

Endosomal TLR signaling is required for anti-nucleic acid and rheumatoid factor autoantibodies in lupus

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Using the *Unc93b1 3d* mutation that selectively abolishes nucleic acid-binding Toll-like receptor (TLR) (TLR3, -7, -9) signaling, we show these endosomal TLRs are required for optimal production of IgG autoAbs, IgM rheumatoid factor, and other clinical parameters of disease in 2 lupus strains, B6-*Fas*^{lpr} and BXSB. Strikingly, treatment with lipid A, an autoAb-inducing TLR4 agonist, could not overcome this requirement. The *3d* mutation slightly reduced complete Freund's adjuvant (CFA)-mediated antigen presentation, but did not affect T-independent type 1 or alum-mediated T-dependent humoral responses or TLR-independent IFN production induced by cytoplasmic nucleic acids. These findings suggest that nucleic acid-sensing TLRs might act as an Achilles' heel in susceptible individuals by providing a critical pathway by which relative tolerance for nucleic acid-containing antigens is breached and systemic autoimmunity ensues. Importantly, this helps provide an explanation for the high frequency of anti-nucleic acid Abs in lupus-like systemic autoimmunity.

autoimmunity | SLE | *Unc93b1* | innate immunity

Systemic lupus erythematosus (SLE) is characterized by autoAbs to nuclear and cytoplasmic material that contain RNA, DNA, or both. AutoAbs typically arise before overt manifestations of disease, and high titers of anti-dsDNA are associated with greater severity (1, 2). Similar findings are observed in lupus-prone mice and, importantly, passively administered anti-DNA mAbs can produce lupus-like immune complex kidney deposits (3). Thus, evidence points to a direct role of anti-nuclear Abs in SLE.

Recent studies show that production of anti-nuclear Abs depends, to varying degrees, on endosomal Toll-like receptors (TLRs) that bind dsDNA (TLR9) or ssRNA (TLR7) (4, 5). Indeed, in vitro experiments have consistently found that either TLR can enhance activation of B cells and dendritic cells (DC) following antigen receptor (B cell receptor, BCR)- or FcγRIIa (FcγRIII in mice)-mediated endocytosis of nucleic acid-containing material or immune complexes (4). Remarkably, chromatin-containing immune complexes, presumably because of combined engagement of BCR and TLR, can stimulate B cells 100-fold more effectively than complexes without nucleic acids (6). On the basis of these findings, Leadbetter et al. (4, 7) proposed the novel hypothesis that these processes might explain the induction and prevalence of anti-nuclear Abs in lupus. Interestingly, these TLRs are not only B cell activators, but also potent inducers in DCs and plasmacytoid (p)DCs of the SLE-promoting type I interferons (IFNs) (8, 9).

When lupus-prone mice were examined, however, TLR9 deficiency had mixed effects on anti-DNA or anti-chromatin and, to a lesser extent, end-organ damage. In MRL-*Fas*^{lpr} mice, lack of TLR9 had different effects on anti-nuclear Ab specificity in different studies, yet overall disease was inexplicably exacerbated (10–12). TLR9-deficiency also enhanced disease in MRL-*Fas*^{wt} (11), B6-*Fas*^{lpr} (13), and mutant *Plcg2*^{+/*Alti5*} mice (14). In striking contrast, lupus-prone FcγRIIB^{-/-} mice expressing a high-affinity anti-DNA transgene had reduced disease (15), although this may be related to dependence on a single trans-

genic anti-DNA specificity. Taken together, despite uncertainty about the role of TLR9 in the production of anti-DNA-related Abs, the majority of studies indicate that TLR9 has an overall lupus-suppressing function.

In contrast, studies have consistently documented a lupus-promoting function for TLR7, most clearly shown by the discovery that an extra copy of TLR7 on the Y chromosome of BXSB mice explained the lupus-enhancing *Yaa* mutation (16–19). Moreover, TLR7-deficient MRL-*Fas*^{lpr} mice had reduced anti-RNP, less lymphoproliferation, and a slightly lower composite renal disease score, but similar levels of anti-nucleosome and anti-dsDNA (12). Thus, in MRL-*Fas*^{lpr} mice, TLR7 appears to play a major role in the induction of RNA-related autoAbs, but only a modest role in overall disease severity. In the tetramethylpentadecane-induced model of lupus, TLR7 is also required for the production of both RNP autoAbs and disease-promoting type I IFNs (20). In contrast, TLR3 deficiency did not significantly affect autoAbs, lymphoproliferation, or glomerulonephritis (GN) in MRL-*Fas*^{lpr} mice (12).

A possible limitation of studying single TLR deletions, however, is that immune complexes of nucleic acid-containing material, such as apoptotic debris, are likely to contain both RNA and DNA. Therefore, Abs to either DNA or RNA could potentially form complexes that activate both TLR7 and TLR9, and deleting any one of these TLRs would provide only partial, if any, inhibition. Thus, assessing the impact of completely blocking all nucleic acid-sensing TLRs on autoAb production and lupus pathogenesis is important. To this end, we studied the induction and progression of lupus-like disease in mice in which endosomal TLR (eTLR) signaling was abolished by the *3d* mutation in *Unc93b1* (21). *Unc93b1* encodes an endoplasmic reticulum (ER)-resident protein that physically associates with TLR3, -7, and -9 and is required for the trafficking of these TLRs from the ER to the endolysosomes where encounter with their cognate ligands occurs (22). We found that the *3d* mutation virtually abolished IgG anti-nuclear Abs and markedly reduced disease in 2 different lupus strains, providing direct evidence that signaling by self-nucleic acid-recognizing TLR is central to the production of autoAbs to nucleic acid-containing material and disease pathogenesis.

Results

The *3d* Mutation Reduces IgG AutoAbs and Lymphoproliferation in B6-*Fas*^{lpr} Mice. To study the effects of blocking TLR3, -7, and -9 in lupus, we backcrossed the *3d* mutation onto B6-*Fas*^{lpr} mice, which develop significant lymphoproliferation and autoAbs, but

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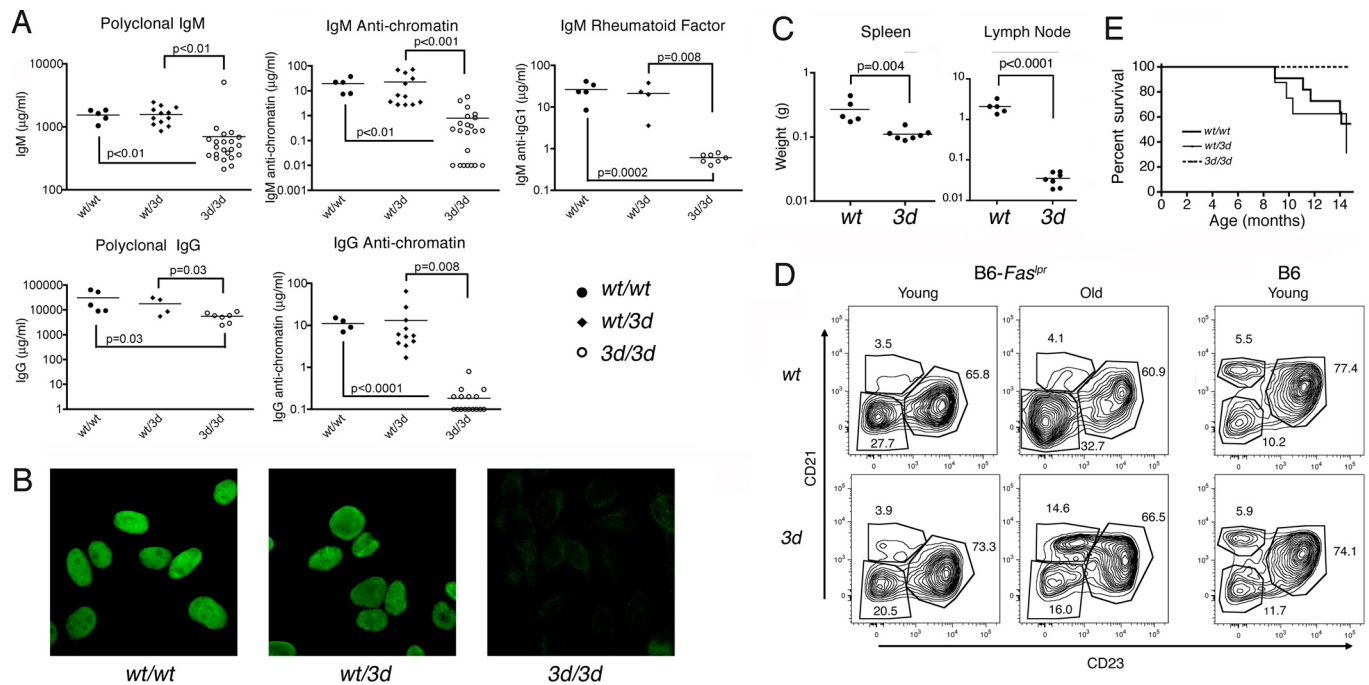


Fig. 1. Immunopathology of B6-*Fas*^{lpr} 3d mice. (A) IgM and IgG polyclonal and autoAbs from 6- to 8-month-old B6-*Fas*^{lpr} wild-type (*wt/wt*), heterozygous (*wt/3d*), and mutant (*3d/3d*) mice determined by ELISA. IgM RF was anti-IgG1, 4–22 mice/group. (B) Representative ANA results from 6- to 7-month-old mice (1/100 dilution), 4–7/group. (C) Lymphoid organ weights: spleen and LN (cervical, axillary, inguinal, and mesenteric) weights from 10-month-old mice, 5–7/group. (D) Representative flow cytometry analyses of splenic B cell subsets in young (1 month) or old (15 month) mice with the *Fas*^{lpr} and 3d mutations. Follicular (CD21^{lo} CD23^{hi}), marginal zone (CD21^{hi} CD23^{lo}), and CD21^{lo} CD23^{lo} populations are gated (see Table S3). (E) Cumulative survival, 8–11/group. $P < 0.04$ for 3d/3d versus *wt/wt* or *wt/3d*.

have limited susceptibility to end-organ injury. Compared with wild-type (*wt*) or heterozygous (*wt/3d*) animals, 3d B6-*Fas*^{lpr} mice had modest suppression of polyclonal IgM and IgG levels, but much greater suppression of IgM and IgG anti-chromatin [Fig. 1A, supporting information (SI) Fig. S1]. Strikingly, while all *wt* B6-*Fas*^{lpr} mice were strongly positive for anti-nuclear Abs (ANA), 6- to 8-month-old 3d mice were all negative, and only 2 of 7 of the 14- to 15-month-old 3d mice had weak ANA staining (Fig. 1B, Table S1). Moreover, these weaker ANAs exhibited atypical patterns: one with essentially metaphase chromosome staining and the other similar to human anti-proliferating cell nuclear antigen (anti-PCNA). Also, another 14- to 15-month-old B6-*Fas*^{lpr} 3d mouse was ANA negative, but had Golgi apparatus staining (Table S1). IgM rheumatoid factor (RF), another autoAb found in high titers in this strain, was also markedly lower in 3d mice (Fig. 1A, Fig. S1). Thus, eTLRs appear to play a critical role in the production of classical anti-nuclear and RF Abs.

The lack of eTLRs in B6-*Fas*^{lpr} 3d mice also significantly suppressed lymphoproliferation with substantial reductions in splenomegaly and especially in the characteristic aggressive lymphadenopathy (Fig. 1C). B6-*Fas*^{lpr} mice do not develop significant GN, but succumb to complications secondary to massive lymphoproliferation. Highlighting the significant enhancement of long-term survival imparted by the effect of 3d mutation on lymphoid hypertrophy, mortality at 14–15 months of age was 45% in *wt/wt*, 50% in *wt/3d*, and 0% in 3d/3d B6-*Fas*^{lpr} ($P < 0.04$ for 3d/3d compared with either *wt/wt* or *wt/3d*, Fig. 1E). Substantial lymphadenopathy, however, was readily detectable in 14- to 15-month-old 3d mice, indicating that the reduction in lymphoproliferation was not from correction of the defective *Fas*-mediated apoptosis *per se*, but from the lack of self and foreign nucleic acid sensing by eTLRs (Table S2).

When T cell subsets in the spleen and lymph node (LN) of young mice (1 month) were characterized, the only major difference was an $\approx 30\%$ reduction in the CD44^{hi} subset of CD8

T cells (Table S2). In contrast, old 3d mice (≈ 15 months) had a lower percentage of T cells in the spleen and in the LNs, reduced percentages of CD4 and CD8 T cells associated with an increased percentage of double negative (DN, CD4⁻CD8⁻) T cells. In both spleen and LNs, reductions in the naive CD62L⁺ subset of CD8⁺ T cells were also observed. Analysis of splenic B cells in young mice showed a significant 27% decrease in the CD21^{lo} CD23^{lo} population, which is expanded in *Fas*^{lpr} mice (Table S3, Fig. 1D). In old 3d mice, this population was reduced even more (77% decrease) and the percentage of CD138⁺ plasma cells was lower (65% decrease). These findings are consistent with suppression of autoimmunity in older 3d B6-*Fas*^{lpr} mice.

Reduced Lupus Pathology in 3d Male BXSB Background Mice. The 3d mutation was next backcrossed to the lupus-prone BXSB background to further examine the effects of blocking nucleic acid-sensing TLRs on systemic autoimmunity. Similar to the B6-*Fas*^{lpr} 3d mice, 3d-deficient male BXSB mice had marked reductions in autoAbs, including IgM and IgG anti-chromatin, -ssDNA, and -dsDNA and IgG anti-ribonuclear protein (RNP) (Fig. 2A and B). The reductions were greatest for IgG autoAbs, and ANAs were undetectable (Table S4). GN in 3- to 4-month-old mice was also suppressed (kidney disease score: *wt*, 2.4 ± 0.7 ; 3d, 0.4 ± 0.4 ; $P = 0.04$, $n = 4$ –5/group) as reflected by a 100% survival in the BXSB 3d group up to 18.6 weeks compared to a median survival of 11.6 weeks in *wt* mice (Fig. 2C). Thus, eTLR signaling was critical for both anti-nuclear production and disease-associated pathology and mortality.

Effect of TLR4 Stimulation of B6-*Fas*^{lpr} 3d Mice on AutoAb Production. To further study the role of eTLRs in the generation of anti-nuclear Abs, we treated 3d mutant B6-*Fas*^{lpr} mice with the TLR4 ligand, lipid A, a nonimmunogenic form of LPS. TLR4 signaling is not affected by the 3d mutation (21) and its engagement, like that of eTLRs, significantly enhances disease in lupus-prone strains,

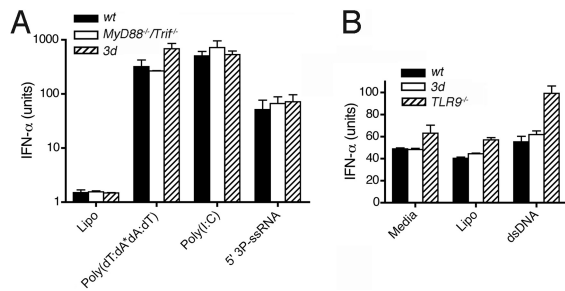


Fig. 5. TLR-independent activation of DCs. (A) DC activation with cytoplasmic dsRNA and 5' (3P)-RNA. Bone marrow (BM)-derived *wt*, *3d*, and *MyD88^{-/-}/Trif^{-/-}* double-knockout DCs were transfected with 250 μ g/mL poly(I:C) alone or by Lipofectamine with 10 μ g/mL poly(dT:dA:dT) or 200 ng 5' (3P)-dsRNA. Poly(I:C) in high concentrations directly enters cells. $P < 0.009$ for all *wt*, *3d*, and *MyD88^{-/-}/Trif^{-/-}* double-knockout DCs transfected with RNA versus Lipofectamine-treated (Lipo) DCs. (B) DC activation with cytoplasmic double-stranded mammalian DNA. DCs from *wt*, *3d*, and *TLR9^{-/-}* mice were transfected with 10 μ g/mL calf thymus DNA. $P < 0.05$ for dsDNA-transfected *wt*, *3d*, and *TLR9^{-/-}* DCs versus Lipo alone DCs. A and B are mean \pm SEM of triplicates.

The TD response was assessed in mice immunized on days 0 and 21 with TNP-KLH plus either of 2 types of adjuvants, alum or sequential complete Freund's adjuvant (CFA) and incomplete Freund's adjuvant (IFA) (CFA/IFA) (Fig. 4B). Both alum and CFA/IFA induced IgM and IgG responses after the initial and recall immunizations in both *wt* and *3d* mice, although Ab concentrations were modestly higher with CFA/IFA, particularly for the IgG isotype. With alum, the overall polyclonal and TNP-specific IgM and IgG levels in *wt* and *3d* mice were very similar after both the first and the second immunizations, indicating no impairment of T cell helper activity. With CFA/IFA, however, levels of Abs in *3d* mice, particularly those of the IgG isotype, were often slightly, but nevertheless significantly ($P < 0.05$), lower than corresponding concentrations in *wt* mice. Thus, a slight reduction in TD humoral response in *3d* mice was detected with CFA, an adjuvant that contains ligands for the nucleic acid-sensing TLRs, but not with alum, which lacks nucleic acids.

To examine the relative roles of *3d* antigen-presenting cells (APCs) and T cells in recall T cell activation, we used antigen (ovalbumin, OVA) plus APCs from either *wt* or *3d* *B6-Fas^{lpr}* mice to stimulate *wt* or *3d* T cells (Fig. 4C). As previously reported (21), *3d* APCs were less effective than *wt* APCs in stimulating *wt* T cells and *3d* T cells, although interestingly, *3d* T cells had slightly greater thymidine incorporation than *wt* T cells regardless of whether the APCs were *wt* or *3d*. Thus, *3d* T cells are fully capable of responding to Ags despite reduced APC function.

3d Mutation Does Not Affect TLR-Independent Nucleic Acid-Induced Cell Activation. Recently described cytoplasmic nucleic acid receptor or signaling molecules, such as retinoic acid-inducible gene I (RIG-I, sensing 5' triphosphate RNA), melanoma differentiation-associated gene 5 (Mda5, sensing dsRNA), absent in melanoma 2 (AIM2, sensing DNA), and stimulator of IFN genes (STING, sensing B-DNA), can also in some cases activate cells to produce proinflammatory responses, such as IFN- α/β , and to upregulate costimulatory molecules (24–32). Because the role of these cytoplasmic sensors in lupus is not known, we sought to determine whether the *3d* mutation affected cytoplasmic nucleic acid recognition. When *wt*, *3d*, *MyD88^{-/-} Trif^{-/-}* double knockout, or *TLR9^{-/-}* DCs were transfected with 5' triphosphate RNA, dsRNA, or mammalian dsDNA, no suppression of TLR-independent cytokine production by the *3d* mutation was detected (Fig. 5A and B). Thus, the effects of the *3d* mutation on

autoAb production and other lupus manifestations cannot be attributed to defective TLR-independent nucleic acid sensing.

Discussion

Herein we show that the *Unc93b1^{3d}* mutation, which abolishes nucleic acid-sensing TLR signaling, markedly suppressed spontaneous anti-nuclear, anti-RNP, and RF Ab production, GN, and mortality in lupus-prone strains. Moreover, this suppression was not overcome by treating mice with lipid A, a TLR4 agonist that promotes autoAbs and lupus. We further show that *3d* had no effect on TI-1 humoral responses or TD Ab responses with alum as the adjuvant, but slight, although significant, reducing effects in CFA/IFA-mediated TD responses. TLR-independent nucleic acid sensing was also unaffected by the *3d* mutation. Thus, eTLRs are largely dispensable for humoral responses to foreign protein antigens (Ags), but are, for all practical purposes, necessary for the generation of autoAbs to nucleic acid and nucleic acid-containing material and IgM RF in systemic autoimmunity. These results definitively demonstrate that the eTLRs play a key and essential role in the pathogenesis of lupus in 2 susceptible strains and, in conjunction with previous studies (6, 7, 12, 18, 20, 33), provide an explanation for the frequent and dominant presence of ANAs in SLE.

Previous studies, showing reduced APC function in *3d* mice, suggested that *Unc93b1* might play a direct role in Ag presentation to both CD8 and to a lesser extent CD4 T cells in addition to trafficking eTLRs (21, 22). Our finding, however, that the *3d* mutation did not impair TD Ab responses when nucleic acid-free alum was the adjuvant indicates that T helper function in these mutant mice is not significantly altered. Thus, reduced *3d* APC activity for *3d* T cells cannot account for the marked suppression of autoAbs in the lupus-prone mice. The finding that eTLR deficiency did not affect TD humoral responses is consistent with a recent study showing that absent TLR signaling in *Myd88/Trif* double-deficient mice did not significantly reduce adjuvant-enhanced Ab responses (34). Nonetheless, we found that the *3d* mutation slightly reduced the humoral response in *3d* mice when CFA was the adjuvant. This suggests that nucleic acids in CFA contribute to the overall adjuvant effect of CFA in *wt* mice, whereas the adjuvanticity of the alum-based mixture is not affected by eTLR deficiency because the response is mediated primarily through the *Nalp3* inflammasome (35).

Despite the normal TD humoral response, our findings confirmed reduced activity of APC from *3d* mice for stimulating *wt* T cells (21, 22). However, we found that the activation of OVA-primed *3d* T cells by *3d* APC was not impaired, and *3d* T cells exhibited greater stimulation than *wt* T cells when activated by *wt* APC. These findings, combined with normal TD humoral responses in *3d* mice (this study) and the lack of T cell population changes in nonautoimmune *3d* mice (21), suggest that *3d* T cells compensate for the slightly reduced function of *3d* APCs. Possible explanations for this are dynamic tuning of T cells for which a large number of different molecular mechanisms have been identified, including cell signaling feedback, level of CD5 expression, sialylation, and miRNA expression (36–40), or modification of the T cell receptor repertoire. We are currently generating T cell receptor transgenic *3d* mice to address this issue. Overall, our findings indicate that eTLRs play a limited, but significant role in determining the overall steady state of APC activity, possibly because of constant exposure of APC to subactivating amounts of nucleic acid-containing material from endogenous sources such as apoptotic cells or from exogenous commensal organisms and dietary substances. This possibility is supported by the observation that the copy number of TLR7 can alter response to self and foreign Ags (18).

Previous studies in lupus-prone mice showed that lack of TLR7 specifically inhibited RNP Ab production, whereas TLR9 deficiency had varying effects on DNA-related Abs (4, 9, 33).

Our finding that complete elimination of eTLR signaling inhibits the specific production of IgG anti-nucleic acid-associated Abs in lupus-prone mice indicates that these autoAb specificities are strongly dependent on TLR engagement. Therefore, the combined data suggest that anti-RNP B cells require TLR7 ligands for activation, whereas anti-DNA B cells can be activated through either TLR7 or TLR9. Thus, the major Ags for anti-RNP B cells most likely contain primarily RNA and little DNA, whereas the major antigenic targets for anti-DNA B cells must contain both DNA and RNA, with only the DNA-related Ags accessible to BCRs (otherwise anti-RNP B cells could take up these targets and be activated by TLR9 in TLR7-deficient mice). Apoptosis-derived blebs and particles, considered to be the major source of self Ag in SLE, contain varying amounts of nucleosomes, cytoplasmic RNA, and RNPs (41, 42). Among these, large apoptotic blebs, known to contain both nucleosomes and RNA, would appear to be a major self Ag for DNA and nucleosome-specific B cells, whereas the major self Ag for RNP-specific B cells may be smaller RNP particles (42).

The critical importance of eTLRs in the production of anti-nucleic acid Abs was strongly supported by 2 key findings. First, atypical ANA staining of 2 of 7 old B6-*Fas^{lpr}* 3d mice was detected only at an age where there were no significant differences in lymphoid hypertrophy and hyperIgG compared to *wt* mice. Second, chronic TLR4 stimulation of 3d mice could not overcome the requirement for eTLRs in autoAb production. TLR4 is the only nonnucleic acid-sensing TLR known to induce type I IFNs and to enhance lupus-like autoimmunity (23). Moreover, TLR4 is similar to the eTLRs in its signaling through both MyD88 (TLR7 and -9) and TRIF (TLR3) and in activating B cells and other APCs (9). Thus, the inability of autoimmune-prone 3d mice to sustain high levels of anti-nuclear Abs and RF in old lupus-prone mice or after TLR4 stimulation must be related to the specific recognition of nucleic acids by eTLRs or less likely by another unique property of these TLRs.

The lipid A treatment initially (day 3) increased concentrations of anti-nuclear Abs and RF in 3d mice commensurate to *wt* mice and consistent with polyclonal B cell activation, but no further increases in autoAbs occurred. Thus, it can be deduced that eTLRs were not required for activation of anti-nucleic acid recognizing B cells and the initial production of autoAbs, but were required for the subsequent amplification of this response. This is consistent with our previously hypothesized 2-phase paradigm of SLE (9), in which the first TLR-independent phase, triggered by activation of pDCs and DCs by apoptotic cell debris and associated nucleic acids, leads to the elaboration of activating cytokines and low levels of autoAb production. For sustained autoAb production and disease, however, a second TLR-dependent amplification phase is required, mediated by the engagement of eTLRs by nucleic acid-containing material taken up either directly via Ag receptors in B cells or as autoAb complexes in pDCs and DCs. This second phase is likely to involve a positive feedback loop that results in substantial magnification of the initial response.

Previous in vitro studies showed that activation of anti-IgG2a RF-expressing AM14 B cells by IgG2a anti-DNA/DNA complexes required TLR9, while activation by IgG2a anti-RNP/RNA complexes required TLR7, consistent with endocytosis of the nucleic acid complexes via the surface-expressed RF and subsequent engagement of eTLRs by their cognate ligand (7, 43). More recent in vivo studies in AM14 IgH-chain transgenic MRL-*Fas^{lpr}* lupus mice showed that direct activation of low-affinity RF B cells was not dependent on T cell help, but required the presence of TLR7 and TLR9 (33). Here, we extend this finding to show that dependence on nucleic acid-sensing TLRs also applies to spontaneous production of RF and, by inference, for RF regardless of affinity. Moreover, we show that, similar to the production of anti-nucleic acid Abs, TLR4 stimulation by lipid A administration in B6-*Fas^{lpr}* 3d

mice cannot bypass the dependence of sustained RF production on nucleic acid-recognizing TLRs.

The 3d mutation reduced, but did not ameliorate the lymphoid hypertrophy associated with defective *Fas^{lpr}*, although the degree of suppression was probably underestimated because of deaths in the more severely affected *wt* group before the final analysis. Examination of the cellular composition of the spleen and lymph nodes revealed some differences suggesting reduced activation of 3d T cells, including less splenic T cells, less CD4 T cells in lymph nodes, and more naive (CD62L⁺) and fewer activated (CD44^{hi}) CD8 T cells in both spleen and lymph node. Another finding was a reduction in the CD21^{lo}CD23^{lo} B cell population in 3d mice, which is abnormally increased in B6-*Fas^{lpr}* mice with age. This, along with the finding of a reduction in the CD138⁺B220⁻, possibly more mature, plasmablasts/plasma cell population, is consistent with a general reduction in overall B cell activation and autoimmunity. Thus, although it is not known to what extent endogenous self-nucleic acid Ags and foreign nucleic acid material might be responsible, we clearly document a major, but not essential, role for eTLRs in the lymphoid hypertrophy that develops in *Fas^{lpr}* mice.

Finally, our findings strongly support the therapeutic targeting of nucleic acid-sensing TLRs in SLE (44) and demonstrate the dramatic benefit of simultaneously inhibiting all such TLR family members. These findings also raise the possibility that eTLR signaling may play a critical role in other autoimmune diseases that have anti-nucleic acid Abs.

Materials and Methods

Mice. C57BL/6 (B6)-*Unc93b1^{3d}* (3d), B6-*Fas^{lpr}*, BXSB, B6-*MyD88^{-/-}*, B6-*Trif^{Δps2/Lps2}*, and B6-*Tlr9^{CpG1,CpG1}* mice were bred and maintained at The Scripps Research Institute. Experiments followed approved Institutional Animal Care and Use Committee protocols. *Fas^{lpr}* and 3d were identified by PCR (45) or sequencing. 3d BXSB mice were N2 or greater, *Yaa⁺*, and fixed for BXSB on chromosome 1 between 19.8 and 174.9 Mb (*D1Mit3*, *D1Mit21*, *D1Mit387*, and *D1Mit206*). Lupus was assessed as described (46). Fifty micrograms of lipid A (Calbiochem) in PBS were given 2 times per week i.p. for the indicated durations.

Immunopathology and Serology. Zinc formalin-fixed and PAS/hematoxylin-stained tissue sections were scored blindly for GN on a 0–4 scale (47). Ab concentrations were measured by ELISA (48). RNP plates were from Inova Diagnostics. ANAs were detected on Hep-2 slides (Bion Enterprises) using 1/100 serum and 1/200 Alexa Fluor 488-goat anti-mouse IgG dilutions (Invitrogen).

Flow Cytometry. Isolated splenic and LN cells, blocked with anti-CD16/CD32, were stained with combinations of dye-conjugated Abs to B220, CD4, CD5, CD8, CD19, CD21, CD23, CD44, CD62L, CD86, CD90.2, CD138, F4/80, IgM, and I-A/I-E (BD Biosciences or Biolegend). Data were acquired on a LSRII (BD Biosciences) and analyzed by Flowjo (Tree Star).

Humoral Responses. For the TI-1 response, mice were immunized once with 50 μg TNP-LPS (Biosearch Technologies) in PBS i.p. and for the TD response twice with 50 μg TNP-KLH on days 0 and 21 either in alum or in CFA on day 0 and in IFA on day 21 (CFA/IFA). Total, anti-TNP3, and anti-TNP15 IgM and IgG were measured by ELISA (49).

T Cell Proliferation. A total of 10⁵ T cells and 6 × 10⁵ APCs (non-T cells), isolated from spleens 10 days after immunization with 100 μg OVA in CFA, were incubated for 4 days with or without 20 μg OVA and harvested 18 h after addition of tritiated thymidine. Data are sample cpm minus media-alone cpm.

TLR-Independent Pathway Induction. BM-derived DCs, generated by culturing BM cells with 10 ng/mL GM-CSF for 7 days and then by CD11c microbead (MACSMiltenyi Biotech) isolation, were transfected with poly(I:C) or poly(dA:dT*dT:dA) (Sigma), 5' (3P)-transcribed RNA (pGEM express positive control, Riboprobe system T7, Promega), or calf-thymus DNA (Sigma) by Lipofectamine (Invitrogen). After 24-h cultures, IFN type I in supernatants was determined (50).

Statistical Analysis. Group comparisons used unpaired 2-tailed *t* tests. Survival was analyzed by Kaplan-Meier plots with a log-rank test.

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