DNA released from dying host cells mediates aluminum adjuvant activity

Thomas Marichal[1,2], Keiichi Ohata[3], Denis Bedoret[1,2], Claire Mesnil[1,2], Catherine Sabatel[1,2], Kouji Kobiyama[3,4], Pierre Lekeux[1,2], Cevayir Coban[3], Shizuo Akira[3], Ken J Ishii[3–5], Fabrice Bureau[1,2,5] & Christophe J Desmet[1,2,5]

Aluminum-based adjuvants (aluminum salts or alum) are widely used in human vaccination, although their mechanisms of action are poorly understood. Here we report that, in mice, alum causes cell death and the subsequent release of host cell DNA, which acts as a potent endogenous immunostimulatory signal mediating alum adjuvant activity. Furthermore, we propose that host DNA signaling differentially regulates IgE and IgG1 production after alum-adjuvanted immunization. We suggest that, on the one hand, host DNA induces primary B cell responses, including IgG1 production, through interferon response factor 3 (Irf3)-independent mechanisms. On the other hand, we suggest that host DNA also stimulates 'canonical' T helper type 2 (T_H2) responses, associated with IgE isotype switching and peripheral effector responses, through Irf3-dependent mechanisms. The finding that host DNA released from dying cells acts as a damage-associated molecular pattern that mediates alum adjuvant activity may increase our understanding of the mechanisms of action of current vaccines and help in the design of new adjuvants.

Since the first report of its adjuvant activity in 1926 (ref. 1), it has been widely believed that alum creates an immunologically inert 'depot' that prolongs antigen exposure or facilitates antigen uptake by antigen-presenting cells (APCs). However, in line with the decades-old observation that alum triggers interleukin (IL)-1β production[2], recent reports indicated that alum may activate caspase 1 (encoded by *Casp1*)[3] through inflammasomes containing the cytosolic receptor Nlrp3 (NLR family, pyrin domain containing 3)[4–8]. Whether the Nlrp3 inflammasome consistently contributes to the adjuvant effect of alum on humoral responses *in vivo*, however, remains unclear[5–10]. Nevertheless, these studies suggested that signaling through specific receptors of innate immunity, commonly referred to as pattern recognition receptors (PRRs), might have a role in the adjuvant activity of alum. Notably, it has also been proposed that alum induces the local accumulation of uric acid, which would act as an endogenous adjuvant molecule boosting T cell responses[11].

It has actually long been known that alum exerts some level of cytotoxicity[12]. Moreover, dying cells release numerous molecules that act as endogenous danger signals, or damage-associated molecular patterns (DAMPs), and which can alert the innate immune system through the activation of various PRR signaling pathways[13,14]. We thus hypothesized that DAMPs released at immunization sites and the subsequent activation of PRR signaling could contribute to the adjuvant effects of alum. Here we report that alum-induced cytotoxicity results in the release of host DNA, which acts as a DAMP mediating the adjuvant activity of alum.

RESULTS

Host DNA boosts adaptive responses after alum immunization

We observed that endotoxin-free ovalbumin (OVA) adsorbed on Imject Alum (alum), administered intramuscularly (i.m.) or intraperitoneally (i.p.), induced the local accumulation of extracellular double-stranded DNA (dsDNA) in mice (**Fig. 1a** and **Supplementary Fig. 1a**), part of which was entrapped in macroscopic alum depots (**Fig. 1b** and **Supplementary Fig. 1b**). This dsDNA release correlated with the death of local cells (**Fig. 1c** and **Supplementary Fig. 1c**). Both alum and aluminum hydroxide gel (Alhydrogel) induced similar DNA release and cell death (**Supplementary Fig. 1d,e**).

Because released host cell DNA may trigger immune responses[13,15–17], we hypothesized that extracellular host DNA could act as an endogenous adjuvant in alum vaccination. Remarkably, i.p. or i.m. injections of OVA mixed with purified mouse genomic DNA (OVA and DNA), in quantities similar to those released by alum, boosted OVA-specific IgM, IgG1 and IgE responses as efficiently as OVA and alum (**Fig. 2a–c** and **Supplementary Fig. 2a–c**). IgG2c production was not significantly induced in these conditions (data not shown). Thus, like alum, host DNA may act as an adjuvant that preferentially boosts T_H2-dependent humoral responses. Digestion of extracellular DNA *in vivo* by treatment with DNase I decreased antigen-specific humoral responses in OVA and alum–treated mice (**Fig. 2d–f**). Furthermore, the transfer of OVA, together with acellular peritoneal lavage fluid from mice immunized i.p. with OVA and alum, triggered OVA-specific IgG1 and IgE production in naive recipient mice, a response

Figure 1 Alum induces cell death and release of host DNA at sites of injection. (**a**) Quantity of free dsDNA in the acellular fraction of the peritoneal lavage fluid of mice treated i.p. with increasing doses of alum, measured over time using quantitative fluorescent dsDNA stain. (**b**) Confocal microscopic imaging of extracellular DNA deposition in alum macroscopic i.p. depots stained with 4′,6-diamidino-2-phenylindole (DAPI). Scale bars, 25 μm. (**c**) Cell death rate in the peritoneal lavage fluid of mice treated i.p. with increasing doses of alum, assessed by staining with 7-aminoactinomycin D (7-AAD) and flow cytometry. $n = 5$ (**a**,**c**). Data are representative of one of three independent experiments. Error bars show means ± s.d. *$P < 0.05$, **$P < 0.01$, ***$P < 0.001$.
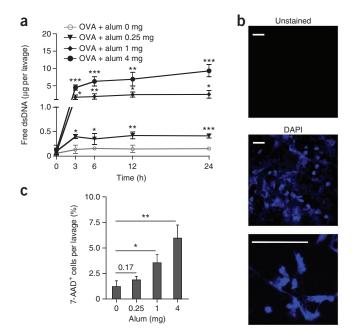

that was abrogated by DNase I treatment of the lavage fluid prior to transfer (**Supplementary Fig. 3a,b**).

We also observed that the proliferation of both adoptively transferred carboxyfluorescein succinimidyl ester (CFSE)-labeled (**Fig. 2g**) and endogenous (**Supplementary Fig. 4**) OVA-specific CD4⁺ T cells was comparable in OVA and DNA– and OVA and alum–treated mice and reduced after treatment of the mice with DNase I (**Fig. 2g** and **Supplementary Fig. 4**). These results thus support the idea that host DNA is a potent endogenous adjuvant molecule that has a role in the induction of humoral and T cell responses by alum.

Alum signals through Tbk1 and Irf3 to boost IgE responses

In vivo, only toll-like receptor 9 (Tlr9) has been implicated in host DNA recognition, under specific pathologic conditions[18,19]. However, we observed that mice deficient in Tlr9 developed humoral responses similar to those of their wild-type (WT) counterparts in response to alum immunization (data not shown).

In addition to Tlr9, other molecules have been identified as putative host DNA receptors *in vitro*[20–27]. These receptors activate

either the inflammasome pathway, leading to mature IL-1β protein expression[21,23,24], or the Irf3 pathway, leading to interferon (IFN)-β1 expression[20,22,25–28]. Whereas OVA and DNA upregulated IFN-β1 secretion similarly to OVA and alum (**Fig. 3a**), OVA and DNA did not induce more IL-1β production than OVA alone, unlike OVA and alum (**Fig. 3b**). This suggested that, after alum immunization, extracellular DNA activated Irf3, but not the inflammasome pathway. By further investigating the general importance of inflammasomes
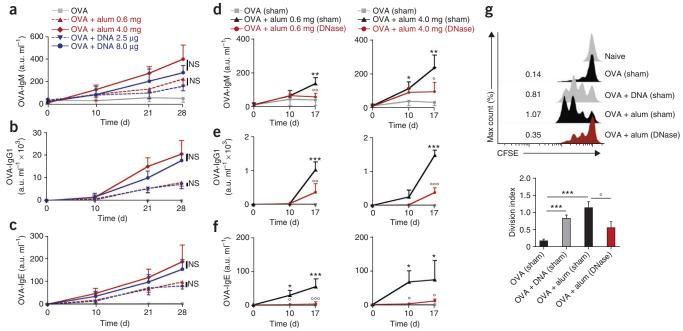

Figure 2 Host DNA released by alum cytotoxicity mediates alum adjuvant activity on humoral and T_H2 cell responses. (**a–f**) Serum titers of OVA-specific IgM (OVA-IgM; **a,d**), IgG1 (OVA-IgG1; **b,e**) and IgE (OVA-IgE; **c,f**). Titers were measured on the indicated days in (**a–c**) mice immunized i.p. with OVA alone, OVA and alum, or OVA and DNA on days 0 and 14, and boosted with OVA on d 21; or in (**d–f**) mice immunized i.p. with OVA or OVA and alum, treated i.p. with DNase I both 3 and 18 h later, and then boosted with OVA 10 d later. (**g**) Proliferation profile (top) and division index (bottom) of adoptively transferred CFSE-labeled OVA-specific CD4⁺ OT-II cells in the bronchial lymph nodes of mice treated i.p. with OVA, OVA and DNA, OVA and alum, or OVA and alum followed by DNase I treatment. Inserted numbers indicate division index values. $n = 5$; data are representative of one of two (**a–c**) or three (**d–g**) independent experiments. Error bars show means ± s.d. *, OVA versus OVA and adjuvant; °, OVA and alum versus OVA and alum followed by DNase I treatment; *,°$P < 0.05$, **,°°$P < 0.01$, ***,°°°$P < 0.001$. a.u., arbitrary unit. NS, not significant.
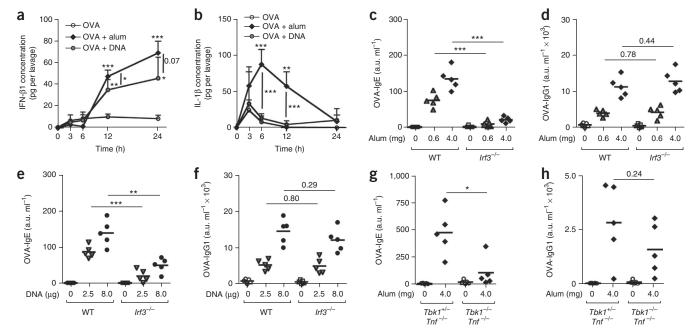
Figure 3 Alum and host genomic DNA trigger type I IFN secretion and IgE responses through activation of the Tbk1-Irf3 axis. (**a,b**) Quantities of (**a**) IFN-β1 and (**b**) IL-1β in the peritoneal lavage fluid over time by ELISA in WT mice treated i.p. with OVA, OVA and DNA, or OVA and alum. (**c–f**) Serum titers of OVA-specific IgE (**c,e**) and IgG1 (**d,f**) measured on day 28 in WT and *Irf3*^{−/−} mice immunized i.p. with OVA, OVA and alum (**c,d**), or OVA and DNA (**e,f**) on days 0 and 14, and then boosted with OVA on day 21. (**g,h**) Serum titers of OVA-specific IgE (**g**) and IgG1 (**h**) measured on day 28 in *Tbk1*^{+/−}/*Tnf*^{−/−} and *Tbk1*^{−/−}/*Tnf*^{−/−} mice immunized i.p. with OVA and alum on days 0 and 14, and then boosted with OVA on day 21. *n* = 5; data are representative of one of three independent experiments. Error bars show means ± s.d. **P < 0.05, **P < 0.01, ***P < 0.001*. a.u., arbitrary unit.

in the adjuvant activity of alum, we observed that Nlrp3-deficient (*Nlrp3*^{−/−}) and Caspase 1–deficient (*Casp1*^{−/−}) mice developed humoral responses comparable to those of WT mice after alum immunization (**Supplementary Fig. 5a,b**).

We next examined whether Irf3 is required for the adjuvant activity of alum. We found that, compared with their WT counterparts, in Irf3-deficient (*Irf3*^{−/−}) mice, i.p. immunization with OVA and alum triggered reduced antigen-specific IgE responses (**Fig. 3c**). Notably, however, the antigen-specific IgG1 serum titers of *Irf3*^{−/−} mice were similar to those of WT mice (**Fig. 3d**). Furthermore, we also saw the dependency of antigen-specific IgE production on Irf3 when we used human serum albumin (HSA) in place of OVA and Alhydrogel in place of alum (**Supplementary Fig. 6a**). We obtained similar results when we administered OVA and alum i.m. rather than i.p. (**Supplementary Fig. 7a**). IgG1 production remained unaffected in *Irf3*^{−/−} mice under these different conditions (**Supplementary Fig. 6b** and **Supplementary Fig. 7b**). We also observed a reduction in the production of antigen-specific IgE, but not of IgG1, when we immunized *Irf3*^{−/−} mice i.p. or i.m. with OVA and DNA in place of OVA and alum (**Fig. 3e,f** and **Supplementary Fig. 7c,d**), supporting the notion that both alum and host DNA induce IgE responses through the same pathway.

Most of the host DNA–activated pathways that lead to Irf3 activation signal through TANK-binding kinase 1 (Tbk1)[20,22,25,28]. Tbk1-deficient mice (*Tbk1*^{−/−}) die *in utero* from massive liver apoptosis, but a combined deficiency for tumor necrosis factor (encoded by *Tnf*) avoids these lethal effects[28,29]. We found that immunized *Tbk1/Tnf* double-knockout mice developed reduced OVA-specific IgE responses (**Fig. 3g**) but similar OVA-specific IgG1 serum titers (**Fig. 3h**) when compared to immunized control *Tbk1*^{+/−}/*Tnf*^{−/−} mice, thereby suggesting a role for the Tbk1-Irf3 axis in the adjuvant activity of alum on IgE responses.

The only currently known mammalian DNA-sensitive pathways able to activate the Tbk1-Irf3 axis involve the sensing of genomic DNA by Sting (a stimulator of IFN genes)[25], and the activation of Ddx58 (DEAD box polypeptide 58) by RNA transcribed from DNA by polymerase III (ref. 22). Both Sting and Ddx58 act upstream of mitochondrial antiviral signaling protein (encoded by *Mavs*) to activate Tbk1 and Irf3 (refs. 22,25). We observed that OVA and alum–immunized *Mavs*^{−/−} mice developed OVA-specific IgE and IgG1 serum titers similar to those of WT mice (data not shown). Zbp1 (Z-DNA binding protein 1, also called Dai) has also been proposed as a cytoplasmic DNA sensor *in vitro*[20], although its role remains unclear *in vivo*[30]. *Zbp1*^{−/−} mice did not show reduced antigen-specific IgE or IgG1 production in response to OVA and alum immunization in comparison with WT mice (data not shown).

Together, these results indicate that alum-induced host DNA release promotes IgG1 and IgE responses through distinct signaling pathways that do not depend on currently identified DNA receptors. They also identify Irf3 as an essential mediator of IgE production after alum immunization.

Irf3 is essential for alum-boosted canonical T$_H$2 responses

Given that IgE isotype switching by B cells is best induced by T cells expressing a T$_H$2 cytokinic profile, we suspected that Irf3 deficiency would have an effect on T$_H$2 cell responses after alum immunization. Indeed, the proliferation of transferred CFSE-labeled OVA-specific CD4[+] cells (**Fig. 4a**), the differentiation of endogenous antigen-specific T$_H$2 cells (**Supplementary Fig. 8a,b**) and the secretion of T$_H$2 cytokines in response to OVA re-stimulation (**Fig. 4b**) were reduced in the bronchial lymph nodes (BLNs) of *Irf3*^{−/−} mice injected i.p. with OVA and DNA or with OVA and alum. Furthermore, in a classical asthma model based on OVA and alum sensitization followed by
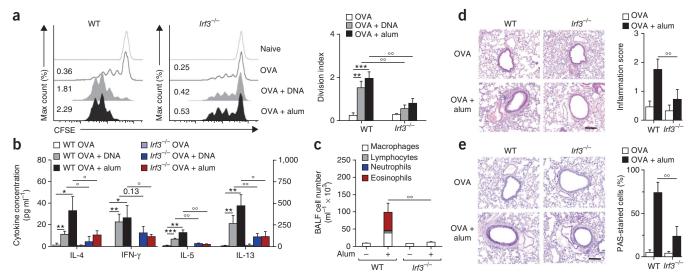
Figure 4 Irf3 is essential for the boosting of canonical T_H2 cells by alum and genomic DNA. (**a**) Proliferation profile (left) and division index (right) of adoptively transferred OVA-specific CD4+ OT-II cells in the BLNs of WT and $Irf3^{-/-}$ mice treated i.p. with OVA, OVA and DNA, or OVA and alum. Inserted numbers indicate division index values. (**b**) Cytokine concentrations in the supernatant of OVA-stimulated BLN cells isolated from WT and $Irf3^{-/-}$ mice treated with OVA, OVA and DNA, or OVA and alum. (**c**–**e**) Assessment of allergic airway inflammation in OVA- or OVA and alum–sensitized WT and $Irf3^{-/-}$ mice challenged with aerosolized OVA. (**c**) Total and differential immune cell counts in the bronchoalveolar lavage fluid (BALF). (**d**) Representative section and inflammatory scores of hematoxylin-eosin staining of lung sections. Scale bar, 50 μm. (**e**) Representative staining and percentage of periodic acid Schiff (PAS)-stained goblet cells per total epithelial cells in randomly selected bronchi. Scale bar, 50 μm. $n = 5$; data are representative of one of two (**a**) or three (**b**–**e**) independent experiments. Error bars show means ± s.d. *, OVA versus OVA and adjuvant; °, WT OVA and adjuvant versus $Irf3^{-/-}$ OVA and adjuvant; *,°$P < 0.05$, **,°°$P < 0.01$, ***$P < 0.001$.

airway challenge with OVA, we observed that $Irf3^{-/-}$ mice were protected from allergic airway inflammation (**Fig. 4c**–**e**), and that their BLN cells showed impaired proliferation and T_H2 cytokine secretion in response to OVA re-stimulation (**Supplementary Fig. 9a,b**). It is noteworthy that $Irf3^{-/-}$ mice, when compared with WT mice, had decreased serum OVA-specific IgE levels but similar OVA-specific IgG1 titers in this model (**Supplementary Fig. 9c,d**).

Previous studies in our laboratory indicated that $Irf3^{-/-}$ mice show normal boosting of T_H2 responses by Irf3-independent adjuvants[31]. Moreover, $Irf3^{-/-}$ T cells proliferate and secrete T_H2 cytokines normally in response to CD3 and CD28 stimulation (**Supplementary Fig. 10a,b**). $Irf3^{-/-}$ mice also develop humoral responses similar to those of WT mice when immunized with Freund's adjuvant (**Supplementary Fig. 11a–c**).

Taken together, these results indicate that $Irf3^{-/-}$ mice are impaired in their ability to boost T_H2 cell responses to alum immunization and to genomic DNA. They further suggest that Irf3-dependent T_H2 cell responses to alum may sustain tissue inflammation but do not affect IgG1 production. The T_H cells implicated in these Irf3-dependent responses would thus functionally correspond to canonical T_H2 cells[32]. In addition, T cell responses, although reduced, were not completely abrogated in $Irf3^{-/-}$ mice (**Fig. 4a,b** and **Supplementary Fig. 8a,b**). This suggests that Irf3-independent pathways also contribute to the adjuvant activity of alum on T cell responses, and that these Irf3-independent T cell responses primarily sustain B cell responses, including IgG1 production.

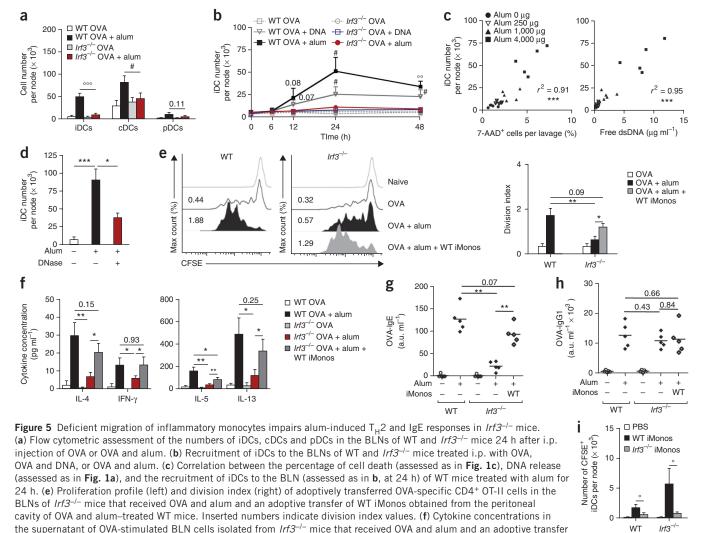## Alum-activated iDCs boost canonical T_H2 responses

The inability of $Irf3^{-/-}$ mice to produce normal T_H2 cell responses suggested the possibility of deficient antigen presentation to T cells. We thus studied the recruitment of dendritic cells and other innate immune cells to alum injection sites and their draining lymph nodes. Although the recruitment of immune cell populations at sites of alum injection did not differ significantly between WT and $Irf3^{-/-}$ mice

(**Supplementary Fig. 12a,b**), there was a reduction in the numbers of inflammatory monocyte (iMono)-derived inflammatory DCs (iDCs) and, to a lesser extent, of conventional DCs (cDCs) in the draining lymph nodes of i.p. or i.m. OVA and alum–treated $Irf3^{-/-}$ mice, whereas numbers of plasmacytoid DCs (pDCs) were not reduced (**Fig. 5a** and **Supplementary Fig. 13a**). Similarly, i.p. or i.m. OVA and DNA treatment induced an Irf3-dependent recruitment of iDCs to the BLNs (**Fig. 5b** and **Supplementary Fig. 13b**). Adding further support to the idea that host DNA acts as a trigger of iDC recruitment upon alum immunization, we observed that the recruitment of iDCs to the BLNs of alum-treated mice strongly correlated with the percentage of cell death and DNA release (**Fig. 5c**), and that it was reduced after DNase I treatment (**Fig. 5d**).

Our observations suggested that Irf3 may be required for T_H2 cell and IgE responses through its role in the normal recruitment of iMonos, the precursors of iDCs[11,33,34], to the lymph nodes that drain alum injection sites. Supporting this hypothesis, we found that the transfer of iMonos isolated from OVA and alum–immunized WT mice to $Irf3^{-/-}$ mice treated with OVA and alum increased the proliferation of transferred CFSE-labeled OVA-specific CD4+ cells (**Fig. 5e**) and the differentiation of endogenous OVA-specific CD4+ T_H2 cells (**Supplementary Fig. 14a,b**). In addition, we found that transferring WT iMonos to OVA and alum–treated $Irf3^{-/-}$ mice increased T_H2-type cytokine production (**Fig. 5f**) and OVA-specific IgE titers (**Fig. 5g**), almost to the levels of OVA and alum–treated WT mice. In contrast, iMono transfer did not affect OVA-specific IgG1 titers (**Fig. 5h**). Notably, we observed that, although the transfer of iMonos from OVA and alum–treated mice was sufficient to induce robust T_H2 cell responses in the draining lymph nodes of naive recipient mice (**Supplementary Fig. 15a–d**), it did not induce any detectable IgG1 or IgE responses (data not shown).

These results therefore indicate that Irf3 is essential for the triggering of iDC recruitment by alum. They also suggest that these iDCs subsequently induce a canonical T_H2 response that is unable to

Figure 5 Deficient migration of inflammatory monocytes impairs alum-induced T$_H$2 and IgE responses in *Irf3*$^{-/-}$ mice. (**a**) Flow cytometric assessment of the numbers of iDCs, cDCs and pDCs in the BLNs of WT and *Irf3*$^{-/-}$ mice 24 h after i.p. injection of OVA or OVA and alum. (**b**) Recruitment of iDCs to the BLNs of WT and *Irf3*$^{-/-}$ mice treated i.p. with OVA, OVA and DNA, or OVA and alum. (**c**) Correlation between the percentage of cell death (assessed as in **Fig. 1c**), DNA release (assessed as in **Fig. 1a**), and the recruitment of iDCs to the BLN (assessed as in **b**, at 24 h) of WT mice treated with alum for 24 h. (**e**) Proliferation profile (left) and division index (right) of adoptively transferred OVA-specific CD4$^+$ OT-II cells in the BLNs of *Irf3*$^{-/-}$ mice that received OVA and alum and an adoptive transfer of WT iMonos obtained from the peritoneal cavity of OVA and alum–treated WT mice. Inserted numbers indicate division index values. (**f**) Cytokine concentrations in the supernatant of OVA-stimulated BLN cells isolated from *Irf3*$^{-/-}$ mice that received OVA and alum and an adoptive transfer of WT iMonos obtained as in **e**. (**g,h**) Serum titers of OVA-specific IgE (**g**) and IgG1 (**h**) in *Irf3*$^{-/-}$ mice treated with OVA and alum, transferred with WT iMonos as in **e** on days 0 and 14, and then boosted with OVA on day 21. As controls, we used WT and *Irf3*$^{-/-}$ mice that received PBS with OVA alone or OVA and alum. (**i**) Numbers of CFSE$^+$ iDCs in the BLNs of OVA and alum–treated WT mice that received CFSE-labeled WT or *Irf3*$^{-/-}$ iMonos 18 h earlier. Control mice received PBS alone. *n* = 5; data are representative of one of four (**a,c**), one of three (**f–i**) or one of two (**b,d,e**) independent experiments. Error bars show means ± s.d. #, WT OVA and adjuvant versus *Irf3*$^{-/-}$ OVA and adjuvant (**a,b**); *,°$P < 0.05$, **,°°$P < 0.01$, ***,°°°$P < 0.001$. a.u., arbitrary unit.

autonomously support humoral responses, but that has the ability to promote IgE responses from independently activated B cell responses.

Alum-induced IL-12p80 promotes iMono migration

Cross-transfer experiments suggested that *Irf3*$^{-/-}$ iMonos had a cell-intrinsic defect that rendered them unable to migrate to the lymph nodes after alum treatment (**Fig. 5i**). In cells isolated from sites of OVA and alum injection, the production of IFN-β1, a hallmark of Irf3 activation, was most prominent in iMonos, as compared with other peritoneal cells (**Supplementary Fig. 16**). We found that IFN-β1 production was abrogated in iMonos from *Irf3*$^{-/-}$ mice, suggesting that Irf3 indeed is activated in iMonos (**Supplementary Fig. 16**). As demonstrated by the use of *Irf9*$^{-/-}$ mice and mice lacking the gene encoding interferon α- and β- receptor 2 (*Ifnar2*$^{-/-}$ mice), type I and type III IFNs did not affect alum-induced iDC recruitment or humoral responses (data not shown).

Although Irf3 deficiency did not seem to affect the overall maturation state of iMonos (a process usually associated with the acquisition

of migratory activity), we noticed that a higher proportion of *Irf3*$^{-/-}$ iMonos highly expressed CCR7, a chemokine receptor crucial for DC migration to the lymph nodes, compared with their WT counterparts (**Fig. 6a**). This suggested that *Irf3*$^{-/-}$ iMonos might be unable to respond to CCR7-activating signals, which would lead to their accumulation. Recent evidence indicates that iMono-derived DCs depend on signaling by homodimers of IL-12p40 (IL-12p80) for their ability to respond to CCR7-activating chemokines[35,36]. We found that alum induced the local production of IL-12p80 in WT mice, and that this production was reduced in *Irf3*$^{-/-}$ mice (**Fig. 6b**). We showed the importance of IL-12p80 expression in regulating iMono migration by treating *Irf3*$^{-/-}$ mice with recombinant IL-12p80, which restored migration of iDCs to the BLNs of alum-injected *Irf3*$^{-/-}$ mice, almost to the levels seen in WT mice (**Fig. 6c**). Moreover, neutralization of IL-12p80 by p40-specific neutralizing antibodies partially attenuated IgE responses and the recruitment of iDCs to the BLNs in OVA and alum–treated WT mice, whereas it did not affect IgG1 responses (**Fig. 6d–f**). Thus, these results suggest that alum-induced iDC activation relies on signaling by IL-12p80.

Figure 6 Alum-induced iMono migration depends on IL-12p40 homodimer signaling. (**a**) Gating strategy for iMonos and flow cytometric analysis of their surface expression of CCR7 and co-stimulatory molecules in OVA and alum–treated WT and $Irf3^{-/-}$ mice. Bottom right, percentage of $CCR7^{hi}$ iMonos in OVA and alum–treated WT and $Irf3^{-/-}$ mice is indicated. (**b**) ELISA measurement of IL-12p70, IL-23 and IL-12p80 in the acellular phase of the peritoneal lavage fluid of WT and $Irf3^{-/-}$ mice treated overnight with OVA or with OVA and alum. #, WT OVA and alum versus $Irf3^{-/-}$ OVA and alum. (**c,d**) Recruitment of iDCs to the BLNs of WT and $Irf3^{-/-}$ mice treated with OVA and alum and recombinant IL-12p80 (rIL12p80) (**c**) or p40-specific neutralizing antibody (**d**). Control OVA- and OVA and alum–treated mice received PBS only. (**e,f**) Serum titers of OVA-specific IgE (**e**) and IgG1 (**f**) measured on day 17 in WT mice treated i.p. with OVA and alum and p40-specific neutralizing antibody, and then boosted with OVA i.p. 10 d later. $n = 5$; data are representative of one of three (**a**) or two (**b**–**f**) independent experiments. Error bars show means ± s.d. *$P < 0.05$, **$P < 0.01$, ***$P < 0.001$. a.u., arbitrary unit.
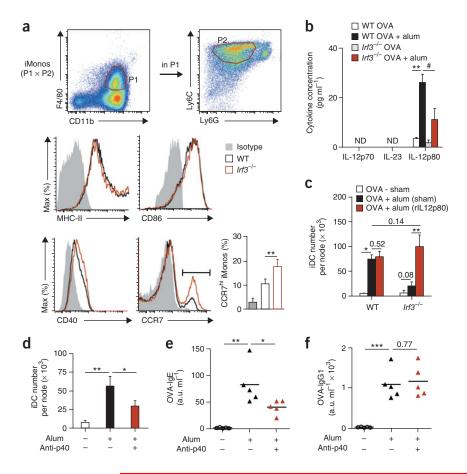
DISCUSSION

Extracellular host DNA has been recently recognized as a DAMP, and it seems to be involved in an increasing number of immune processes and diseases[16,19,37–39]. In this study, we show that host DNA released at sites of injection by dying cells can mediate the adjuvant effect of alum on adaptive responses. Our data thus support the idea that DAMPs have a key role in the adjuvant activity of alum and that, like any other efficient adjuvant[40,41], alum acts mainly by triggering innate immunity. In this regard, it should also be noted that uric acid, a major metabolite of nucleic acids and a potential DAMP when crystallized, has been suggested to have a role in iDC recruitment and T cell activation after alum immunization[11].

It has recently been proposed that alum immunization may induce functionally distinct T_H cell subsets comprising T follicular helper (T_{FH}) cells, which would be responsible for B cell help, and lymphoid and emigrant T_H cells, which would be responsible for other effector functions[42]. Another report suggested that IL-4–producing T_{FH} cells may be distinguished functionally from canonical T_H2 cells[32]. These IL-4–producing T_{FH} cells, also referred to as $T_{FH}2$ cells[43], would be endowed with high B cell helper activity and would promote IgG1 responses, but they would be poor inducers of classical T_H2 cell effector functions in peripheral tissues[32]. Canonical T_H2 cells, on the contrary, would be specialized in mediating T_H2 cell effector functions in peripheral tissues[32].

In line with these studies, our data allow us to propose a model in which the contribution of host DNA to the adjuvant activity of alum would comprise both an Irf3-independent and an Irf3-dependent component (**Supplementary Fig. 17**). In the Irf3-independent component of the response, host DNA induces IgG1 production through the promotion of T_H responses endowed with high B cell helper activity. These T_H cells would thus functionally relate to $T_{FH}2$ cells. In the Irf3-dependent component, iDCs activated by host DNA trigger T_H2 cell responses that mediate effector T_H2 cell functions in peripheral tissues but lack the ability to stimulate antibody production on their own. These T_H cells

would thus functionally relate to canonical T_H2 cells. They would also promote IgE production from independently induced B cell responses. Future detailed phenotypic and functional characterization of these T_H cell subtypes should offer new perspectives on the mechanisms driving the production of specific antibody responses.

Our data suggest that alum-activated iDCs preferentially induce canonical T_H2 cell responses. In other settings, iMono-derived iDCs preferentially induce other types of T_H responses, such as T_H1 responses[44]. The functional specialization of iDCs upon alum immunization most probably originates from the T_H2 microenvironment that develops at alum injection sites[10]. In contrast with a previous report[11], we observed that impairment of iDC function does not affect IgG1 production. This disagreement probably comes from the fact that the general depletion of CD11c+ cells used to target iDCs in the previous study also depleted other CD11c+ APCs. Given the diversity of lymph node APCs[45] and the current lack of molecular tools for manipulating them specifically *in vivo*, the identification of the APC(s) responsible for initiating IgG1 responses will be a stimulating challenge for future research.

It is traditionally thought that IgG1 and IgE responses are sequentially boosted by a common IL-4-producing T_H cell response. Our data, however, support the idea that IgE and IgG1 responses are uncoupled and that they are regulated by functionally distinct T_H cell mechanisms. Understanding how alum-induced DNA release stimulates $T_{FH}2$-related T cell help for IgG1 production could thus have a promising influence on the design of new vaccine formulations devoid of canonical T_H2 cell and IgE bias. We foresee that this breakthrough might come from the identification of the DNA sensors triggered by alum.

In conclusion, we suggest that the DNA released by dying host cells is an important DAMP mediating the adjuvant activity of alum.

This knowledge may help in improving or developing current and experimental adjuvant formulations, which will be key to the success of future vaccination strategies.

METHODS

Methods and any associated references are available in the online version of the paper at http://www.nature.com/naturemedicine/.

Note: Supplementary information is available on the Nature Medicine website.

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ONLINE METHODS

Mice. *Tbk1/Tnf* DKO, *Tlr9*^{−/−} and *Zbp1*^{−/−} mice have been described elsewhere[28,46,47]. We purchased *Irf3*^{−/−} (ref. 48) and *Irf9*^{−/−} mice[49] from RIKEN BioResource Center. *Nlrp3*^{−/−} and *Casp1*^{−/−} mice[50,51] were a kind gift of V. Dixit (Genentech). *Ifnar2*^{−/−} and *Mavs*^{−/−} mice were obtained as described previously[30]. OVA-specific, MHC II–restricted, T cell receptor–transgenic OT-II (H-2b; C57BL/6 background) mice were obtained from The Jackson Laboratory. All transgenic mice except *Tbk1/Tnf* double-knockout and *Zbp1*^{−/−} mice were backcrossed, in total, for 10 to 16 generations onto a C57Bl/6 background. *Tbk1/Tnf* double-knockout and *Zbp1*^{−/−} mice were on a 129/Ola X C57Bl/6 background. *Tbk1*^{+/−}/*Tnf*^{−/−} littermates were used as control for *Tbk1/Tnf* DKO mice. All mice were bred and housed in institution-specific pathogen-free facilities. Age-paired groups of females were used at 8–12 weeks of age. All experimental procedures and protocols were approved by the Institutional Animal Care and Use Committees of the University of Liège and of the Research Institute for Microbial Diseases, Osaka University, Japan.

Immunizations. Endotoxin-free (endotoxin <1 EU mg^{−1}) OVA was from Hyglos. Unless otherwise indicated, we immunized mice i.p. or i.m. with 10 µg of OVA alone or adsorbed on 4 mg of Imject Alum (OVA and alum; Pierce Biochemicals). For immunizations with genomic DNA, we purified genomic DNA from mouse tissue using High Pure PCR Template Preparation kits (Roche), and, unless otherwise indicated, we immunized mice with 10 µg of OVA mixed with 8 µg of DNA (OVA and DNA). The endotoxin content of the DNA preparations, as measured by Hyglos using the Limulus Amoebocyte Lysate assay, was always <1 EU mg^{−1}. Unless otherwise indicated, we immunized mice i.p. or i.m. on days 0 and 14 with 10 µg of OVA alone, of OVA and DNA or of OVA and alum, and then boosted them i.p. on day 21 with 20 µg of OVA. Alternatively, we replaced OVA with HSA (Sigma), and alum by Alhydrogel (Sigma).

***In vivo* DNase I treatment.** We injected mice i.p. with 2,000 IU of DNase I (Roche) in 100 µl of HBSS (Lonza) 3 and 18 h after immunization.

Measurement of antigen-specific antibody titers. For OVA- and HSA-specific IgG1 and OVA-specific IgM and IgG2c detection, we incubated diluted sera from immunized mice on ELISA plates coated with antigen at 100 µg ml^{−1}. We detected bound IgG1, IgM and IgG2c using horseradish peroxidase (HRP)-conjugated mouse IgG1-, IgM- and IgG2c-specific antibodies (Southern Biotechnology) followed by incubation with tetramethyl benzidine and measurement by spectrophotometry. For OVA- and HSA-specific IgE detection, we incubated diluted sera on ELISA plates coated with mouse IgE-specific antibodies (BD Biosciences) at 2 µg ml^{−1}. We detected bound IgE using biotinylated antigen followed by streptavidin peroxidase (Zymed) incubation. We calculated antibody titers by plotting the serum dilution that gave half-maximal signal. When no signal was detected, we assigned a titer of 2.

Antibodies. A detailed description of the antibodies used in this study is available in the **Supplementary Methods**.

Assessment of OVA-specific T cell activation. We assessed the proliferation of transferred CFSE-labeled OT-II cells, as described previously[52], 3 d after immunization. We assessed proliferation and IL-4 expression of endogenous OVA-specific CD4^{+} cells as described previously[31], except that we cultured cells for 5 d with or without OVA (OVA grade V, Sigma; 50 µg ml^{−1}).

Flow cytometry and cell sorting. We incubated single cell suspensions with 2.4G2 Fc-receptor antibodies to reduce nonspecific binding before staining. We carried out staining reactions at 4 °C. We did flow cytometry on a FACScanto II instrument (BD Biosciences) and analyzed results using FlowJo software (Tree Star). We used a FACSAria instrument (BD Biosciences) for sorting F4/80^{int} CD11b^{+}Ly6C^{+}Ly6G^{−} iMonos.

IL-12p40 homodimer ELISA, complementation and neutralization experiments. We measured IL-12p40 homodimer concentrations by ELISA as previously described[53]. Briefly, we used mouse recombinant IL-12p80 (Biolegend) as a standard. We used the standards for mouse IL-12p40, IL-12p70 and IL-23 (R&D Systems) to test the specificity of the ELISA. For complementation and neutralization experiments, we immunized mice with OVA and alum and, 6 and 18 h later, treated them with rIL-12p80 (1 µg in 100 µl of PBS) or low endotoxin azide-free–purified IL-12p40-specific antibodies (Biolegend) (500 µg in 100 µl of PBS), respectively.

Statistical analysis. Data are presented as means ± s.d. We estimated the differences between mean values by using two-tailed pairwise Student's *t* tests after Anderson-Darling tests for assessment of normality of the data distributions. We calculated correlation coefficients using Pearson's two-tailed tests after assessment of data distribution normality. In all figures, we use asterisk (*) and degree (°) symbols to indicate significant differences observed when comparing indicated groups (*,° $P < 0.05$; **,°° $P < 0.01$; and ***,°°° $P < 0.001$).

Additional methods. Detailed methodology is described in the **Supplementary Methods**.

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