

Supplemental Data

Patients with *CD3G* mutations reveal a role for human CD3 γ in T_{reg} diversity and suppressive function

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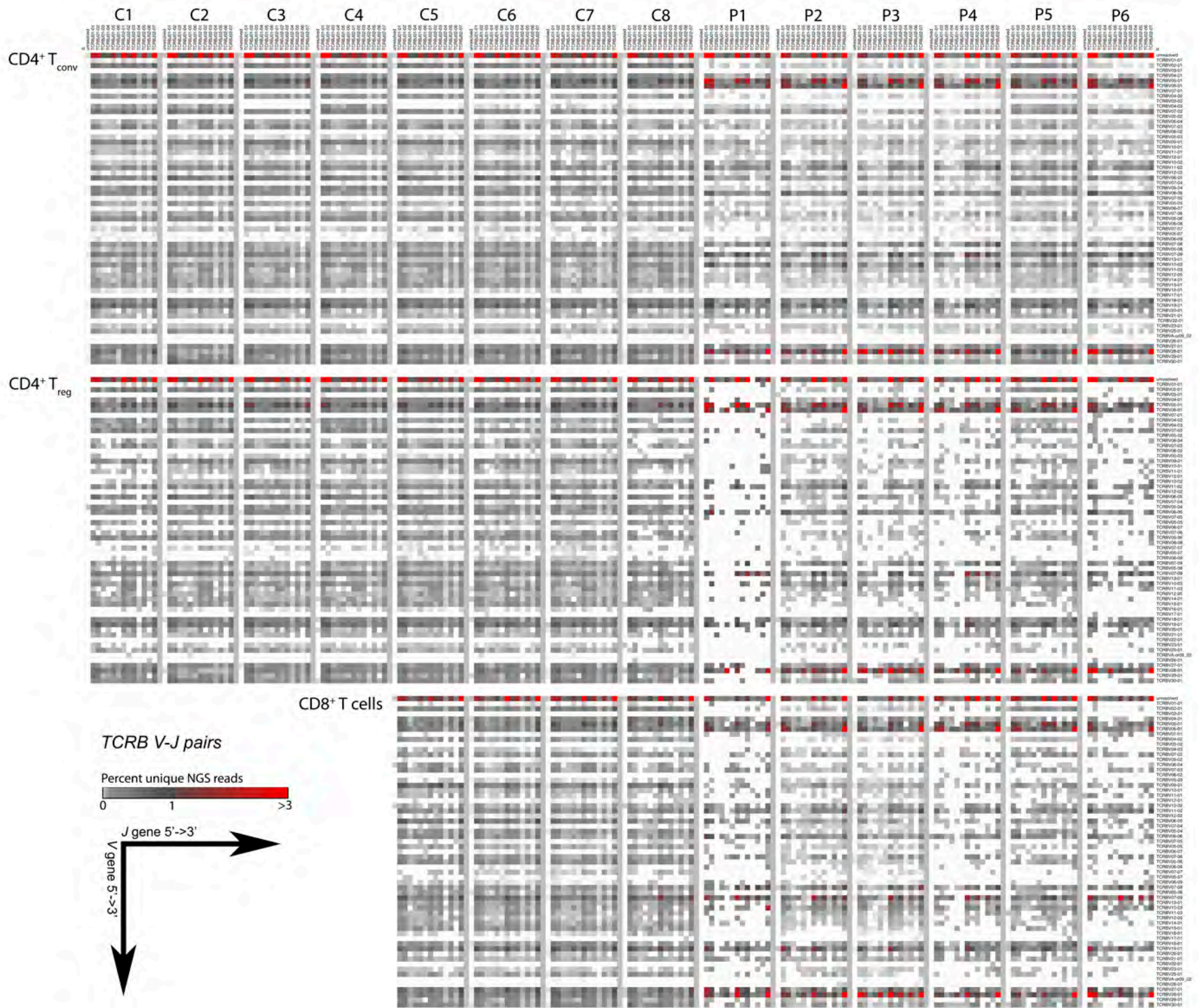
Table S1 – Characteristics of autoimmunity in CD3G-mutated patients

Patient #	Autoimmune manifestation	Age at onset	Treatment	Age at the time of the study	Outcome	Current age (years)
P1	Enteropathy AIHA GLILD	15 months 21 months 21 months	Rituximab, sirolimus	27 months	MUD-HSCT at 32 months, deceased at 47 months (GvHD, infections)	N/A
P2	Subclinical hypothyroidism (positive TG and TPO Ab)	11 years	none	19 years	Actively followed, no treatment needed	20
P3	Hypothyroidism	5 years	Thyroid hormone replacement	6 years	Still under hormone replacement therapy	7
P4	Evans syndrome Autoimmune hepatitis Nephropathy ALPS	12 months 5 years 5 years 12 years	Steroids, cyclosporine A, high dose IVIG	15 years	Multiple relapses of Evans syndrome, persistent hepatomegaly and splenomegaly	18
P5	Hypothyroidism	10 years	Thyroid hormone replacement	23 years	Still under hormone replacement therapy	24
P6	Vitiligo Subclinical hypothyroidism (positive TG and TPO Ab)	12 months 18 years	none	34 years	Actively followed, no treatment needed	36

Ab, antibody; AIHA, autoimmune hemolytic anemia; ALPS, autoimmune lymphoproliferative syndrome; GLILD, granulomatous lymphocytic interstitial lung disease; GvHD, graft versus host disease; IVIG, intravenous immunoglobulins; MUD-HSCT, hematopoietic stem cell transplantation from a matched unrelated donor; TG, thyroglobulin; TPO, thyroid peroxidase

N/A: not applicable

Figure S1 - High throughput sequencing analysis of TRB repertoire in CD4⁺ T_{conv}, CD4⁺ T_{reg}, and CD8⁺ T cell subsets. Heat maps showing Vβ and Jβ pairing in unique TRB reads of CD4⁺ T_{conv} (top), CD4⁺ T_{reg} (middle) or CD8⁺ T cells (bottom) from all controls and all CD3G-mutated patients included in the study.



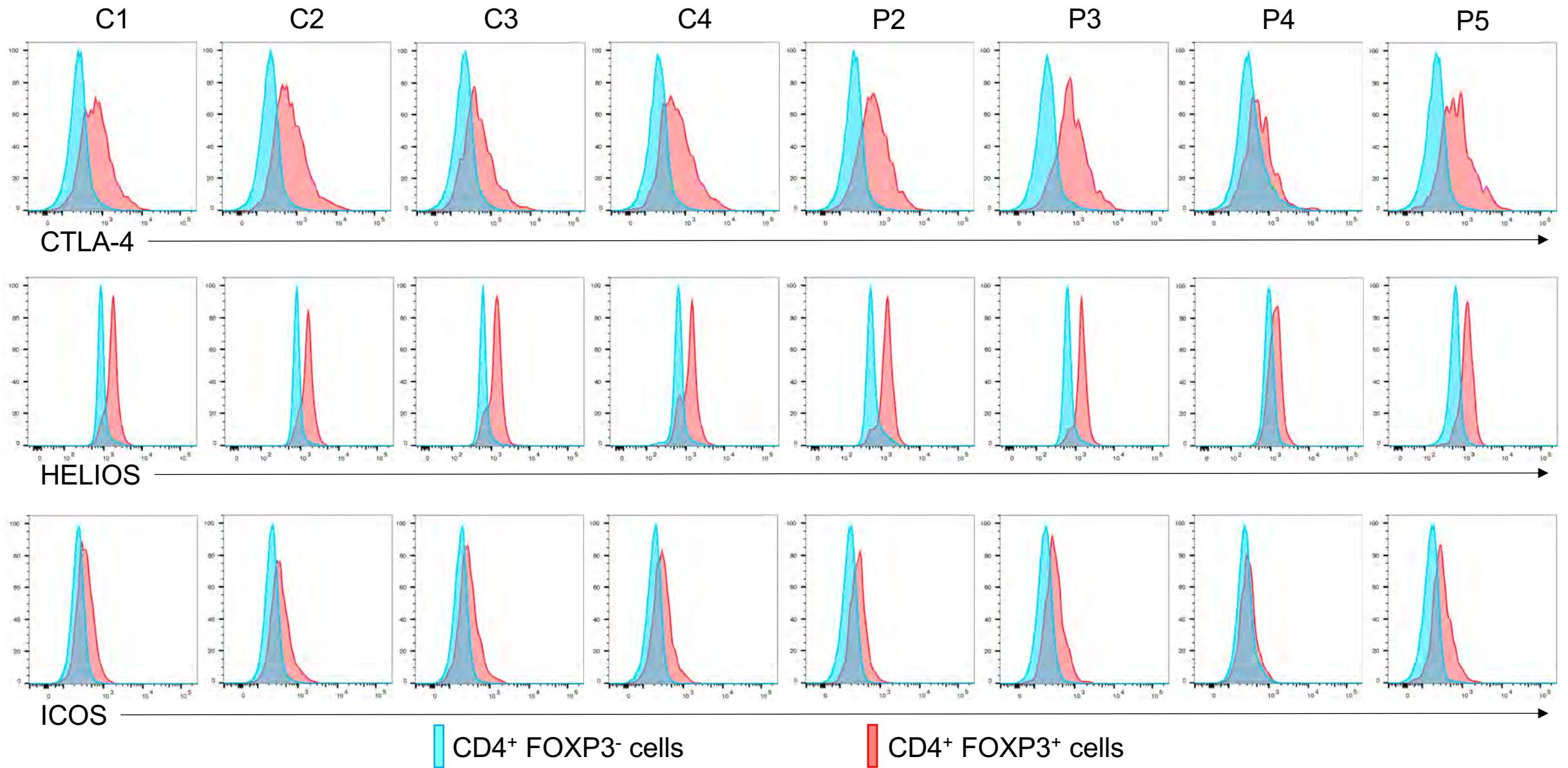


Figure S2 – Overlay of expression of CTLA-4, HELIOS, and ICOS on CD4⁺ FOXP3⁻ T conventional (T_{conv}) cells (blue) and CD4⁺ FOXP3⁺ regulatory T (T_{reg}) cells (red) from controls (C1-C4) and patients (P2-P5). Results demonstrate that T_{reg} cells from both controls and patients show increased expression of CTLA-4, HELIOS, and ICOS as compared to T_{conv} cells.