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DOI:  
[10.1073/pnas.1503315112](https://doi.org/10.1073/pnas.1503315112)

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*Document Version*  
Publisher's PDF, also known as Version of record

*Citation for published version (Harvard):*  
Barone, F, Nayar, S, Dias De Campos, J, Withers, D, Toellner, K-M, Fisher, B, Bowman, S, Cloake, T, Zhang, Y, Fouser, L, Rangle-Moreno, J, de la Luz Garcia-Hernandez, M, Randall, TD, Lucchesi, D, Bombardieri, M, Pitzalis, C, Buckley, C & Luther, SA 2015, 'IL-22 regulates lymphoid chemokine production and assembly of tertiary lymphoid organs', *National Academy of Sciences. Proceedings*, vol. 112, no. 35, pp. 11024-11029. <https://doi.org/10.1073/pnas.1503315112>

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IL-22 regulates tertiary lymphoneogenesis, Francesca Barone et al, *Proceedings of the National Academy of Sciences*, Sep 2015, 112 (35) 11024-11029; DOI: 10.1073/pnas.1503315112

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# IL-22 regulates lymphoid chemokine production and assembly of tertiary lymphoid organs

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Edited by Jason G. Cyster, University of California, San Francisco, CA, and approved July 31, 2015 (received for review February 19, 2015)

**The series of events leading to tertiary lymphoid organ (TLO) formation in mucosal organs following tissue damage remain unclear. Using a virus-induced model of autoantibody formation in the salivary glands of adult mice, we demonstrate that IL-22 provides a mechanistic link between mucosal infection, B-cell recruitment, and humoral autoimmunity. IL-22 receptor engagement is necessary and sufficient to promote differential expression of chemokine (C-X-C motif) ligand 12 and chemokine (C-X-C motif) ligand 13 in epithelial and fibroblastic stromal cells that, in turn, is pivotal for B-cell recruitment and organization of the TLOs. Accordingly, genetic and therapeutic blockade of IL-22 impairs and reverses TLO formation and autoantibody production. Our work highlights a critical role for IL-22 in TLO-induced pathology and provides a rationale for the use of IL-22-blocking agents in B-cell-mediated autoimmune conditions.**

IL-22 | tertiary lymphoid organs | chemokines | Sjogren's syndrome | autoimmunity

Tertiary lymphoid organs (TLOs) are organized clusters of immune cells that preferentially form in autoimmune diseases such as Sjogren's syndrome and Hashimoto thyroiditis (1). TLOs' cellular compartments, spatial organization, vasculature, and function are similar to those of secondary lymphoid organs (SLOs), providing a local hub for autoreactive B-cell proliferation and affinity maturation. TLOs appear to contribute to disease progression and to the emergence of malignant B clones responsible for lymphoma development (2). Within TLOs, the ectopic expression of lymphoid chemokines has been shown to correlate with the size or degree of organization of lymphoid aggregates and with the production of autoantibodies (3, 4). In the case of malignant transformation, lymphoid chemokine expression increases during disease progression (5). The pathways regulating chemokine expression in SLOs during embryonic life have been largely described and mainly involve the engagement of lymphotoxin  $\beta$  receptor on a family of gp38<sup>+</sup> lymphoid tissue-organizer stromal cells (6). At sites of TLO formation, lymphotoxin and lymphotoxin-producing cells appear to be dispensable for the early production of lymphoid chemokines required during SLO formation, raising the possibility that signals other than lymphotoxin can regulate the formation of TLOs, stimulating the production of lymphoid chemokines during inflammation (7, 8). Some of these signals have been identified in the family of the IL-23/IL-17 cytokines (9), but other cytokines have been advocated in TLO formation; thus different molecules may play differential roles depending on site and nature of the etiological agent (10).

IL-22, a member of the IL-10 cytokine superfamily, regulates mucosal responses to danger and wound healing (11). IL-22

promotes tissue repair, inducing epithelial cell proliferation and survival, in both physiological and pathological conditions (12–15). Recently, an association between IL-22 expression and autoimmune B-cell activation has been proposed (16). In Sjogren's syndrome, serum levels of IL-22 correlate with clinical manifestations including autoantibody production (17). Furthermore, B-cell-depleting treatment modulates salivary gland expression of IL-22, thus suggesting a potential functional relationship between IL-22 expression, B-cell infiltration, and local pathology (18). Previously we have demonstrated that direct cannulation of murine salivary glands with a replication-deficient adenovirus induces the formation of organized T-cell and B-cell aggregates, local expression of lymphoid chemokines, and the enzyme aicda [activation-induced cytidine deaminase (AID)] required for affinity maturation and isotype switching in B cells (19). Exploiting this model, we tested the hypothesis that IL-22 is involved in the generation of a humoral response within the TLOs (19). Here we demonstrate that early expression of IL-22 is instrumental for lymphoid chemokine production by stromal cells that, in turn,

## Significance

Ectopic clusters of immune cells that mimic the structure and function of secondary lymphoid organs are defined as tertiary lymphoid organs (TLOs). They have been observed at sites of chronic inflammation for decades, but their formation and function have remained enigmatic. TLOs are thought to contribute to disease pathogenesis by promoting autoreactive lymphocyte survival and autoantibody production. In this study we identify a novel role for the cytokine IL-22 in TLO development and biology. We provide evidence that IL-22 expression within TLOs is instrumental for the production of the lymphoid chemokines, chemokine (C-X-C motif) ligand 13 and chemokine (C-X-C motif) ligand 12, which in turn orchestrate B-cell clustering, lymphoid aggregation, and autoantibody production. Our data provide a strong rationale for targeting IL-22 in TLO-associated autoimmune diseases.

Author contributions: F.B. and C.D.B. designed research; S.N., J.C., T.C., D.R.W., Y.Z., and D.L. performed research; K.-M.T., L.F., B.F., S.B., J.R.-M., M.d.I.L.G.-H., T.D.R., M.B., and C.P. contributed new reagents/analytic tools; F.B., S.N., D.R.W., Y.Z., and D.L. analyzed data; and F.B., S.A.L., and C.D.B. wrote the paper.

The authors declare no conflict of interest.

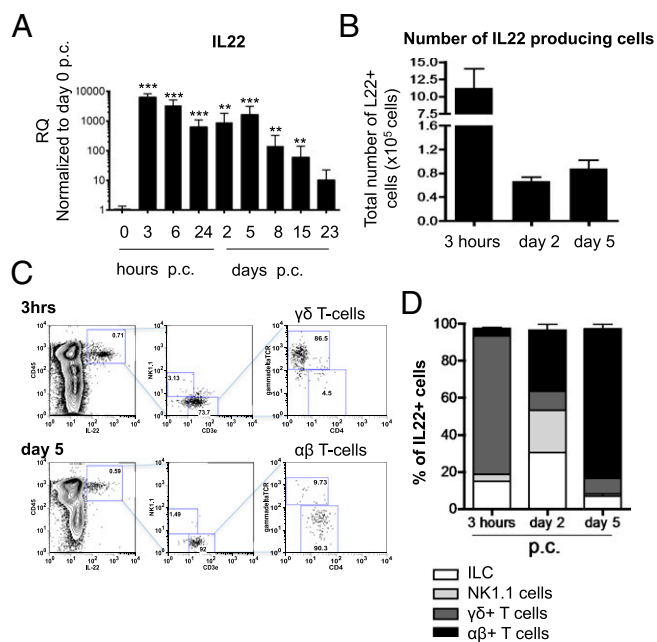
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This article contains supporting information online at [www.pnas.org/lookup/suppl/doi:10.1073/pnas.1503315112/-DCSupplemental](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1503315112/-DCSupplemental).



**Fig. 1.** Source of IL-22 in TLO formation. (A) Quantitative RT-PCR analysis of mRNA transcript for *IL22* in WT mice on day 0, at 3, 6, and 24 h p.c., and on days 2, 5, 8, 15, and 23 p.c., normalized to  $\beta$ -actin. Relative expression (RQ) was calibrated to 0 p.c. salivary glands.  $**P < 0.01$ ;  $***P < 0.001$  versus day 0. Results represent two or three experiments with four glands analyzed per group. (B) Absolute number of IL-22<sup>+</sup> cells in the CD45<sup>+</sup> population from salivary glands at 3 h p.c. and on days 2 and 5 p.c. (C) Representative dot plots identifying IL-22-producing cells in  $\gamma\delta$ <sup>+</sup> T cells (CD3e<sup>+</sup>gdTCR<sup>+</sup>NK1.1<sup>-</sup>CD4<sup>-</sup>) at 3 h p.c. and in  $\alpha\beta$ <sup>+</sup> T cells (CD3e<sup>+</sup>gdTCR<sup>-</sup>NK1.1<sup>-</sup>CD4<sup>+</sup>) on day 5 p.c. Data present the mean  $\pm$  SD of two different experiments with three salivary glands per experiment. (D) Distribution of IL-22<sup>+</sup> cells in the CD45<sup>+</sup> population at 3 h p.c. and on days 2 and 5 p.c. Data present the mean  $\pm$  SD from two different experiments with three salivary glands per experiment.

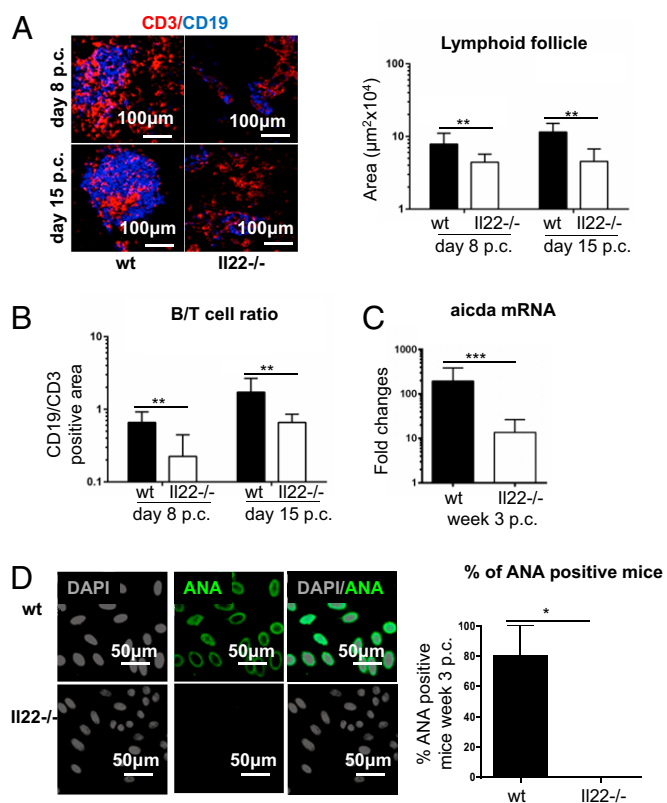
regulates B-cell aggregation. This work highlights a novel role for IL-22 in lymphoid chemokine expression and provides a mechanistic link between deranged mucosal responses and autoantibody production.

## Results

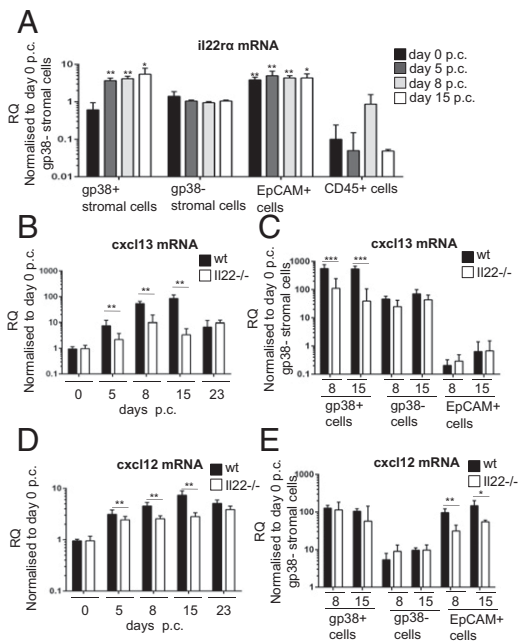
**Induction of IL-22-Expressing Cells in Murine Salivary Gland TLOs.** To determine whether IL-22 is produced in cannulated salivary glands, we analyzed mice early post viral cannulation (p.c.). In WT mice we detected early up-regulation of IL-22 mRNA transcript at 3 h p.c., with a second peak of expression at day 5 p.c. (Fig. 1A). These data were confirmed at the protein level by flow cytometry analysis of hematopoietic cells isolated from cannulated salivary glands. IL-22-producing cells were identified within the CD45<sup>+</sup> cell population by 3 h p.c., with the largest source of IL-22 [both in terms of percentage of positive cells and mean fluorescence index (MFI)] deriving from T cells at all time points examined (Fig. 1B–D and Fig. S1). At 3 h p.c. IL-22 production derived largely from  $\gamma\delta$  T cells, but by day 5 p.c.  $\alpha\beta$  T lymphocytes were the main producers of this cytokine (Fig. 1C). Other cell populations responsible for early IL-22 production were innate lymphoid cells (ILCs) and natural killer (NK) cells (Fig. 1D).

**TLO Formation Is Impaired in the Absence of IL-22.** To evaluate the consequence of IL-22 deficiency in TLO formation and autoantibody production, we moved our analysis to *IL22*<sup>-/-</sup> mice. First, resting salivary glands from *IL22*<sup>-/-</sup> and WT mice were evaluated for the presence of potential anatomical or structural differences that could interfere with virus infectivity. No

differences were found between WT and *IL22*<sup>-/-</sup> mice in resting condition by histological examination and flow cytometry analysis of the salivary gland epithelial component (Fig. S2A–C). Moreover, antiviral responses measured by ELISA showed no significant differences that could interfere with the establishment of the model of TLO formation in *IL22*<sup>-/-</sup> mice (Fig. S2D). Histological examination revealed that *IL22*<sup>-/-</sup> mice develop significantly smaller salivary gland lymphocytic aggregates than WT mice (Fig. 2A). Aggregates in the *IL22*<sup>-/-</sup> mice were characterized by a defect in B-cell accumulation (shown as a decreased B-cell/T-cell ratio) and follicular organization (Fig. 2A and B) as well as a strikingly lower expression of *aicda* transcripts (Fig. 2C). This defect was functionally associated with suppression in autoantibody production (Fig. 2D), suggesting that the lack of IL-22 impairs B-cell accumulation, organization, and maturation within TLOs. Interestingly, this defect was not observed in SLOs, where *aicda* expression was maintained post



**Fig. 2.** Lack of IL-22 results in defective TLO formation and decreased autoantibody production. (A, Left) Salivary gland staining for CD3 (red) and CD19 (blue) in WT and *IL22*<sup>-/-</sup> mice at days 8 and 15 p.c. (Right) Lymphoid aggregate area calculated at days 8 and 15 p.c. for WT (black bars) and *IL22*<sup>-/-</sup> (white bars) mice. Data present the mean  $\pm$  SD from two different experiments with two or three mice (four or six salivary glands) per group;  $**P < 0.01$ . (B) Ratio between the CD19<sup>+</sup> and CD3<sup>+</sup> area at days 8 and 15 p.c. in WT (black bars) and *IL22*<sup>-/-</sup> (white bars) mice. Data present the mean  $\pm$  SD from two different experiments (four or six salivary glands);  $**P < 0.01$ . (C) mRNA transcripts of the *aicda* gene (normalized to  $\beta$ -actin), in week 3 p.c. salivary glands from WT and *IL22*<sup>-/-</sup> mice. RQ values are relative to day 0 p.c. *aicda* mRNA; data represent the mean  $\pm$  SD of three independent experiments with four to six salivary glands analyzed in each experiment.  $***P < 0.001$ . (D, Left) Autoantibody detection on Hep2 cells from WT and *IL22*<sup>-/-</sup> mice at week 3 p.c. (nuclear staining in gray, autoantibody reactivity in green) (serum dilution 1:80). ANA, antinuclear antibody. (Right) Detection of autoantibody reactivity (percentage of mice positive for autoantibody at 1:80 dilution or more) in WT and *IL22*<sup>-/-</sup> mice at day 23 p.c. Data present the mean  $\pm$  SD of two experiments with five cannulated mice;  $*P < 0.05$ ; unpaired *t* test.



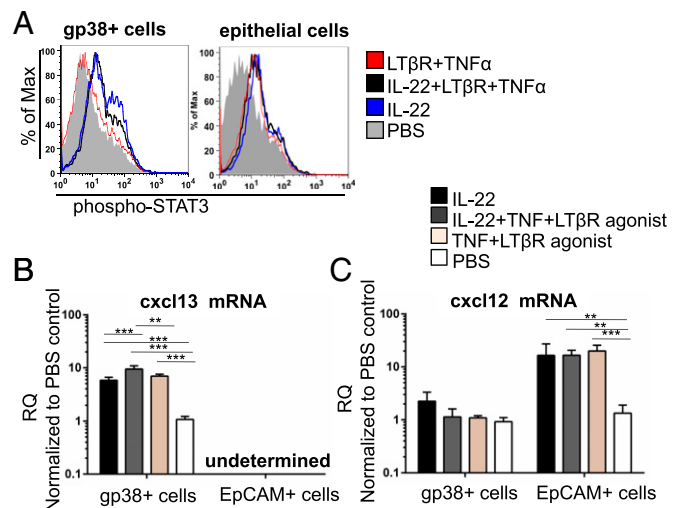
**Fig. 3.** IL-22-deficient mice fail to induce CXCL13 and CXCL12 expression within stromal cells of TLOs. (A) Expression of *I122ra* mRNA in FACS-sorted CD45<sup>+</sup>EpCAM<sup>-</sup>CD31<sup>-</sup>gp38<sup>+</sup> cells (black bars) in comparison with CD45<sup>+</sup>EpCAM<sup>-</sup>CD31<sup>-</sup>gp38<sup>-</sup> cells (white bars), CD45<sup>+</sup>EpCAM<sup>+</sup> epithelial cells (dark gray bars), and CD45<sup>-</sup> cells (light gray bars). *I122ra* mRNA transcripts were assessed by quantitative RT-PCR and normalized to  $\beta$ -actin. RQ was calculated with day 0 CD45<sup>+</sup>EpCAM<sup>-</sup>CD31<sup>-</sup>gp38<sup>-</sup> cells as calibrator. Results represent the mean  $\pm$  SD of two independent experiments with four biological replicates. \**P* < 0.05; \*\**P* < 0.01. (B and D) Quantitative PCR analysis of *cxcl13* and *cxcl12* mRNA transcripts obtained from salivary glands at days 5, 8, 15, and 23 p.c. from *I122*<sup>-/-</sup> (white bars) and WT (black bars) mice. Transcripts were normalized to *pdgfrb* mRNA; RQ values were calculated with day 0 salivary gland used as calibrator. \*\**P* < 0.01. Data present the mean  $\pm$  SD of two independent experiments with two or three mice per group (four or six salivary glands per experiment). (C and E) mRNA expression of *cxcl13* and *cxcl12* mRNA on FACS-sorted CD45<sup>+</sup>EpCAM<sup>-</sup>CD31<sup>-</sup>gp38<sup>+</sup>, CD45<sup>+</sup>EpCAM<sup>-</sup>CD31<sup>-</sup>gp38<sup>-</sup>, and EpCAM<sup>+</sup> epithelial cells from WT (black bars) and *I122*<sup>-/-</sup> (white bars) mice at days 8 and 15 p.c. (normalized to  $\beta$ -actin), calculated with day 0 CD45<sup>+</sup>EpCAM<sup>-</sup>CD31<sup>-</sup>gp38<sup>-</sup> cells used as calibrator. Data present the mean  $\pm$  SD of three biological replicates. \**P* < 0.05, \*\**P* < 0.01; \*\*\**P* < 0.001.

immunization (Fig. S3A). To evaluate the potential contribution of IL-17 expression to the overall effect observed in the *I122*<sup>-/-</sup> mice, quantitative PCR was performed on cannulated WT and *I122*<sup>-/-</sup> mice. Preserved IL-17 up-regulation was observed in *I122*<sup>-/-</sup> mice p.c., suggesting that the effect on TLO formation in the *I122*<sup>-/-</sup> mice is not dependent on IL-17 (Fig. S3B).

***I122*<sup>-/-</sup> Mice Fail to Induce Chemokine (C-X-C Motif) Ligand 13 and Chemokine (C-X-C Motif) Ligand 12 Expression Within TLOs.** To dissect the mechanism by which IL-22 affects B-cell accumulation and response in TLOs, we first evaluated the distribution of IL-22 receptor  $\alpha$  (IL-22R $\alpha$ ) on isolated cell populations from virus-cannulated salivary glands. As expected, IL-22R $\alpha$  expression was restricted to nonhematopoietic (CD45<sup>-</sup>) cells (Fig. 3A). IL-22R $\alpha$  mRNA expression was found on epithelial cells and on a nonendothelial, nonepithelial stromal cell population positive for gp38 [CD45<sup>-</sup>epithelial cell adhesion molecule (EpCAM)<sup>-</sup>CD31<sup>-</sup>gp38<sup>+</sup>], previously demonstrated to produce lymphoid chemokines during inflammation (20). We therefore hypothesized that IL-22R $\alpha$  expression on stromal cells might contribute to chemokine (C-X-C motif) ligand 13 (CXCL13) expression in our model, accounting for the decreased B-cell

recruitment observed in *I122*<sup>-/-</sup> mice. Indeed, total mRNA transcript and protein expression for CXCL13 were significantly decreased in *I122*<sup>-/-</sup> mice compared with WT mice (Fig. 3B and Fig. S4). These data were confirmed on sorted gp38<sup>+</sup> stromal cells that showed a significant decrease in the transcript for CXCL13 in *I122*<sup>-/-</sup> mice (Fig. 3C). In contrast, we observed no significant difference between WT and *I122*<sup>-/-</sup> mice in CXCL13 expression on sorted epithelial cells or gp38<sup>-</sup> stromal cells (Fig. 3C). Next, we investigated another B-cell attractant, chemokine (C-X-C motif) ligand 12 (CXCL12). Whole-gland mRNA transcript analysis revealed impaired CXCL12 expression in *I122*<sup>-/-</sup> mice (Fig. 3D). However, the decrease in CXCL12 expression was caused by the failure of CXCL12 up-regulation by epithelial cells rather than by gp38<sup>+</sup> stromal cells (Fig. 3E). Taken together, these findings demonstrate that IL-22 promotes differential CXCL13 and CXCL12 expression in gp38<sup>+</sup>/EpCAM<sup>-</sup> stromal cells and epithelial stromal cells within TLOs. These findings also were confirmed in *I122ra*<sup>-/-</sup> mice that showed decreased expression of the lymphoid chemokines CXCL13 and CXCL12 p.c., phenocopying the observation in *I122*<sup>-/-</sup> mice (Fig. S4 B and C).

**IL-22 Differentially Induces CXCL13 and CXCL12 Expression in Fibroblastic and Epithelial Cells in Vitro.** To confirm a direct role for IL-22 upstream of CXCL13 and CXCL12 production, we isolated gp38<sup>+</sup> stromal cells and EpCAM<sup>+</sup> epithelial cells from WT cannulated mice. Incubation of these cells with recombinant IL-22 protein induced the phosphorylation of Stat-3 (downstream of IL-22R $\alpha$  signaling) (Fig. 4A). Recombinant IL-22 led to up-regulation of CXCL13 in gp38<sup>+</sup> fibroblasts but not in epithelial cells (Fig. 4B). In contrast, IL-22 significantly stimulated CXCL12 expression in epithelial cells but not in fibroblasts



**Fig. 4.** IL-22 induces differential chemokine expression in fibroblasts and epithelial cells. (A) STAT3 phosphorylation assessed by flow cytometry in CD45<sup>+</sup>EpCAM<sup>-</sup>CD31<sup>-</sup>gp38<sup>+</sup> stroma and EpCAM<sup>+</sup> epithelial cells isolated from salivary glands at day 2 p.c. and stimulated in vitro with IL-22 (blue lines), IL-22, TNF- $\alpha$  and LT $\beta$ R agonist (black lines), TNF- $\alpha$  and LT $\beta$ R agonist (red lines), or PBS (gray lines). Two independent experiments were performed with three replicates per experiment. (B and C) Quantitative PCR analysis of *cxcl13* and *cxcl12* mRNA from CD45<sup>+</sup>EpCAM<sup>-</sup>CD31<sup>-</sup>gp38<sup>+</sup> stromal and EpCAM<sup>+</sup> epithelial cells isolated from salivary glands at day 2 p.c. stimulated in vitro with IL-22 (black bars), IL-22, TNF- $\alpha$ , and LT $\beta$ R agonist (gray bars), TNF- $\alpha$  and LT $\beta$ R agonist (light brown bars), or PBS (white bars). *cxcl13* and *cxcl12* mRNA transcripts were normalized to  $\beta$ -actin mRNA. Results are presented as RQ values compared with PBS-treated gp38<sup>+</sup> stromal cells (B) or epithelial cells (C) as calibrator. Data present the mean  $\pm$  SD of two independent experiments with three technical replicates per experiment. \*\**P* < 0.01; \*\*\**P* < 0.001.

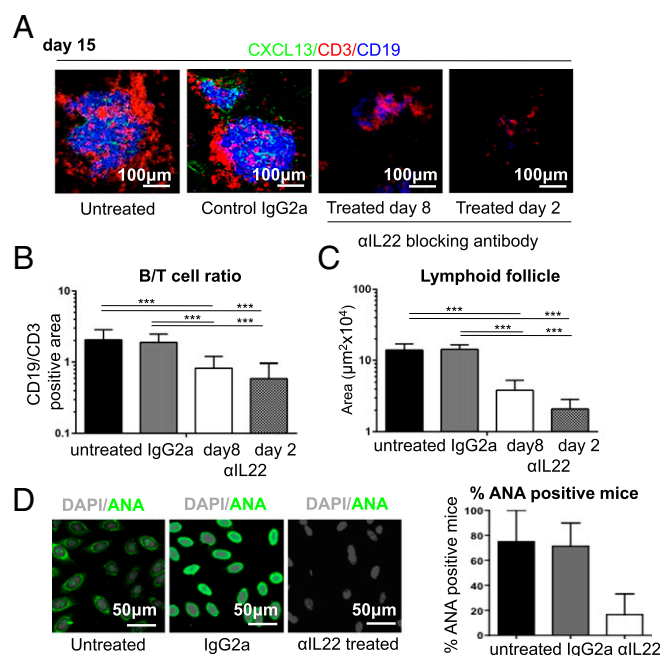
(Fig. 4C). TNF and lymphotoxin, which are known to activate chemokine expression by stromal cells, also induced Stat3 phosphorylation as well as inducing efficient CXCL13 and CXCL12 expression in a manner comparable to IL-22 on its own (Fig. 4B and C). Taken together, these data provide a molecular explanation for the defective B-cell accumulation and aberrant TLO formation observed in the *Il22*<sup>-/-</sup> mice, suggesting a novel role for IL-22 upstream of CXCL12 and CXCL13 expression in inflammatory conditions at mucosal sites.

**Therapeutic Blockade of IL-22 Inhibits Stromal Cell CXCL13 and CXCL12 Expression and Reduces Autoantibody Production.** We next explored whether therapeutic antibody blockade of IL-22 could induce regression of TLOs, once formed, and abolish autoantibody production, thus reproducing the phenotype observed in *Il22*<sup>-/-</sup> mice. We treated cannulated mice with anti-IL-22 antibody blocking starting at either day 2 or day 8 p.c. Immunofluorescence analysis on day 15 p.c. revealed defective lymphoid aggregate formation, both in terms of TLO size (Fig. 5A) and B-cell accumulation (Fig. 5A–C). This effect was more pronounced in mice that were treated from day 2 p.c. than in mice in which treatment began on day 8 p.c. (Fig. 5A–C). Quantitative RT-PCR analysis of the salivary glands from treated mice showed a strong reduction in the production of both CXCL13

and CXCL12 (Fig. S5). This defect was accompanied by suppression of *aicda* mRNA transcripts (Fig. S5) that coincided with a significant decrease in autoantibody production (Fig. 5D), supporting the concept that IL-22-blocking agents might be therapeutically effective in clinical practice.

## Discussion

IL-22 is known to regulate mucosal homeostasis and promote epithelial repair following tissue damage (21, 22). Recently, IL-22 has been implicated in malignant epithelial cell proliferation (23), but its role in chronic inflammation remains enigmatic (15). Although observational studies have proposed a relationship between mucosal IL-22 expression and humoral immunity in humans (18, 24) a direct functional role for IL-22 in ectopic B-cell activation and TLO formation has not been established. In this study we explored the hypothesis that IL-22 regulates TLO development and autoantibody production, influencing the release of lymphoid chemokines such as CXCL13 and CXCL12 that are critical for TLO assembly. We used a recently described disease model in which cannulation of the salivary glands with a replication-deficient adenovirus leads to the formation of TLOs with breakage of local tolerance and autoantibody production (19). Here we demonstrate that absence of IL-22 results in strongly reduced B-cell accumulation within TLOs and significant reduction in autoantibodies. This defective autoimmune response correlates with a failure to induce the B-cell chemoattractants CXCL13 and CXCL12 by fibroblasts and epithelial cells, respectively. The role of IL-22 in a model of lymphoid organ development has been described previously (25). Ota et al. (25) reported the ability of IL-22 to restore the organization of gut-associated lymphoid tissue in mice lacking lymphotoxin signals. Nonetheless, the authors interpreted their results as suggesting that IL-22 expression is under the control of lymphotoxin (25). Here we provide evidence that, in vitro, IL-22 can regulate the production of lymphoid chemokines in fibroblasts, independently of lymphotoxin. IL-22 is known to regulate RANKL expression on synovial fibroblasts from patients with rheumatoid arthritis via activation of p38 MAPK/NF- $\kappa$ B or JAK-2/STAT-3 signaling (26). Similarly, in our model we demonstrated that treatment of isolated salivary gland fibroblasts and epithelial cells with recombinant IL-22 was sufficient to induce STAT3 phosphorylation and production of CXCL13 and CXCL12 in these two stromal cell populations. Our work and that of Ota et al. (25) do not exclude the possibility that in vivo, during the dynamic formation of TLOs, IL-22 and lymphotoxin synergistically contribute to chemokine production, thus coregulating the enlargement, organization, and maintenance of the inflammatory aggregates. Strikingly, the induction of IL-22 expression occurs within hours of salivary gland cannulation and in different innate immune cell populations. Because *Il22*<sup>-/-</sup> mice are characterized by profound defects in early TLO maturation, with significant loss in lymphoid chemokine expression, we propose that first resident innate cells and later  $\gamma\delta$  and  $\alpha\beta$  T cells, sensing tissue damage, produce IL-22 to activate stromal cell expression of B-cell chemokines, thereby allowing the development of B-cell-rich infiltrates and autoantibody production. The recruitment of IL-22-producing retinoid-related orphan receptor  $\gamma$  t (ROR $\gamma$ t)<sup>+</sup> natural killer p46 (NKp46)<sup>-</sup> CD3<sup>-</sup> CD4<sup>+</sup> lymphoid tissue inducer cells in a CXCL13 transgenic model that mimics isolated lymphoid follicle formation in the gut has been described (27). Taken together, these data suggest that IL-22 plays a regulatory role in the early phases of lymphoid tissue development at mucosal sites. Transgenic expression of lymphoid chemokines is known to induce TLO formation (8, 28–30). However, TLOs can form in the absence of ROR $\gamma$ t lymphoid tissue inducer cells (9, 31, 32). This observation suggests that cytokines present within the inflammatory microenvironment provide alternative pathways of activation, different from those



**Fig. 5.** In vivo blockade of IL-22 reverses TLO formation and inhibits autoantibody production. (A) Lymphocytic aggregates from WT cannulated salivary glands from untreated mice, control IgG-treated mice, and mice treated with anti-IL-22 blocking antibody from day 8 or day 2 p.c. analyzed at day 15 p.c. for CD3 (red), CD19 (blue), and CXCL13 (green). (B) The ratio between the area covered by CD19<sup>+</sup> B cells and CD3<sup>+</sup> T cells at day 15 p.c. for the conditions above. Data present the mean  $\pm$  SD of two independent experiments with two or three mice (four or six salivary glands) per group. \*\*\* $P$  < 0.001. (C) Lymphoid follicle size for conditions above calculated as described in *Methods* for aggregates at day 15 p.c. Data present the mean  $\pm$  SD of two independent experiments with two or three mice (four or six salivary glands) per group. \*\*\* $P$  < 0.001. (D, Left) Immunofluorescent detection of autoantibody on Hep2 cells showing nuclear reactivity in sera at day 23 p.c. (nuclear staining in gray, autoantibody reactivity in green) (dilution 1:80). (Right) Graph summarizing autoantibody reactivity. The data present the mean  $\pm$  SD of two experiments with four cannulated mice (biological replicates expressed as the percentage of mice positive for autoantibody at 1:80 dilution or more).

conventionally associated with physiological lymphoid tissue formation. A role for IL-17 has been shown in two models of TLO formation in the lungs (9, 33). Peters et al. (34) also established the requirement for gp38<sup>+</sup> Th17 cells for the development of experimental allergic encephalomyelitis in the brain. We demonstrated *in vitro* and *in vivo* that IL-22 is sufficient to induce chemokine expression. Moreover in *Il22*<sup>-/-</sup> mice the signal for IL-17 is unaffected, thus excluding any significant contribution of IL-17 in the defect we observed in *Il22*<sup>-/-</sup> mice. Nonetheless, we cannot exclude a combined effect of IL-17 and IL-22 in TLO development at other sites and under different conditions. It has been elegantly demonstrated that CXCL13 expression is necessary and sufficient, both in the embryonic life and at ectopic sites, for the establishment of lymphoid follicles (28, 29) and the regulation of functional germinal centers (35). CXCL12 plays a role both in germinal center development and plasma cell attraction (30, 36). More recently, CXCL12 expression in nonepithelial stromal cells also has been implicated in B-cell recruitment to bronchial-associated lymphoid tissue (33). Carefully dissecting the source of lymphoid chemokines within TLOs, we have demonstrated that IL-22 exerts an unexpected differential role in the induction of these two chemokines on different stromal cell populations. On EpCAM<sup>+</sup>gp38<sup>+</sup> fibroblasts, IL-22 stimulation induces CXCL13 expression, both independently from and additively with TNF- and lymphotoxin  $\beta$  (LT $\beta$ )-receptor signals (known regulators of CXCL13 expression) (37). Conversely, on EpCAM<sup>+</sup>gp38<sup>+</sup> epithelial cells, IL-22R $\alpha$  engagement by IL-22 is able to increase CXCL12 but not CXCL13 production. Autoantibody production is the hallmark of many autoimmune diseases. Autoantibodies have been shown to precede disease onset, define prognosis, and stratify response to therapy (38–40). There is convincing evidence that AID expression on locally activated B cells within TLOs is able to support class-switch recombination and somatic hypermutation, thus sustaining affinity maturation of autoreactive B cells to tissue autoantigens (41, 42). Strikingly, therapeutic blockade of IL-22 by function-blocking antibodies is sufficient to impair TLO formation, by preventing organized B-cell aggregation and significantly affecting autoantibody production. In this context it is important to highlight the absence of a defect in *aicda* up-regulation in SLOs, further emphasizing the differences between SLO and TLO biology. Recently it has been shown that exposure of salivary gland epithelial cells to anti-Ro/SSA and La/SSB antibodies can induce IL-22 production (43). In agreement with these *in vitro* observations, there is evidence that B-cell depletion following treatment with rituximab leads to decreased expression of IL-22 in the salivary glands (18). This report suggests the possibility of a feed-forward loop involving IL-22 expression, survival of autoreactive B cells, and autoantibody production. Such an amplificatory loop, dependent on IL-22, also may be relevant to the well-known but poorly understood association between Sjogren's syndrome and the predisposition to B-cell mucosal-associated lymphoma (41). Previously we have shown that lymphoma-associated epithelial cells preferentially express CXCL12, potentially contributing to ductal infiltration and B-cell survival (5). It is tempting to speculate that, just as excess production of IL-22 leads to uncontrolled epithelial proliferation and tumor development in the intestine (42), overproduction of IL-22 in exocrine glands leads to excessive CXCL12 and CXCL13 production, contributing to B-cell persistence in TLOs and lymphoma development. The relationship between TLO persistence and cancer development is well recognized; for example, in the gut the formation of mucosa-associated lymphoid tissue lymphoma has been associated with excessive CXCL13 expression (44). Our data suggest that IL-22, a critical cytokine activated in response to mucosal danger signals, provides a previously unidentified functional link between mucosal autoimmunity, TLO development, and cancer. Our findings suggest

that targeting the IL-22 pathway might be an effective treatment in autoimmune diseases characterized by TLO formation. A translational program involving the use of anti-IL-22 in Sjogren's syndrome is currently in development in the Rheumatology Unit at the University of Birmingham.

## Methods

**Mice and Salivary Gland Cannulation.** C57BL/6 mice were purchased from Harlan. *Il22*<sup>-/-</sup> mice were generated at Lexicon Genetics in collaboration with Pfizer. *Il22Ra1*<sup>-/-</sup> mice were sourced from the European Mouse Mutant Archive, Sanger Center, United Kingdom. Mice were maintained in the Biomedical Service Unit at the University of Birmingham according to Home Office and local ethics committee regulations (University of Birmingham). Submandibular glands of 8- to 12-wk-old female C57BL/6 and knockout mice were intraductally cannulated with 10<sup>8</sup>–10<sup>9</sup> pfu of luciferase-encoding replication-defective adenovirus (Ad5) under anesthesia, as described (19). Animals were culled at specific time points by terminal anesthesia and cardiac puncture. Salivary glands and whole blood were harvested. Mice were excluded from the analysis on the basis of suboptimal virus delivery (Fig. S6) (19).

**In Vivo Blocking of Salivary Glands with Anti-IL-22.** Rat anti-mouse IL-22 Ab-03 was used as described (45). Starting at either day 2 or day 8 p.c., mice were administered a dose of 200  $\mu$ g of anti-IL-22 antibody via i.p. injection followed by injections three times per week.

**Histology and Immunofluorescence.** Seven-micrometer sections of salivary gland were cut by cryosectioning and were stained with directly conjugated or unconjugated antibodies (*SI Flow Cytometry Antibodies*). Sections were mounted using Prolong Gold Antifade reagent (Invitrogen Life Technologies). Images were acquired on a Zeiss LSM 510 laser-scanning confocal head with a Zeiss Axio Imager Z1 microscope; LSM510 Image Examiner Software was used to process the images.

**Isolation of Stromal Cells.** Harvested salivary glands were cut into small pieces and tissue digested for 40 min at 37 °C with stirring in 1.5 mL RPMI 1640 medium containing collagenase D (3.7 mg/mL) (Roche), DNase I (30  $\mu$ g/mL) (Sigma), and 2% (vol/vol) FCS. The suspension was gently pipetted at 15-min intervals to break up cell aggregates. Remaining fragments were digested further for 20 min at 37 °C with medium containing collagenase dispase (3.7 mg/mL) and DNase I (30  $\mu$ g/mL). EDTA was added to a final concentration of 5 mM; then cells were passed through a 70- $\mu$ m mesh, washed twice, and resuspended in RPMI 1640 medium containing 10% (vol/vol) FCS.

**Isolation of Leukocytes and in Vitro Stimulation for Cytokine Production.** After digestion, cells were resuspended in DMEM [with 10% (vol/vol) FCS, 1% Glutamine-Penicillin-Streptomycin (GPS) solution, 1% nonessential amino acids (NEAA), 1% HEPES, and 50  $\mu$ M  $\beta$ -mercaptoethanol] for *in vitro* stimulation culture. Isolated cells were incubated with 50 ng/mL phorbol-12-myristate-13-acetate, 750 ng/mL ionomycin, 10  $\mu$ g/mL Brefeldin A (all from Sigma Aldrich), 10  $\mu$ g/mL recombinant IL-23 (BioLegend), and 10  $\mu$ L of 10<sup>8</sup>–10<sup>9</sup> pfu of adenovirus for 4 h at 37 °C.

**Flow Cytometry.** Single-cell suspensions were incubated with antibodies on ice for 30 min in PBS (with 0.5% BSA and 2 mM EDTA) with mixtures of antibodies (*SI Flow Cytometry Antibodies*). Intracellular staining was performed according to the manufacturer's protocol (BD Cytofix/Cytoperm). After surface staining with mixtures of desired antibodies, cells were washed in PBS (with 0.5% BSA and 2 mM EDTA), resuspended in 150  $\mu$ L Cytofix/Cytoperm (BD Pharmingen), and incubated overnight at 4 °C. Cells were washed twice with the BD PermWash Buffer; then intracellular antibody was added, and cells were incubated at 4 °C for 30–40 min. Cells were washed twice, resuspended, and then analyzed using a Cyan-ADP (Dako) with forward/side scatter gates set to exclude nonviable cells. Data were analyzed with FlowJo software (Tree Star). For cell sorting, stained cells were sorted using MoFlo-XDP (Beckman Coulter, Inc.). Sorted cell purity exceeded 96%.

**In Vitro Stimulation of Sorted Stromal Cells with Recombinant Cytokines.** Isolated stromal cell subsets were resuspended at the same cell density in 500  $\mu$ L of DMEM [with 10% (vol/vol) FCS, 1% GPS, 1% NEAA, 1% HEPES, and 50  $\mu$ M  $\beta$ -mercaptoethanol] in 48-well plates. Recombinant cytokines were reconstituted in Dulbecco's phosphate-buffered saline (DPBS) (Sigma Aldrich). Cells were incubated with 2  $\mu$ g/mL of recombinant IL-22 (Peprotech) alone, 2  $\mu$ g/mL LT $\beta$ R agonist antibody (Novus Biologicals) plus 500 ng/mL recombinant TNF- $\alpha$  (Peprotech), 2  $\mu$ g/mL IL-22  $\mu$ g/mL plus LT $\beta$ R agonist

antibody plus 500 ng/mL TNF- $\alpha$ , or DPBS alone (Sigma Aldrich) for 24 h at 37 °C. Cells were harvested after 24 h and taken for quantitative PCR analysis.

**RNA Isolation and Quantitative PCR.** Total RNA was isolated from salivary glands with an RNeasy mini kit (Qiagen) and reverse-transcribed using the high-capacity reverse transcription cDNA synthesis kit (Applied Biosystems) on a Techne 312 Thermal Cycler PCR machine. Quantitative RT-PCR (Applied Biosystems) was performed on cDNA samples for *cxc13*, *cxc12*, *aicda*, *il22ra*, *il22*, and *il17* mRNA expression.  $\beta$ -actin and *pdgfrb* were used as an endogenous control. PCR products were detected using an ABI PRISM 7900HT instrument and were analyzed with the Applied Biosystems SDS software (SDS 2.3) as described (46).  $C_T$  values above 34 were not accepted, nor were technical replicates with more than two cycle differences between them.

**Detection of Autoantibodies and Antiviral Response.** Hep-2 cells cultured on slides (IMMCO Diagnostics) were incubated with murine serum diluted in PBS

(1:80). Sera were incubated for 1 h at room temperature, washed twice in PBS, and incubated with FITC anti-mouse IgG (Sigma) for 30 min. Slides were washed in PBS, mounted, and analyzed blindly by two different investigators. Antiviral titer was calculated as already described (19).

**Statistical Analysis.** Statistical significance was determined by Generalized Estimating Means (GEE analysis) followed by Sidak's significance test. In Figs. 3 A and C and 4 B and C statistical significance was determined by two-way ANOVA. In Fig. 2D an unpaired *t* test was used for statistical significance.

**ACKNOWLEDGMENTS.** We thank Dr. T. Cupedo and Dr. M. Coles for useful discussions during manuscript preparation, Dr. P. Nightingale for professional statistical advice, and the Biomedical Services Unit facility at the University of Birmingham. F.B. and D.R.W. are funded by the Wellcome Trust. C.D.B. is supported by Arthritis Research UK Grant 19791.

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