first is the overexpression of PD-L1 on naïve T cells which provides a general mechanism for how constitutively active STAT1 blocks the development of the T<sub>H</sub>17 lineage. The second feature, accelerated B-cell apoptosis that may result in progressive B-cell lymphopenia and humoral immunodeficiency, further broadens the clinical phenotype associated with gain-of-function *STAT1* mutations.

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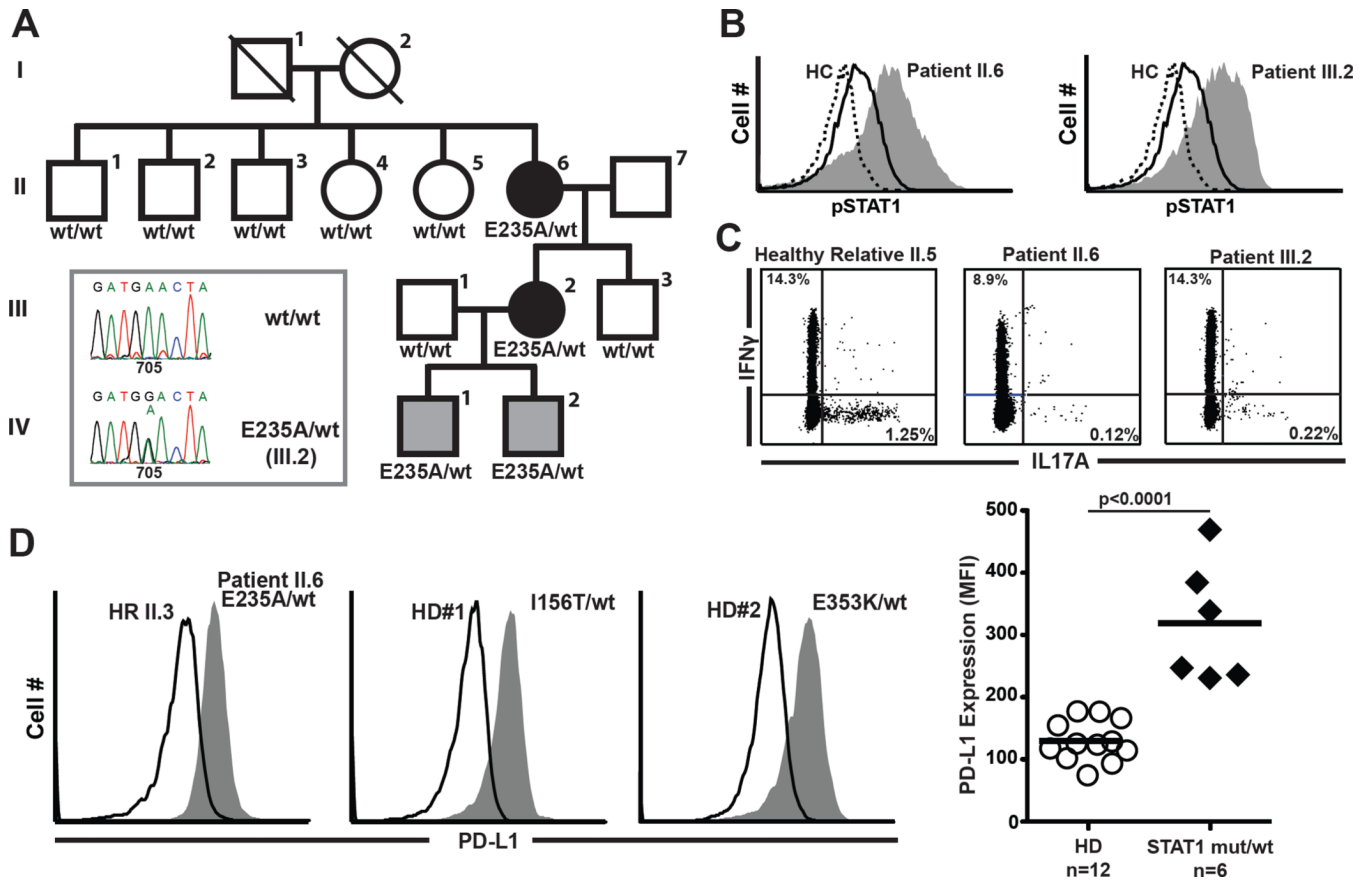
## Abbreviations

<b>STAT1</b>	Signal transducer and activator of transcription 1
<b>PD-1</b>	Programmed cell death protein 1
<b>PD-L1</b>	Programmed cell death protein ligand 1
<b>CMC</b>	Chronic mucocutaneous candidiasis

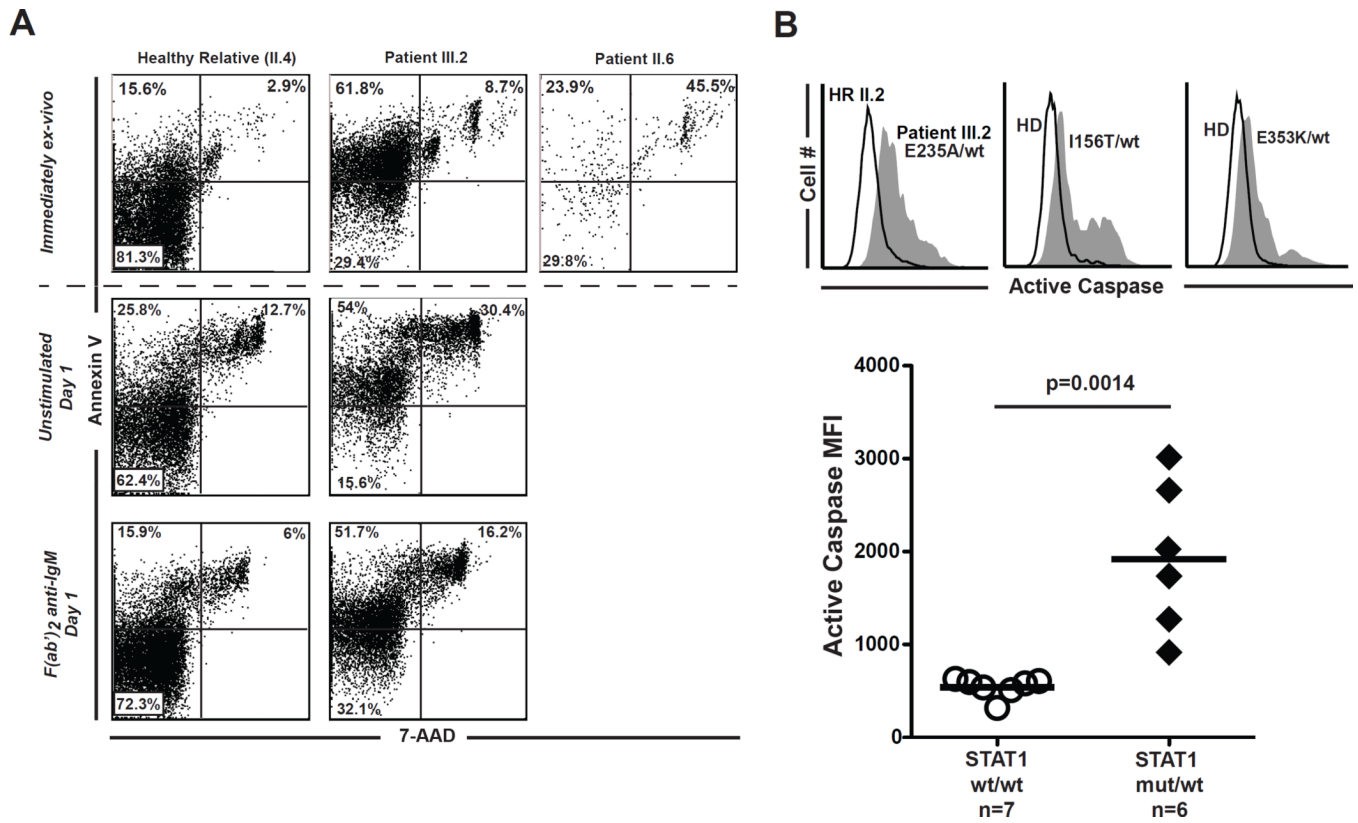
## References

- van de Veerdonk FL, Plantinga TS, Hoischen A, Smeekens SP, Joosten LA, Gilissen C, et al. *STAT1* mutations in autosomal dominant chronic mucocutaneous candidiasis. *N Engl J Med*. 2011 Jul 7; 365(1):54–61. [PubMed: 21714643]
- Liu L, Okada S, Kong XF, Kreins AY, Cypowij S, Abhyankar A, et al. Gain-of-function human *STAT1* mutations impair IL-17 immunity and underlie chronic mucocutaneous candidiasis. *The Journal of Experimental Medicine*. 2011 Aug 1; 208(8):1635–1648. [PubMed: 21727188]
- Milner JD, Brechley JM, Laurence A, Freeman AF, Hill BJ, Elias KM, et al. Impaired T(H)17 cell differentiation in subjects with autosomal dominant hyper-IgE syndrome. *Nature*. 2008 Apr 10; 452(7188):773–776. [PubMed: 18337720]
- Hirahara K, Ghoreschi K, Yang XP, Takahashi H, Laurence A, Vahedi G, et al. Interleukin-27 Priming of T Cells Controls IL-17 Production In trans via Induction of the Ligand PD-L1. *Immunity*. 2012 Jun 29; 36(6):1017–1030. [PubMed: 22726954]
- Chin YE, Kitagawa M, Su WC, You ZH, Iwamoto Y, Fu XY. Cell growth arrest and induction of cyclin-dependent kinase inhibitor p21 WAF1/CIP1 mediated by *STAT1*. *Science*. 1996 May 3; 272(5262):719–722. [PubMed: 8614832]

6. Sironi JJ, Ouchi T. STAT1-induced apoptosis is mediated by caspases 2, 3, and 7. *J Biol Chem*. 2004 Feb 6; 279(6):4066–4074. [PubMed: 14623896]
7. Niitsu N, Higashihara M, Honma Y. Human B-cell lymphoma cell lines are highly sensitive to apoptosis induced by all-trans retinoic acid and interferon-gamma. *Leuk Res*. 2002 Aug; 26(8):745–755. [PubMed: 12191570]

**FIG 1.**

Gain-of-function *STAT1* mutations increase PD-L1 expression. **A**, Pedigree with patients (black) and carriers (gray) heterozygous at *STAT1* nucleotide 705. **B**, Increased *STAT1* phosphorylation in IL-21 treated T-cells from patients (filled), compared to controls (solid) and unstimulated (dashed). **C**, Reduced T<sub>H</sub>17-cell frequencies in E235A *STAT1* patients. **D**, PD-L1 overexpression on CD3<sup>+</sup>CD4<sup>+</sup>CD62L<sup>+</sup>CD45RO<sup>-</sup> cells from subjects with *STAT1* mutations (filled) versus controls (solid).

**FIG 2.**

Gain-of-function *STAT1* mutations promote B-cell death. **A**, Annexin V and 7-AAD staining of CD19<sup>+</sup> cells immediately ex-vivo (top row) and after 24 hours in culture with (middle row) or without rescue with F(ab')<sub>2</sub>-anti-IgM (bottom row). **B**, Active caspase staining of CD19<sup>+</sup> cells in subjects heterozygous for the indicated *STAT1* mutations (filled) vs. related and unrelated controls (solid).