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Defective lymphoid organogenesis underlies the immune deficiency caused by a heterozygous S32I mutation in IκBα

Jana L. Mooster,1,3 Severine Le Bras,1,3 Michel J. Massaad,1,3 Haifa Jabara,1,3 Juhan Yoon,1,3 Claire Galand,1,3 Balthasar A. Heesters,2,6 Oliver T. Burton,1,3 Hamid Mattoo,7 John Manis,4,5 and Raif S. Geha1,3

1Division of Allergy and Immunology and 2Program in Cellular and Molecular Medicine, Boston Children’s Hospital, Boston, MA 02115
2Department of Pediatrics, 3Division of Transfusion Medicine, and 4Department of Pathology, Harvard Medical School, Boston, MA 02115
5Department of Medical Microbiology, University Medical Center Utrecht, 3584 CX Utrecht, Netherlands
6Department of Cancer Research, Massachusetts General Hospital, Boston, MA 02114
7Center for Cancer Research, Massachusetts General Hospital, Boston, MA 02114

Patients with ectodermal dysplasia with immunodeficiency (ED-ID) caused by mutations in the inhibitor of NF-κB α (IκBα) are susceptible to severe recurrent infections, despite normal T and B cell numbers and intact in vitro lymphocyte function. Moreover, the outcome of hematopoietic stem cell transplantation (HSCT) in these patients is poor despite good engraftment. Mice heterozygous for the IκBα S32I mutation found in patients exhibited typical features of ED-ID. Strikingly, the mice lacked lymph nodes, Peyer’s patches, splenic marginal zones, and follicular dendritic cells and failed to develop contact hypersensitivity (CHS) or form germinal centers (GCs), all features not previously recognized in patients and typical of defective noncanonical NF-κB signaling. Lymphotixin β receptor (LTβR)–driven induction of chemokines and adhesion molecules mediated by both canonical and noncanonical NF-κB pathways was impaired, and levels of p100 were markedly diminished in the mutant. IκBα mutant →Rag2−/−, but not WT→IκBα mutant, bone marrow chimeras formed proper lymphoid organs and developed CHS and GCs. Defective architectural cell function explains the immunodeficiency and poor outcome of HSCT in patients with IκBα deficiency and suggests that correction of this niche is critical for reconstituting their immune function.

Abbreviations used: AD, autosomal dominant; BAFF, B cell activating factor; CBC, complete blood count; CHS, contact hypersensitivity; EDA, ectodysplasin A; ED-ID, ectodermal dysplasia with immunodeficiency; ES, embryonic stem; FDC, follicular DC; GC, germinal center; HSCT, hematopoietic stem cell transplantation; IκB, inhibitor of NF-κB; IKK, IκB kinase; IκK, IκB essential modulator; NIK, NF-κB–inducing kinase; OXA, oxazolone; P, Peyer’s patch; PNA, peanut agglutinin; qPCR, quantitative PCR; T, T dependent; T, T independent.

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polyubiquitination and proteasome processing into p52, allowing the p52:RelB dimer to translocate into the nucleus and activate the transcription of genes involved in lymphoid organogenesis (Dejardin et al., 2002; Seymour et al., 2006). There is cross talk between the canonical and noncanonical NF-κB pathways. NIK-dependent activation of IKKα can cause IκBα phosphorylation and degradation, p50:p65 regulates the expression of p100 and RelB, and p100 regulates RelA/p65-containing complexes (Matsushima et al., 2001; Tücker et al., 2007; Shihi et al., 2011; Chen et al., 2013).

Ectodermal dysplasia with immunodeficiency (ED-ID) is characterized by sparse hair, conical teeth, reduced number of sweat glands, and susceptibility to severe infections (Kere et al., 1996; Srivastava et al., 2001; Orange and Geha, 2003; Picard et al., 2011). The X-linked form of ED-ID is caused by hypomorphic mutations in NEMO (Puel et al., 2004; Hanson et al., 2008). Autosomal-dominant (AD) ED-ID is caused by heterozygous hypermutmorphic mutations in the IκBα gene NFKBIA (Courtois et al., 2003; Kawai et al., 2012). Six mutations in IκBα, S32I, W11X, E14X, Q9X, M37X, and S36Y, have been identified in AD ED-ID (Courtois et al., 2003; Jansen et al., 2004; McDonald et al., 2007; Lopez-Granados et al., 2008; Ohnishi et al., 2012; Schimke et al., 2013). In each case, the mutation impairs phosphorylation-driven degradation of the mutant protein, resulting in the sequestration of NF-κB in the cytoplasm (Courtois et al., 2003; McDonald et al., 2007; Kawai et al., 2012). In both forms of ED-ID, activation of the canonical NF-κB pathway is impaired, resulting in ED caused by defective signaling downstream of the IL-1 receptor, impaired TLR responses, and decreased in vitro B cell response to CD40 ligation (Orange et al., 2005). The severity of the disease correlates with the degree of NF-κB impairment (Orange and Geha, 2003).

Two aspects of the disease phenotype of patients affected by IκBα deficiency have long been a puzzle. The patients suffer from severe, recurrent, and potentially fatal infections despite having normal or elevated T and B cell numbers and intact in vitro T cell function (Courtois et al., 2003; Jansen et al., 2004; McDonald et al., 2007; Kawai et al., 2012). The outcome of hematopoietic stem cell transplantation (HSCT) in these patients is poor in spite of good engraftment of donor lymphoid cells. Of three patients treated with HSCT, only one with the S32I IκBα mutation has survived, but continues to suffer from recurrent infections despite excellent donor lymphoid cell engraftment (Dupuis-Girod et al., 2006; Cancrini, C., personal communication). We have created an IκBα S32I knock-in mouse model of AD ED-ID to gain insights into the disease. The IκBα mutant mouse recapitulates many of the ectodermal and immune abnormalities found in patients with ED-ID. Strikingly, the mutant completely lacked LNs and Peyer’s patches (PPs), and its spleen lacked follicles, marginal zones (MZs), MZ B cells, and follicular DCs (FDCs) and failed to form germinal centers (GCs), all features not previously recognized in patients with ED-ID and typical of defective noncanonical NF-κB signaling. The levels of p100 and noncanonical NF-κB signaling in response to LTβR ligation were decreased in the IκBα mutant. Analysis of BM radiation chimeras demonstrated that the defective lymphoid organogenesis in the IκBα mutant is caused by a defect in nonhematopoietic cells, thus explaining the poor outcome of HSCT in patients with IκBα deficiency.

RESULTS
Mice heterozygous for the S32I mutation in IκBα have ED and impaired IκBα phosphorylation and degradation

The strategy for the generation and identification of the heterozygous IκBα S32I mutant (IκBα mutant) mice is shown in Fig. S1. IκBα mutant mice were born at the normal Mendelian ratio but were significantly smaller in size and weight than their WT littermates (Fig. 1, A and B) and had a 50% survival rate at 8 wk compared with 100% for WT littermates (Fig. 1 C). IκBα mutant mice are missing their third molars, lack guard hairs, and have hypoplastic eccrine glands (Fig. 1, D–F), a phenotype observed in mice with disruption of the Eda gene, mutated in patients with X-linked anhidrotic ED (Srivastava et al., 2001).

Immunoblotting cannot distinguish between WT IκBα and the S32I mutant proteins. We sought evidence for the expression of the mutant protein in heterozygous IκBα mutant mice by examining the susceptibility of IκBα to phosphorylation and degradation after stimulation of fibroblasts with IL-1β. IκBα phosphorylation was significantly weaker in fibroblasts from mutant mice compared with WT littermates (Fig. 1 G). IκBα was mostly degraded by 15 min and completely degraded by 30 min in WT fibroblasts. In contrast, there was markedly less IκBα degradation in the mutant fibroblasts. Similar results were obtained when the fibroblasts were stimulated with TNF and LPS, two other well-known activators of the canonical NF-κB pathway (not depicted).

BMDCs were differentiated from BM cells with GM-CSF and IL-4 and used to examine the response to TLR ligation in a homogeneous population of cells. BMDCs from IκBα mutant mice secreted significantly less TNF in response to TLR1/2, TLR4, TLR7, and TLR9 ligands (Fig. 1 H). Furthermore, up-regulation of Vcam1 (Vascular adhesion molecule 1) and Icam1 (Intercellular adhesion molecule 1) gene expression and of VCAM1 surface expression after TNF stimulation, which are dependent on canonical NF-κB signaling (Dejardin et al., 2002; Winning et al., 2010), was deficient in MEFs from IκBα mutant mice (Fig. 1, I and J). Collectively, these results indicate that the IκBα mutant mice we generated represent a faithful model of ED-ID caused by heterozygous IκBα S32I mutation.

Defective secondary lymphoid organogenesis in IκBα mutant mice

The gross appearance, weight, and cellularity of the thymus and the distribution of thymocyte subsets into CD4+CD8+ double-negative cells, CD4+CD8+ double-positive cells, and CD4+ or CD8+ single-positive cells were comparable in IκBα mutant mice and WT littermates (not depicted). BM cellularity and distribution of (B220+IgM–CD43+) pre-B, (B220+IgM–CD43+) pre-B, (B220+IgM–) immature, and
Figure 1. **IκBα** mutant mice have ED, impaired IκBα processing, and deficient TLR response. (A) IκBα mutant mouse and WT littermate photographed at 3 wk of age. Data are representative of >20 mice per group. (B and C) Growth (B) and Kaplan-Meier survival (C) curves of IκBα mutant mice and WT littermates weighed every 3–4 d and observed daily. Data were derived from 34 mutant mice and 19 WT littermates weighed. (D–F) Photographs of mandibular bones (D) and fur (E) and H&E staining of footpad sections (F) in 6 wk-old IκBα mutant mouse and WT littermate. Red arrows in D show the missing third molars in the mutant mice. Data are representative of four or more mice per group in three independent experiments. Bar, 100 μm. (G) Immunoblot of fibroblast lysates using antibodies to phospho-IκBα or IκBα. Actin was used as a loading control. Pooled results of IκBα phosphorylation at 5 min using three mice per group in three independent experiments. (H) TNFα as measured by ELISA in supernatants of BMDCs stimulated with the TLR ligands Pam3Csk4 (TLR1/2), LPS (TLR4),loxoribine (TLR7), and CpG oligonucleotide (TLR9). Data are representative of four mice per group in two independent experiments for Pam3Csk4 and LPS and four mice per group in three independent experiments for loxoribine and CpG. (I and J) qPCR analysis of Vcam1 and Icam1 mRNA (I) and flow cytometric analysis of VCAM1 surface expression (J) after TNF stimulation in MEFs from IκBα mutant mice and WT controls. Results in I are expressed relative to unstimulated WT MEFs. Data are representative of three mice per group in three independent experiments. Circles and columns represent means, and bars represent SD in B, C, and G–I. *, P < 0.05; **, P < 0.01; ***, P < 0.001.
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(B220+IgM+) mature B cell subsets were comparable in IkBα mutant mice and WT littermates (not depicted).

Gross examination revealed no visible cervical, axillary, inguinal, or mesenteric LNs in >15 IkBα mutant mice examined (Fig. 2A and not depicted). No LN tissue could be detected in hematoxylin and eosin (H&E)-stained serial sections of the inguinal fat pad in IkBα mutant mice (not depicted). To confirm the absence of LNs, we injected mice in the footpad with Evan’s blue dye. Blue-colored lymphatic vessels were readily apparent in both mutant and WT mice. Blue-colored LNs could be detected in the popliteal, inguinal, and paraaortic area in WT mice, but were absent in the mutants (Fig. 2B and not depicted). Examination of H&E-stained sections of the small intestine revealed no detectable PPs in seven WT littermates examined (Fig. 2C and not depicted). The numbers of circulating lymphocytes were significantly increased in the IkBα mutant compared with 7.1 ± 2.3 PPs in seven WT littermates (Fig. 2D). The number of splenic T cells and B cells was greater in IkBα mutant mice than WT littermates, but the difference was not statistically significant (Fig. 3B). The number of splenic T cells and the distribution of CD4+ and CD8+ subsets were comparable between the IkBα mutant mice and WT littermates (Fig. 3C). Analysis of B cell subsets revealed comparable percentages of B220+CD93+ transitional B cells and B220+CD23+CD21+ follicular cells in the two groups, but drastically reduced percentages of B220+CD23−CD21+ MZ B cells in mutant spleens (Fig. 3D and F). The percentages of MZ B cells in the blood were comparable in the mutant and WT littermates (12.1 ± 1.0% in the mutant vs. 10.6 ± 1.8% in WT littermates; n = 3). Given the severe decrease in splenic MZ B cells in the mutant, this result suggests a retention defect but does not rule out a defect in development of MZ cells. IkBα mutant mice had increased numbers of CD11b+F4/80+ macrophages but normal numbers of CD11b+Ly6G+ granulocytes and CD3−NK1.1+ cells and CD11c+ DCs compared with WT littermates (not depicted). The percentages of CD11c+MHCII+ B220+ plasmacytoid DCs and CD11c+MHCII+CD11b+CD4+ and CD11c+MHCII+CD11b+CD8+CD4− myeloid DCs were comparable between the two groups, whereas the percentage of CD11c+MHCII+CD11b−CD8+ lymphoid DCs was significantly reduced in the mutant (9.3 ± 0.9% in the mutant vs. 14.8 ± 1.0% in WT littermates; n = 3; P < 0.05).

IkBα mutant mice had a severely disorganized splenic architecture. H&E staining revealed virtual absence of follicles (Fig. 3G). Immunofluorescence analysis revealed that MOMA+ macrophages were present in IkBα mutant mice but failed to form the ring structure typical of a MZ (Fig. 3H). The number of Macd1+ cells, which represent the endothelial cells that line the marginal sinus, was decreased, and the organization of these cells was disrupted in the IkBα mutant (Fig. 3I). FDCM1+ FDCs were absent from the spleens of the mutant (Fig. 3J).

Figure 2. IkBα mutant mice lack LNs and PPs. (A and B) Photograph of mesenteric area (A) and popliteal LNs (B) in an IkBα mutant mouse and WT littermate. Mice in B were injected with Evan’s blue. The normal position of LNs is outlined by a dotted circle. Arrows indicate the lymphatic vessels. (C and D) Longitudinal sections of the small intestine (C) and transverse sections of the head at the level of the nasal cavities (D) were stained with H&E. Bars, 1,000 µm. Data in A–D are representative of a minimum of three mice per group in three independent experiments.

Normal number of lymphoid tissue inducer (LTI) cells but defective response to LTβR ligation and TNF in the IkBα mutant

Lymphorganogenesis is initiated in the embryo by the interaction of LTi cells with stromal organizer cells, which express LTβR (Rennert et al., 1996; Mebius, 2003; Bénézech et al., 2010). FACS analysis of intestine from embryonic day (E) 17 embryos revealed the presence of higher percentages and numbers of CD45−CD3−CD4−IL-7R+ LTi cells in IkBα mutant mice compared with WT littermates (Fig. 4, A and B), indicating that the IkBα mutation did not interfere with the development of LTi cells.

LTi cells express the LTβR ligand LTα1β2, whereas stromal cells express LTβR. LTβR signaling in stromal cells is essential for lymphoid organogenesis (Fu and Chaplin, 1999). Both the canonical and noncanonical NF-κB pathways are
activated by LTβR and are important for the expression of adhesion molecules and chemokines that drive lymphoid organogenesis, with the noncanonical pathway playing a major role (Miyawaki et al., 1994; Koike et al., 1996; Shinkura et al., 1999; Dejardin et al., 2002; Lo et al., 2006; Vondenhoff et al., 2009). The absence of secondary lymphoid structures in IκBα mutant mice prompted us to examine LTβR signaling in these mice. We stimulated MEFs, as surrogates for stromal cells, with the agonistic LTβR mAb AFH6 and examined the induction of Vcam1 and Icam1 expression, as readouts of canonical NF-κB activation (Dejardin et al., 2002), Cxcl10 (C-X-C motif chemokine 10) expression, as a readout of predominantly canonical NF-κB activation (Hoffmann et al., 2003; Caposio et al., 2007; Shultz et al., 2009), and Madcam1.
expression, as a readout of noncanonical NF-κB pathway activation downstream of LTβR (Ganeff et al., 2011). Expression of all these genes after stimulation with LTβR mAb AFH6 was significantly reduced in MEFs from IkBα mutant mice compared with WT controls (Fig. 4 C and not depicted).

Induction of the chemokine genes Ccl19 (C-C motif chemokine 19), Cxcl12, and Cxcl13 after LTβR signaling is important for lymphoid organogenesis and is dependent on the noncanonical NF-κB pathway (van de Pavert et al., 2009; van de Pavert and Mebius, 2010). Because LTβR failed to up-regulate the expression of these genes in WT MEFs (not depicted), we examined their baseline expression in the spleen. Expression of all three chemokine genes was significantly reduced in spleens of IkBα mutant mice compared with WT littermates (Fig. 4 D).

Expression in cell lines of the genes that encode for the noncanonical NF-κB family members p100 and RelB have been reported to be regulated by the canonical NF-κB pathway (Liptay et al., 1994; Bren et al., 2001). The level of p100, but not RelB, was significantly reduced in B cells from IkBα mutant mice (Fig. 4, E and F). NIK levels were comparable in B cells from IkBα mutant mice and WT controls (not depicted). These findings suggest that the S32I IkBα mutation impairs signaling via the noncanonical NF-κB pathway because it results in diminished expression of p100.
Normal intrinsic T cell function but impaired delayed contact hypersensitivity (CHS) in IκBα mutant mice

T cell proliferation upon stimulation with immobilized anti-CD3 with and without soluble anti-CD28 and IL-2 secretion in response to immobilized anti-CD3 were comparable in mutant mice and WT littermates (Fig. 5, A and B). CD28 ligation increased IL-2 secretion by the mutant T cells, albeit to a significantly lower extent than in WT T cells (Fig. 5 B).
The differentiation of naive CD4$^+$ T cells in vitro under polarizing conditions into Th2 cells was comparable in the mutant and WT littermates, but differentiation into Th1 and Th17 cells was impaired in the mutant (Fig. 5 C). These findings indicate that TCR-driven T cell proliferation, IL-2 secretion, and in vitro Th2 cell differentiation were normal in the mutant, but in vitro Th1 and Th17 cell differentiation were significantly impaired.

Figure 6. B cell function in IκBα mutant mice. (A and B) Proliferation determined by $[^{3}H]$thymidine incorporation into DNA (A) and IgM and IgG production determined by ELISA in the supernatants (B) of purified splenic B cells stimulated for 3 d with LPS. Data are representative of a minimum of four mice per group in three experiments. (C) Aicda and Prdm1 mRNA expression in B cells stimulated with 10 µg/ml LPS measured by qPCR. Data are representative of three mice per group in three experiments. (D) Proliferation by purified splenic B cells stimulated with anti-μ, anti-CD40 mAb, LPS, or TGFβ in the presence of IL-4 determined by $[^{3}H]$thymidine incorporation into DNA. Data are representative of a minimum of four mice per group in three experiments. (E) IgA production by purified splenic B cells stimulated with LPS + TGFβ, as determined by ELISA of supernatants. Data are representative of a minimum of four mice per group in four experiments. (F and G) B cell survival mediated by BAFF determined by analysis of trypan blue-negative cells (F) and IgG1 production determined by ELISA in the supernatants (G) by purified naive splenic B cells stimulated with BAFF in the absence or presence of IL-4 for 4 (F) or 7 (G) d. Data are representative of three mice per group in three experiments. (H) Serum Ig levels from unimmunized 6-8-wk-old mice. Data represent the mean of duplicate determinations on sera from the number of mice indicated. (I) Small intestine sections were stained by immunofluorescence for IgA, CD45, and DAPI. The dashed line delineates the edge of the PP. Bars, 100 µm. Data are representative of two independent experiments with one mouse per group. Lines and columns represent mean, and bars represent SD. *, P < 0.05; **, P < 0.01; ***, P < 0.001.
After i.p. immunization with OVA, T cells in splenocyte cultures from \(1k \beta \alpha\) mutant mice proliferated normally and secreted IL-2 robustly in response to OVA stimulation (Fig. 5 D). In contrast to the response to i.p. immunization, which involves interaction in the spleen between antigen-presenting DCs and T cells, the T cell response to cutaneously introduced antigen involves interaction in the skin-draining LN between DCs that have captured antigen in the skin and recirculating antigen-specific T cells. \(1k \beta \alpha\) mutant mice sensitized with the hapten oxazolone (OXA) developed significantly less ear swelling with markedly less cellular infiltration and expressed significantly less \(\text{Ilf} 3\) and II4 mRNA than WT controls (Fig. 5, E and F). Because \(1k \beta \alpha\) mutant mice have partial intrinsic T cell defects, the absence of LNs likely plays a major role in the failure of these mice to mount a CHS response.

**B cell function is modestly decreased in vitro but severely impaired in vivo in \(1k \beta \alpha\) mutant mice**

Proliferation and IgG secretion by purified total splenic B cells in response to stimulation with LPS were modestly, but significantly, reduced in the mutant, whereas IgM secretion was not significantly reduced (Fig. 6, A and B). \(\text{Aicda}\) (activation-induced cytidine deaminase) expression in LPS-stimulated B cells from the mutant was comparable with WT controls, but \(\text{Prlm} 1\) (PR domain zinc finger protein 1) expression was significantly decreased (Fig. 6 C). The proliferation of splenic B cells from \(1k \beta \alpha\) mutant mice in response to stimulation with anti-IgM + IL-4, anti-CD40 + IL-4, LPS + IL-4, and LPS + TGF\(\beta\) was modestly, but significantly, decreased compared with B cells from WT littermates, as measured by \(\text{H}^{3}\)thymidine incorporation (Fig. 6 D) and CFSE dye dilution (not depicted). IgM, but not IgG, production was significantly decreased in response to LPS + IL-4 and anti-CD40 + IL-4 (not depicted), suggesting that IL-4 may overcome some of the hyporesponsiveness of the B cells to LPS alone. LPS + TGF\(\beta\) stimulation caused significantly less IgA production by mutant B cells than by WT B cells (Fig. 6 E). BAFF, which activates primarily the noncanonical NF-\(\kappa\)B pathway (Claudio et al., 2002), promotes B cell survival and drives modest IgG1 isotype switching in naive B cells (Batten et al., 2000; Castighi et al., 2005). The survival of naive B cells in the presence of BAFF and their IgG1 production in response to BAFF alone or with added IL-4 were significantly reduced in \(1k \beta \alpha\) mutant mice compared with WT littermates (Fig. 6, F and G). This is consistent with the impaired noncanonical NF-\(\kappa\)B signaling in the mutant.

Serum IgM levels were comparable in mutant mice and WT littermates, whereas serum IgG levels were significantly decreased, and serum IgA was nearly absent in the mutants (Fig. 6 H). IgA\(^{+}\) plasma cells were absent from the lamina propria of mutant mice (Fig. 6 I).

The antibody responses to the type I T-independent (TI) antigen TNP-LPS, the type II TI antigen TNP-Ficoll, and the T-dependent (TD) antigens OVA and TNP-KLH were all severely diminished in \(1k \beta \alpha\) mutant mice compared with WT controls (Fig. 7, A–C; and not depicted). The IgG antibody response to TD antigens requires GC formation (Rajewsky, 1996). OVA immunization resulted in the robust development of GCs in the spleens of WT mice as indicated by staining for peanut agglutinin (PNA; Fig. 7 D). In contrast, GC development was severely deficient in \(1k \beta \alpha\) mutant mice. OVA immunization caused a significant increase in the percentage of B220\(^{+}\)PNA\(^{+}\) and B220\(^{+}\)Fas\(^{+}\)GL7\(^{+}\) GC B cells in the spleens of WT mice but not \(1k \beta \alpha\) mutant mice (Fig. 7, E and F).

**Reconstitution of \(1k \beta \alpha\) mutant mice with WT BM fails to correct their secondary lymphoid organs and immune defects**

To determine whether the immune phenotype of \(1k \beta \alpha\) mutant mice was intrinsic to lymphocytes, we examined immune function in WT→\(1k \beta \alpha\) mutant chimeras. The same pool of BM cells from CD45.1\(^{+}\) WT mice was used to reconstitute irradiated CD45.2\(^{+}\) \(1k \beta \alpha\) mutant mice and WT littermates. 8 wk after BM transplantation, the two chimeras had comparable numbers of lymphocytes in the blood, with virtually all B cells and >80% of T cells of donor origin (not depicted). Gross examination revealed that LNs and PPs were present in WT→WT control chimeras (Fig. 8, A and B). In contrast, they were not detectable in WT→\(1k \beta \alpha\) mutant chimeras. Splenic cellularity and the percentages of CD3\(^{+}\), CD4\(^{+}\), CD8\(^{+}\), and B220\(^{+}\) cells were comparable in the two chimeras (not depicted). However, follicles and FDCs, which were present in the spleens of control chimeras, were absent from WT→\(1k \beta \alpha\) mutant chimeras (Fig. 8 C), and the percentage of MZ B cells was distinctly decreased in the WT→\(1k \beta \alpha\) mutant chimeras (Fig. 8 D). These findings strongly suggest that nontransferable stromal cells are responsible for the lack of LNs, PPs, splenic follicles, and FDCs and for the decrease in MZ B cells in \(1k \beta \alpha\) mutant mice.

Evaluation of CHS revealed significantly decreased ear swelling, inflammation, and cytokine expression in WT→\(1k \beta \alpha\) mutant chimeras compared with control chimeras (Fig. 8, E–G). WT→\(1k \beta \alpha\) mutant chimeras had impaired antibody responses to TNP-Ficoll and OVA compared with control chimeras (Fig. 8, H and I). OVA immunization resulted in the development of GCs and a significant increase in the percentage of B220\(^{+}\)PNA\(^{+}\) GC B cells in the spleens of control chimeras, but not WT→\(1k \beta \alpha\) mutant chimeras (Fig. 8, J and K). The failure of reconstitution with WT BM cells to correct the immune defects in \(1k \beta \alpha\) mutant mice indicates that nonhematopoietic cells are major contributors to the immune deficiency in \(1k \beta \alpha\) mutant mice.

**Secondary lymphoid organs, CHS, and GC formation, but not B cell function, are restored in \(1k \beta \alpha\) mutant→\(Rag^{2/−}\) chimeras**

\(Rag^{2/−}\) mice possess LTi cells, and the cross talk between LTi cells and stromal cells is intact in these mice (Fu and Chaplin, 1999). To determine whether some of the immune defects in \(1k \beta \alpha\) mutant mice are intrinsic to hematopoietic cells, we generated \(Rag^{2/−}\) chimeras. LNs were present and comparable in size, cellularity, and distribution of T and B cells in \(1k \beta \alpha\) mutant→\(Rag^{2/−}\) chimeras and WT→\(Rag^{2/−}\) control chimeras (Fig. 9, A and B). PPs and splenic follicles
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Figure 7. Impaired antibody responses in IκBα mutant mice. (A–C) TNP-specific serum Ig measured by ELISA in mice immunized with TNP-LPS (A) and TNP-Ficoll (B) and OVA-specific serum Ig measured by ELISA in mice immunized with OVA (C). Data represent the mean of duplicate determinations on sera from four to seven mice per group. (D) Spleen sections from immunized mice were stained by histochemistry with anti-B220 or PNA. Bars, 500 μm. Data are representative of six mice per group in three independent experiments. (E and F) Analysis of PNA+ and Fas+GL7+ B cells among the B220+ population of splenic B cells of OVA-immunized mutant mice and WT controls (E) and percentages of these subpopulations in unimmunized and OVA-immunized mice of both groups (F), as analyzed by flow cytometry. Data are representative of three to six mice per group in three independent experiments. Imm, immunization. Lines and columns represent mean, and bars represent SD in A–C and F. *, P < 0.05; **, P < 0.01; ***, P < 0.001.

and FDCs were also present and comparable in size in the two chimeras (Fig. 9, C and D). Furthermore, MZ B cells were present in IκBα mutant→Rag2−/− chimeras, although their number was less than in control chimeras (Fig. 9 E). These findings indicate that the defective lymphoid organogenesis in IκBα mutant mice was not caused by an intrinsic defect in hematopoietic cells.

IκBα mutant→Rag2−/− chimeras mounted a CHS response to OVA comparable with that of control chimeras, as measured by ear swelling, cellular infiltration, and Ifng and Il4 mRNA expression (Fig. 10, A–C). These findings strongly suggest that the lack of CHS in the IκBα mutants is caused by their lack of LNs.

Despite the presence of splenic follicles, FDCs, and MZ B cells, the antibody responses to TNP-LPS, TNP-Ficoll, and OVA were all significantly lower in IκBα mutant→Rag2−/− chimeras than in control chimeras (Fig. 10, D–F). After OVA immunization, the chimeras formed splenic GCs that were comparable in size and number (Fig. 10 G) and had a comparable increase in the percentage of B220+PNA+ cells (Fig. 10 H). Quantitative PCR (qPCR) analysis performed on B220+PNA+ cells revealed that mRNA levels of Aicda and Bcl6, which are expressed by GC B cells, were comparable in the two chimeras. In contrast, mRNA levels of Prdm1 and Xbp1 (X-box–binding protein 1), which drive plasma cell differentiation, were significantly lower in the IκBα mutant→Rag2−/− chimeras than in control chimeras (Fig. 10 I). These results suggest that defective GC formation in IκBα mutant mice is caused by a defect in nonhematopoietic architectural cells rather than in hematopoietic cells. Nevertheless, the B cells from the mutant have an intrinsic defect in terminal differentiation into plasma cells.

**DISCUSSION**

We have generated mice heterozygous for the S32I mutation in IκBα that recapitulate the known features of ED-ID patients who carry the same mutation. Strikingly, these mice had severely defective lymphoid organogenesis caused by a defect in nontransferable architectural cells, a finding not previously appreciated in patients, which may explain their susceptibility to infection despite normal lymphocyte number and in vitro function and their poor response to HSCT.
Figure 8. Secondary lymphoid organs and immune function in WT→IκBα mutant chimeras. (A) Photograph of inguinal fat pad with the vessel junction. The dotted circle indicates the LN at the vessel junction in the fat pad. (B) H&E staining of small intestine sections. The arrow points to a PP in the WT mouse. (C) Spleen sections were stained with H&E (left) and by immunofluorescence for FDCM1, IgD, and IgM (right). (D) Percentage of CD21+CD23− MZ B cells among B220+ cells in the spleen, as determined by flow cytometry (n = 4–6 mice per group). (E–G) Increase in ear thickness measured by a micrometer (E), representative H&E-stained sections (F), and Ifng and Il-4 mRNA expression determined by qPCR (G) 24 h after ear challenge with OXA or vehicle (n = 3–5 mice per group). (H and I) Serum TNP-specific (H) and OVA-specific (I) antibody measured by ELISA in mice immunized with TNP-Ficoll (H) and OVA (I), respectively (n = 5 mice per group). (J) Spleen sections from immunized mice stained by histochemistry with anti-B220 mAb or PNA. Bars: (B) 500 μm; (C, left), F, and J 200 μm; (C, right) 100 μm. (K) Percentage of PNA+ B cells among the B220+ population of splenic B cells determined by flow cytometry (n = 3–5 mice per group). Data in A–C and J are representative of three to six mice. Columns and bars in D, E, G–I, and K represent mean and SD. *, P < 0.05; **, P < 0.01; ***, P < 0.001.
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The IκBα mutant mouse we generated had ectodermal defects similar to those present in EDA- and EDA-R–deficient patients and mice and in patients with ED-ID (Srivastava et al., 2001; Laurikka et al., 2002; Claus et al., 2008; Hanson et al., 2008; Kawai et al., 2012). Growth, weight gain, and survival were diminished in the mutant, like in patients with hypermorphic IκBα mutations (Courtois et al., 2003; Janssen et al., 2004; McDonald et al., 2007; Lopez-Granados et al., 2008; Ohnishi et al., 2012; Yoshioka et al., 2013). Activation of the canonical NF-κB pathway was deficient in these mice, as in the patients (Courtois et al., 2003; McDonald et al., 2007). This was indicated by impaired IκBα phosphorylation and degradation in fibroblasts after stimulation with IL-1, TNF, and LPS. The mutant also showed reduced cytokine production by BMDCs in response to TLR stimulation and decreased up-regulation of ICAM1 and VCAM1 expression by MEFs in response to TNF. Thus, the heterozygous IκBα mutant mice we generated represent a faithful model of humans with hypermorphic IκBα mutations.

Secondary lymphoid organogenesis was severely defective in the IκBα mutant mouse. The mutant completely lacked LNs and PPs, and its spleen lacked follicles, MZs, MZ B cells, and FDCs. This was surprising because secondary lymphoid organogenesis is relatively well preserved in p50-deficient mice, which only lack inguinal LNs (Lo et al., 2006), and partially preserved in mice deficient in TNF, TNFR1 (TNF receptor 1), RANK (receptor activator of NF-κB), or RANKL (RANK ligand), which signal through the canonical NF-κB pathway (Pasparakis et al., 1996, 1997; Körner et al., 1997; Matsumoto et al., 1997; Dougall et al., 1999; Kong et al., 1999; Kim et al., 2000). Instead, the phenotype of IκBα mutant mice closely resembles that of mice deficient in RelB or NIK and mice deficient in LTα, LTβ, or LTβR, which signal through both the canonical and noncanonical NF-κB pathways (Miyawaki et al., 1994; Banks et al., 1995; Weih et al., 1995; Koike et al., 1996; Koni et al., 1997; Fütterer et al., 1998; Weih et al., 2001; Tumanov et al., 2003; Seymour et al., 2006).

Defective lymphoid organogenesis has not been previously appreciated in patients with ED-ID. However, tonsils and cervical LNs have been noted to be absent in at least one of the patients with the S32I IκBα mutation (Dupuis-Girod et al., 2006). Circulating lymphocyte counts were significantly elevated in IκBα mutant mice, as has been noted in patients with IκBα deficiency (Courtois et al., 2003; Janssen et al., 2004; McDonald et al., 2007; Ohnishi et al., 2012) and in mice that lack LNs and PPs (De Togni et al., 1994; Alimzhanov et al., 1997; Fütterer et al., 1998).

Lymphoid organogenesis involves activation of both the canonical and noncanonical NF-κB pathways through LTαβLTβR interactions between LTI cells and stromal organizer cells (Mebius, 2003; Vondenhoff et al., 2009; van de Pavert and Mebius, 2010; Roosendaal and Mebius, 2011). LTI cells were present in the IκBα mutant, consistent with previous observations in mice with defects in the canonical NF-κB pathway (Alcamo et al., 2002). As expected, MEFs from the mutant failed to up-regulate Vcam1, Lami1, and Cxcl10 mRNA expression in response to LTβR ligation, which depends on the canonical NF-κB pathway (Dejardin et al., 2002; Caposio et al., 2007). In addition, they failed to up-regulate the expression of Madcam1, which depends on the noncanonical pathway downstream of
for p100 processing into its active product p52 (Dejardin et al., 2002), the impaired noncanonical NF-κB pathway activity in the mutant is likely caused by the decreased p100 levels. In addition to the control of p100 level by the canonical pathway, there is cross talk at other levels between the canonical and noncanonical NF-κB pathways (Matsushima et al., 2001; Shih et al., 2011). The complexity of this talk may account for the variable effects of deficiency in individual components of the canonical NF-κB pathway on lymphoid LTβR (Ganeff et al., 2011). Furthermore, splenocytes from the IκBα mutant expressed significantly less mRNA for the chemokine genes Ccl19, Cxcl12, and Cxcl13, the expression of which depends on the noncanonical NF-κB pathway (Dejardin et al., 2002). We demonstrated that the levels of p100 were markedly reduced in cells from the mutant. This is consistent with previous observations that canonical NF-κB signaling controls p100 levels in cell lines (Liptay et al., 1994; Dejardin et al., 2002; Shih et al., 2011). Because canonical NF-κB activation is not required

Figure 10. Defective immune function in IκBα mutant→Rag2−/− chimeras. (A–C) Increase in ear thickness measured by a micrometer (A), representative H&E-stained sections (B), and Ifng and Il-4 mRNA expression determined by qPCR (C) 24 h after ear challenge with OXA or vehicle. Data in A–C are representative of five to seven mice per group in three experiments. (D–F) Serum TNP-specific (D) and OVA-specific (F) antibody measured by ELISA in mice immunized with TNP-LPS (D), TNP-Ficoll (E), and OVA (F), respectively. Data in D–F are representative of a mean of two determinations on sera from 5–10 mice per group. (G) Spleen sections from immunized mice stained by histochmotery with anti-B220 or PNA. Bars: (B) 200 μm; (G) 25 μm. (H) Percentage of PNA+ B cells among the B220+ population of splenic B cells (n = 3 mice per group) determined by flow cytometry. (I) Aicda, Bcl6, Prdm1, and Xbp1 mRNA expression in B220+PNA+ cells from immunized mice measured by qPCR. Data in G–I are representative of three mice per group in three experiments. Columns and bars in A, C–F, H, and I represent mean and SD. *, P < 0.05; **, P < 0.01; ***, P < 0.001.
organogenesis. Thus, although lymphoid organogenesis is absent in Rela<sup>−/−</sup> mice rescued from lethality by breeding on the Tnfr1<sup>−/−</sup> background (Alcamo et al., 2002), it is relatively unaffected in p50-deficient Nfkbi<sup>−/−</sup> mice. NEMO-deficient patients are known to possess LNs (Orange et al., 2004; Imamura et al., 2011). A recent study demonstrates increased baseline levels of NIK and increased activation of the noncanonical pathway in NEMO-deficient MEFs caused by loss of the IKK complex (Gray et al., 2014). Although not discussed in that study, p100 levels were low in these MEFs. Increased NIK levels and activity in NEMO deficiency may compensate for the low levels of p100. In contrast, NIK levels were not increased in MEFs from the IκBα mutant mice.

Thymocyte development and splenic T cell numbers were normal in the IκBα mutant. Moreover, T cell proliferation and IL-2 secretion in response to TCR ligation with anti-CD3 or antigen stimulation after i.p. immunization were normal in the mutant. However, up-regulation of IL-2 secretion by CD28 ligation and in vitro differentiation of naive T cells into Th1 and Th17, but not Th2 cells, were impaired. The CHS response, which largely depends on T cells (Wang et al., 2000; Gorbachev and Fairchild, 2004), was markedly deficient in the mutant. Failure to mount a CHS response also occurs in Ltbr<sup>−/−</sup> mice (Kernert et al., 2001) and is likely caused by the absence of LNs, where contact between hapten-specific T cells and hapten-bearing DCs takes place. This is supported by the observation that the CHS response was reconstituted in IκBα mutant→Rag2<sup>−/−</sup> chimeras, which developed LNs that were normal in size and cellularity.

The IκBα mutant had normal B cell development in the BM and normal B cell numbers in the spleen, but had a severe deficiency in splenic MZ B cells. Proliferation of mutant B cells to LPS was normal, but proliferation to response to BCR, CD40, and TLR4 ligation in the presence of IL-4 was modestly reduced, suggesting a role for IκBα in B cell proliferation supported by IL-4. B cells from the mutant secreted less IgG after LPS stimulation and less IgA after LPS + TGFβ stimulation than control B cells. Expression of Aicda was normal, but expression of Pdml1 was reduced in LPS-stimulated B cells from the mutant, suggesting a defect in terminal B cell differentiation. B cell survival and IgG1 isotype switching in response to BAFF, which activates NF-κB primarily via the canonical pathway, was impaired in the mutant. Serum IgG levels were decreased in the mutant mice, and serum IgA and intestinal IgA<sup>+</sup> cells were virtually undetectable. IgA levels are low in patients with the IκBα S32I mutation (Courtois et al., 2003; Janssen et al., 2004). Serum IgA is virtually absent in Ltbr<sup>−/−</sup> and NIK-deficient mice (Shinkura et al., 1999), and its level is decreased in Nfkbi<sup>−/−</sup> mice (Snapper et al., 1996) and Rela<sup>−/−</sup>/Tnfr1<sup>−/−</sup> mice, which have impaired canonical signaling (Alcamo et al., 2002). Thus, defects in both canonical and noncanonical pathways may underlie the absence of IgA in the mutant.

Antibody responses to TI and TD antigens were severely deficient in the mutant. The defective response to TNP-Ficoll is consistent with the virtual absence of MZ B cells, which play a critical role in the antibody response to type II TI antigens (Baláz et al., 2002; Pillai et al., 2005). Remarkably, despite intact proliferation of splenic T cells in response to OVA stimulation ex vivo, the antibody response of the mutant to i.p. immunization with OVA was virtually absent and was accompanied by a total failure to form GCs. None of the antibody responses were restored in IκBα mutant→Rag2<sup>−/−</sup> chimeras, despite the restoration of MZ B cells and GC formation with generation of normal numbers of PNA<sup>+</sup> B cells, suggesting that an intrinsic defect in terminal B cell differentiation exits in the mutant. This was supported by the observation that the expression by PNA<sup>+</sup> cells of Pdm1 and Xbps1, which encode two transcription factors required for plasma cell differentiation, was significantly reduced in the IκBα mutant→Rag2<sup>−/−</sup> chimeras. Thus, the defective antibody response of IκBα mutant mice is two-tiered, involving defective lymphoid organogenesis and an intrinsic defect in terminal B cell differentiation.

Reconstitution of the IκBα mutant with WT BM cells failed to restore lymphoid organogenesis and to correct the immune defects in irradiated IκBα mutant recipients, despite excellent chimerism in the T and B cell lineages. This finding recapitulates the poor results of allogeneic HSCT in patients with IκBα mutations despite the engraftment of donor lymphocytes. Notably, despite 100% chimerism in lymphoid cells, the single patient with the S32I IκBα mutation who has survived HSCT continues to suffer from recurrent infections requiring gamma globulin replacement and has persistent lymphocytosis, no visible tonsillar tissue, and no palpable LNs even during infection (Dupuis-Girod et al., 2006; Cancrini, C., personal communication).

Our findings suggest that defects in architectural nonhematopoietic cells, likely a mesenchymally derived stromal cell, underlie the immune deficiency in patients with ED-ID caused by IκBα mutations despite the engraftment of donor lymphocytes. Strategies that correct defective architectural cell function, such as implantation of stromal cell–embedded biocompatible scaffolds to generate LNs (Okamoto et al., 2007), may be valuable in patients with IκBα deficiency.

**MATERIALS AND METHODS**

**Generation of heterozygous IκBα S32I knock-in mice.** The arms of the IκBα S32I targeting vector were amplified by PCR from genomic DNA of the C57 embryonic stem (ES) cell line derived from the 129Sv mouse strain. The 2.8-kb 5′ and 3.1-kb 3′ arms were cloned into the pKS-neo-DTAIII vector (a gift from the laboratory of M. Greenberg, Harvard Medical School, Boston, MA), and the nucleic acid G94T mutation was introduced by site-directed mutagenesis (Agilent Technologies) into exon 1 in the 5′ arm, resulting in the amino acid S32I mutation (Fig. S1 A). The linearized targeting construct was electroporated into C57 ES cells, which were then selected in 0.4 μg/ml G418 and 10 mg/ml ganciclovir. An ES cell clone containing a disrupted allele in the amino acid S32I mutation (Fig. S1 A). The linearized targeting construct was electroporated into C57 ES cells, which were then selected in 0.4 μg/ml G418 and 10 mg/ml ganciclovir. An ES cell clone containing a disrupted allele in the amino acid S32I mutation who has survived HSCT continues to suffer from recurrent infections requiring gamma globulin replacement and has persistent lymphocytosis, no visible tonsillar tissue, and no palpable LNs even during infection (Dupuis-Girod et al., 2006; Cancrini, C., personal communication).

Our findings suggest that defects in architectural nonhematopoietic cells, likely a mesenchymally derived stromal cell, underlie the immune deficiency in patients with ED-ID caused by IκBα mutations and explain the poor outcome of allogeneic HSCT in these patients. Strategies that correct defective architectural cell function, such as implantation of stromal cell–embedded biocompatible scaffolds to generate LNs (Okamoto et al., 2007), may be valuable in patients with IκBα deficiency.

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cages with trimethoprim/sulfamethoxazole and enrofloxacin added to the water to prevent infection. Procedures were performed in accordance with the Animal Care and Use Committee of the Children's Hospital Boston.

**Dermal fibroblast isolation and culture.** Primary dermal fibroblasts were isolated as in McDonald et al. (2007) and grown in RPMI media plus 10% FCS (HyClone) plus t-glutamine, penicillin, and streptomycin (Invitrogen). Fibroblasts were stimulated with media, 10 ng/ml IL-1, 10 ng/ml TNF (R&D Systems), or 1 mg/ml LPS (Sigma-Aldrich) for the indicated times.

**Western blotting.** Western blotting was performed as previously described (McDonald et al., 2007) using anti-phospho IkBα (Cell Signaling Technology), anti-mouse full-length IkBα (Abcam), and anti-actin (Sigma-Aldrich) according to the manufacturers' recommendations. Immunoblotting was also performed using anti-p100/p52, RelB, and NIK (Cell Signaling Technology). Sheep anti-mouse horseradish peroxidase–conjugated and sheep anti-rabbit horseradish peroxidase–conjugated secondary antibodies were obtained from GE Healthcare. Quantification was performed using the public domain ImageJ/FJIL software from the National Institutes of Health.

**BMDC isolation and TLR ligand stimulation.** BM was flushed from the femur and tibia of 6–8-wk-old mice, plated in DMEM (Gibco) plus 10% FCS and plus t-glutamine, penicillin, and streptomycin.

**VCAM1 surface staining.** MEFs were treated for 24 h with 10 ng/ml TNF, dissociated with trypsin/EDTA, and stained with anti-VCAM1 (eBioscience).

**LN detection by Evan’s blue injection.** LN detection was performed as described in Harrell et al. (2008).

**H&E staining.** Samples of inguinal fat pad, small intestine, nasal cavity, thymus, spleen, and LN were fixed in formalin for 24 h, and then H&E staining of paraffin-embedded cut sections was performed by the Rodent Histopathology and Specialized Histopathology Services cores at the Dana Farber/Harvard Cancer Center (Boston, MA) or by Histo-Scientific Research Laboratories (HSRL, Mt. Jackson, Virginia).

**Complete blood counts (CBCs).** Blood was collected from the facial vein, and CBC analysis was performed by the Animal Research at Children’s Hospital (ARCH) facility.

**FACS analysis.** Single-cell suspensions from spleen, thymus, or BM of 4–6-wk-old mice were stained with FITC, PE, PE-Cy7, APC, PerCP-Cy5.5, Alexa Fluor 700, or APC–eFluor 780–labeled mAbs and analyzed on the FACSCanto (BD). Antibodies to B220, CD3, IgM, CD21, CD23, CD93, CD43, CD44, GL-7, CD11b, Ly6G, F4/80, NK1.1, CD45, IL-7R, CD45.1, and CD45.2 were from eBioscience, mAb to Fas was from BD, and FITC-conjugated PNA was from Sigma-Aldrich.

**Immunofluorescence.** Immunofluorescence was performed as described in Cariappa et al. (2009) and Kranich et al. (2008).

**LTI cell isolation from embryonic intestine.** Embryonic intestines were isolated from E17.5 fetuses, incubated in Liberase (Roche) for 1 h at 37°C, and then passed through a 70-µm filter. A portion of the embryos was reserved for genotyping.

**qPCR.** MEFs were stimulated with 2 µg/ml anti-LTβR agonist antibody AFH6 (a gift from J. Browning, Biogen Idec, Cambridge, MA) or 10 ng/ml TNF (R&D Systems) for 24 h, and RNA was extracted using the TRIzol (Invitrogen) method. cDNA was synthesized using Script (Bio-Rad Laboratories), and qPCR was performed using TaqMan probes and the 7300 Real-time PCR System (Applied Biosystems).

**Purification and stimulation of splenic T and B cells.** T cells were purified from spleen suspensions using the pan-T cell isolation kit II (Miltenyi Biotec), and 100,000 per 96 wells were plated, along with 300,000 irradiated (3,000 rad) non-T cell APCs purified by the CD90.2 kit (Miltenyi Biotec) using RPMI supplemented with penicillin/streptomycin, Hepes (Gibco), sodium pyruvate, and β-mercaptoethanol. 200,000 cells in 96-well plates were plated for proliferation assays, and 250,000 per 24-well plates for in vitro Ig analysis. Cells were stimulated with 0.1–10 µg/ml LPS (Sigma-Aldrich) or 10 µg/ml LPS, 10 µg/ml anti-m-IL-4 (Jackson Immunoresearch Laboratories, Inc.), or 500 ng/ml anti-CD40 (eBioscience), alone or in combination with 40 ng/ml IL-4 (R&D Systems). Tritiated thymidine (PerkinElmer) was added at day 3, and incorporation was assayed after 15–18 h.

**Serum Ig levels and antibody responses.** 5 ng/ml IL-12 (BioLegend) and 10 µg/ml anti–IFN-γ neutralizing antibodies were added. For Th1 cell differentiation, 40 ng/ml IL-23 (R&D Systems), 1 ng/ml TGF-β (BioLegend), 10 µg/ml anti–IL-4 (11B11; BioLegend), and 10 µg/ml anti–IL-10 (BioLegend) neutralizing antibodies were added. For Th1 cell differentiation, 5 ng/ml IL-12 (BioLegend) and 10 µg/ml anti–IL-4 neutralizing mAbs (11B11; BioLegend) were added.

**In vitro T helper cell differentiation.** For Th0 conditions, naïve CD4+ T cells were sorted by MACS and were stimulated for 5 d with plate-bound antibodies against CD3 (145-2C11; 2 µg/ml; eBioscience) and CD28 (PV-1; 2 µg/ml; BioLegend). For Th2 cell differentiation, 50 ng/ml IL-4 (BioLegend) and 10 µg/ml neutralizing anti–IFN-γ mAb (XM1.2; BioLegend) were added. For Th17 cell differentiation, 40 ng/ml IL-6 (BioLegend), 40 ng/ml IL-23 (R&D Systems), 1 ng/ml TGFB-β (BioLegend), 10 µg/ml anti–IL-10 (11B11; BioLegend), and 10 µg/ml anti–IL-4 (BioLegend) neutralizing antibodies were added. For Th1 cell differentiation, 5 ng/ml IL-12 (BioLegend) and 10 µg/ml anti–IL-4 neutralizing mAbs (11B11; BioLegend) were added.
CHS to OXA. Mice were sensitized by the application of 100 μl of 2% OXA (Sigma-Aldrich) in ethanol on day 1 to shaved abdominal skin. On day 5, the left ear was challenged with 10 μl of 1% OXA, and the right ear was challenged with vehicle (ethanol) on dorsal and ventral surfaces. Ear thickness was measured after 24 h with a micrometer (Mitutoyo).

BM reconstitution of mutant mice. Adult healthy mutant mice and their WT littermates received two 550-cGy (550Cesium source) doses of total body irradiation 3 h apart. BM was collected from CD45.1 mice (Charles River), and 10^6 viable BM cells were injected i.t. into the mice. Reconstitution was assessed by flow cytometry using antibodies to CD45.1 (donor) and CD45.2 (recipient).

BM reconstitution of Rag2^-/- mice. BM was collected from 6-wk-old mutant mice or their WT littermates. Flow cytometry was performed to assure the CD34^+ population was comparable between WT and mutant BM. Adult Rag2^-/- recipient mice received a single 250-cGy dose of total body irradiation. Viability of the BM cells was examined by trypan blue dye exclusion, and 10^6 viable BM cells were injected i.t. into the Rag2^-/- mice.

PNA B cell sorting. 10 d after immunization with OVA plus alum, spleens from Rag2^-/- BM chimeras were harvested and stained for B220 and PNA. Double-positive cells were collected at the Children's Hospital cell sorting core.

Statistical analysis. Statistical analysis of data was performed with Prism software (GraphPad Software) using the Student's t test or ANOVA.

Online supplemental material. Fig. S1 shows the Nkbia targeting construct and screening strategy. Online supplemental material is available at http://www.jem.org/cgi/content/full/jem.20140979/DC1.

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by classical NF-κB key regulator of osteoclastogenesis, lymphocyte development and lymph-


