

Table S1. Genetic and clinical features of the patients

Patient	Syndrome ^a	Gene	Mutations	Age (yr)	Sex	Origin	Infections					References
							<i>Pneumococcus</i>	<i>Staphylococcus</i>	<i>Salmonella</i> ^b	<i>Mycobacteria</i> ^c	<i>Candida</i>	
1	MPPBI	<i>IRAK4</i>	E402X	11	M	Spain	+	–	–	–	–	Ku et al. ^d
2	MPPBI	<i>IRAK4</i>	1-1096_40+23del	11	M	Israel	+	–	–	–	–	Ku et al.
3 ^e	MPPBI	<i>IRAK4</i>	M1V/1188+520A>G	3	F	Slovenia	–	–	–	–	–	Ku et al.
4	MPPBI	<i>IRAK4</i>	1189-1G>T/1188+520A>G	10	M	Hungary	+	–	–	–	–	Ku et al.
5	MPPBI	<i>IRAK4</i>	Q293X	33	F	UK	+	–	–	–	–	Ku et al.
6	MPPBI	<i>IRAK4</i>	Q293X	28	M	Canada	+	–	–	–	–	Ku et al.
7	MPPBI	<i>MYD88</i>	L93P/R196C	4	F	Turkey	+	–	–	–	–	unpublished data
8	MPPBI	<i>MYD88</i>	R196C	16	F	Portugal	+	–	+	–	–	unpublished data
9	MPPBI	<i>MYD88</i>	R196C	10	M	Portugal	+	+	+	–	–	unpublished data
10	MSMD	<i>IL12B</i>	297del8	7	M	Tunisia	–	–	<i>Se</i>	–	–	This report
11	MSMD	<i>IL12B</i>	297del8	24	M	Tunisia	–	–	–	–	–	This report
12	MSMD	<i>IL12RB1</i>	1791+2T>G	12	F	Spain	–	–	–	<i>Mtb</i>	–	Caragol et al. ^f
13	MSMD	<i>IL12RB1</i>	1791+2T>G	20	F	Spain	–	–	<i>Se</i>	<i>Mtb</i>	–	Caragol et al.
14	MSMD	<i>IL12RB1</i>	1791+2T>G	22	F	Spain	–	–	–	–	–	Caragol et al.
15	MSMD	<i>IL12RB1</i>	628-644dup	12	M	Turkey	–	–	–	BCG	–	Tanir et al. ^g
16	MSMD	<i>IL12RB1</i>	628-644dup	3	M	Turkey	–	–	–	–	+	Tanir et al.

17	MSMD	<i>IL12RB1</i>	Q32X	12	F	France	–	–	–	BCG	–	Fieschi et al ^h
18	MSMD	<i>IL12RB1</i>	K305X	29	F	Morocco	–	–	<i>St</i>	BCG	–	Fieschi et al.
19	MSMD	<i>IL12RB1</i>	700+362_1619-944del	11	F	Israel	–	–	–	–	–	Scheuerman et al. ⁱ
20	MSMD	<i>IL12RB1</i>	C198R	15	M	Turkey	–	–	–	BCG	–	Lichtenauer-Kaligis et al. ^j
21	MSMD	<i>IL12RB1</i>	R173P	14	M	Turkey	–	–	<i>Se</i>	–	–	This report
22	MSMD	<i>IL12RB1</i>	1745-46delinsCA/ 1483+182_1619-1073del	37	F	France	–	–	+	BCG	–	Fieschi et al.
23	MSMD	<i>IL12RB1</i>	C198R	8	F	Turkey	–	–	–	–	–	This report
24	MSMD	<i>IL12RB1</i>	C198R	4	M	Turkey	–	–	+	BCG	–	This report
25	MSMD	<i>IL12RB1</i>	Y367C	8	M	Cameroon	–	–	<i>Sd</i>	BCG	–	Fieschi et al.
26	MSMD	<i>IL12RB1</i>	1791+2T>G	24	F	Sri Lanka	–	–	–	BCG	–	Fieschi et al.
27	CE	<i>TGFB1</i>	R218C	31	F	France	–	–	–	–	–	Campos-Xavier et al. ^k
28	CE	<i>TGFB1</i>	R218C	62	F	France	–	–	–	–	–	Campos-Xavier et al.
29	CE	<i>TGFB1</i>	R218C	53	M	France	–	–	–	–	–	Campos-Xavier et al.
30	MLS	<i>TGFBR1</i>	K333Q	7	F	France	–	–	–	–	–	This report
31	MLS	<i>TGFBR2</i>	R537C	34	M	France	–	–	–	–	–	Mizuguchi et al. ^l
32	MLS	<i>TGFBR2</i>	C394W	41	F	France	–	–	–	–	–	This report
33	MLS	<i>TGFBR2</i>	C394W	14	F	France	–	–	–	–	–	This report
34	MLS	<i>TGFBR2</i>	C394W	10	F	France	–	–	–	–	–	This report
35	AD-HIES	<i>STAT3</i>	V463del	34	F	France	–	+	–	–	–	This report

36	AD-HIES	STAT3	V463del	8	M	France	+	+	-	-	+	This report
37	AD-HIES	STAT3	V463del	9	F	France	-	+	-	-	-	This report
38	AD-HIES	STAT3	K709E	17	M	France	-	+	-	-	+	This report
39	AD-HIES	STAT3	T412S	19	F	France	-	+	-	-	-	This report
40	AD-HIES	STAT3	V463del	37	F	Pakistan	-	+	-	-	+	This report
41	AD-HIES	STAT3	V463del	9	M	Pakistan	+	+	-	-	+	This report
42	AD-HIES	STAT3	K642E	36	M	France	-	+	-	-	+	This report
43	AD-HIES	STAT3	R382W	28	F	France	-	+	-	-	+	This report
44	AD-HIES	STAT3	R382Q	19	M	Turkey	-	+	-	-	+	This report
45	AD-HIES	STAT3	R382W	21	F	France	+	+	-	-	+	This report
46	AD-HIES	STAT3	R382W	16	M	Algeria	+	+	-	-	+	This report
47	AD-HIES	STAT3	R382W	23	M	France	+	+	-	-	+	This report
48	AD-HIES	STAT3	V463del	28	M	France	-	+	-	-	+	This report
49	AD-HIES	STAT3	N472D	17	M	France	-	+	-	-	+	This report
50	AD-HIES	STAT3	I665N	43	F	France	-	+	-	-	+	This report

^aShown are Mendelian predisposition to pyogenic bacterial infections (MPPBI), Mendelian susceptibility to mycobacterial diseases (MSMD), Camurati-Engelmann (CE) disease, Marfan-like syndromes (MLS), and AD-HIES.

^bInfections caused by *Salmonella enteritidis* (*Se*), *Salmonella typhimurium* (*St*), and *Salmonella dublin* (*Sd*).

^cInfections caused by Bacille Calmette-Guerin (BCG) or by *Mycobacterium tuberculosis* (*Mtb*).

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^ePatient 3 suffered from invasive infection caused by *Pseudomonas aeruginosa*.

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Table S2. Percentage of CCR6-positive CD4 T cells in controls and STAT-3–deficient patients ex vivo

Patients and controls^a	Age (yr)	CCR6⁺ CCR4⁺ CD4⁺ T cells (%)	CCR6⁺ CCR4⁻ CD4⁺ T cells (%)
P 35	7	4.4	0.5
P 36	9	5.7	0.5
P 37	34	6.7	1.1
P 38	16	8.7	7
P 46	16	8.9	1.4
Other patient ^b	7	4.7	2.4
Other patient ^b	15	3.1	0.2
Other patient ^b	21	4.9	0.5
C 1	5	7.8	1.5
C 2	7	6.8	6.5
C 3	7	6.6	8.8
C 4	7	11.1	6.2
C 5	12	16.6	15.2
C 6	13	6.8	12.2
C 7	16	8.4	18.5
C 8	unknown	8.3	11.8
C 9	unknown	10.9	20.3

^aEight STAT-3–deficient patients (P) and nine healthy controls (C) were studied.

^bThese patients, not described in Table S1, were not studied for IL-17 production.

Table S3. Percentage of CD4- and CD8-positive T cells in controls and patients ex vivo

Patient	Gene	Age (yr)	Lymphocytes ($\times 10^9$ per μl)	CD4⁺ (%)	CD8⁺ (%)
2	<i>IRAK4</i>	11	4.8	63	17
4	<i>IRAK4</i>	10	1.9	45	25
6	<i>IRAK4</i>	28	1.3	46	20
17	<i>IL12RB1</i>	12	Not done	35	19
18	<i>IL12RB1</i>	29	1.8	23	37
38	<i>STAT3</i>	17	1.3	37	21
39	<i>STAT3</i>	19	Not done	38	33
40	<i>STAT3</i>	37	2.9	41	23
41	<i>STAT3</i>	9	3.4	31	12
42	<i>STAT3</i>	36	0.8	46	20
43	<i>STAT3</i>	28	3	28	21
45	<i>STAT3</i>	21	2.6	40	34
46	<i>STAT3</i>	16	3.3	43	22
47	<i>STAT3</i>	23	1.5	35	24
49	<i>STAT3</i>	17	Not done	35	19
50	<i>STAT3</i>	43	1	39	21