

# Loss of c-REL but not NF- $\kappa$ B2 prevents autoimmune disease driven by FasL mutation

LA O'Reilly<sup>\*1,2</sup>, P Hughes<sup>1,3</sup>, A Lin<sup>1</sup>, P Waring<sup>4</sup>, U Siebenlist<sup>5</sup>, R Jain<sup>1,2</sup>, DHD Gray<sup>1,2</sup>, S Gerondakis<sup>6,7</sup> and A Strasser<sup>\*1,2,7</sup>

FASL/FAS signaling imposes a critical barrier against autoimmune disease and lymphadenopathy. Mutant mice unable to produce membrane-bound FASL (*FasL* <sup>$\Delta m/\Delta m$</sup> ), a prerequisite for FAS-induced apoptosis, develop lymphadenopathy and systemic autoimmune disease with immune complex-mediated glomerulonephritis. Prior to disease onset, *FasL* <sup>$\Delta m/\Delta m$</sup>  mice contain abnormally high numbers of leukocytes displaying activated and elevated NF- $\kappa$ B-regulated cytokine levels, indicating that NF- $\kappa$ B-dependent inflammation may be a key pathological driver in this multifaceted autoimmune disease. We tested this hypothesis by genetically impairing canonical or non-canonical NF- $\kappa$ B signaling in *FasL* <sup>$\Delta m/\Delta m$</sup>  mice by deleting the *c-Rel* or *NF- $\kappa$ B2* genes, respectively. Although the loss of NF- $\kappa$ B2 reduced the levels of inflammatory cytokines and autoantibodies, the impact on animal survival was minor due to substantially accelerated and exacerbated lymphoproliferative disease. In contrast, a marked increase in lifespan resulting from the loss of c-REL coincided with a striking reduction in classical parameters of autoimmune pathology, including the levels of cytokines and antinuclear autoantibodies. Notably, the decrease in regulatory T-cell numbers associated with loss of c-REL did not exacerbate autoimmunity in *FasL* <sup>$\Delta m/\Delta m$</sup> *c-rel*<sup>-/-</sup> mice. These findings indicate that selective inhibition of c-REL may be an attractive strategy for the treatment of autoimmune pathologies driven by defects in FASL/FAS signaling that would be expected to circumvent many of the complications caused by pan-NF- $\kappa$ B inhibition.

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Apoptotic death of autoreactive lymphocytes constitutes a major mechanism of immunological tolerance. Defects in the deletion of autoreactive T and B lymphocytes, either in primary lymphoid organs<sup>1–3</sup> or in the periphery,<sup>2</sup> can lead to autoimmune disease. The 'death receptor' FAS (CD95, APO-1), and its ligand FASL (CD95L, APO-1L), has critical roles in the removal of autoreactive lymphocytes in the periphery,<sup>4</sup> and in contraction of immune responses and lymphocyte homeostasis.<sup>5</sup> Accordingly, loss-of-function mutations in these genes cause lymphadenopathy and systemic lupus erythematosus (SLE)-like multiorgan autoimmune disease in mice<sup>6</sup> and humans.<sup>7</sup>

FAS, which is expressed on the surface of many cell types, allows FASL-mediated killing of pathogen-infected or damaged cells,<sup>4</sup> whereas its ligand (FASL) is thought to be present only on activated T cells and NK cells.<sup>4</sup> Upon deposition in the plasma membrane, membrane-bound FASL (mFASL) can also be processed and shed as secreted FASL (sFASL).<sup>4</sup> The generation of mice that lack mFASL but express sFASL (*FasL* <sup>$\Delta m/\Delta m$</sup> ) demonstrated unambiguously that only the former is essential for FAS-mediated cell killing, activation-induced T-cell death (AICD) and lymphocyte homeostasis.<sup>8</sup> Although *FasL* <sup>$\Delta m/\Delta m$</sup>  mice lack functional mFASL, they can still produce sFASL, which can engage FAS, an interaction

postulated to trigger non-apoptotic signals,<sup>4,9</sup> which promote inflammation and cytokine production.<sup>8,10</sup> Consistent with this hypothesis, *FasL* <sup>$\Delta m/\Delta m$</sup>  mice develop SLE-like kidney disease, autoimmune allergic dermatitis and histiocytic sarcoma with higher incidence and accelerated rate compared with *FasL*<sup>*gld/gld*</sup> mice,<sup>8</sup> which carry a mutation that impairs the function of both mFASL and sFASL.

The onset of disease in *FasL* <sup>$\Delta m/\Delta m$</sup>  mice is preceded by an elevation in immune cells with activated nuclear factor- $\kappa$ B (NF- $\kappa$ B) that coincides with elevated serum levels of NF- $\kappa$ B-regulated inflammatory cytokines.<sup>8</sup> The NF- $\kappa$ B family of transcription factors, comprising homodimers and heterodimers of the five related proteins RELA, RELB, c-REL, NF- $\kappa$ B1 (p105/p50) and NF- $\kappa$ B2 (p100/p52), regulates the development, survival and function of innate and adaptive immune cells.<sup>11,12</sup> Different NF- $\kappa$ B dimers retained in a latent cytoplasmic state by various I $\kappa$ B inhibitory proteins are differentially activated by the canonical and non-canonical arms of the NF- $\kappa$ B pathway in response to distinct upstream signals (Supplementary Figure S1).<sup>13</sup> Antigen receptors and inflammatory cytokines, such as TNF, IL-1 $\beta$  and IL-6, are among the many signals that activate the canonical NF- $\kappa$ B transcription factors (dimers of c-REL or RELA and p50/NF- $\kappa$ B1<sup>11,12</sup>) through IKK $\beta$ -dependent proteasome-mediated

<sup>1</sup>Molecular Genetics of Cancer Division, The Walter and Eliza Hall Institute of Medical Research, Parkville 3052, Victoria, Australia; <sup>2</sup>Department of Medical Biology, The University of Melbourne, Parkville 3010, Victoria, Australia; <sup>3</sup>Department of Nephrology, The Royal Melbourne Hospital, Parkville 3052, Victoria, Australia; <sup>4</sup>Department of Pathology, The University of Melbourne, Parkville 3010 Victoria, Australia; <sup>5</sup>Laboratory of Molecular Immunology, National Institute of Allergy and Infectious Diseases, NIH, Bethesda, MD, USA and <sup>6</sup>Australian Centre for Blood Diseases and Department of Clinical Hematology, Monash University Central Clinical School, Melbourne 3004, Victoria, Australia

\*Corresponding author: LA O'Reilly or A Strasser, Molecular Genetics of Cancer Division, The Walter and Eliza Hall Institute of Medical Research, 1G Royal Parade, Parkville 3052, Victoria, Australia. Tel: +61 3 9345 2555, Fax: +61 3 9347 0852. E-mail: oreilly@wehi.edu.au or strasser@wehi.edu.au

<sup>7</sup>These authors share senior authorship.

**Abbreviations:** ANA, antinuclear autoantibodies; GN, glomerulonephritis; sFASL, secreted FASL; mFASL, membrane-bound FASL; mTEC, medullary thymic epithelial cells; AIRE, autoimmune regulator

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degradation of I $\kappa$ B inhibitory proteins (Supplementary Figure S1). Conversely, certain TNF receptor family members (e.g., BAFF-R) trigger the non-canonical NF- $\kappa$ B pathway through IKK $\alpha$ -induced proteolysis of p100/NF- $\kappa$ B2 bound to RELB, thereby generating dimers of p52/NF- $\kappa$ B2 and RelB (Supplementary Figure S1).<sup>11,12</sup> Canonical and non-canonical NF- $\kappa$ B transcription factors generally control different sets of genes,<sup>14</sup> although a subset of genes is co-regulated by the two arms of the NF- $\kappa$ B pathway.

Abnormally elevated NF- $\kappa$ B activity features in various human autoimmune diseases, including SLE and rheumatoid arthritis,<sup>12,15,16</sup> where it is thought to drive pathology by regulating the proliferation, survival and effector activities of immune cells.<sup>16–19</sup> The functions that NF- $\kappa$ B-regulated transcriptional programs exert in the immune system, combined with our findings that leukocytes in *FasL $\Delta m/\Delta m$*  mice have abnormally elevated NF- $\kappa$ B activity before disease onset, led us to investigate whether NF- $\kappa$ B signaling is critical for this disease. Due to the profound deleterious consequences associated with systemic NF- $\kappa$ B inhibition,<sup>20</sup> we focused on investigating the impact of selective loss of c-REL or NF- $\kappa$ B2 on autoimmune pathology in *FasL $\Delta m/\Delta m$*  mice. These NF- $\kappa$ B transcription factors were chosen because of their selective function in the canonical or non-canonical NF- $\kappa$ B pathways, and because they are mainly required for immune cell function and not for embryogenesis or hematopoietic cell development. Mice lacking c-REL, which is mainly expressed in hematopoietic cells,<sup>21</sup> have no major abnormality but exhibit defective activation of B and T lymphocytes, macrophages and DC that collectively contribute to reduced antibody and cytokine production.<sup>12,22,23</sup> In *nfk2 $\beta$ <sup>-/-</sup>* mice, T-cell and myeloid cell development and functions are intact, but reduced mature B-cell numbers due to their impaired survival, combined with germinal center defects, result in reduced humoral immune responses.<sup>24,25</sup> Interestingly, with age, *nfk2 $\beta$ <sup>-/-</sup>* mice exhibit multiorgan leukocyte infiltrates and mild autoimmune disease.<sup>18,26</sup> Our study demonstrates that loss of c-REL prevents autoantibody production in *FasL $\Delta m/\Delta m$*  mice and thereby substantially prolongs their lifespan. This suggests that selective targeting of c-REL may represent a promising strategy for the treatment of systemic autoimmune diseases.

## Results

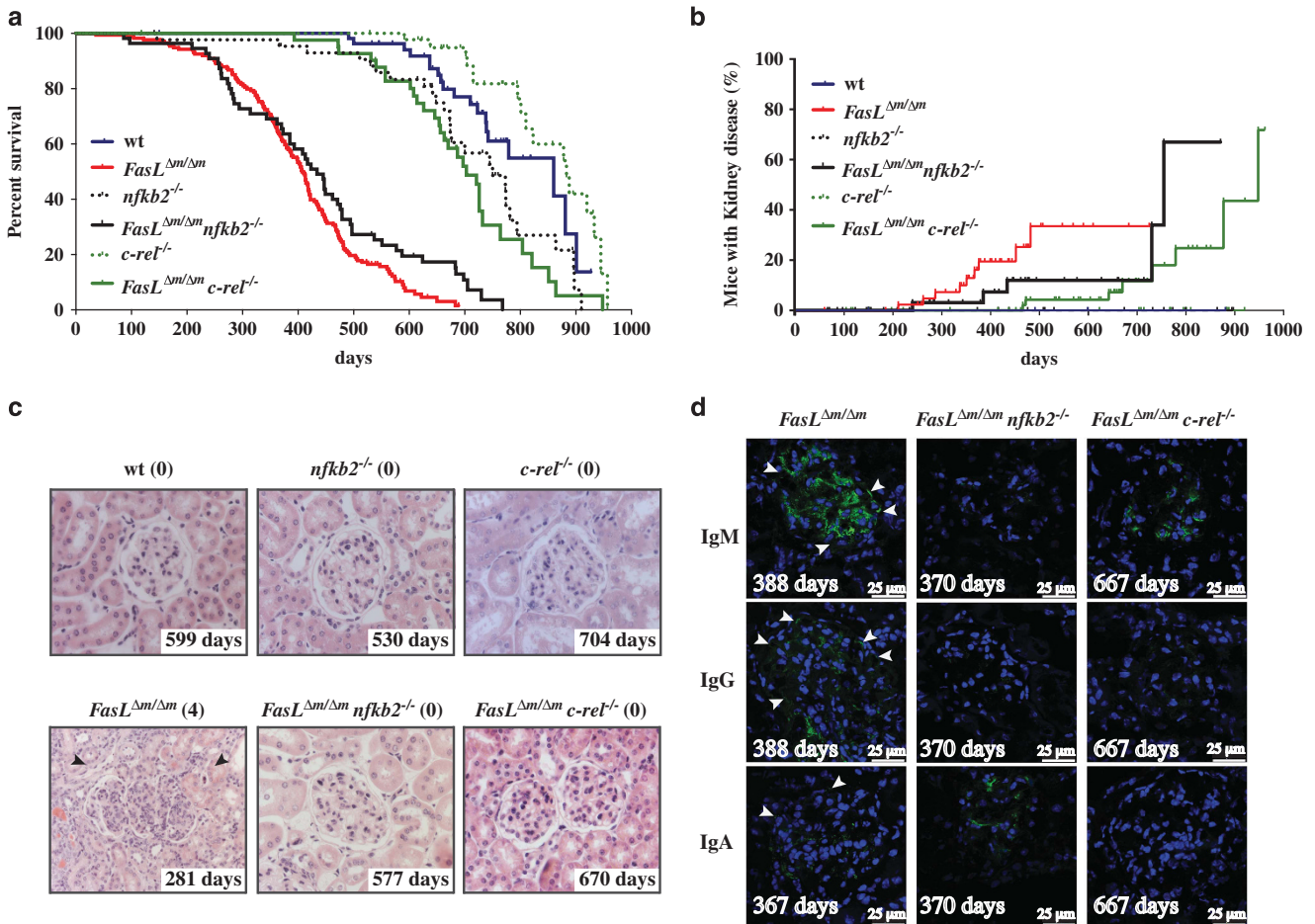
**Loss of c-REL extensively prolongs the lifespan of *FasL $\Delta m/\Delta m$*  mice, whereas loss of NF- $\kappa$ B2 has only a modest impact.** To investigate the roles of the canonical or non-canonical NF- $\kappa$ B signaling pathways in the lymphadenopathy and autoimmune disease caused by defects in FASL, we intercrossed *FasL $\Delta m/\Delta m$*  animals with *c-rel<sup>-/-</sup>* or *nfk2 $\beta$ <sup>-/-</sup>* mice. Both the lifespan and pathology of the *FasL $\Delta m/\Delta m$ c-rel<sup>-/-</sup>* and *FasL $\Delta m/\Delta m$ nfk2 $\beta$ <sup>-/-</sup>* compound mutants were compared with cohorts of wt, *FasL $\Delta m/\Delta m$* , *c-rel<sup>-/-</sup>* and *nfk2 $\beta$ <sup>-/-</sup>* mice for a period of 2 years. Although the absence of NF- $\kappa$ B2 or c-REL alone causes various immune defects,<sup>12,18</sup> these animals had lifespans comparable to wt mice (wt versus *nfk2 $\beta$ <sup>-/-</sup>* and wt versus *c-rel<sup>-/-</sup>* both  $P=NS$ , Figure 1a). Loss of NF- $\kappa$ B2 only modestly ( $P=0.0344$ ) prolonged the

lifespan of *FasL $\Delta m/\Delta m$*  mice, whereas this was greatly extended by loss of c-REL ( $P<0.0001$ , Figure 1a).

The morbidity and premature death of *FasL $\Delta m/\Delta m$*  mice are due to lymphadenopathy and SLE-like autoimmune glomerulonephritis (GN, Figure 1b).<sup>8</sup> Histological examination (Figure 1c) and immunostaining for glomerular immune complex deposition indicated that the incidence of severe autoimmune kidney damage was significantly reduced in both *FasL $\Delta m/\Delta m$ c-rel<sup>-/-</sup>* and *FasL $\Delta m/\Delta m$ nfk2 $\beta$ <sup>-/-</sup>* mice, and the damaging immunoglobulin deposition on the glomerular capillary loops (Figure 1d, arrows) in these animals was now mainly confined to mesangial cells (Figure 1d). Impairment of renal function (pathology grading 3–4) was modest and delayed in *FasL $\Delta m/\Delta m$ nfk2 $\beta$ <sup>-/-</sup>* mice (*FasL $\Delta m/\Delta m$ nfk2 $\beta$ <sup>-/-</sup>* versus *FasL $\Delta m/\Delta m$* ,  $P<0.05$ ). Loss of c-REL had a more profound impact on GN-associated autoimmune pathology ( $P=0.0002$ ), manifest as substantial reductions in hypercellularity, capillary loop obliteration and fibrinoid necrosis (Figures 1b and c).

**NF- $\kappa$ B2 and c-REL are required for inflammatory cytokine production in *FasL $\Delta m/\Delta m$*  mice.** Before the onset of SLE-like autoimmune disease (~5 months), *FasL $\Delta m/\Delta m$*  mice harbor high levels of circulating inflammatory cytokines.<sup>8</sup> We examined the levels of 31 cytokines and chemokines and found that TNF, IL-17A/F, KC (CXCL1) and MIP-1 $\beta$  levels were elevated in *FasL $\Delta m/\Delta m$*  mice at 5 months of age (Figure 2a), whereas IFN $\beta$  remained largely undetectable by ELISA (Supplementary Figure S2), although type I interferons have been implicated previously in the GN in *Fas<sup>lpr/lpr</sup>* mutant mice.<sup>27</sup> NF- $\kappa$ B2 deficiency blunted the excess production of most cytokines (except KC) in *FasL $\Delta m/\Delta m$*  mice (Figure 2a). As *FasL $\Delta m/\Delta m$*  mice aged, IL-1 $\beta$  and IL-6 levels increased significantly, whereas in aged *FasL $\Delta m/\Delta m$ nfk2 $\beta$ <sup>-/-</sup>* mice (283–683 days) all cytokines tested, with the exception of IL-17A/F and KC, remained at wt levels (Figure 2b). Remarkably, none of the cytokines or chemokines tested were elevated in *FasL $\Delta m/\Delta m$ c-rel<sup>-/-</sup>* mice, even when aged (Figures 2a and b). This indicates that c-REL and, to a lesser extent, NF- $\kappa$ B2 are required for accumulation of the cytokines and chemokines associated with autoimmune pathology in *FasL $\Delta m/\Delta m$*  mice.

**NF- $\kappa$ B2 restrains lymphadenopathy in *FasL $\Delta m/\Delta m$*  mice.** Defects in AICD of T cells are a major cause of lymphadenopathy in mice and humans deficient in FASL/FAS function and are associated with the accumulation of 'unusual' TCR $\alpha\beta^+$  CD3 $^+$ CD4 $^-$ CD8 $^-$ B220 $^+$  (hereafter called DN) T cells.<sup>6,7</sup> *FasL $\Delta m/\Delta m$*  mice display lymphadenopathy and splenomegaly commencing from ~3 months of age (Figures 3a and b). Surprisingly, loss of NF- $\kappa$ B2 greatly accelerated lymphadenopathy in *FasL $\Delta m/\Delta m$*  mice (Figures 3a and b) with a fourfold increase in DN T cells compared with *FasL $\Delta m/\Delta m$*  controls (Figure 3c). By 3 months, these DN T cells comprised the majority of leukocytes in the lymph nodes ( $81.9 \pm 2.8\%$ ) and spleens of *FasL $\Delta m/\Delta m$ nfk2 $\beta$ <sup>-/-</sup>* mice ( $76.7 \pm 1.1\%$ , Figure 3c), resulting in a >10-fold increase in total lymph node and splenic cellularity compared with *FasL $\Delta m/\Delta m$*  mice (Figure 3e). In contrast, loss of c-REL had no impact on the onset or magnitude of lymphadenopathy and DN T-cell numbers in *FasL $\Delta m/\Delta m$*  mice (Figures 3a–e). To assess whether the enhanced lymphadenopathy of *FasL $\Delta m/\Delta m$ nfk2 $\beta$ <sup>-/-</sup>* mice was



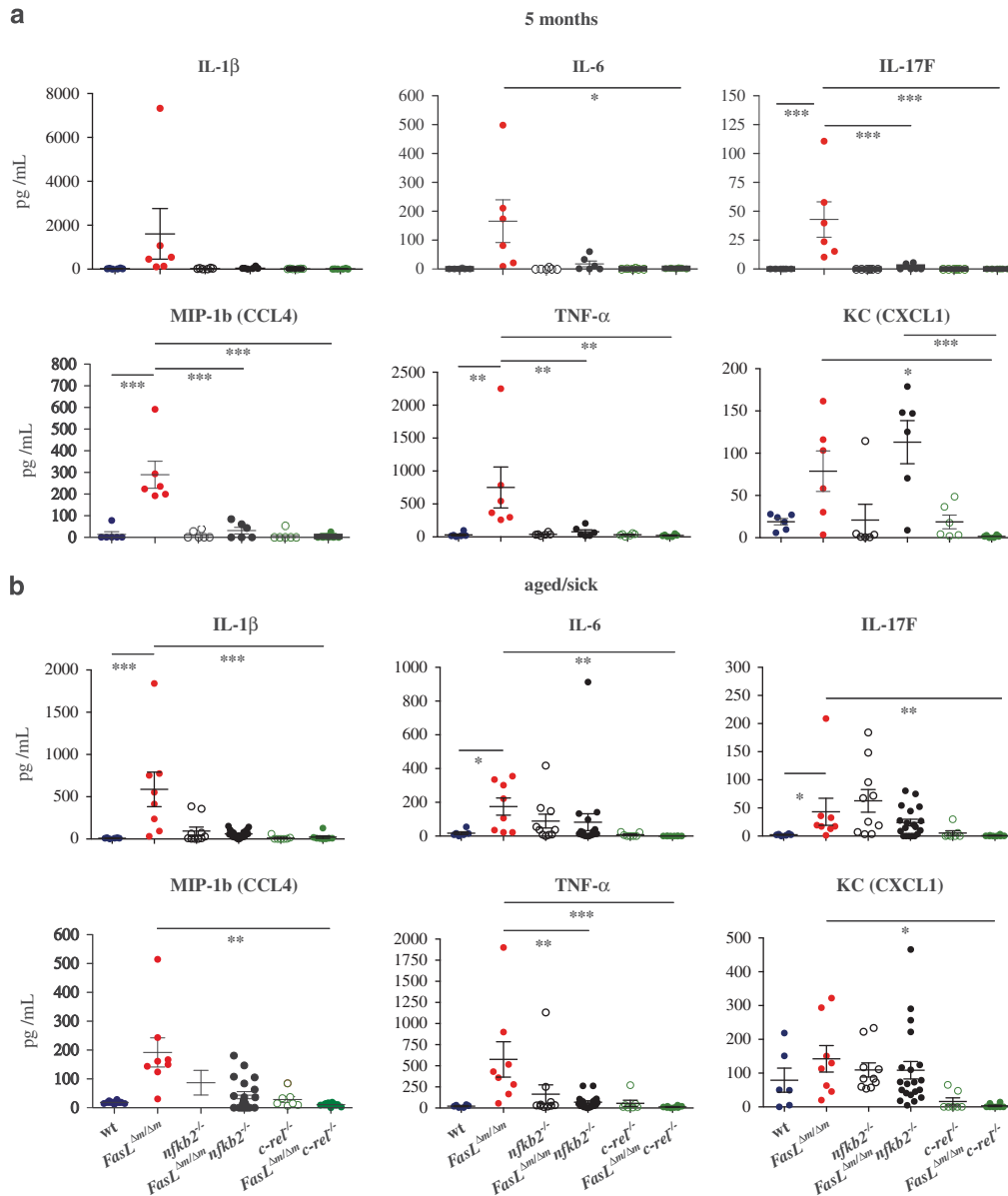
**Figure 1** Loss of NF- $\kappa$ B2 or c-REL both prolonged the lifespan of *FasL*<sup>Δm/Δm</sup> mutant mice and delayed or even prevented fatal SLE-like GN in these animals. (a) Kaplan-Meier survival curves for wt ( $n = 54$ , mean survival 860 days, blue line), *FasL*<sup>Δm/Δm</sup> ( $n = 174$ , mean survival 407 days, red line), *nfk2*<sup>-/-</sup> ( $n = 43$ , mean survival 750 days, dotted black line), *FasL*<sup>Δm/Δm</sup>*nfk2*<sup>-/-</sup> ( $n = 54$ , mean survival 434 days, solid black line), *c-rel*<sup>-/-</sup> ( $n = 46$ , mean survival 883 days, dotted green line) and *FasL*<sup>Δm/Δm</sup>*c-rel*<sup>-/-</sup> ( $n = 40$ , mean survival 703 days, solid green line) mice (wt versus *FasL*<sup>Δm/Δm</sup>;  $P < 0.0001$ ; *FasL*<sup>Δm/Δm</sup> versus *FasL*<sup>Δm/Δm</sup>*nfk2*<sup>-/-</sup>;  $P = 0.0344$ ; *FasL*<sup>Δm/Δm</sup> versus *FasL*<sup>Δm/Δm</sup>*c-rel*<sup>-/-</sup>;  $P < 0.0001$ ; wt versus *nfk2*<sup>-/-</sup>;  $P = \text{NS}$ ; wt versus *FasL*<sup>Δm/Δm</sup>*c-rel*<sup>-/-</sup>;  $P = \text{NS}$ ). (b) Graphical representation of the incidence of severe autoimmune kidney disease (GN score  $\geq 3$ ) in mice of the indicated genotypes (*FasL*<sup>Δm/Δm</sup> versus *FasL*<sup>Δm/Δm</sup>*nfk2*<sup>-/-</sup>;  $P < 0.05$ , *FasL*<sup>Δm/Δm</sup> versus *FasL*<sup>Δm/Δm</sup>*c-rel*<sup>-/-</sup>;  $P = 0.0002$ , lines colored as for (a), wt ( $n = 4$ ), *FasL*<sup>Δm/Δm</sup> ( $n = 44$ ), *nfk2*<sup>-/-</sup> ( $n = 22$ ), *FasL*<sup>Δm/Δm</sup>*nfk2*<sup>-/-</sup> ( $n = 40$ ), *c-rel*<sup>-/-</sup> ( $n = 17$ ), *FasL*<sup>Δm/Δm</sup>*c-rel*<sup>-/-</sup> ( $n = 56$ )). (c) Representative H&E-stained sections of the kidneys from mice of the indicated genotypes were examined for pathological changes, such as hypercellularity, cellular crescents, dilated tubules or sclerotic glomeruli. Arrows indicate fibrinoid necrosis. Magnification  $\times 40$ . (d) Representative confocal photomicrographs of frozen sections (eight kidneys per genotype analyzed) stained for the presence of IgA-, IgG- or IgM-containing immune complexes (green) in glomeruli. Nuclei are revealed by staining with DAPI (blue). Arrows indicate Ig deposition on the glomerular basement membrane (scale bars represent 25  $\mu\text{m}$ )

intrinsic to the hematopoietic system, bone marrow chimeras were generated. C57BL/6-Ly5.1 (wt) mice reconstituted with hematopoietic stem/progenitor cells (HSPCs) from *FasL*<sup>Δm/Δm</sup> *nfk2*<sup>-/-</sup> mice exhibited enhanced lymphadenopathy and elevated numbers of DN T cells compared with mice reconstituted with *FasL*<sup>Δm/Δm</sup> HSPCs (Supplementary Figure S3a and c). This indicates that loss of NF- $\kappa$ B2 in hematopoietic cells was sufficient to enhance the lymphadenopathy caused by defects in FASL.

**Loss of c-REL or NF- $\kappa$ B2 reduces total antibody and autoantibody levels in *FasL*<sup>Δm/Δm</sup> mice.** *FasL*<sup>Δm/Δm</sup> mice have elevated levels of antibodies of all Ig isotypes (Figure 3f and O'Reilly et al.<sup>8</sup>). In contrast, mice lacking NF- $\kappa$ B2 or c-REL have reduced levels of Ig isotypes requiring Ig heavy chain class switching (Figure 3f and Franzoso et al.<sup>25</sup>, Köntgen

et al.<sup>28</sup>). With the exception of IgA in *FasL*<sup>Δm/Δm</sup>*nfk2*<sup>-/-</sup> mice, loss of either NF- $\kappa$ B2 or c-REL in *FasL*<sup>Δm/Δm</sup> mice reduced all Ig isotype levels to those of wt mice (Figure 3f). High titers of antinuclear autoantibodies (ANA) producing a homogenous, nuclear staining pattern are a hallmark of SLE and have been observed in *FasL*<sup>Δm/Δm</sup> mice (Figures 3g–i and O'Reilly et al.<sup>8</sup>). Although the absence of NF- $\kappa$ B2 significantly reduced ANA levels in *FasL*<sup>Δm/Δm</sup> mice ( $P < 0.005$ ; Figures 3g–i), there was no difference in the Ig isotypes among the ANA (Supplementary Figure S4). Remarkably, the sera of *FasL*<sup>Δm/Δm</sup>*c-rel*<sup>-/-</sup> mice remained ANA negative (Figures 3g–i).

To determine whether lymphocytic tissue infiltration, another feature of the autoimmune pathology in *FasL*<sup>Δm/Δm</sup> mice,<sup>8</sup> was associated with the production of organ-specific autoantibodies, we assessed the reactivity of sera from sick

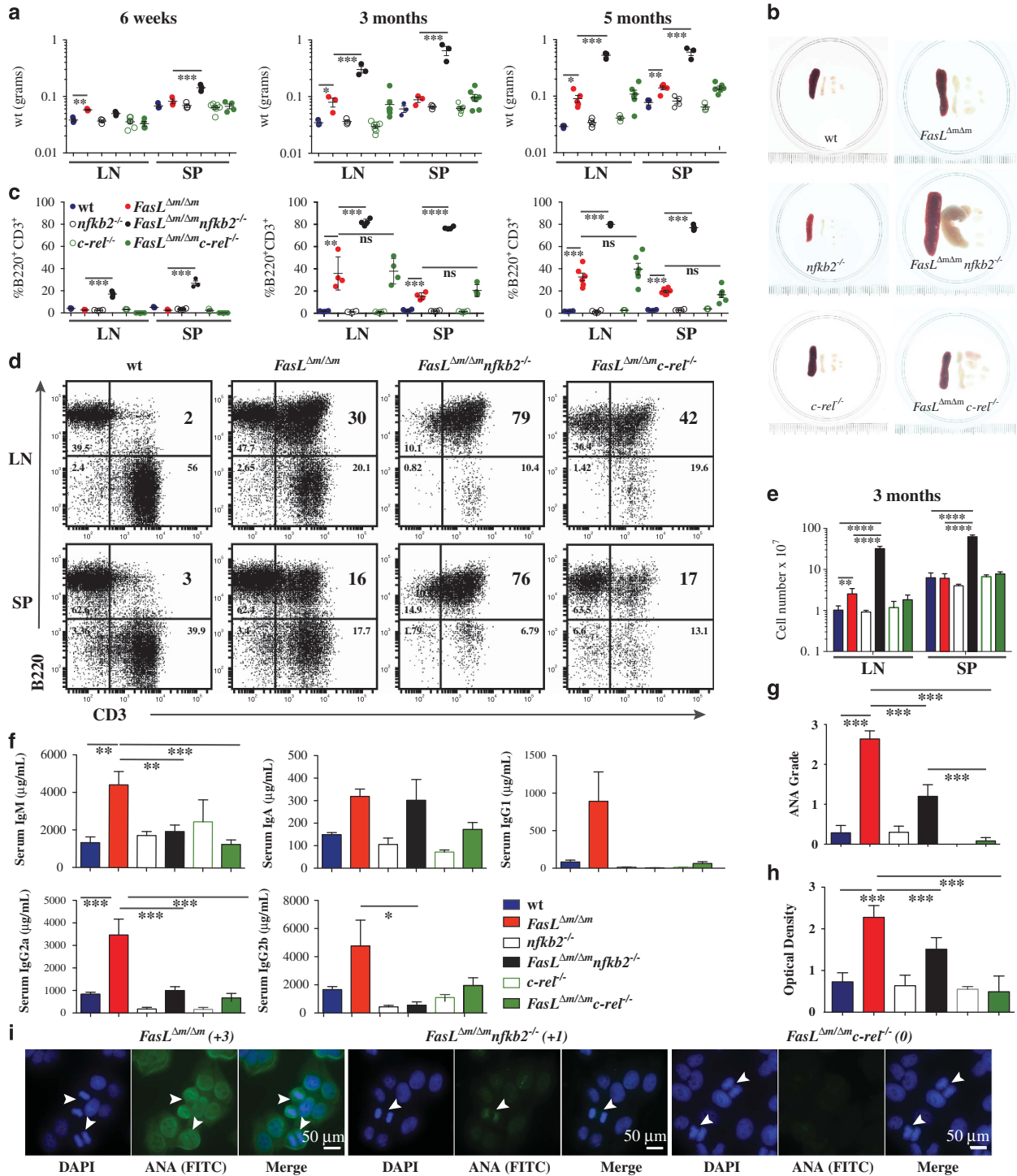


**Figure 2** Impact of loss of c-REL or NF- $\kappa$ B2 on the levels of pro-inflammatory cytokines and chemokines in *FasL* $\Delta m/\Delta m$  mice. (a) The levels of 31 cytokines and chemokines (indicated in the Materials and Methods) were measured in the sera of mice of the indicated genotypes by the multiplex system at 5 months when animals still appeared healthy or (b) when they were terminally ill or aged. The levels of IL-1 $\beta$ , IL-6, IL-17F, CCL4, TNF- $\alpha$  and CXCL1 are shown. Each dot represents a single mouse; the bar indicates the average; the error bars represent S.E.M.

animals against a panel of tissues from *rag-1* $^{-/-}$  mice. Sick *FasL* $\Delta m/\Delta m$  mice harbored autoantibodies (IgM, IgA or IgG) against lung epithelia, thyroid follicles, gastric parietal cells, submandibular gland and pancreatic acinar cells (Figure 3f). No such autoantibodies were detected in the sera from similarly aged wt or *c-rel* $^{-/-}$  mice. Sera from *nfkb2* $^{-/-}$  mice stained certain organs at a low level, and, although *FasL* $\Delta m/\Delta m$  *nfkb2* $^{-/-}$  animals generally had lower autoantibody levels than *FasL* $\Delta m/\Delta m$  mice, the persistent elevation of gastric autoantibodies may be due to the sustained serum IgA levels (Figure 3f). *FasL* $\Delta m/\Delta m$  *c-rel* $^{-/-}$  mice had a substantial reduction in IgM, IgA or IgG autoantibodies, particularly those directed against the submandibular gland and thyroid,

compared with *FasL* $\Delta m/\Delta m$  animals (Supplementary Figure S5, Table 1).

**The roles of NF- $\kappa$ B2 and c-REL in autoimmune pathology in *FasL* $\Delta m/\Delta m$  mice.** To further understand how NF- $\kappa$ B2 and c-REL influence the early phase of autoimmune pathology in *FasL* $\Delta m/\Delta m$  mice, we examined the lungs, liver, salivary glands and pancreas at 5 months. Although lung disease was not apparent in *FasL* $\Delta m/\Delta m$  mice, severe peribronchial and perivascular lymphocytic infiltrates were prominent in *FasL* $\Delta m/\Delta m$  *nfkb2* $^{-/-}$  mice and constituted the main pathology (Figures 4a and b). The infiltrates were predominantly composed of DN T cells with a polyclonal TCR $\beta$  variable



**Figure 3** Loss of NF- $\kappa$ B2 accelerated and enhanced lymphadenopathy and splenomegaly in *FasL<sup>Δm/Δm</sup>* mice and loss of NF- $\kappa$ B2 and c-REL both reduced hypergammaglobulinaemia and ANA in these animals. (a) Weights of the lymph nodes (axillary, brachial, inguinal and mesenteric) and spleens from mice of the indicated genotypes at the ages indicated. (b) Representative photomicrographs of the lymph nodes (axillary, brachial, inguinal and mesenteric) and spleen from mice of the indicated genotypes, all at ~6 months of age. Scale bar, each line represents 1 mm. (c) The percentages of the 'unusual' DN T cells in the lymph nodes of mice of the indicated genotypes were measured by FACS analysis. In a and c, each dot represents a single mouse and the figure legend in c also applies to a. (d) Representative flow cytometric images of the DN T cells (numbers indicate percentage of cells in each quadrant). (e) Graph of total cell numbers for the indicated organs and the indicated genotypes of mice at 3 months of age ( $n=3-4$  mice/genotype). (f) The levels of antibodies of the different Ig isotypes in the sera of the mice of the indicated genotypes (age 5 months). Data represent mean  $\pm$  S.E.M. ( $n=6$ /genotype). (g) ANA levels in the sera of mice of the indicated genotypes (5 months) were quantified by staining of human HEp-2 epithelial cells and scored according to a scale of fluorescence intensity on a scale of 0 (no fluorescence) to 3+ (maximal fluorescence). (h) Graphical representation of ANA levels in the sera of mice of the indicated genotypes quantified by ELISA ( $n=7-12$  mice/genotype). (i) Pictorial examples of ANA quantification in the sera of mice of the indicated genotypes; arrows indicate staining of nuclear components in mitotic figures. Immunofluorescence intensity score is indicated in brackets; ANA (green) and nuclei (blue) are revealed by staining with DAPI. Positive control sera were from *Fas<sup>gld/gld</sup>* mice. Values in graphs represent mean  $\pm$  S.E.M., \* $P<0.05$ , \*\* $P<0.001$ , \*\*\* $P<0.0001$

**Table 1** Percentage of mice with organ-specific autoantibodies against the tissues indicated

Strain	Salivary glands	Pancreas	Lacrimal glands	Lung	Liver	Thyroid	Stomach	n	Mean age
wt (aged)	0	0	20	0	0	ND	0	5	384 ± 55
<i>FasL</i> <sup><math>\Delta m/\Delta m</math></sup> (sick)	75	63	50	100	100	100	100	8	369 ± 97
<i>nfk2</i> <sup><math>-/-</math></sup> (aged)	25	25	13	38	25	0	25	8	392 ± 24
<i>FasL</i> <sup><math>\Delta m/\Delta m</math></sup> <i>nfk2</i> <sup><math>-/-</math></sup> (sick)	25	38	63	75	13	0	100	8	291 ± 79
<i>FasL</i> <sup><math>\Delta m/\Delta m</math></sup> <i>c-rel</i> <sup><math>-/-</math></sup> (aged)	0	0	27	0	0	0	12	8	435 ± 61

chain repertoire (Supplementary Figure S6), demonstrating their nonmalignant status.

Unlike sick *FasL* <sup>$\Delta m/\Delta m$</sup> *nfk2* <sup>$-/-$</sup>  mice, which exhibited extensive lymphoid infiltrates and damage in the lungs, liver, pancreas and salivary glands, similarly aged sick *FasL* <sup>$\Delta m/\Delta m$</sup>  ( $P < 0.0001$ ) and *FasL* <sup>$\Delta m/\Delta m$</sup> *c-rel* <sup>$-/-$</sup>  mice ( $P < 0.0001$ ) displayed milder pathology (Figure 4c). Aged or sick *nfk2* <sup>$-/-$</sup>  mice only exhibited mild organ leukocyte infiltration (*nfk2* <sup>$-/-$</sup>  versus wt,  $P < 0.0001$ ; *nfk2* <sup>$-/-$</sup>  versus *c-rel* <sup>$-/-$</sup>  mice,  $P < 0.05$ ). Many of the CD4<sup>+</sup>8<sup>-</sup> and CD4<sup>+</sup>8<sup>+</sup> T cells from the organs of aged or sick *FasL* <sup>$\Delta m/\Delta m$</sup> *nfk2* <sup>$-/-$</sup>  mice were CD69<sup>+</sup> (Figure 4d), indicating recent activation, possibly by self-antigens. Sick *FasL* <sup>$\Delta m/\Delta m$</sup> *nfk2* <sup>$-/-$</sup>  mice (but none of the animals of the other genotypes) also suffered from multiorgan necrotizing vasculitis, an inflammatory disease characterized by damaged vessel walls, involving the kidney and occasionally other organs (Figure 4e, Supplementary Table S1).

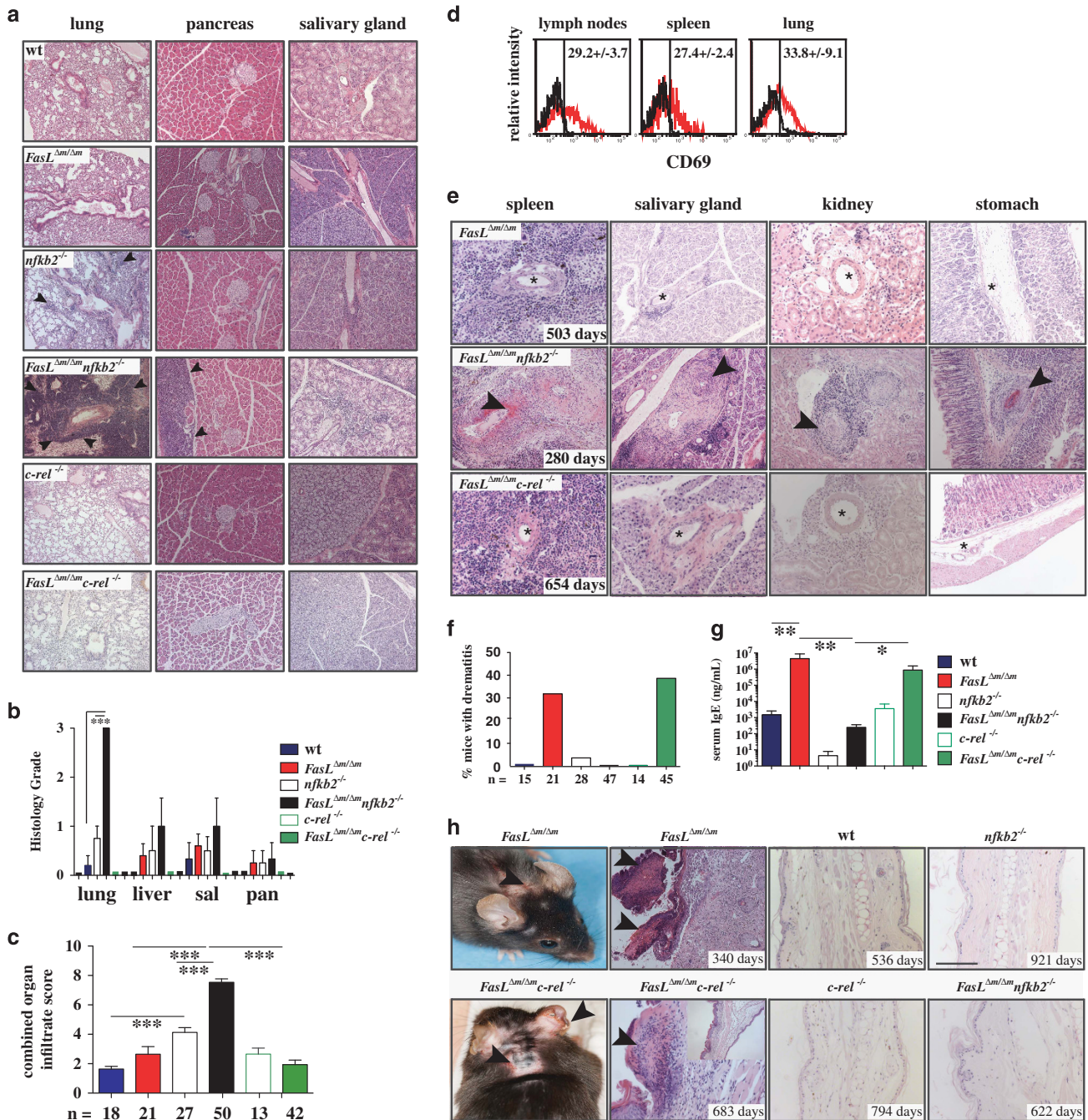
*FasL* <sup>$\Delta m/\Delta m$</sup>  mice develop autoimmune dermatitis (32%), which presents as reddening, thickening and scabbing of the ears and neck (Figure 4h and O'Reilly et al.<sup>8</sup>). Interestingly, *FasL* <sup>$\Delta m/\Delta m$</sup> *nfk2* <sup>$-/-$</sup>  mice did not develop autoimmune dermatitis, whereas *FasL* <sup>$\Delta m/\Delta m$</sup> *c-rel* <sup>$-/-$</sup>  mice showed a similar incidence of this disease (39%), albeit with delayed onset compared with the *FasL* <sup>$\Delta m/\Delta m$</sup>  animals (660 ± 134 days versus 435 ± 86 days) (Figures 4f and h). Dermatitis in *FasL* <sup>$\Delta m/\Delta m$</sup>  and *FasL* <sup>$\Delta m/\Delta m$</sup> *c-rel* <sup>$-/-$</sup>  mice was associated with elevated serum IgE levels (Figure 4g and O'Reilly et al.<sup>8</sup>), a feature shared with humans suffering from this autoimmune pathology.<sup>29</sup> These findings show that loss of c-REL merely delayed autoimmune dermatitis in *FasL* <sup>$\Delta m/\Delta m$</sup>  mice, whereas loss of NF- $\kappa$ B2 abrogated this pathology.

Regulatory T cells (Tregs) have a key role in the suppression of autoimmunity and modulation of the intensity and quality of immune responses to foreign antigens.<sup>30</sup> Of relevance to this study, reductions in Treg numbers or impaired Treg function have been implicated in murine and human autoimmune dermatitis.<sup>31,32</sup> To ascertain whether this was the case in *FasL* <sup>$\Delta m/\Delta m$</sup>  mice and whether this was influenced by the absence of NF- $\kappa$ B2 or c-REL, Tregs (CD4<sup>+</sup>CD25<sup>+</sup>Foxp3<sup>+</sup>) were enumerated in lymphoid tissues. Treg numbers in *FasL* <sup>$\Delta m/\Delta m$</sup>  and *nfk2* <sup>$-/-$</sup>  mice were normal (Figures 5a and b) and even elevated in the *FasL* <sup>$\Delta m/\Delta m$</sup> *nfk2* <sup>$-/-$</sup>  mice (Figure 5b). However, consistent with a previous report<sup>33</sup> showing that loss of c-REL resulted in a substantial reduction in Tregs, both the percentages and absolute numbers of Tregs in *FasL* <sup>$\Delta m/\Delta m$</sup> *c-rel* <sup>$-/-$</sup>  mice were markedly reduced compared with *FasL* <sup>$\Delta m/\Delta m$</sup>  and wt animals (Figures 5a and b). These results, combined with the impact that NF- $\kappa$ B2 or c-REL loss had on the phenotype in *FasL* <sup>$\Delta m/\Delta m$</sup>  mice, indicate that while Treg defects are not likely to be critical for the development of

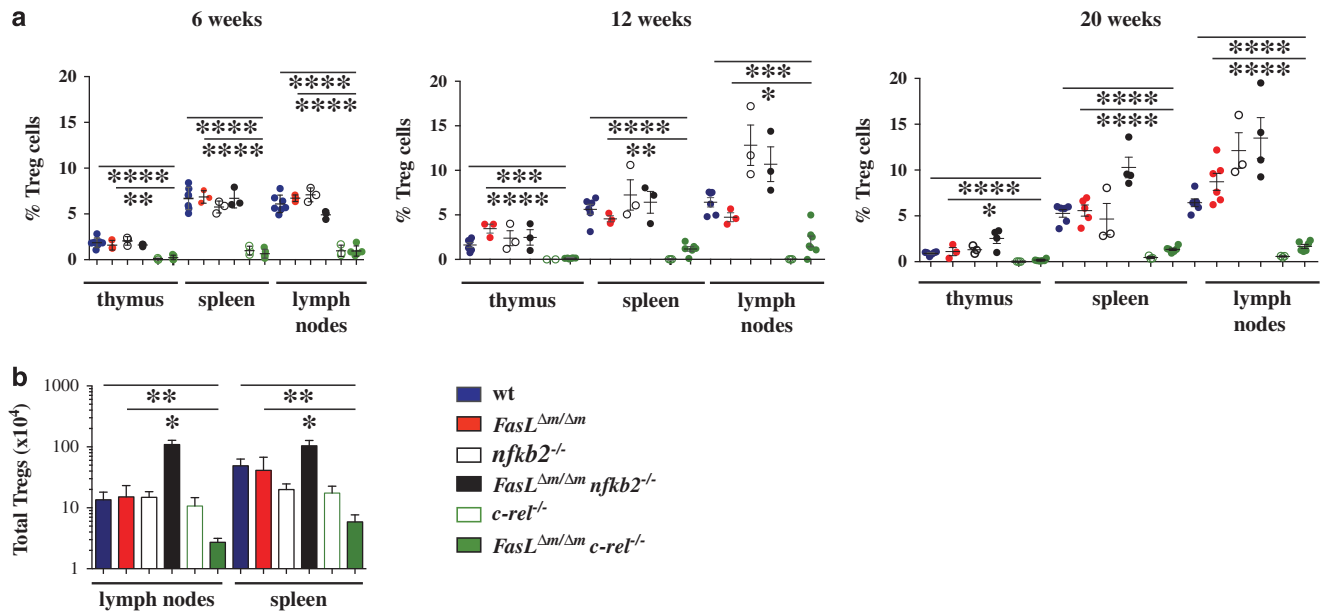
autoimmune pathology in these animals reduction in Treg numbers may explain in part the dermatitis seen in the *FasL* <sup>$\Delta m/\Delta m$</sup> *c-rel* <sup>$-/-$</sup>  mice.

**NF- $\kappa$ B2 but not mFASL or c-REL is required for the development of AIRE<sup>+</sup> mTECs.** The epithelial components of the thymic stroma have critical roles in promoting central tolerance to tissue-specific antigens.<sup>34</sup> A reduction in the number or function of medullary thymic epithelial cells (mTECs) is linked to T-cell-dependent inflammatory diseases triggered by autoreactive T cells that escape negative selection.<sup>34,35</sup> Both the canonical and non-canonical NF- $\kappa$ B pathways have been implicated in tolerogenic processes mediated by the thymic stroma,<sup>36</sup> with the non-canonical pathway reported to be important for the differentiation and/or survival of mTECs.<sup>34</sup> NF- $\kappa$ B2 deficiency was associated with a loss of CD80<sup>+</sup> mTEC, but not the reduction of mTECs that express the autoimmune regulator (AIRE).<sup>18</sup> We conducted a detailed analysis of the thymic microenvironment to examine whether mTEC defects might be responsible for the organ-specific autoimmunity observed in *FasL* <sup>$\Delta m/\Delta m$</sup>  mice or for the alterations in the *FasL* <sup>$\Delta m/\Delta m$</sup>  disease profile exerted by loss of NF- $\kappa$ B2 or c-REL (summarized in Table 2).

Thymi from 6- or 20-week-old *FasL* <sup>$\Delta m/\Delta m$</sup>  mice stained for mTEC markers (keratin-5 (K5), UEA-1 and AIRE) displayed a medullary architecture comparable to that of age-matched wt mice, although the levels of UEA-1 may be lower (Figures 6a and b). The expression of mTEC markers in the thymi of *c-rel* <sup>$-/-$</sup>  and *FasL* <sup>$\Delta m/\Delta m$</sup> *c-rel* <sup>$-/-$</sup>  mice was also normal (Figures 6a and b). In contrast, thymi from *nfk2* <sup>$-/-$</sup>  and *FasL* <sup>$\Delta m/\Delta m$</sup> *nfk2* <sup>$-/-$</sup>  mice exhibited reduced UEA-1 expression (from 6 weeks) and an abnormal distribution of K5<sup>+</sup> mTEC (from 20 weeks, Figures 6a and b). Flow cytometric analysis indicated that the total numbers of epithelial cells were similar across all genotypes (Figure 6c) and the total numbers of cortical TEC (cTEC) and mTEC subsets expressing high or low levels of MHCII (mTEC<sup>hi</sup> and mTEC<sup>low</sup>) were comparable between wt, *FasL* <sup>$\Delta m/\Delta m$</sup> , *c-rel* <sup>$-/-$</sup>  and *FasL* <sup>$\Delta m/\Delta m$</sup> *c-rel* <sup>$-/-$</sup>  mice (Figure 6d). However, the numbers of mTEC<sup>hi</sup> cells were substantially diminished in thymi of *nfk2* <sup>$-/-$</sup>  and *FasL* <sup>$\Delta m/\Delta m$</sup> *nfk2* <sup>$-/-$</sup>  mice (Figure 6e), reflected by an almost complete absence of AIRE<sup>+</sup> TECs (~10% of wt numbers). Interestingly, *FasL* <sup>$\Delta m/\Delta m$</sup>  mice also displayed elevated CD4<sup>+</sup>CD8<sup>+</sup> thymocyte numbers (Figure 6f). In summary, although *FasL* <sup>$\Delta m/\Delta m$</sup>  mice display many features of autoimmune disease, they have a normal TEC composition, indicating that mFASL has no major role in TEC development and survival. Loss of AIRE expressing mTECs, however, correlates with impaired negative selection and multiorgan autoimmunity<sup>34</sup> and this may provide an explanation for the mild age-related autoimmunity



**Figure 4** *FasL* <sup>$\Delta m/\Delta m$</sup> *nfk2*<sup>-/-</sup> mice developed more severe lymphoproliferative disease compared with *FasL* <sup>$\Delta m/\Delta m$</sup>  mice. (a) Representative photomicrographs of the lung, pancreas and salivary glands from mice of the indicated genotypes (5 months); arrows indicate areas of infiltration;  $n=4$ /genotype. (b) Summary (mean  $\pm$  S.E.M.) of the incidence of lymphocytic infiltration into the indicated organs of mice of the indicated genotypes (5 months of age), graded 0–3+, 3–5 mice/genotype;  $***P<0.0001$ . (c) Graphical depiction (mean  $\pm$  S.E.M.) of the combined scores for lymphocytic infiltration (lung, liver, pancreas, salivary and stomach), each scored 0–3+ from the indicated mice when sick/aged, from wt ( $555 \pm 176$  days), *FasL* <sup>$\Delta m/\Delta m$</sup>  ( $436 \pm 98$  days), *nfk2*<sup>-/-</sup> ( $627 \pm 182$  days), *FasL* <sup>$\Delta m/\Delta m$</sup> *nfk2*<sup>-/-</sup> ( $405 \pm 181$  days), *c-rel*<sup>-/-</sup> ( $790 \pm 84$  days) and *FasL* <sup>$\Delta m/\Delta m$</sup> *c-rel*<sup>-/-</sup> ( $603 \pm 116$  days) mice,  $***P<0.0001$ ,  $n$  = number of mice analyzed. (d) Expression of the activation marker CD69 on CD4<sup>+</sup> T cells (percentage of positive cells indicated) from the indicated organs from *FasL* <sup>$\Delta m/\Delta m$</sup> *nfk2*<sup>-/-</sup> mice (red line) and wt mice (black line);  $n=3$ /genotype. (e) Representative photomicrographs depicting necrotizing vasculitis in mice of the indicated genotypes. Arrows indicate areas of necrotizing vasculitis; \*indicates vasculature. Magnification  $\times 20$ . (f) Graphical representation of the percentages of mice of the indicated genotypes with autoimmune dermatitis (ears and neck);  $n$  = number of mice per genotype analyzed. (g) Serum IgE levels measured by ELISA when the animals were afflicted with dermatitis or otherwise sick. (h) Representative images, depicting autoimmune dermatitis either macroscopically or microscopically. Magnification  $\times 10$  or  $\times 40$ ; scale bar  $128 \mu\text{m}$



**Figure 5**  $CD4^+$  Tregs were not a limiting factor in the autoimmune disease of  $FasL^{\Delta m/\Delta m}$  mice. (a) The percentages of Treg ( $CD4^+CD25^+Foxp3^+$ ) were determined by flow cytometric analysis of the thymus, spleen and lymph nodes of mice of the indicated genotypes. Each dot represents a single mouse ( $n=3-4$ /genotype); bars indicate the average. (b) Absolute numbers of Tregs ( $CD4^+CD25^+Foxp3^+$ ) determined by flow cytometry ( $n=4$ /genotype at 3 months of age). The legend in b also applies to a. Error bars represent S.E.M., \* $P<0.05$ , \*\* $P<0.001$ , \*\*\* $P<0.0001$

**Table 2** Summary of the effects that loss of NF- $\kappa$ B2 or c-REL exerts on abnormalities in  $FasL^{\Delta m/\Delta m}$  mice

Genotype	Lifespan	Cytokine production	Lymphadenopathy splenomegaly	Organ infiltration	Serum IgG	ANA	Dermatitis
$FasL^{\Delta m/\Delta m}$ <sup>a</sup>	Reduced	Elevated	Present	Present	Elevated	Present	Present
$FasL^{\Delta m/\Delta m} nfkb2^{-/-}$ <sup>b</sup>	Minimal effect	No effect	Accelerated	Accelerated	Reduced	Reduced	Absent
$FasL^{\Delta m/\Delta m} c-rel^{-/-}$ <sup>b</sup>	Enhanced	Reduced	No effect	Reduced	Reduced	Reduced	Present

<sup>a</sup>Compared with wt. <sup>b</sup>Compared with  $FasL^{\Delta m/\Delta m}$

in  $nfkb2^{-/-}$  mice<sup>18,26</sup> and the exacerbated organ-specific lymphocyte infiltration observed in  $FasL^{\Delta m/\Delta m}nfkb2^{-/-}$  animals.

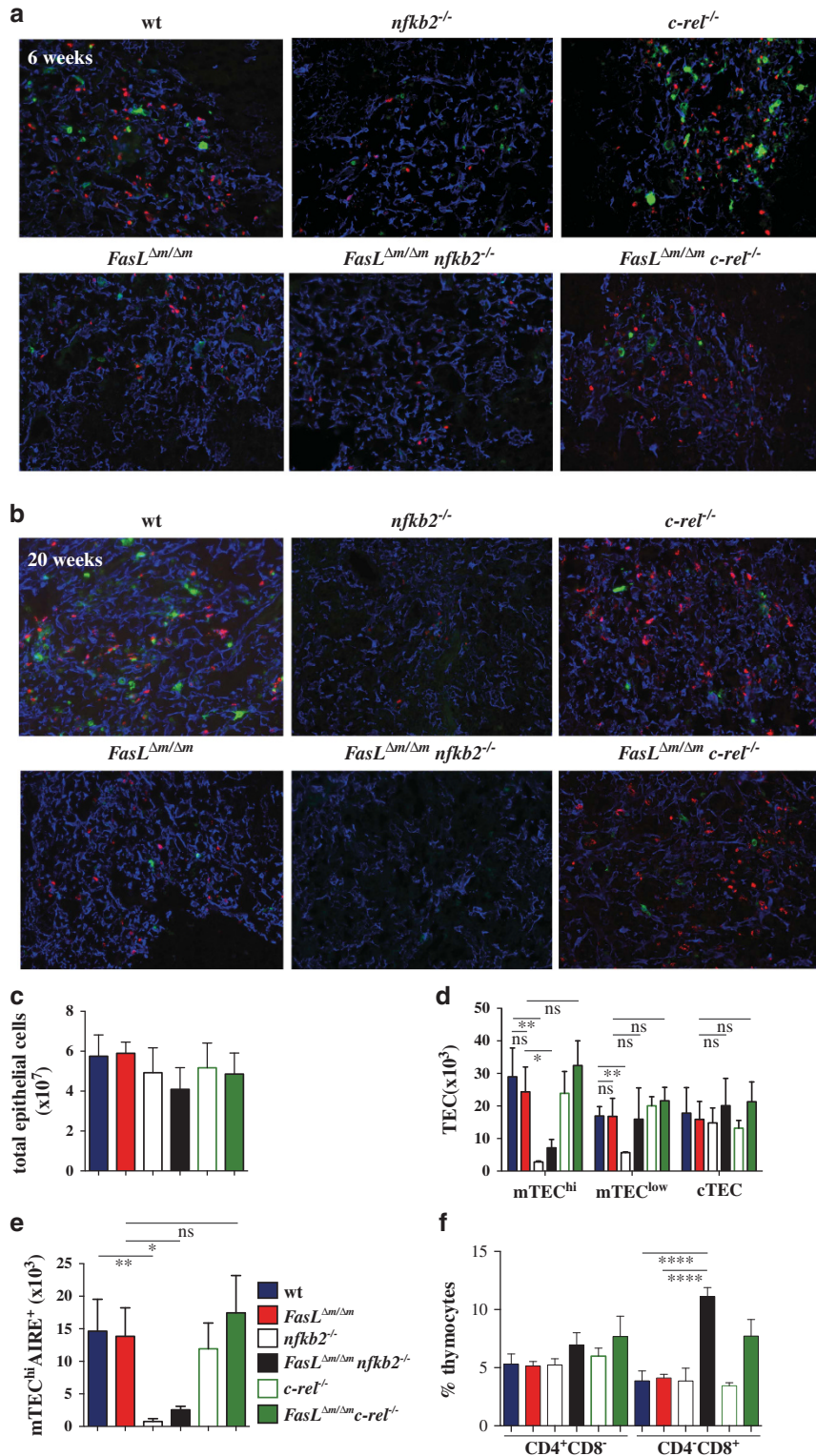
## Discussion

The many roles of canonical and non-canonical NF- $\kappa$ B signaling in the immune system identify constituents of these pathways as potential therapeutic targets for inflammatory and autoimmune diseases. Indeed, the immunosuppressive and anti-inflammatory activity of glucocorticoids, a treatment of choice for many autoimmune diseases, is at least in part due to attenuated NF- $\kappa$ B signaling.<sup>20</sup> However, as NF- $\kappa$ B signaling regulates a wide variety of biological processes both within and outside the immune system, prolonged blockade of overall NF- $\kappa$ B activity, not surprisingly, can cause severe adverse effects.<sup>37</sup> Accordingly, we assessed whether targeting individual NF- $\kappa$ B transcription factors would be a superior strategy for treating autoimmune disease.  $FasL^{\Delta m/\Delta m}$  mice were used as a model to determine the impact that the loss of c-REL (canonical pathway) or NF- $\kappa$ B2 (non-canonical pathway) had on their multifaceted autoimmune pathology. The largely hematopoietic restricted expression of c-REL<sup>21</sup> offers the added advantage of primarily targeting immune functions.<sup>22</sup>

Notably, targeting the canonical pathway in T cells using a proteasome degradation resistant form of I $\kappa$ Ba(I $\kappa$ B) afforded only minor protection from autoimmune disease in  $FasL^{gld/gld}$  mice.<sup>38</sup> However, given that NF- $\kappa$ B signaling was only diminished, not abrogated, combined with the essential loss of FAS function in B cells in this autoimmune disease,<sup>39</sup> the full impact of completely inhibiting a particular NF- $\kappa$ B transcription factor on autoimmune disease caused by defects in FASL/FAS remains unclear. Our study is the first detailed, long-term analysis of the roles that specific NF- $\kappa$ B family members have in the autoimmune disease caused by defects in FASL/FAS.

With respect to non-canonical NF- $\kappa$ B signaling, loss of NF- $\kappa$ B2 had contrasting impacts on different aspects of the  $FasL^{\Delta m/\Delta m}$  autoimmune disease. Autoantibody-mediated pathologies (dermatitis and severe GN) were reduced but the lymphoproliferative disease was enhanced, causing respiratory difficulties and multiorgan vasculitis. This establishes that NF- $\kappa$ B2-dependent signals restrain proliferation and/or survival of the lymphocytes that accumulate to excess as a consequence of defective FASL/FAS-mediated apoptosis. Our studies using chimeric mice indicate that NF- $\kappa$ B2 activity within the hematopoietic system is mainly responsible for restraining the proliferation and/or survival of these T lymphocytes, although non-hematopoietic functions regulated





**Figure 6** *FasL*<sup>Δm/Δm</sup> mice had normal mTEC numbers with normal AIRE expression but loss of NF- $\kappa$ B2 caused abnormalities in the thymic stroma. (a) Representative confocal images of thymi from mice of the indicated genotypes at 6 weeks. Sections were stained for UEA-1 (green), AIRE (red) and keratin-5 (blue); *n* = 3/genotype/time point. (b) Representative confocal images of thymi from mice of the indicated genotypes at 20 weeks as for (a). (c) Graphical representation of total thymic cellularity from mice of the indicated genotypes at 20 weeks of age; *n* = 4/genotype. (d) Graphical depiction of flow cytometric analysis of TECs from mice of the indicated genotypes at 20 weeks of age, as defined; mTEC<sup>hi</sup> (CD45<sup>-</sup> Ep-CAM<sup>+</sup>class II MHC<sup>hi</sup> Ly51<sup>-</sup>), mTEC<sup>low</sup> (CD45<sup>-</sup> Ep-CAM<sup>+</sup>class II MHC<sup>low</sup> Ly51<sup>-</sup>), cTEC (CD45<sup>-</sup> Ep-CAM<sup>+</sup>class II MHC<sup>+</sup> Ly51<sup>+</sup>); *n* = 4/genotype. (e) Graphical depiction of flow cytometric analysis of mTEC<sup>hi</sup>AIRE<sup>+</sup> mTECs (CD45<sup>-</sup> Ep-CAM<sup>+</sup>class II MHC<sup>hi</sup> Ly51<sup>-</sup> AIRE<sup>+</sup>) from mice of the indicated genotypes at 20 weeks of age. (f) Percentages of thymocytes that are CD4 single positive (CD4<sup>+</sup> CD8<sup>-</sup>) or CD8 single positive (CD4<sup>-</sup> CD8<sup>+</sup>) from mice of the indicated genotypes at 3 months of age; *n* = 4/genotype. \**P* < 0.05, \*\**P* < 0.005

by NF- $\kappa$ B2 may contribute. The NF- $\kappa$ B2 precursor p100, a protein with I $\kappa$ B-like activity,<sup>40</sup> limits NF- $\kappa$ B-dependent T-cell proliferation by preventing the nuclear activation of p50-NF- $\kappa$ B1/p65 heterodimers.<sup>41</sup> The absence of this inhibitory p100-NF- $\kappa$ B2 protein may be a mechanism that contributes to the increased accumulation of T cells in *FasL $\Delta m/\Delta m$ nfkB2 $^{-/-}$*  mice. The escape of T cells with higher affinity autoreactive TCRs due to impaired negative selection as a consequence of reduced numbers of AIRE<sup>+</sup> mTECS might also contribute to the enhanced accumulation of DN T cells in *FasL $\Delta m/\Delta m$ nfkB2 $^{-/-}$*  mice. Finally, the reduction in autoantibody-mediated aspects of *FasL $\Delta m/\Delta m$*  autoimmune disease associated with NF- $\kappa$ B2 loss can be explained by reduced BAFF-R survival signals in B cells and/or germinal center defects, both of which involve p52-NF- $\kappa$ B2/RELB dimers.<sup>19</sup> Overall, these findings suggest that, while systemic therapeutic targeting of NF- $\kappa$ B2 could alleviate autoantibody-induced pathology, this benefit would be negated by enhanced lymphoproliferative disease.

Our study also provides insight into the role of NF- $\kappa$ B2 in autoimmune dermatitis. Unlike *FasL $\Delta m/\Delta m$*  mice that develop IgE-dependent autoimmune dermatitis, the *FasL $\Delta m/\Delta m$ nfkB2 $^{-/-}$*  mice were protected. Consistent with NIK, the apical kinase in the non-canonical NF- $\kappa$ B2-dependent pathway<sup>42</sup> being critical for isotype switching to IgE,<sup>42</sup> our aged *nfkB2 $^{-/-}$*  mice had ~100-fold lower serum IgE levels compared with wt controls, and even in *FasL $\Delta m/\Delta m$ nfkB2 $^{-/-}$*  mice the serum IgE levels were still below those found in wt mice. Given that topical delivery of NF- $\kappa$ B decoy oligodeoxynucleotides alleviates murine atopic dermatitis,<sup>43</sup> our findings indicate that topical NF- $\kappa$ B2 inhibitors may represent a promising strategy for treating IgE-dependent autoimmune dermatitis. In contrast to a prior report, which showed that isotype switching to IgE was c-REL dependent,<sup>44,45</sup> we found that IgE levels were not markedly reduced by loss of c-REL on either a wt or a *FasL $\Delta m/\Delta m$*  background. The reason for these differences remains to be determined.

This report demonstrates that c-REL loss substantially prolongs the lifespan of *FasL $\Delta m/\Delta m$*  mice by preventing autoantibody-mediated pathology. Although c-REL is critical for normal T-cell activation through a combination of T-cell intrinsic and extrinsic processes,<sup>12,22,28</sup> lymphadenopathy was unaffected by loss of c-REL. This indicates that the accumulation of DN T cells is driven by other NF- $\kappa$ B proteins or is NF- $\kappa$ B independent. This disjunction between extended lifespan despite persistence of the lymphoproliferative component of the autoimmune disease elicited by defects in FASL/FAS is consistent with studies of gene-targeted mice lacking FAS in select cell populations.<sup>39</sup> Loss of c-REL has also been shown to prevent pathology in mouse models of collagen-induced arthritis,<sup>46</sup> streptozotocin-induced diabetes<sup>47</sup> and autoimmune encephalomyelitis.<sup>48</sup> These outcomes can be explained by the involvement of Th1 cytokines and/or IL-17 in these diseases, because their production and the differentiation of Th1,<sup>49</sup> as well as Th17<sup>48</sup> effector T cells, are c-REL-dependent. The fact that c-REL is also critical for cytokine production by macrophages, dendritic cells and NKT cells<sup>12</sup> is also likely to contribute to the reduced pathology in *FasL $\Delta m/\Delta m$ c-rel $^{-/-}$*  mice. Although we were unable to detect IFN $\beta$  in the sera of sick *FasL $\Delta m/\Delta m$* , *FasL $\Delta m/\Delta m$ nfkB2 $^{-/-}$*  and *FasL $\Delta m/\Delta m$ c-rel $^{-/-}$*  mice, others have reported that introduction of a

loss-of-function mutation of the type I interferon receptor decreased glomerular immune complex deposition in *Fas $^{lpr/lpr}$*  mutant mice.<sup>27</sup> However, no data on animal lifespan were provided and these mice were on a mixed genetic background; notably, genetic background greatly affects pathology in *Fas $^{lpr/lpr}$*  mice. Thus, the role of type I interferons in SLE-like disease in mice with defects in FASL or FAS remains unclear.

A requirement for c-REL in antigen receptor-induced T- and B-cell activation<sup>23</sup> is the most likely explanation for why the reduction in Treg numbers, caused by the loss of c-REL,<sup>33</sup> neither exacerbates autoimmune disease in *FasL $\Delta m/\Delta m$ c-rel $^{-/-}$*  mice nor instigates autoimmunity in *c-rel $^{-/-}$*  mice.<sup>23</sup> Interestingly, mice deficient for PELI1,<sup>50</sup> a ubiquitin E3-ligase that controls c-REL levels, develop an autoimmune phenotype similar to those observed in *FasL $\Delta m/\Delta m$*  mice, accompanied by nuclear accumulation of c-REL and excessive canonical NF- $\kappa$ B activation. As PELI1 appears to be critical for the maintenance of T-cell tolerance, it would be interesting to test whether c-REL dysregulation, specifically in T cells, is responsible for the SLE-like autoimmune disease in *FasL $\Delta m/\Delta m$*  mice.

In conclusion, our studies using *FasL $\Delta m/\Delta m$*  mice demonstrate that the canonical and non-canonical NF- $\kappa$ B pathways have critical, yet distinct roles in autoimmune and lymphoproliferative pathology. Although autoimmune diseases caused by inherited mutations in FASL or FAS (ALPS patients) are rare, subtle abnormalities in this apoptotic pathway arising from gene polymorphisms are likely to contribute to autoimmune disease predisposition more generally.<sup>51</sup> Combined with the recent discovery of links between polymorphisms in *c-Rel* and increased susceptibility to certain human autoimmune diseases,<sup>23</sup> our findings suggest that targeting c-REL constitutes an attractive strategy to alleviate or prevent autoimmune pathology, while avoiding the major side effects associated with pan-NF- $\kappa$ B inhibition.

## Materials and Methods

**Mouse.** All animal experiments were approved by the Walter and Eliza Hall Institute Animal Ethics Committee. *FasL $\Delta m/\Delta m$*  mice were generated on a C56BL/6 genetic background.<sup>8</sup> *nfkB2 $^{-/-}$* <sup>25</sup> and *c-rel $^{-/-}$*  mice<sup>22,28</sup> were generated on a mixed C57BL/6x129SV background and backcrossed onto a C57BL/6 background for >10 generations. *FasL $\Delta m/\Delta m$*  mice were intercrossed with *nfkB2 $^{-/-}$*  or *c-rel $^{-/-}$*  mice. The resultant *FasL $\Delta m/wt$ nfkB2 $^{+/-}$*  or *FasL $\Delta m/wt$ c-rel $^{+/-}$*  mice were intercrossed to produce *FasL $\Delta m/\Delta m$ nfkB2 $^{-/-}$*  or *FasL $\Delta m/\Delta m$ c-rel $^{-/-}$*  mice. Control wt, *FasL $\Delta m/\Delta m$* , *nfkB2 $^{-/-}$*  and *c-rel $^{-/-}$*  mice used in this study were maintained within the same colony.

**Histological analysis and scoring.** Mice were killed by CO<sub>2</sub> asphyxiation. Organs were removed and fixed in 80% Histochoice (Amresco, Solon, OH, USA) or 10% buffered formalin and paraffin embedded. Histological examination was performed by staining with hematoxylin and eosin (H&E). H&E-stained sections of the kidneys were examined for evidence of GN and scored on a scale of 0–4 (0 = normal; 1 = minor mesangial hypercellularity; 2 = moderate hypercellularity; 3 = severe hypercellularity; 4 = severe hypercellularity with obliteration and thickening of all capillary loops or the presence of fibrinoid necrosis or glomerular crescents). Lymphoid infiltration into organs (lung, liver, pancreas, submandibular gland and stomach) was assessed by examining H&E-stained sections and graded 0–3 (0 = none; 1 = occasional small perivascular foci (age related); 2 = more dense, well-defined perivascular and periductal foci; 3 = extensive infiltrate with parenchymal destruction). Organs were also examined and scored for the presence of necrotizing vasculitis. All photomicrographs were acquired using a  $\times 10$ /NA 0.3,  $\times 20$ /NA 0.50 or  $\times 40$ /NA 0.75 objective lens attached to an Axioplan 2 microscope (Carl Zeiss, North Ryde, NSW, Australia).

**Immunofluorescent staining and confocal microscopy.** The presence of autoantibodies against classical target tissues in autoimmune diseases (salivary gland, lacrimal gland, eye, kidney, liver, lung, pancreas, stomach and thyroid) was assessed by staining frozen tissue sections (5  $\mu$ m, air dried and fixed in acetone) of *rag-1*<sup>-/-</sup> mice (to eliminate endogenous Ig, a source of background staining) with sera (1/100 dilution) from aged or sick mice. Staining was detected with FITC conjugated goat antibodies against mouse IgG, IgA or IgM (Cappel (MP Biochemicals), Santa Ana, CA, USA). To stain for immune complex deposits, the kidneys from aged or sick mice were snap-frozen in isopentane, sectioned, acetone fixed, and blocked with PBS/2% FCS, followed by staining with FITC-coupled goat antibodies specific to mouse IgM, IgG or IgA (Southern Biotechnology, Birmingham, AL, USA). All immunofluorescent staining was analyzed using a Leica laser scanning (SP2) confocal microscope.

Frozen sections of thymi from 6- or 20-week-old mice were cut at -20 °C on a cryostat (5  $\mu$ m), air dried and fixed in acetone. Sections were incubated with primary mAbs in 0.5%FCS/PBS for 30 min in a moist chamber, washed three times in PBS and incubated, while protected from light, with the appropriate secondary Abs plus DAPI, followed by washing. Primary Abs included biotin-conjugated anti-JEA-1 (Vector Laboratories, Burlingame, CA, USA), rabbit anti-keratin-5 (MK5, Covance, Biologend, San Diego, CA, USA) and Alexa 405-conjugated anti-AIRE. Detection was achieved using Alexa Fluor 555-conjugated goat anti-rabbit IgG antibodies or FITC-conjugated Streptavidin (both from Invitrogen, Carlsbad, CA, USA). Staining was analyzed using a Zeiss LSM5 line scanning confocal microscope (Carl Zeiss).

ANA in sera from aged or sick mice was detected by staining (1/100 serum dilution) of slides coated with HEp-2 human epithelial cells according to the manufacturer's instructions (Immuno Concepts, Sacramento, CA, USA) and detected with FITC-conjugated rabbit antibodies against all mouse Igs (Cappel (MP Biochemicals)) or FITC-conjugated rabbit antibodies selective for mouse IgM, IgG or IgA (Southern Biotechnology). ANA levels were quantified according to the brightness of fluorescence intensity on a scale of 0 (no fluorescence) to 3+ (maximum fluorescence intensity). Serum ANA levels were also measured by ELISA (The Binding Site, Birmingham, UK) according to the manufacturer's instructions.

**Hematopoietic reconstitution.** Ablation of the hematopoietic compartment was achieved by exposing mice to two doses of 5.5 Gy of  $\gamma$ -irradiation from a <sup>60</sup>Co source, 2 h apart. Irradiated mice were reconstituted by intravenous (i.v.) injection of 10<sup>6</sup> total bone marrow cells in a 100  $\mu$ l volume of PBS. Reconstituted mice were maintained on 2 mg/ml neomycin-supplemented water for 21 days.

**Flow cytometric analysis of thymic stromal cell populations.** This procedure was adapted from previously described protocols.<sup>52,53</sup> Thymi were processed individually for digestion as described.<sup>53</sup> First, thymi were cleaned of fat and connective tissue, the lobes separated and the capsule nicked with fine scissors. Thymi were then individually agitated in 10 ml RPMI-1640 medium for several minutes to gently flush out thymocytes using a wide-bore pipette. The remaining thymic tissue was then incubated at 37 °C for 15 min in 1 ml of Liberase/DNAase I (0.5 units and 0.1%, respectively, both from Roche, Germany) in RPMI-1640 medium with regular agitation, as detailed.<sup>54</sup> Fragments were allowed to settle and the supernatant was collected (fraction 1), kept on ice and the digestion process repeated with the settled thymic fragments. After a second digestion (fraction 2), the remaining aggregates were centrifuged, and resuspended in PBS with 5 mM EDTA, 2% FCS and 0.02% Na<sub>2</sub>S<sub>2</sub>O<sub>3</sub>. Cells were then filtered through a 100  $\mu$ m mesh and counted using a Countless automated cell counter (Invitrogen). For phenotypic analysis, 1.5  $\times$  10<sup>7</sup> cells were stained.

**Flow cytometric analysis.** Single-cell suspensions of the spleen and lymph nodes (pooled axillary, brachial, inguinal and mesenteric) were stained as described<sup>1</sup> with FITC-, R-PE- or APC-conjugated mAbs specific for mouse cell surface markers: RB6-8C5: anti-Gr-1, MI/70: anti-Mac-1, M3/84.6.34: anti-Mac-2, Ter119, F4/80, T24.31.2: anti-Thy-1, GK1.5: anti-CD4, 53.6.72: anti-CD8, RA3-6B2: anti-CD45R-B220 and PK136: anti-NK1.1 (BD Biosciences, Franklin Lakes, NJ, USA). The vital dye PI (1  $\mu$ g/ml) was included and data were acquired on a FACScan (Becton Dickinson, Franklin Lakes, NJ, USA) or LSR II (Becton Dickinson). Tregs were detected after fixation and permeabilization (eBiosciences) by staining with anti-CD25-R-PE, anti-CD4-FITC and anti-Foxp3-APC antibodies, followed by data acquisition on a LSR II (Becton Dickinson). Single-cell suspensions were prepared from the lungs by digestion for 20 min at room temperature with collagenase/DNAase (7 mg collagenase (Worthington, Lakewood, NJ, USA) in 1 ml volume of 0.1% w/v DNAase1 (Roche)) treated with EDTA as described,<sup>55</sup> and then

stained with an R-PE-coupled antibody to Thy-1 plus a panel of biotinylated antibodies specific to different TCRV $\beta$  regions followed by staining with FITC-coupled streptavidin (BD-Pharmingen).

Phenotypic analysis of thymic stromal cells by flow cytometry was performed using the following antibody conjugates: PerCP-cy5.5 anti-CD45.2 (clone 30F11, Biologend, San Diego, CA, USA) and PE anti-Ly51 (Biologend). APC anti-AIRE, FITC anti-MHC class II (H-2A/E, clone M5 114/15.2) and APC-cy7 anti-EpCAM (clone G8.8a) were produced in-house.

**Measurement of serum immunoglobulin levels.** Serum immunoglobulin levels were determined by ELISA using sheep anti-mouse Ig antibodies (Silenus Laboratories, Australia) as a capture reagent and developed with mouse Ig isotype-specific goat antibodies conjugated to horseradish peroxidase (Southern Biotechnology). Purified myeloma proteins were used as standards (Sigma, MO, USA).

**Analysis of serum cytokine and chemokine levels.** Cytokine and chemokine levels in sera were measured by using Bio-Plex Pro mouse cytokine 23-plex (Bio-Rad, Gladesville, NSW, Australia) and Bio-Plex Pro mouse cytokine Th17-plex magnetic bead immunoassays on a Bio-Rad Bio-Plex instrument, following the manufacturer's instructions. The cytokines measured included the following: CD40L, IL-1 $\alpha$ , IL-1 $\beta$ , IL-2, IL-3, IL-4, IL-5, IL-6, IL-9, IL-10, IL-12(P40), IL-12(p70), IL-13, IL-17A/F, IL-21, IL-22, IL-23(p19), IL-31, IL-33, eotaxin, G-CSF, GM-CSF, IFN- $\gamma$ , KC (CXCL1), MCP-1, MIP-1 $\alpha$ , MIP-1 $\beta$ , MIP-3 $\alpha$ , RANTES and TNF $\alpha$ . Serum IFN $\beta$  levels were measured using the VeriKine-HS (PBL Assay Science, Piscataway, NJ, USA) Mouse IFN- $\beta$  serum ELISA kit, according to the manufacturer's instructions.

**Statistical analysis.** Statistical analysis was performed using Student's *t*-test, log rank (Mantel-Cox) test for animal survival curves, one-way analysis of variance using Tukey's comparison test to compare multiple groups or the Mann-Whitney test where appropriate.

### Conflict of Interest

The authors declare no conflict of interest.

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