

## Low Penetrance, Broad Resistance, and Favorable Outcome of Interleukin 12 Receptor $\beta 1$ Deficiency: Medical and Immunological Implications

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## Abstract

The clinical phenotype of interleukin 12 receptor  $\beta 1$  chain (IL-12R $\beta 1$ ) deficiency and the function of human IL-12 in host defense remain largely unknown, due to the small number of patients reported. We now report 41 patients with complete IL-12R $\beta 1$  deficiency from 17 countries. The only opportunistic infections observed, in 34 patients, were of childhood onset and caused by weakly virulent *Salmonella* or *Mycobacteria* (Bacille Calmette-Guérin -BCG- and environmental *Mycobacteria*). Three patients had clinical tuberculosis, one of whom also had salmonellosis. Unlike salmonellosis, mycobacterial infections did not recur. BCG inoculation and BCG disease were both effective against subsequent environmental mycobacteriosis, but not against salmonellosis. Excluding the probands, seven of the 12 affected siblings have remained free of case-definition opportunistic infection. Finally, only five deaths occurred in childhood, and the remaining 36 patients are alive and well. Thus, a diagnosis of IL-12R $\beta 1$  deficiency should be considered in children with opportunistic mycobacteriosis or salmonellosis; healthy siblings of probands and selected cases of tuberculosis should also be investigated. The overall prognosis is good due to broad resistance to infection and the low penetrance and favorable outcome of infections. Unexpectedly, human IL-12 is redundant in protective immunity against most microorganisms other than *Mycobacteria* and *Salmonella*. Moreover, IL-12 is redundant for primary immunity to *Mycobacteria* and *Salmonella* in many individuals and for secondary immunity to *Mycobacteria* but not to *Salmonella* in most.

Key words: *Mycobacteria* • *Salmonella* • immunodeficiency • interleukin 12 receptor • interferon  $\gamma$

## Introduction

Mendelian susceptibility to mycobacterial disease (MSMD; Mendelian inheritance in man [MIM] 209950) (1) is a rare syndrome which results in predisposition to clinical disease caused by poorly virulent mycobacterial species such as the Bacillus Calmette-Guérin (BCG)\* vaccines (2) and nontuberculous environmental *Mycobacteria* (EM; reference 3). Patients are also susceptible to virulent *Mycobacterium tuberculosis* (1). Unlike patients with classic immunodeficiencies, these patients rarely present other unusual infectious diseases, with the exception of extra-intestinal nontyphoid salmonellosis, which affects less than half of them. Histopathological and clinical heterogeneity are suggestive of genetic heterogeneity. Some sporadic and most familial cases suggest autosomal recessive heredity, but the syndrome segregates in an autosomal dominant (4) or X-linked recessive (5) pattern in some families.

Five disease-causing autosomal genes have been found, and allelic heterogeneity accounts for the existence of nine defined disorders, all of which result in impaired IFN- $\gamma$ -mediated immunity (1). Null recessive mutations in the IFN- $\gamma$ -receptor ligand-binding chain (IFN- $\gamma$ R1)-encoding gene (*IFNGR1*) abolish either receptor expression (6, 7) or the binding of surface-expressed receptors to IFN- $\gamma$  (8). Partial recessive (9) and dominant (4) IFN- $\gamma$ R1 deficiencies have also been described. Different recessive mutations in the gene of the IFN- $\gamma$  signaling chain (IFN- $\gamma$ R2), *IFNGR2*, are responsible for complete (10) or partial (11) IFN- $\gamma$ R2 deficiency. A dominant mutation in *STAT1*, resulting in partial STAT-1 (signal transducer and activator of transcription-1) deficiency and impaired cellular responses to IFN- $\gamma$ , has also been identified in other kindreds (12).

\*Abbreviations used in this paper: BCG, Bacille Calmette-Guérin; EM, environmental *Mycobacteria*; SSCP, single-stranded conformation polymorphism.

The remaining two genetic defects result in normal cellular responses to IFN- $\gamma$ , but abnormal IL-12-dependent production of IFN- $\gamma$ . A few children are homozygous for null mutations in *IL12B*, encoding the p40 subunit of IL-12 (13–15). Null recessive *IL12RB1* mutations have been identified in other patients with IL-12 receptor  $\beta 1$  chain deficiency (15–21). These studies (13–21) only addressed case reports or small cohorts of patients, making it difficult to estimate the range of pathogenic microorganisms and the severity of the clinical course. These issues are of both clinical and immunological importance, given the central role attributed to IL-12 in Th1 development and immunity to various pathogens (22–24). Here, we describe the *IL12RB1* genotype and the cellular and clinical phenotypes of 41 patients with IL-12R $\beta 1$  deficiency.

## Materials and Methods

**Subjects, Kindreds, and Statistical Methods.** We investigated 120 unrelated patients, including five previously described patients (16, 19, 20). Our study was conducted according to the principles expressed in the Helsinki Declaration, with informed consent obtained from each patient or the patient's family. The proportion of infection-free individuals, survival, and penetrance were calculated by the Kaplan-Meier method, and the differences between curves were assessed by the log-rank test. All calculations were performed with the Lifetest procedure of SAS version 8.2 (SAS).

**Cell Culture and Stimulation.** Epstein-Barr virus-transformed lymphoblastoid cell lines (EBV-B cell lines) were cultured as described previously (14). Whole blood samples were diluted 1:2 in RPMI 1640 (Invitrogen) and infected by incubation with live *M. bovis* BCG at a multiplicity of infection of 20:1, alone or with recombinant IL-12p70 (20 ng/ml; R&D Systems), for 48 h. Alternatively, PBMCs were cultured in RPMI 1640 supplemented with 10% heat-inactivated pooled human AB serum, and acti-

vated by incubation with PHA P 1/700 (Bacto PHA-P; Becton Dickinson) for 72 h. T cell blasts were restimulated every 48 h with IL-2 (40 IU/ml; Chiron Corp.).

**ELISA and Flow Cytometry.** Cell culture supernatants were assayed for IFN- $\gamma$  by ELISA, according to the kit manufacturer's recommendations (Pelikn Compact, CLB). IFN- $\gamma$  concentration was calculated per 1 million PBMCs. T cell blasts and/or EBV-B cells were first incubated with an IL-12R $\beta$ 1-specific mouse IgG1 mAb (24E6), an IL-12R $\beta$ 1-specific rat IgG2a mAb (2B10), or isotypic control mAbs, then with a biotinylated rat anti-mouse Ab or a biotinylated mouse anti-rat Ab, and finally with streptavidin-PE (all reagents were obtained from BD Biosciences/Becton Dickinson). Signals were analyzed with a FACScan<sup>TM</sup> machine, using CELLQuest<sup>TM</sup> software (Becton Dickinson).

**DNA and RNA Extraction, cDNA Synthesis, and PCR Amplification.** Genomic DNA and total RNA were extracted from EBV-transformed B cells or T cell blasts, as described previously (14). RNA was reverse transcribed in the presence of oligo (dT) with Superscript II reverse transcriptase (Invitrogen; reference 14). The *IL12RB1* cDNA, coding exons and flanking intron regions were amplified using pairs of primers and PCR conditions available upon request.

**Single-stranded Chain Polymorphism and Sequencing.** Single-stranded chain polymorphism (SSCP) analysis and PCR were performed with pairs of intron primers flanking each *IL12RB1* exon, under conditions available upon request. PCR products were sequenced by dideoxynucleotide termination with nested primers (available upon request) and the BigDye terminator kit. PCR products were sequenced on an ABI Prism 3100 apparatus, and analyzed with Sequencing Analysis software (Applied Biosystems).

## Results

We investigated 120 unrelated probands, including 100 with MSMD syndrome (disseminated BCG or EM disease with or without salmonellosis), and 20 with only nontyphoid extraintestinal salmonellosis, as we recently described a child with salmonellosis as the sole infectious disease and IL-12p40 deficiency (14).

**Detection of *IL12RB1* Mutations and Intrafamilial Segregation.** The 17 *IL12RB1* coding exons and flanking intron regions were analyzed by SSCP. 25 patients displayed a unique SSCP pattern, each for only one exon (Kindreds 1–9, 11–16, 18–26, 29; Table I and Fig. 1). Three additional patients (17.II.2, 27.II.2, 28.II.3) displayed unique SSCP patterns for two separate exons, and in one patient (10.II.2), exons 8 to 13 were not amplified. All coding exons were sequenced in these 29 patients with abnormal SSCP patterns. Introns 7 and 13 were also sequenced in patient 10.II.2, and introns 12 and 13 in patient 19.II.1 (owing to the identification of a cDNA of low molecular weight, see below). The patients were found to be homozygous, or compound heterozygous for nonsense ( $n = 4$ ), splice ( $n = 4$ ), and missense ( $n = 6$ ) mutations, small insertions ( $n = 1$ ), large deletions ( $n = 2$ ), and deletions/insertions ( $n = 4$ ; a total of 21 mutant alleles; Table I). None of the mutations were found in 50 unrelated healthy individuals from the corresponding ethnic group. We then amplified and sequenced the full-length *IL12RB1* cDNA coding region for the 29 probands. *IL12RB1* transcripts of low

molecular weight were found in patients 10.II.2 and 19.II.1. In patients homozygous for splice mutations, no full-length mRNAs were found. All parents tested were heterozygous for one *IL12RB1* mutation. Only 12 of the 61 known siblings could not be genotyped (designated “E?” in Fig. 1). Two had died of proven BCG (26.II.1) or EM (4.II.2) infection and therefore probably carried the disease-associated *IL12RB1* genotype. 10 of the 49 siblings tested, from nine kindreds, carried two mutant *IL12RB1* alleles, raising the number of affected siblings to 12. Altogether, we report 41 individuals from 29 kindreds demonstrated ( $n = 39$ ) or thought ( $n = 2$ ) to be homozygous or compound heterozygous for *IL12RB1* mutations, including 29 probands and 12 siblings.

**Impaired Surface Expression and Function of IL-12R $\beta$ 1.** We assessed IL-12R $\beta$ 1 expression on the surface of T cell blasts (kindreds 9, 14, 15, 22, 25), EBV-transformed B cells (kindreds 8, 11, 19, 20, 29), or both (kindreds 3, 6, 10, 12, 16, 17, 23, 24, 26–28), from 21 of the 24 newly diagnosed probands, by flow cytometry with specific antibodies. Consistent with the five patients previously reported (16, 19, 20), no IL-12R $\beta$ 1 molecules were detected on the surface of cells from all but one patient. Patient 10.II.2, carrying a large in-frame deletion, presented an expression of the  $\beta$ 1 chain, both on T cell blasts and EBV-transformed B cells, detectable with one monoclonal antibody (24E6), but not with another (2B10; data not depicted). Cells from the three remaining probands (kindreds 13, 18, 21) were not tested for IL-12R $\beta$ 1 expression, but were homozygous for mutations found in other kindreds (Table I). The cells from all 21 patients tested, including the proband from kindred 10 with residual IL-12R $\beta$ 1 expression produced less IFN- $\gamma$  than the “travel” and “local” laboratory controls in response to BCG (mean of 76 versus 664 and 4,109 pg/ml), and production was not increased by IL-12 (mean of 87 versus 22,432 and 65,636 pg/ml; Fig. 2). Some patients were not tested with this assay (kindreds 4, 7, 12, 13, 14, 18, 22, 29). However, kindreds 4, 7, 14, 18, and 22 carry the same mutation as other patients (Table I). We and others have demonstrated that T cells homozygous for the *IL12RB1* missense mutation R213W (kindred 2) are complemented by transfection with a wild-type *IL12RB1* allele (20, 21). Overall, we have demonstrated complete IL-12R $\beta$ 1 deficiency in 29 patients (absence of detectable IL-12R $\beta$ 1 surface expression,  $n = 28$ , absence of response to IL-12,  $n = 27$ , or both,  $n = 27$ ).

**Onset and Outcome of Infections in IL-12R $\beta$ 1-deficient Patients.** We first analyzed the 34 individuals who developed case-definition opportunistic infections caused by weakly virulent microorganisms (BCG, EM, and nontyphoid *Salmonella*; Table I). 25 of these 34 patients had been inoculated with live BCG in childhood. Only 18 developed disseminated BCG disease, within six months of inoculation (Fig. 3 a). No recurrence of BCG disease was observed in the 16 patients who survived the initial BCG infection. Eight patients suffered EM disease. EM disease began before the age of six years, in all but two patients (4.II.1 and 7.II.5) who developed *M. avium* and *M. fortuitum* infections at the

**Table I.** Genotypes and Clinical Phenotypes of Patients with IL-12 Receptor  $\beta 1$  Deficiency

Kindred	Patients <sup>a</sup>	Mutations <sup>b</sup>	Origin <sup>c</sup>	Follow-up <sup>d</sup>	Age <sup>e</sup> (yr)	BCG <sup>f</sup>	EM <sup>f</sup>	Mtb <sup>f</sup>	Salmonella <sup>f</sup>
1	II.2	K305X	Morocco	Alive	23	D	–	–	<i>S. typhimurium</i>
2	II.1	R213W	Morocco	Alive	26	Resistant	–	GI tract	–
	II.3			Alive	15	D	–	–	<i>S. typhimurium</i>
3	II.3	Y367C	Cameroon	Alive	2	D	–	–	<i>S. dublin</i>
4	II.1	1623_1624delinsTT	Cyprus	Alive	33	Resistant	<i>M. avium</i> <i>M. triplex</i>	–	<i>S. enteritidis</i>
	II.2			Deceased	8	nv	<i>M. avium</i>	–	–
	II.3			Alive	27	Resistant	–	–	<i>S. enteritidis</i>
5	II.3	783+1G>A	Turkey	Alive	15	D	–	–	–
	II.4			Alive	12	nv	–	–	–
6	II.2	783+1G>A	Turkey	Alive	14	D	–	–	<i>S. enteritidis</i>
	II.3			Alive	9	D	–	–	–
7	II.5	R173P	Turkey	Alive	14	Resistant	<i>M. chelonae</i>	–	–
8	II.2	R173P	Turkey	Alive	9	D	–	–	<i>S. enteritidis</i>
9	II.3	557_563delins8	Turkey	Alive	12	D	–	–	<i>S. enteritidis</i>
10	II.2	700+362-1619-944del	Israel	Alive	4	nv	–	–	<i>S. group D</i>
11	II.2	1190-1G>A	Saudi Arabia	Alive	3	D	–	–	–
12	II.2	C186S	Qatar	Alive	7	D	–	–	<i>S. enteritidis</i>
13	II.3	C186S	Qatar	Alive	7	Resistant	–	–	<i>S. spp</i>
	II.4			Alive	4	Resistant	–	–	<i>S. typhimurium</i>
14	II.2	1791+2T>G	Iran	Alive	8	nv	–	–	<i>S. enteritidis</i>
15	II.2	S321X	Pakistan	Alive	19	Resistant	–	–	<i>S. enteritidis</i>
16	II.1	1791+2T>G	Sri-Lanka	Alive	18	D	–	–	–
17	II.1	[Q32X+1623_1624delinsTT]	France	Alive	18	Resistant	–	–	–
	II.2			Alive	13	nv	<i>M. genavense</i>	–	<i>S. enteritidis</i>
18	II.1	Q376X	France	Alive	30	Resistant	–	–	<i>S. dublin</i>
19	II.1	[1745_1746insCA+1483+182-1619-1073del]	France	Alive	31	D	–	–	<i>S. dublin</i>
20	II.1	Q32X	France	Alive	6	D	–	–	–
21	II.1	Q32X	Belgium	Alive	16	nv	–	–	–
	II.2			Deceased	7	nv	<i>M. avium</i>	–	–
22	II.1	1623_1624delinsTT	Germany	Deceased	3.5	nv	<i>M. avium</i>	–	–
23	II.1	1623_1624delinsTT	Germany	Alive	10	D	–	–	<i>S. enteritidis</i>
	II.2			Alive	7	nv	–	–	–
24	II.1	1791+2T>G	Spain	Alive	16	nv	–	–	–
	II.2			Alive	14	nv	–	Lungs	<i>S. enteritidis</i>
	II.3			Alive	7	nv	–	D	–
25	II.1	1791+2T>G	Spain	Alive	4	nv	<i>M. avium</i>	–	<i>S. enteritidis</i>
26	II.1	549+2T>C	Bosnia-Herzegovina	Deceased	4	D	–	–	–
	II.4			Alive	6	nv	<i>M. spp</i>	–	–
27	II.2	[1440_1447delins16+Q171P]	Slovakia	Deceased	2	D	–	–	–
28	II.3	[1007_1008delinsG+Q171P]	Slovakia	Alive	3	D	–	–	–
29	II.1	L77P	Brazil	Alive	24	D	–	–	<i>S. typhimurium</i>

<sup>a</sup>Children 3.II.2, 5.II.1, 6.II.1, 7.II.1 and 9.II.2 (Fig. 1) died during their first year of life of unknown causes; they were not available for genetic analysis (no DNA available), and thus were not included in the present series of patients as their phenotype and genotype were unclear. In contrast, two patients with the known phenotype of BCG and/or EM infection were included despite their genotype not being known (4.II.2 and 26.II.1).

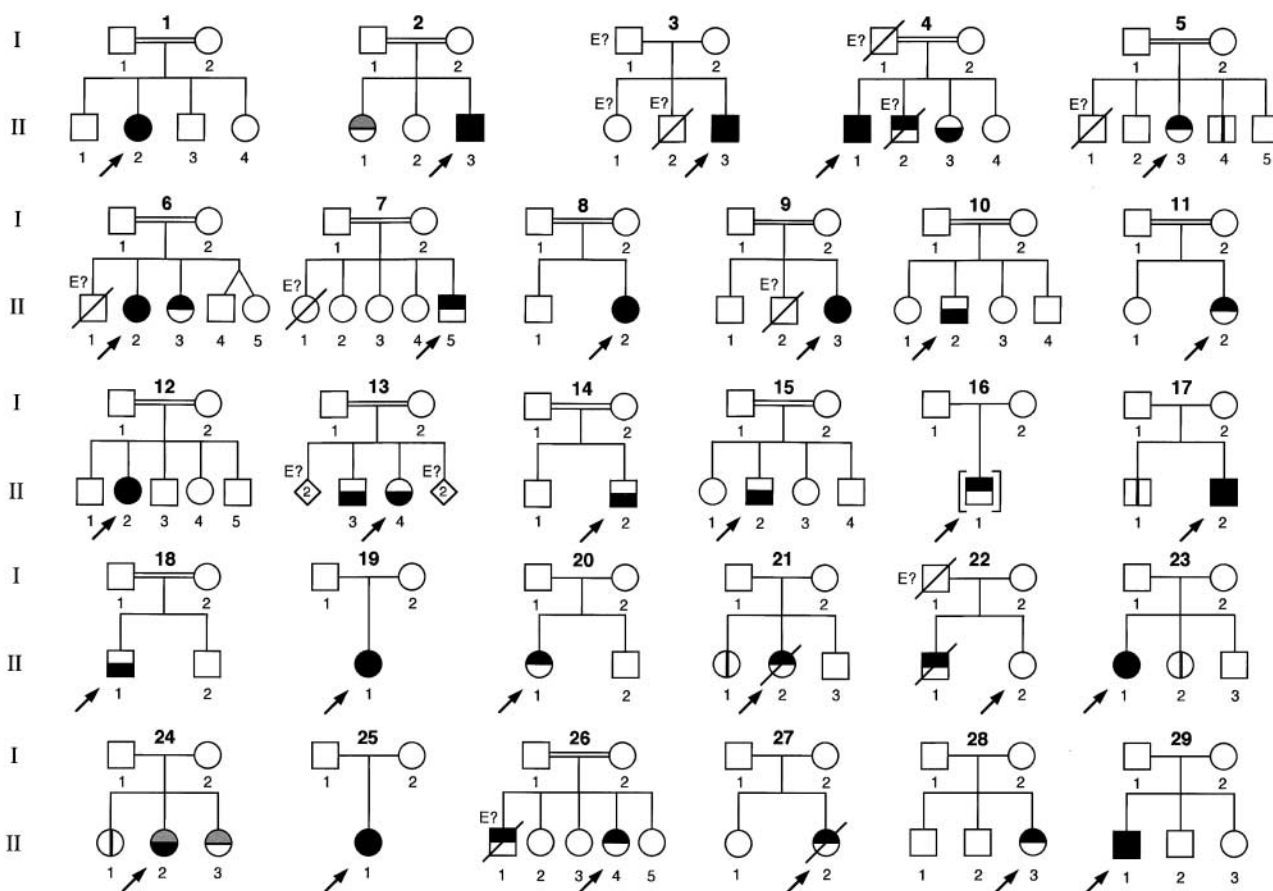
<sup>b</sup>Mutation nomenclature follows the recommendations of Antonarakis and den Dunnen (reference 52). Mutation Q214R initially found in patient 4.II.1 (16), was subsequently found to be a polymorphism. “Delins” indicates mutations combining the deletion and insertion of nucleotides.

<sup>c</sup>The clinical features of kindreds 1, 4, and 5 were initially reported in 1998 (reference 16), kindred 2 in 2001 (reference 20), kindred 4 in 1995 (reference 3) and 2000 (reference 18), kindred 5 in 1988 (reference 53), kindred 7 in 2001 (reference 19), and kindred 21 in 1997 (reference 54). The countries of residence in some cases differ from the countries of origin: kindred 1 lives in France, kindreds 4 and 15 live in the United Kingdom, kindreds 5 and 6 living in Sweden and originating from Turkey are of the Kurd ethnic group, kindred 9 originating from Turkey lives in Germany, kindred 10 originating from and living in Israel is of the Bedouin ethnic group, kindred 14 lives in USA, kindred 16 lives in Switzerland.

<sup>d</sup>Deceased or alive.

<sup>e</sup>Age at death or at the time of writing this report.

<sup>f</sup>BCG, Bacille Calmette-Guérin; D, Disseminated; EM, Environmental *Mycobacteria*; Mtb, *Mycobacterium tuberculosis*; nv, not vaccinated with BCG; Resistant, no adverse reaction to BCG vaccination; *M. spp*, patient 26.II.4 was not BCG-vaccinated and suffered from chronic disseminated granulomatous disease that responded to empirical antimycobacterial treatment (rifampicin, isoniazid, ethambutol, amikacin); no mycobacterial species were visible or cultured, suggesting atypical mycobacteriosis. Patient 10.II.2 also had *Kingella kingae* infections, and patient 29.II.1 *Paracoccidioides brasiliensis* infections. Patient 4.II.1, previously reported not to have been vaccinated (references 3, 16, and 18), had in fact been inoculated with BCG at 5 yr of age. Sera from 26, 25, and 22 patients were tested for Ab against non-typhoid *Salmonella*, *Toxoplasma gondii*, and cytomegalovirus, respectively. The detailed clinical features of all patients will be reported elsewhere.

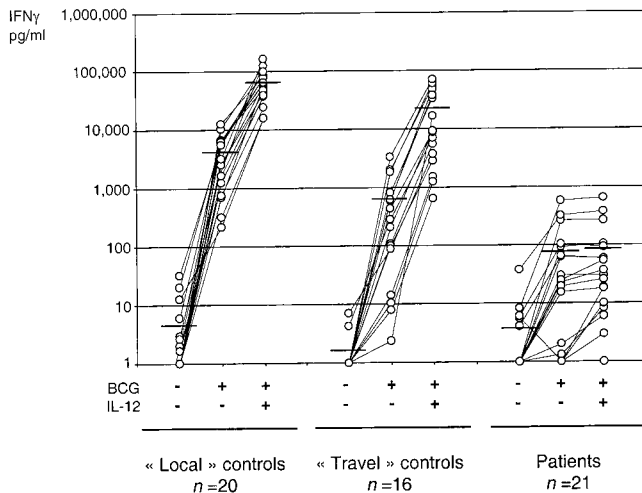


**Figure 1.** Pedigrees of 29 kindreds with IL-12 receptor  $\beta 1$  deficiency. Each kindred is designated by a capital letter (1–29), each generation by a roman numeral (I–II), and each individual by an Arabic numeral (from left to right). Symbols are partitioned in two parts by a horizontal line: the upper part indicates infections with *Mycobacteria* (in black, patients with BCGosis or atypical mycobacteriosis; in gray, patients with tuberculosis); the lower part indicates infections with *Salmonella* (in black, nontyphoid salmonellosis). The proband are indicated by an arrow. Individuals whose genetic status could not be evaluated are indicated by the symbol “E?”. Asymptomatic individuals carrying two mutant *IL12RB1* alleles are represented by a vertical line.

ages of 17 and 10 yr, respectively. One of these two patients (4.II.1), unlike the others, suffered a second EM disease, caused by *M. triplex*. Finally, disseminated nontyphoid salmonellosis occurred in 21 of 34 patients. Salmonellosis was the only infectious disease for seven patients. First *Salmonella* infection occurred at or before the age of 12 yr. Eight patients had recurrent salmonellosis (8/21, 38%), involving the same serotype, whereas BCG and EM infections were not recurrent. Nevertheless, we found high titers of *Salmonella*-specific antibodies in the sera of 4 deficient patients with no history of clinical salmonellosis. Only five of the 34 infected IL-12R $\beta 1$ -deficient patients died due to BCG ( $n = 2$ ) or EM ( $n = 3$ ; a mortality rate among symptomatic patients of only 15% [5/34]) (Fig. 3 b). All deaths occurred before 8 yr of age. No death was attributable to salmonellosis. Thus, infections caused by weakly virulent *Mycobacteria* and *Salmonella* in IL12R $\beta 1$ -deficient patients generally have a childhood onset and a favorable clinical outcome, particularly once patients have entered their teens.

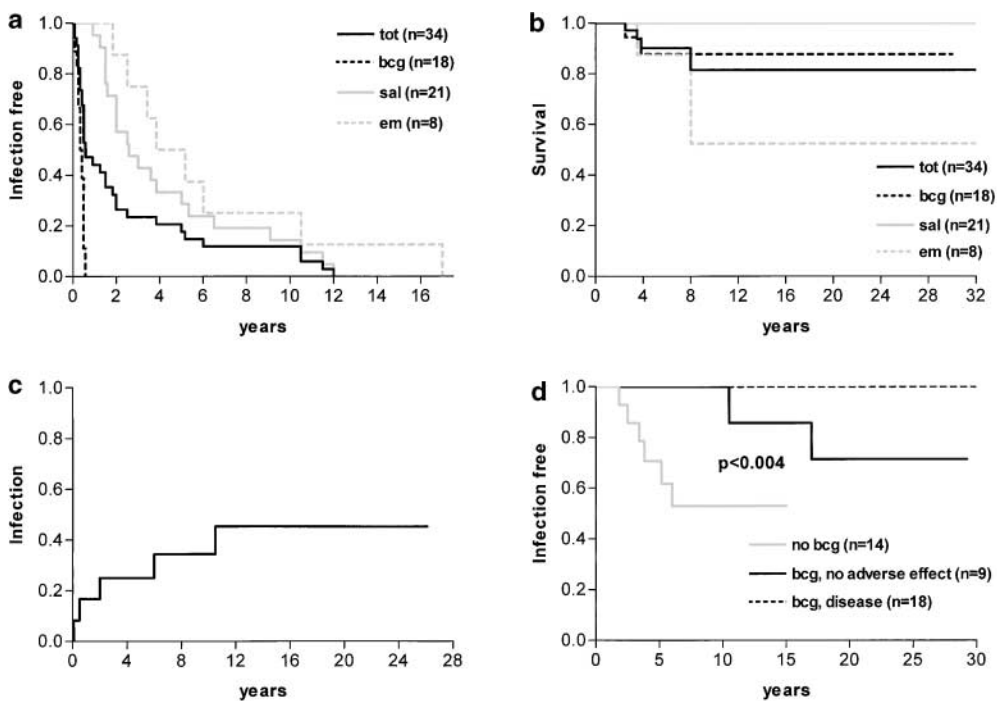
**Clinical Penetrance and Impact of BCG on Clinical Phenotype.** We determined the clinical penetrance of IL12R $\beta 1$  deficiency, by excluding all probands (Fig. 3 c). Five (42%)

of the 12 affected siblings were completely free of unusual infections at their last follow-up visit (mean duration of follow-up: 13.8 yr, range: 7–18 yr). None of the other 7 siblings had more than one infection. Two of the six siblings who had been vaccinated had disseminated BCG disease. Three additional siblings developed infections with weakly virulent microorganisms, EM in one case and *Salmonella* in the other two. Overall, the penetrance of opportunistic infections was estimated to be 45.3% (confidence interval [CI] 95%: 14.2%–75.4%) when calculated with survival analysis techniques to account for differences in follow-up period (Fig. 3 c). Finally, tuberculosis occurred in 2 other siblings (2.II.1, 24.II.3). We then determined the impact of BCG on the clinical phenotype for the 41 patients described herein. BCG inoculation or disease had no impact on the occurrence (and age at onset) of *Salmonella* infection, or the occurrence of tuberculosis. In contrast, none of the 18 patients with BCG disease developed EM disease (interval between onset of BCG disease and last follow-up: 3–31 yr, mean 12.8 yr). Only two (4.II.1 and 7.II.5) of the nine patients resistant to BCG (BCG inoculation without BCG disease) suffered EM disease, with late onset of the



**Figure 2.** Impaired cellular response to interleukin-12. Production of IFN- $\gamma$  by whole blood cells from 20 healthy “local” positive controls (fresh blood), from 16 healthy “travel” positive controls and from 21 patients, either unstimulated (–) or stimulated with BCG alone or with BCG plus recombinant IL-12p70. Fresh, heparinized blood from 7 patients and all “travel” controls were shipped to Paris within 12 to 48 h, where the experiments were performed. Blood from four patients and all “local” control was drawn in Paris and readily stimulated. The supernatants were harvested after 48 h of activation for cytokine quantification by ELISA. The horizontal bars represent the arithmetic mean of the values.

disease. Six (43%) of the 14 patients who had not been inoculated with BCG suffered EM disease (Fig. 3 d), with early onset of the disease. The difference in age at onset of EM disease between the three groups of patients was highly significant ( $P < 0.004$ ; Fig. 3 d). This difference was particularly clear if patients with BCG disease were compared with patients not inoculated with BCG ( $P < 0.004$ ). The



**Figure 3.** Epidemiological features of IL-12R $\beta$ 1 deficiency. First onset (a) and outcome (b) of case-definition opportunistic infectious diseases in 34 deficient patients, according to infections: BCG (broken black line), EM (broken gray line), *Salmonella* (solid gray line), and all 3 infections (solid black line). (c) Penetration of case-definition infectious diseases in the 12 IL-12R $\beta$ 1-deficient siblings (excluding all probands). (d) Variations in onset of EM disease among the 41 deficient patients, who had been vaccinated with BCG and suffered BCG disease (resistance to BCG, solid black line,  $n = 9$ ), or who had not been vaccinated with BCG (solid gray line,  $n = 14$ ).

difference between patients resistant to BCG and nonvaccinated patients was not significant ( $P = 0.09$ ). Finally, the difference in incidence of EM disease between BCG-inoculated patients (with or without BCG disease) and nonvaccinated patients was highly significant ( $P < 0.002$ ).

## Discussion

We report here the first large series of patients with complete IL-12R $\beta$ 1 deficiency, including 41 deficient individuals from 29 unrelated kindreds (Table I and Fig. 1). The kindreds originate from 17 countries, in Africa, America, Asia, and Europe. Seven other unrelated patients reported elsewhere (15, 17, 18, 21) raise the total number of cases to 48, from 36 kindreds, in 20 countries. The cellular phenotype of IL-12-receptor  $\beta$ 1 chain-deficient patients is uniform: all patients tested failed to respond to IL-12. The relative contributions to the clinical phenotype of the lack of response of NK cells (16) and of T lymphocytes (15–18, 21) to IL-12 are unclear. The receptor for IL-23, another IFN- $\gamma$ -inducing cytokine, contains the IL-12R $\beta$ 1 subunit (25), suggesting that patients with IL-12R $\beta$ 1 deficiency probably also suffer from IL-23 receptor deficiency.

The results obtained in this study confirm that IL-12R $\beta$ 1 deficiency is principally associated with mycobacterial diseases (Table I). If we include the seven patients reported elsewhere (15, 17, 18, 21), 33 of the 48 patients have had BCG ( $n = 22$ ) or EM ( $n = 11$ ) disease. We also report three patients from two kindreds with *M. tuberculosis* disease. Thus, a diagnosis of IL-12 receptor deficiency should be considered in patients with unusually severe forms of tuberculosis (20). Mice with a disrupted *IL12B* gene (26) are susceptible to BCG (27), *M. tuberculosis* (27–

29) and, to a lesser extent, to *M. avium* (30–32). Our results extend these observations, and highlight the crucial role of IL-12 in immunity to BCG, several EM and *M. tuberculosis* in humans.

However, IL-12R $\beta$ 1-deficient patients are less susceptible to *Mycobacteria* than was initially thought (15–21). 9 of 27 children inoculated with BCG did not suffer adverse effects. Only 8 of the 41 patients (19%) in this series, and 11 of the 48 (23%) including the cases reported elsewhere (17, 18, 21), developed EM infections. Excluding the probands, only 2 of 6 siblings vaccinated with BCG and 1 of 12 IL-12R $\beta$ 1-deficient siblings developed BCG and EM disease, respectively. Paradoxically, these data indicate that most IL-12R $\beta$ 1-deficient patients do not have the original phenotype of BCG/EM disease or salmonellosis. Healthy siblings of probands should therefore be investigated. IL-12 is surprisingly redundant in protective immunity against primary infection by weakly virulent *Mycobacteria* in a majority of individuals.

In our survey, the age at onset of EM disease differed between patients with and without BCG disease. If we include the other IL-12R $\beta$ 1- and IL-12p40-deficient cases (13–15, 17, 18, 21), there are 63 patients in total, 36 of whom had BCG disease. None of these 36 patients developed EM disease, while among the 27 patients who did not develop any BCG disease 12 developed EM disease ( $P < 10^{-6}$ , Fisher's exact test). This indicates that BCG disease protects against subsequent EM illness in patients with IL-12 deficiency or IL-12R deficiency. Antibiotic treatment against BCG probably played a minor role in preventing EM disease, as it was discontinued after a few years in all patients. Moreover, the impact of BCG inoculation itself (with or without BCG disease) was also significant in our series. Similarly, EM disease was protective in itself. In contrast, BCG inoculation and EM disease had no effect on the incidence of salmonellosis. IL-12 is thus redundant for adaptive, secondary immunity to *Mycobacteria* in most, if not all, patients.

The prognosis of IL-12R $\beta$ 1 deficiency is good due to the low clinical penetrance of primary mycobacterial infection, acquired protective immunity to secondary mycobacterial infection, and the lack of life-threatening nonmycobacterial infections. Only 5 of our 41 (12%) IL-12R $\beta$ 1-deficient patients died of BCG or EM infection, before the age of 8 yr. The overall mortality rate for all known IL-12R $\beta$ 1-deficient patients is only 15% (7/48), increasing to 21% (13/63) if we include IL-12p40-deficient children (13–15, 17, 18, 21). Outcome improved with age, and our patients (now aged 2–33 yr) are healthy and off all treatment. These results should help us to predict clinical outcome for individual patients. They also indicate that immune mechanisms progressively develop ways of compensating for the lack of IL-12-mediated immunity.

21 of the 41 patients (51%) in our series had had extra-intestinal infections caused by *Salmonella*, as had 4 of the other 7 IL-12R $\beta$ 1-deficient patients (15, 17, 18, 21) and 5 of the 15 IL-12p40-deficient patients (a total of 30 out of 63, 48%; references 13–15). Of clinical interest, five chil-

dren presented salmonellosis as the only infectious disease. IL-12 has been shown to play a major role in host defense against *Salmonella* in IL-12p40-KO mice (33, 34). Our observation extends this findings in humans. However, not all *Salmonella*-infected patients developed clinical disease (Table I). IL-12 was redundant in primary immunity against *Salmonella* in some patients. Salmonellosis recurrence was frequent (8/21, 38%), suggesting that IL-12 is required for secondary immunity to *Salmonella* in many patients. These data suggest that different immune mechanisms control secondary immunity to *Salmonella* and *Mycobacteria*.

There were no other unusually severe infections in our 41 patients, as in the other seven patients (15, 17, 18, 21; Table I). CMV infection had been well controlled in at least 14 patients (Table I). Other common viruses did not cause severe disease in our patients. One patient suffered benign infection with *Kingella kingae*. There were no other diseases caused by common bacteria, including macrophage-infecting species such as *Listeria* and *Nocardia*. One IL-12-deficient patient had one episode of nocardiosis (14). *Paracoccidioides brasiliensis* infection in one curable IL-12R $\beta$ 1-deficient patient had a symptomatic yet benign course, and oral candidiasis was diagnosed in a few patients. Other fungi represented no threat to IL-12R $\beta$ 1-deficient patients. *Toxoplasma gondii* infection had an uneventful course in at least five patients. Other common opportunistic protozoa were not pathogenic.

The number of kindreds investigated from various genetic backgrounds and exposed to a variety of environmental conditions, together with the number of siblings affected, ruling out a major ascertainment bias, make it possible to conclude that mycobacteriosis and salmonellosis are the most frequent infectious diseases in patients with IL-12R $\beta$ 1 deficiency. Our study does not exclude the possibility that other infectious or non-infectious diseases may occur in other patients with this genetic defect, or in our patients later in life. The apparent resistance of IL-12R $\beta$ 1-deficient patients to most microorganisms nevertheless contrasts with results obtained in mice. Indeed, IL12p40 KO mice are susceptible to almost all microorganisms tested (35–51). The lack of a requirement for IL-12 and probably IL-23 for protective immunity against most microorganisms in humans may reflect the natural (1), as opposed to experimental, course of most infections in man.

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