

Role of Lymphotoxin- α in Cigarette Smoke-Induced Inflammation and Lymphoid Neogenesis

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Abstract

In chronic obstructive pulmonary disease (COPD), chronic inflammation is accompanied by peribronchial lymphoid aggregates. Lymphotoxin- α (LT α), crucial in secondary lymphoid organogenesis, may be involved in lymphoid neogenesis.

We examined cigarette smoke (CS)-induced pulmonary lymphoid neogenesis and inflammation *in vivo* in LT α knockout (LT $\alpha^{-/-}$) and wild-type (WT) mice and studied expression of lymphoid chemokines by lung fibroblasts *in vitro*.

T-cell numbers (in bronchoalveolar lavage fluid (BALF) and lungs) and lymphoid aggregate numbers were significantly higher in air-exposed LT $\alpha^{-/-}$ mice than in WT animals, and increased upon chronic CS exposure in both genotypes. In contrast, local IgA responses upon chronic CS exposure were attenuated in LT $\alpha^{-/-}$ mice. CXCL13 and CCL19 mRNA in total lung and CXCL13 protein level in BALF increased upon CS exposure in WT, but not in LT $\alpha^{-/-}$ mice. *In vitro* lymphotoxin- β receptor (LT β R) stimulation induced CXCL13 and CCL19 mRNA in WT lung fibroblasts. Furthermore, *in vitro* exposure to cigarette smoke extract (CSE) up-regulated CXCL13 mRNA expression in WT, but not in LT β R $^{-/-}$, lung fibroblasts.

In this murine model of COPD, CS induces pulmonary expression of lymphoid chemokines CXCL13 and CCL19 in a LT α β -LT β R-dependent fashion. However, LT α is not required for CS-induced pulmonary lymphocyte accumulation and neogenesis of lymphoid aggregates.

Key words: chronic obstructive pulmonary disease, cigarette smoking, cytokines and chemokines, lymphotoxin alpha, mouse models

Introduction

Chronic obstructive pulmonary disease (COPD) is characterized by a slowly progressive airflow limitation that is poorly reversible (1). COPD is one of the leading causes of death with cigarette smoking as main risk factor (2). The molecular and cellular mechanisms responsible for the development of COPD are poorly understood. For that reason, a murine smoke-model of COPD was developed, showing a pulmonary pathology comparable to the disease in COPD patients (3), including inflammation and remodeling of the small airways as well as destruction of the lung parenchyma, emphysema.

Recently we (4), as well as others (5), reported the appearance of lymphoid aggregates in murine lung tissue upon chronic cigarette smoke (CS) exposure. Similar lymphoid follicles with clearly delineated T- and B-cell areas can be seen in lung sections of patients with severe COPD (6;7). The underlying mechanisms and functional properties of lymphoid follicle formation in the lung remain to be elucidated.

Lymphotoxin- α (LT α), essential in organogenesis of secondary lymphoid organs, may induce the development of tertiary lymphoid tissue. LT α knockout (LT $\alpha^{-/-}$) mice have no lymph nodes or Peyer's patches and aberrant splenic architecture (8;9), while site-specific LT α expression in transgenic animals causes local inflammation with a cellular composition and organization similar to those of lymph nodes (10).

LT α exists as a soluble homotrimer (LT α_3) or as a membrane-bound heterotrimer with lymphotoxin β (LT $\alpha_1\beta_2$). LT α_3 , produced by activated T-cells and early B-cells, is functionally redundant with TNF- α (11). LT $\alpha_1\beta_2$ on activated T- and B-lymphocytes and natural killer (NK) cells, binds exclusively to the TNFR-like LT β receptor (LT β R), on stromal and myeloid lineage cells (12). The LT $\alpha\beta$ -LT β R pathway is crucial in lymphoid organ development and triggers the expression of lymphoid chemokines, such as CXCL12,

CXCL13, CCL19 and CCL21 (13;14). Furthermore, this $LT\alpha\beta$ - $LT\beta R$ pathway has been implicated in the thymic emigration of ($V\alpha 14$) Invariant Natural Killer T (iNKT) cells (15).

To reveal the *in vivo* role of $LT\alpha$ in CS-induced inflammation, pulmonary emphysema, lymphoid neogenesis and mucosal IgA production, we subjected wild-type (WT) and $LT\alpha^{-/-}$ mice to subacute (4 weeks) or chronic (24 weeks) CS exposure. Moreover, we studied *in vitro* the expression of lymphoid chemokines by lung fibroblasts under baseline conditions and upon stimulation with agonistic $LT\beta R$ Ab and/or cigarette smoke extract (CSE).

Materials and Methods

Animals

Homozygous male C57Bl/6 $LT\alpha^{-/-}$ ($LT\alpha^{tm1Dch}$) mice and C57Bl/6 WT mice (8 weeks old) were obtained from The Jackson Laboratory (Bar Harbor, ME, USA) (8). The local Ethics Committee for animal experimentation of the faculty of Medicine and Health Sciences (Ghent, Belgium) approved all *in vivo* manipulations.

Smoke exposure

Mice (n = 8 per group) were exposed to cigarette smoke (CS), as described previously (16). Briefly, groups of 8 mice were exposed whole body to the tobacco smoke of 5 cigarettes (Reference Cigarette 2R4F without filter, University of Kentucky, Lexington, KY, USA), four times a day with 30 minutes smoke-free intervals, 5 days a week for 4 weeks (subacute exposure) or 24 weeks (chronic exposure). During the exposure an optimal smoke:air ratio of 1:6 was obtained. The control groups were exposed to air. Carboxyhemoglobin in serum of smoke-exposed mice reached a non-toxic level of $8.3 \pm 1.4\%$ (compared with $1.0 \pm 0.2\%$ in air-exposed mice, n = 7 for both groups), similar to carboxyhemoglobin blood concentrations of human smokers (17).

Bronchoalveolar lavage (BAL)

24 hours after the last exposure, mice were euthanized with an overdose of pentobarbital and BAL fluid was collected, as described previously (16). A total cell count was performed in a Bürker chamber, and differential cell counts (on at least 400 cells) were performed on cytocentrifuge preparations using standard morphologic criteria after May-Grünwald-Giemsa

staining. Flow cytometric analysis of BAL cells was performed to enumerate dendritic cells (DCs) and CD4⁺ and CD8⁺ T lymphocytes.

Preparation of lung single-cell suspensions

After rinsing of pulmonary and systemic circulation, the left lung was used for histology, and the right lung for the preparation of a single cell suspension, as detailed previously (16). Cell counting was performed with a Z2 Beckman Coulter particle counter (Beckman Coulter, Ghent, Belgium).

Labelling of BAL cells and single cell-suspensions for flow cytometry

The following mAbs were used to identify mouse dendritic cell (DC) populations: anti-CD11c-APC (HL3) and anti-I-A^b-PE (AF6-120.1). We discriminated between macrophages and myeloid DCs using the methodology described by Vermaelen and Pauwels (18). The following mAbs were used to stain mouse T cell subpopulations: anti-CD4-FITC (GK1.5), anti-CD8-FITC (53-6.7), anti-CD3-APC (145-2C11) and anti-CD69-PE (H1.2F3). Using anti-CD19-PE (1D3) and anti-CD11c, B lymphocytes were identified as the CD11c-low and CD19-positive population. All mAbs were obtained from BD Pharmingen. iNKT cells were stained with anti-CD3 and PE-conjugated CD1d tetramer loaded with α -galactosylceramide (α GalCer) (19). In a last step before analysis, cells were incubated with 7-aminoactinomycin D (or viaprobe; BD Pharmingen) to check cell viability. Flow cytometry data acquisition was performed on a FACScaliburTM running CellQuestTM software (BD Biosciences). FlowJo software was used for data analysis (TreeStar Inc., Ashland, OR, USA).

Histology

The left lung was fixated by intratracheal infusion of fixative (4% paraformaldehyde), as described previously (16). The lung lobe was embedded in paraffin and cut into 3- μ m transversal sections. Photomicrographs were captured using a Zeiss KS400 image analyzer platform (KS400, Zeiss, Oberkochen, Germany).

Quantification of pulmonary emphysema

To evaluate pulmonary emphysema, we determined the enlargement of the alveolar spaces by measuring the mean linear intercept (L_m), as described previously (4;16). Using image analysis software (ImageJ 1.34s) a 100 x 100- μ m grid was placed over images of hematoxylin&eosin-stained lung sections, acquired and scored in a blinded fashion. The total length of each line of the grid divided by the number of alveolar intercepts gives the average distance between alveolated surfaces, the L_m .

Morphometric quantification of lymphoid infiltrates

To evaluate the presence of lymphoid infiltrates in lung tissues, sections obtained from formalin-fixed, paraffin-embedded lung lobes were subjected to an immunohistological CD3/B220 double-staining as described previously (4). Infiltrates in the proximity of airways and blood vessels were counted. Dense accumulations of at least 50 cells were defined as lymphoid aggregates. Counts were normalized for the number of bronchovascular bundles per lung section.

Immunohistochemistry for CXCL13

Paraffin sections were incubated with primary anti-CXCL13 Ab (R&D Systems, Minneapolis, MN, USA), followed by biotinylated rabbit anti-goat-Ab from the Vectastain Elite ABC kit

(Vector Laboratories, Burlingame, CA, USA). After incubation with streptavidin-horseradish peroxidase, slides were coloured with diaminobenzidine (Dako, Carpinteria, CA, USA) and counterstained with Mayer's hematoxylin (Sigma-Aldrich, St. Louis, MO, USA).

ELISA

Commercially available ELISA kits were used to determine CXCL13 and CCL21 protein levels in BALF (R&D Systems) as well as IgM (ZeptoMetrix, Buffalo, NY, USA) and IgA (Alpha Diagnostic International, San Antonio, TX, USA) titers in serum and BALF. Secretory-IgA (S-IgA) was measured with a sandwich ELISA, developed in the laboratory of co-author CP. BALF samples were assayed using a polyclonal goat antibody to rat secretory component (SC), cross-reactive with mouse SC, as capture antibody and biotinylated anti-mouse IgA (Sigma-Aldrich) as detection antibody. S-IgA data were expressed as corrected OD, in the absence of a murine standard.

RT-PCR analysis

Total lung RNA was extracted with the RNeasy Mini Kit (Qiagen, Hilden, Germany). RNA from cultured cells was extracted with the ChargeSwitch Total RNA Cell Kit (Invitrogen Corp, Carlsbad, CA, USA). Expression of CXCL13, CCL19, CCL20, CXCR5 and CCR7 mRNA, relative to hypoxanthine guanine phosphoribosyltransferase (hppt) mRNA, was analyzed with the Assays-on-Demand Gene Expression Products (Applied Biosystems, Foster City, CA, USA). RT-PCR was performed on a LightCycler 480 Instrument (Roche Diagnostics, Basel, Switzerland) with murine leukemia virus RTase (Applied Biosystems) under previously described conditions (4).

Lung fibroblast culture

Lungs from a WT and a $LT\beta R^{-/-}$ mouse (20) (C57Bl/6) were digested, as described above, into single-cell suspensions. Cells were seeded in DMEM supplemented with 10% FBS, *L*-glutamine and penicillin/streptomycin (all from Gibco BRL, Invitrogen Corp) and incubated in a humidified 37°C incubator with 5% CO₂. 24 hrs after plating, non-adherent cells were removed by medium change. Cells were passaged at subconfluency. At passage 5, fibroblastic phenotype was checked on Lab-Tek chamber slides (Nalge Nunc, Rochester, NY, USA). In addition to positive staining of the cells for vimentin with monoclonal anti-vimentin Ab (clone VIM 13.2, Sigma-Aldrich), fibroblast morphology was confirmed by inverted phase-contrast microscopy.

Stimulation of cultured lung fibroblasts

At passage 6, cells were plated onto 24-well plates at a density of 4×10^4 cells/well and grown to confluency. Cells were stimulated for 48 hrs with 1 ml culture medium, isotype control (2 µg/ml), agonistic $LT\beta R$ Ab (2 µg/ml), 5% cigarette smoke extract (CSE), or the combination of agonistic $LT\beta R$ Ab with CSE. 100% CSE was prepared as described previously (21). Agonistic $LT\beta R$ Ab consisted of a 9:1 mix of 4H8 and 3C8 (Carl F. Ware, La Jolla Institute, San Diego, CA, USA). Isotype control consisted of a 9:1 mix of rat IgG_{2a} (A110-2) and rat IgG₁ (R3-34) (BD Pharmingen).

Statistical analysis

Reported values are expressed as mean \pm SEM. Statistical analysis was performed with Sigma Stat software (SPSS 15.0 Inc, Chicago, IL, USA) using nonparametric tests (Kruskall-Wallis; Mann-Whitney *U*). A *p* value ≤ 0.05 was considered significant.

Results

Modulation of CS-induced inflammation in BAL fluid of $LT\alpha^{-/-}$ mice

Subacute and chronic CS exposure increased the number of total cells, alveolar macrophages, dendritic cells (DCs), neutrophils and lymphocytes in the BAL fluid of both WT and $LT\alpha^{-/-}$ mice, compared to air-exposed animals (Fig. 1). Neutrophilic inflammation was significantly attenuated in $LT\alpha^{-/-}$ mice at both time points (Fig. 1 *D*). In contrast, baseline levels of BAL lymphocytes, more specifically both $CD4^{+}$ and $CD8^{+}$ T-cells, were significantly higher in air-exposed $LT\alpha^{-/-}$ mice than in WT mice, and increased further upon CS exposure (Fig. 1, *E* and *F*).

CS-induced pulmonary inflammation and emphysema

The CS-induced increase in lung macrophages and DCs was comparable between WT and $LT\alpha^{-/-}$ mice (Fig. 2, *A* and *B*). Baseline B- and T-cell numbers were higher in the lungs of air-exposed $LT\alpha^{-/-}$ mice versus WT mice. Contrary to the B-cell numbers, T-cell numbers increased strongly upon chronic CS exposure in the lung compartment of both WT and $LT\alpha^{-/-}$ mice (Fig. 2, *C* and *D*).

In both WT and $LT\alpha^{-/-}$ mice, chronic CS exposure induced significantly higher numbers of activated $CD4^{+}CD69^{+}$ and $CD8^{+}CD69^{+}$ T-cells. Yet again, both cell types were significantly increased in air-exposed $LT\alpha^{-/-}$ mice in comparison with WT mice (Fig. 2, *E* and *F*). These steady state differences in lymphocyte numbers, between air-exposed WT and $LT\alpha^{-/-}$ mice, were confirmed in the short term experiment (data not shown).

The L_m increased significantly upon chronic CS exposure in WT mice (air, $34.55 \pm 0.35 \mu\text{m}$ vs. smoke, $37.35 \pm 0.71 \mu\text{m}$; 8,1% increase; $p = 0,037$) as in $LT\alpha^{-/-}$ mice (air, $33.71 \pm 0.50 \mu\text{m}$

vs. smoke, $36.71 \pm 0.82 \mu\text{m}$; 8,9% increase; $p = 0,009$). There was no significant difference in L_m between CS-exposed WT and $LT\alpha^{-/-}$ mice ($p = 0,505$).

Increase of iNKT cells in the lung upon subacute CS exposure is $LT\alpha$ -dependent

iNKT cell numbers were determined through high specificity binding to CD1d-tetramer loaded with αGalCer (19) (Fig. 3 A). Subacute CS exposure caused a significant increase of iNKT cells in WT but not in $LT\alpha^{-/-}$ mice (Fig. 3, B and C). In contrast, chronic CS exposure did not change iNKT cell numbers in the WT and knockout mice, but $LT\alpha^{-/-}$ mice had lower percentages of iNKT cells compared to WT mice (Fig. 3, D and E).

$LT\alpha$ is not required for lymphoid neogenesis upon chronic CS exposure

Lymphoid aggregates were scarce in lung sections of air-exposed WT mice (Fig. 4, A and B), whereas the lungs of air-exposed $LT\alpha^{-/-}$ mice were strongly infiltrated with lymphocytes. (Fig. 4, A and D). Chronic (24 weeks) CS exposure significantly increased the number of peribronchial and perivascular lymphoid aggregates in the lungs of both WT and $LT\alpha^{-/-}$ mice (Fig. 4, A-E). Lymphoid aggregates were absent in the lungs of WT mice after subacute (4 weeks) exposure to air or CS (data not shown). In contrast, lungs of air-exposed $LT\alpha^{-/-}$ mice already showed a high degree of lymphocyte infiltration, but short term CS exposure did not induce additional aggregates (# aggregates / # bronchovascular bundles: air-exposed $LT\alpha^{-/-}$ mice: $0,31 \pm 0,059$, CS-exposed $LT\alpha^{-/-}$ mice: $0,28 \pm 0,054$; means \pm SEM, $n = 5$ animals/group).

Local and systemic IgA response upon chronic CS exposure is attenuated in $LT\alpha^{-/-}$ mice

Serum and BALF IgA levels were significantly diminished in $LT\alpha^{-/-}$ mice compared to WT animals, both at baseline in air-exposed mice and upon CS exposure for 24 weeks (Fig. 5). In serum, the relative increase in IgA levels upon chronic CS exposure was larger in the $LT\alpha^{-/-}$ mice compared to WT (Fig. 5A), whereas in BALF, the opposite was observed (Fig. 5B). Chronic CS exposure increased secretory IgA (S-IgA) levels in BALF of WT mice, while BALF S-IgA levels in $LT\alpha^{-/-}$ mice were significantly lower, both in air- and CS-exposed animals (Fig. 5C), indicating impairment of S-IgA responses. In contrast, IgM levels were higher in serum and BALF of $LT\alpha^{-/-}$ mice, both at baseline and upon CS exposure (Fig. 5, D and E).

$LT\alpha$ affects chemokine expression upon CS exposure

mRNA expression of CXCL13 and CCL19 was significantly increased in total lung after CS exposure in WT animals, whereas this increase could not be seen in $LT\alpha^{-/-}$ mice (Fig. 6, A and B). mRNA levels of CCL20/MIP-3 α , an attractant for immature DCs and lymphocytes, were increased in lungs of WT mice upon CS exposure (Fig. 6C). Baseline MIP-3 α expression was higher in $LT\alpha^{-/-}$ mice compared to WT mice, with a strong tendency to increase after chronic CS exposure ($p = 0,076$).

Total lung mRNA levels of CXCR5 and CCR7, the receptors for CXCL13 and CCL19 respectively, were higher at baseline in $LT\alpha^{-/-}$ mice (CXCR5: WT = air, 0.28 ± 0.045 vs. smoke, 0.36 ± 0.065 ; $LT\alpha^{-/-}$ mice = air, 2.09 ± 0.52 vs. smoke, 1.74 ± 0.26 ; CCR7: WT = air, 0.39 ± 0.038 vs. smoke, 0.73 ± 0.086 ; $LT\alpha^{-/-}$ mice = air, 1.29 ± 0.19 vs. smoke, 1.33 ± 0.16). CCR7 mRNA expression increased upon subacute CS exposure in WT, but not in $LT\alpha^{-/-}$ mice.

CXCL13 protein levels in BALF were up-regulated upon subacute and chronic CS exposure, again only in WT mice (Fig. 6D). Immunohistochemistry in WT mice revealed CXCL13 expression in CS-induced lymphoid aggregates (Fig. 6E). CS exposure did not affect protein levels of total CCL21 in BAL fluid of both WT and $LT\alpha^{-/-}$ mice (data not shown).

In vitro expression of CXCL13 and CCL19 in lung fibroblasts is $LT\beta R$ -mediated

To reveal the mechanism by which CXCL13 and CCL19 are induced in the lung, we stimulated cultured WT and $LT\beta R^{-/-}$ lung fibroblasts with agonistic $LT\beta R$ Ab and/or CSE.

In WT lung fibroblasts, *in vitro* stimulation with agonistic $LT\beta R$ Ab or CSE resulted respectively in a 60-fold and 5-fold increase in expression of CXCL13 mRNA. Furthermore, the combination of $LT\beta R$ Ab with CSE induced a 100-fold up-regulation in CXCL13 mRNA levels (Fig. 7A). A similar CCL19 mRNA response was seen upon stimulation with $LT\beta R$ Ab or $LT\beta R$ Ab with CSE combined, but there was no significant effect of CSE exposure alone (Fig. 7C). The different stimulations did not affect CXCL13 and CCL19 mRNA expression in $LT\beta R^{-/-}$ lung fibroblasts (Fig. 7, B and D), however, baseline mRNA levels of both CXCL13 and CCL19 were higher compared to WT lung fibroblasts (data not shown).

Discussion

Using *in vivo* studies with WT and $LT\alpha^{-/-}$ mice, we have demonstrated for the first time that CS-induced pulmonary lymphocyte accumulation and lymphoid neogenesis do not require $LT\alpha$. Moreover, $LT\alpha^{-/-}$ mice are not protected against the development of CS-induced emphysema. In contrast, $LT\alpha$ is required for CS-induced up-regulation, but not for baseline expression, of the lymphoid chemokines CXCL13 and CCL19. We have shown *in vitro* that the up-regulation of CXCL13 and CCL19 mRNA levels in lung fibroblasts is $LT\beta R$ -mediated. Furthermore, CSE itself is sufficient to induce CXCL13 expression in cultured lung fibroblasts via $LT\beta R$.

In accordance with previous reports (9), the lungs of air-exposed $LT\alpha^{-/-}$ mice were strongly infiltrated with B- and T-cells. Moreover, we found high steady state numbers of $CD4^+$ and $CD8^+$ T-cells in BALF of air-exposed $LT\alpha^{-/-}$ mice compared to WT mice, in line with earlier research (22). Extensive serology proved all animals to be free of a wide range of viruses and bacteria. Most likely, the lack of lymph nodes in $LT\alpha^{-/-}$ mice causes an accumulation of lymphocytes in organs such as the lungs and the liver, as described before (9).

Lymphoid chemokines are critical for neogenesis and organisation of lymphoid tissue (23;24). The lymphoid chemokines CXCL13 and CCL21 are greatly downregulated in the spleen, but not in the lungs of $LT\alpha^{-/-}$ mice (13;25;26), possibly causing a reversed chemokine gradient and consequent homing of lymphocytes to the lung. We found baseline mRNA levels of CXCL13, CCL19 and CCL20 to be elevated in lungs of air-exposed $LT\alpha^{-/-}$ mice compared to WT mice, which further explains increased recruitment of B- and T-cells to the lungs.

In WT mice, CS exposure increases the expression of CXCL13 and CCL19, a B-cell and T-cell attractant respectively. Furthermore, the presence of CXCL13 in pulmonary lymphoid aggregates of smoke-exposed WT mice indicates a role for this chemokine in ongoing organisation. Whereas $LT\alpha$ is not required for the baseline expression of CXCL13 and

CCL19, it is essential for the CS-induced up-regulation of these lymphoid chemokines. In secondary lymphoid tissue organogenesis, a positive feedback loop mechanism has been described: CXCL13 induces $LT\alpha_1\beta_2$ expression on attracted B-cells through CXCR5 and subsequent stimulation of $LT\beta R$ on stromal cells, which in turn up-regulates CXCL13 production (27;28). Indeed, we found cultured lung fibroblasts to be responsive to $LT\beta R$ stimulation through enhanced CXCL13 and CCL19 expression. More importantly, CSE induced *in vitro* CXCL13 expression in WT, but not in $LT\beta R^{-/-}$ lung fibroblasts. In line with a previous report on $LT\beta$ expression in stromal cells (29), we detected in these fibroblasts a discreet expression of $LT\beta$, which was up-regulated by CSE, only in WT fibroblasts (data not shown). CSE may thus contain or induce $LT\beta R$ ligands, implicating the $LT\beta R$ pathway in CS-induced CXCL13 expression in the lung. However, we can not exclude a general non-responsiveness of $LT\beta R^{-/-}$ lung fibroblasts towards the different stimuli. In figure 8 we propose a mechanism for lymphoid chemokine expression in the lung upon CS exposure.

Interestingly, CXCR5 and CCR7, the receptors for CXCL13 and CCL19 respectively, appear to be important for lymphoid neogenesis in chronic inflammatory autoimmune disease (30). In contrast, the inability of $LT\alpha^{-/-}$ mice to generate a CXCL13- and CCL19-response does not prevent the development of pulmonary lymphoid aggregates upon chronic CS exposure, suggesting other chemokines are involved. We demonstrated that CCL20, a chemokine that attracts immature DCs and lymphocytes, was higher in lungs of air-exposed $LT\alpha^{-/-}$ mice compared to WT mice, and increased further upon chronic CS exposure. CCL20 is thus possibly involved in both steady-state and CS-induced lymphocyte recruitment and lymphoid aggregate formation in $LT\alpha^{-/-}$ mice.

In WT mice, chronic CS exposure induces lymphoid neogenesis with germinal centre formation (5); the latter is known to be $LT\alpha$ -dependent (31). Accordingly, baseline and CS-induced IgA levels in BALF and serum were significantly attenuated in $LT\alpha^{-/-}$ mice compared

to WT animals, whereas $LT\alpha^{-/-}$ mice also showed higher baseline IgM titers in BALF and serum, consistent with a putative Ig isotype switching deficiency. WT mice generated a strong IgA response in BALF upon chronic CS exposure. Similarly, IgA appears to be increased in bronchial secretions of chronic bronchitis patients (32;33). However, when IgA levels are corrected for albumin – to exclude diffusion effects – there is in fact a decrease of indices of local active transport of IgA across the epithelium (32;33), which could relate to reduced expression of epithelial polymeric immunoglobulin receptor/transmembrane secretory component in COPD airways (34). In CS-exposed WT mice, IgA transport towards the airway lumen appeared intact, whereas the strongly reduced S-IgA concentrations in BALF of $LT\alpha^{-/-}$ animals could relate to decreased local IgA synthesis (general defect in class switching to IgA, which could also underlie reduced serum IgA levels) and/or to decreased trans-epithelial transport. Altogether, our B-cell data could indicate impairment in class switch recombination mechanisms in $LT\alpha^{-/-}$ mice, as previously suggested by others (35).

Currently, there is considerable controversy on the role of iNKT cells in airway disease (36;37), a cell population that is profoundly diminished in the periphery of $LT\alpha$ deficient mice (15). Therefore, iNKT cell frequencies were monitored during the course of CS exposure. iNKT cell percentages were significantly reduced in lungs of $LT\alpha^{-/-}$ mice in the chronic experiment. Moreover, subacute CS-exposure increased pulmonary iNKT cell numbers in a $LT\alpha$ -dependent fashion. This is in line with the modulating role of iNKT cells between innate and adaptive immunity, since at this time point (4 weeks), immune responses shift from innate to adaptive in this experimental model of COPD (16). When the adaptive immune system has kicked in, iNKT cell numbers are no longer different between air and CS exposed lungs of WT mice. Corresponding with data from COPD patients (37), our findings do not indicate a role for iNKT cells in the chronic disease model.

In COPD patients, lymphoid aggregates have been correlated with disease severity (6;7). Hypothesizing that the pathogenesis of COPD has an autoimmune component, pulmonary lymphoid aggregates could add to the perpetuating nature of COPD and enhance production of auto-antibodies in the germinal centres. On the other hand, ectopic lymphoid tissue could very well have a local protective role against infections (7). Although $LT\alpha$ polymorphisms are not directly associated with COPD (38;39), potential indirect effects on inflammatory mediators should not be excluded.

In conclusion, we have demonstrated that CS-induced pulmonary lymphoid neogenesis does not require $LT\alpha$. CS induces the pulmonary expression of lymphoid chemokines CXCL13 and CCL19 in a $LT\alpha\beta$ - $LT\beta R$ -dependent fashion. This murine model of COPD will be useful to further characterize the developing mechanisms and function of CS-induced lymphoid neogenesis.

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Figure Legends

FIGURE 1. Total BAL cells and cell differentiation in BAL fluid of wild-type (WT) and $LT\alpha^{-/-}$ mice after subacute (4 weeks) or chronic (24 weeks) exposure to air or cigarette smoke: *A*, total BAL cells; *B*, macrophages; *C*, dendritic cells; *D*, neutrophils; *E*, $CD4^{+}$ T-cells and *F*, $CD8^{+}$ T-cells (*A* and *D* were calculated on cytospins; *B*, *C*, *E* and *F* were enumerated by flow cytometry). Results are expressed as mean \pm SEM. $n = 8$ animals/group; *, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$.

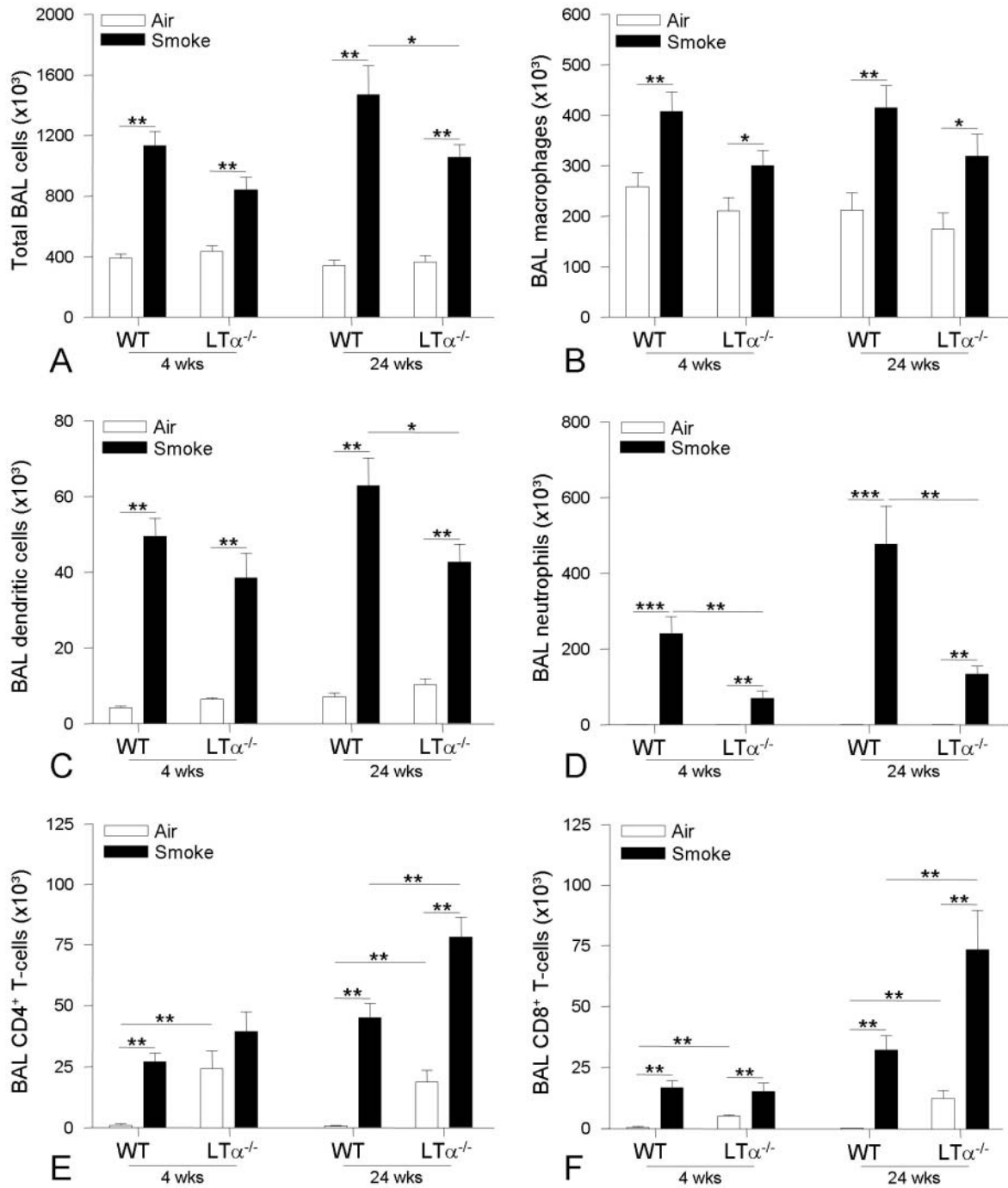


Figure 1.

FIGURE 2. Cell differentiation in the lungs of wild-type (WT) and $LT\alpha^{-/-}$ mice after chronic (24 weeks) exposure to air or cigarette smoke: *A*, macrophages; *B*, dendritic cells; *C*, B-cells; *D*, T-cells; *E*, activated $CD4^{+}$ T-cells and *F*, activated $CD8^{+}$ T-cells (all cell types were

enumerated by flow cytometry). Results are expressed as mean \pm SEM. $n = 8$ animals/group;

*, $p < 0.05$; **, $p < 0.01$.

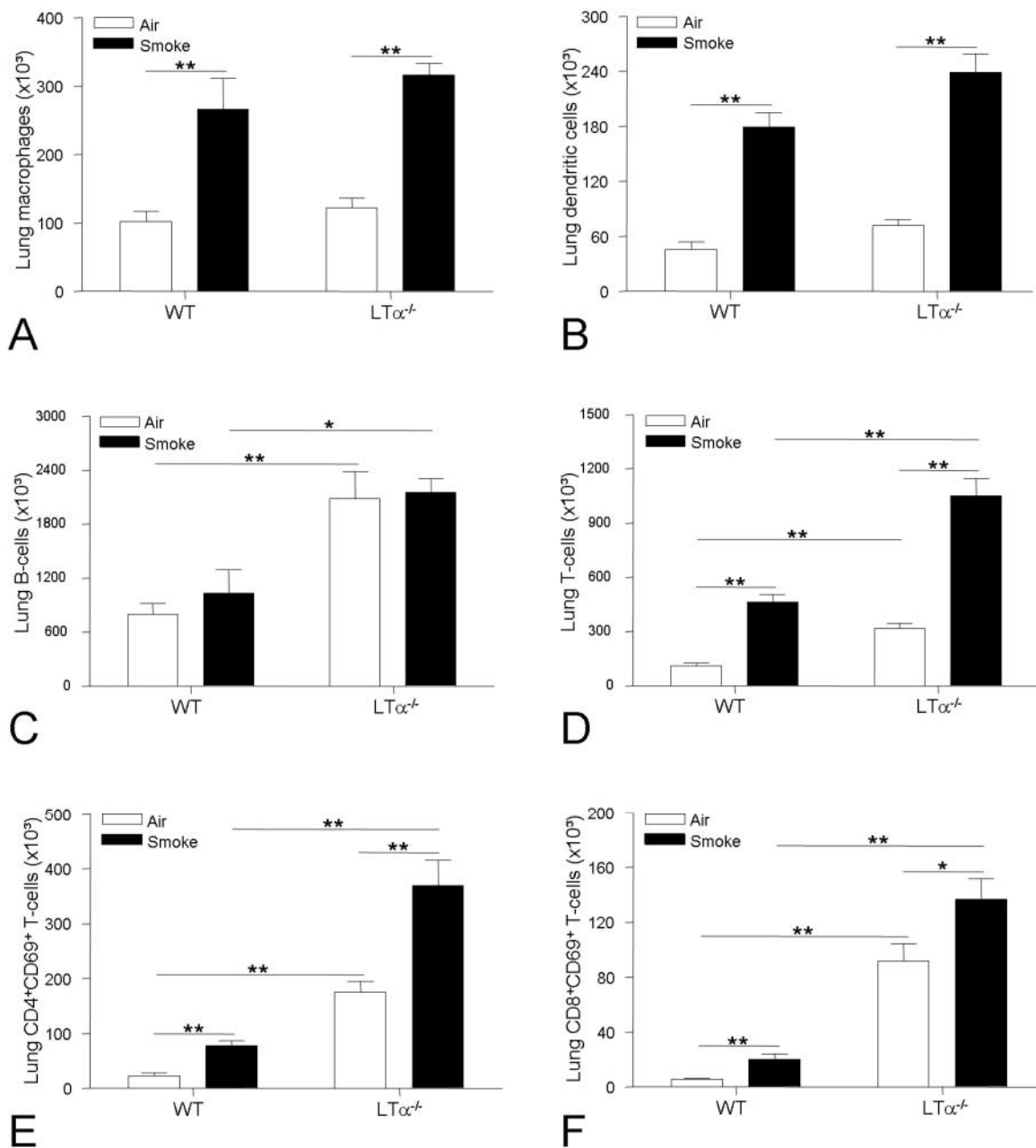


Figure 2.

FIGURE 3. Invariant Natural Killer T (iNKT) cells in lung digests of wild-type (WT) and $LT\alpha^{-/-}$ mice after subacute (4 weeks) or chronic (24 weeks) exposure to air or cigarette smoke, enumerated by flow cytometry. *A*, iNKT cells were stained with APC-conjugated anti-CD3

antibody and PE-conjugated CD1d-tetramer loaded with α -galactosylceramide (α -GalCer).

Results are expressed as percentage of total cells (B and D) and as absolute cell number (C

and E), mean \pm SEM. $n = 8$ animals/group; *, $p < 0.05$.

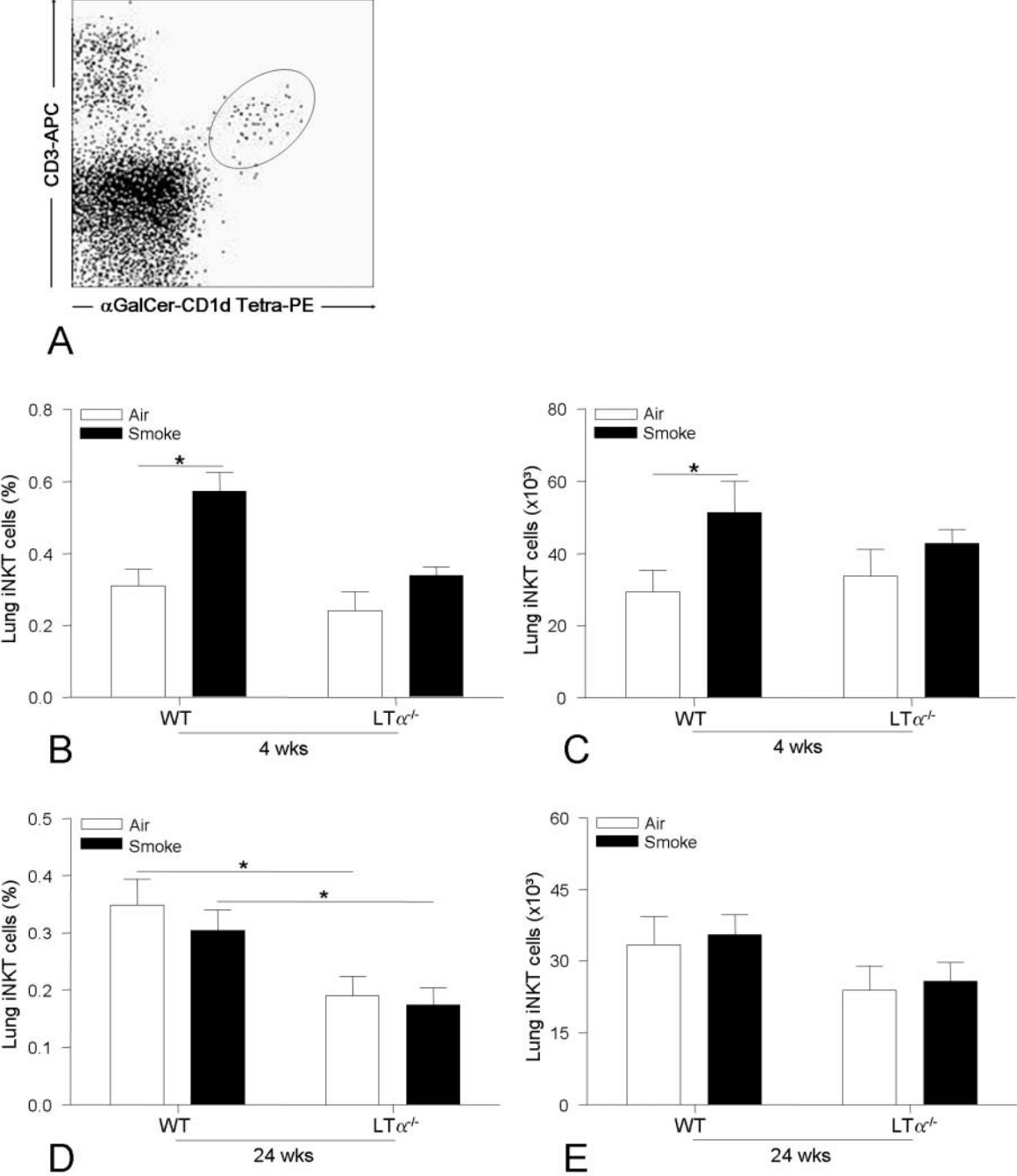


Figure 3.

FIGURE 4. Lymphoid aggregates in lung tissue of wild-type (WT) and $LT\alpha^{-/-}$ mice after chronic (24 weeks) exposure to air or cigarette smoke (CS). *A*, Quantification of peribronchial and perivascular lymphoid aggregates. Results are expressed as mean \pm SEM. $n = 8$ animals/group; **, $p < 0.01$; ***, $p < 0.001$. *B-E*, Photomicrographs of lymphoid aggregates in CD3/B220 immuno-stained lung tissue of air- and smoke-exposed WT and $LT\alpha^{-/-}$ mice at 24 weeks (magnification, $\times 100$; brown = CD3 positive cells; blue = B220 positive cells). *B*, air-exposed WT mice; *C*, CS-exposed WT mice; *D*, air-exposed $LT\alpha^{-/-}$ mice; and *E*, CS-exposed $LT\alpha^{-/-}$ mice. Scale bars = 100 μm .

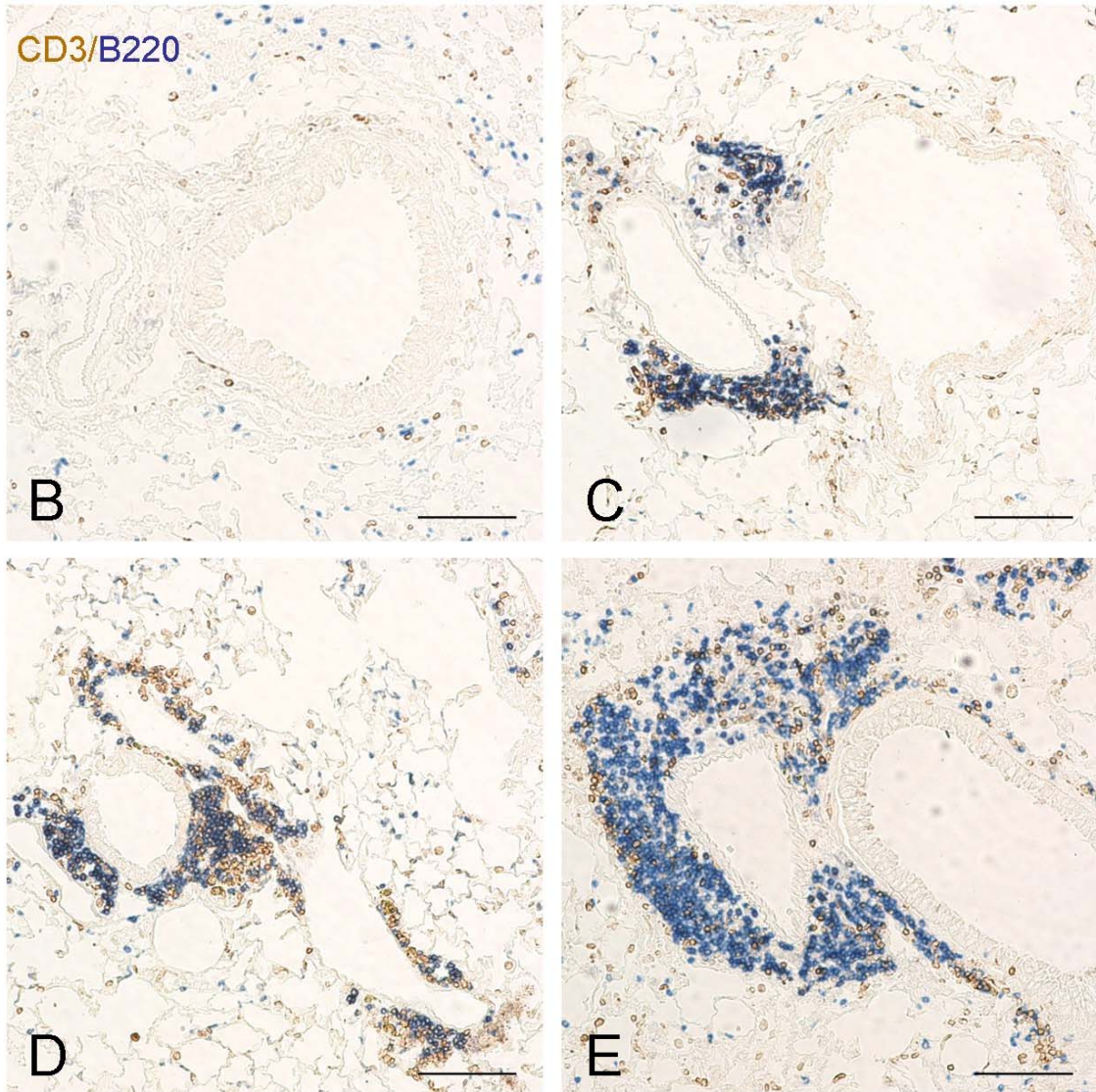
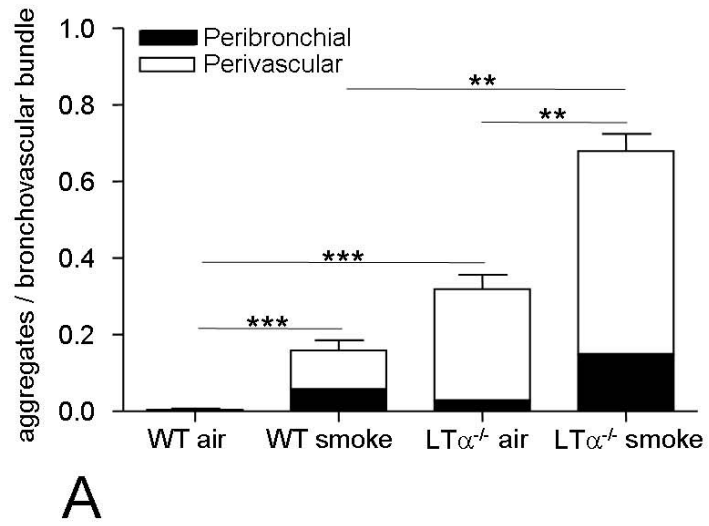


Figure 4.

FIGURE 5. Immunoglobulin A (IgA) levels in serum (*A*), total IgA (*B*) and secretory IgA (*C*) in BAL fluid of wild-type (WT) and $LT\alpha^{-/-}$ mice after chronic (24 weeks) exposure to air or cigarette smoke, as measured by ELISA. IgA titers are expressed as $\mu\text{g/ml}$ and secreted-IgA (S-IgA) levels as corrected optical density (OD) (mean \pm SEM). $n = 8$ animals/group; *, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$. Immunoglobulin M (IgM) levels in serum (*D*) and BAL fluid (*E*) of WT and $LT\alpha^{-/-}$ mice after subacute (4 weeks) exposure to air or cigarette smoke, as measured by ELISA. Results are expressed as $\mu\text{g/ml}$ (serum IgM) and ng/ml (BALF IgM), mean \pm SEM. $n = 8$ animals/group; *, $p < 0.05$; **, $p < 0.01$.

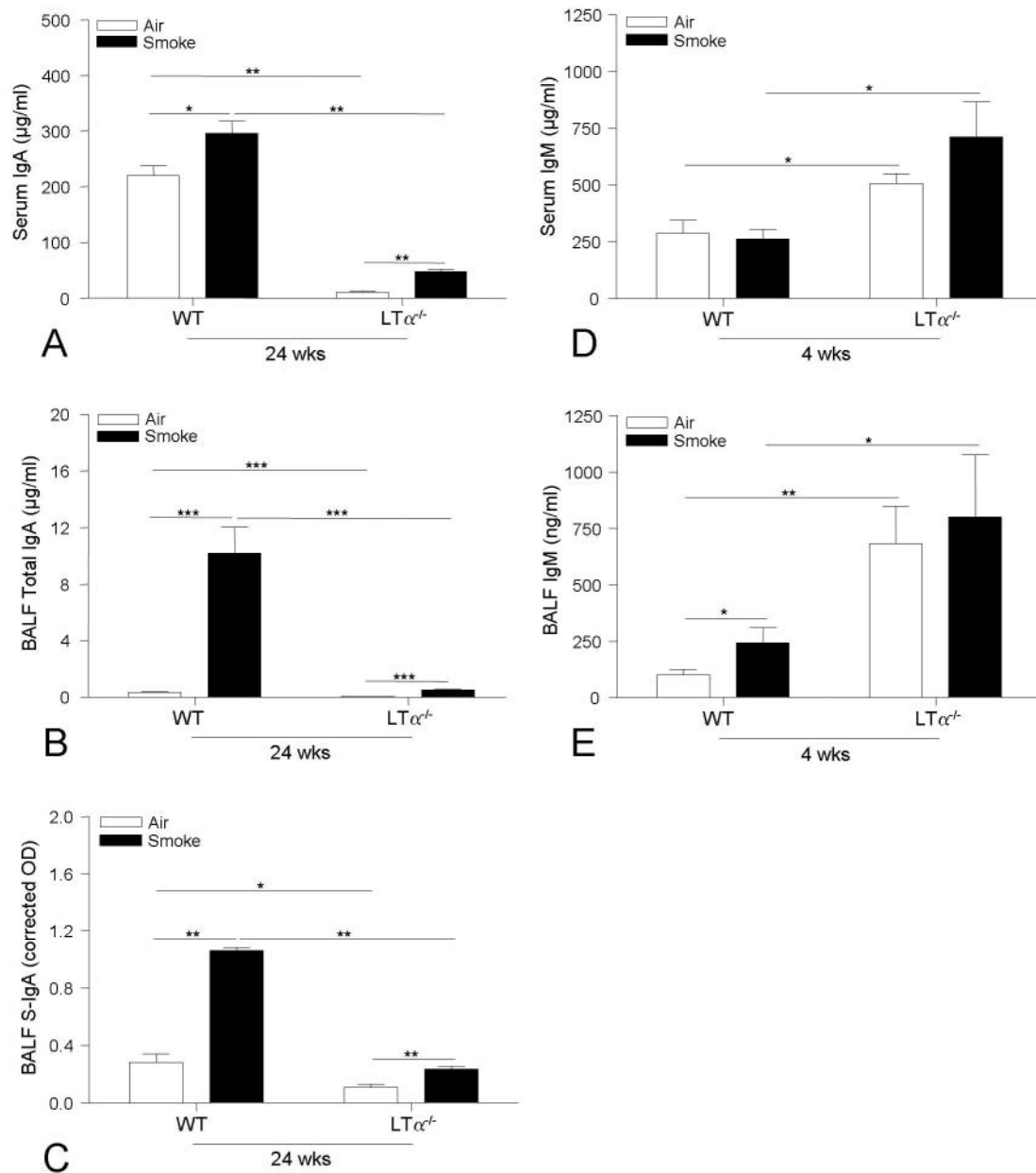


Figure 5.

FIGURE 6. Measurement of chemokines in BAL fluid and lung tissue. mRNA levels of CXCL13 (A), CCL19 (B) and CCL20 (C) in total lung tissue of wild-type (WT) and $LT\alpha^{-/-}$ mice after subacute (4 weeks) or chronic (24 weeks) exposure to air or cigarette smoke, as measured by RT-PCR. Results are expressed as ratio with *hprt* mRNA (mean \pm SEM). $n = 5$ animals/group; *, $p < 0.05$; **, $p < 0.01$. D, Protein levels of CXCL13 in BAL fluid of WT and $LT\alpha^{-/-}$ mice after subacute (4 weeks) or chronic (24 weeks) exposure to air or cigarette

smoke, as measured by ELISA. Results are expressed as pg/ml (mean \pm SEM). $n = 8$ animals/group; *, $p < 0.05$; **, $p < 0.01$. *E*, Photomicrographs of a CXCL13-positive lymphoid aggregate in a smoke-exposed WT animal at 24 weeks. Magnification, x100: scale bar = 100 μ m; magnification, x400: scale bar = 20 μ m.

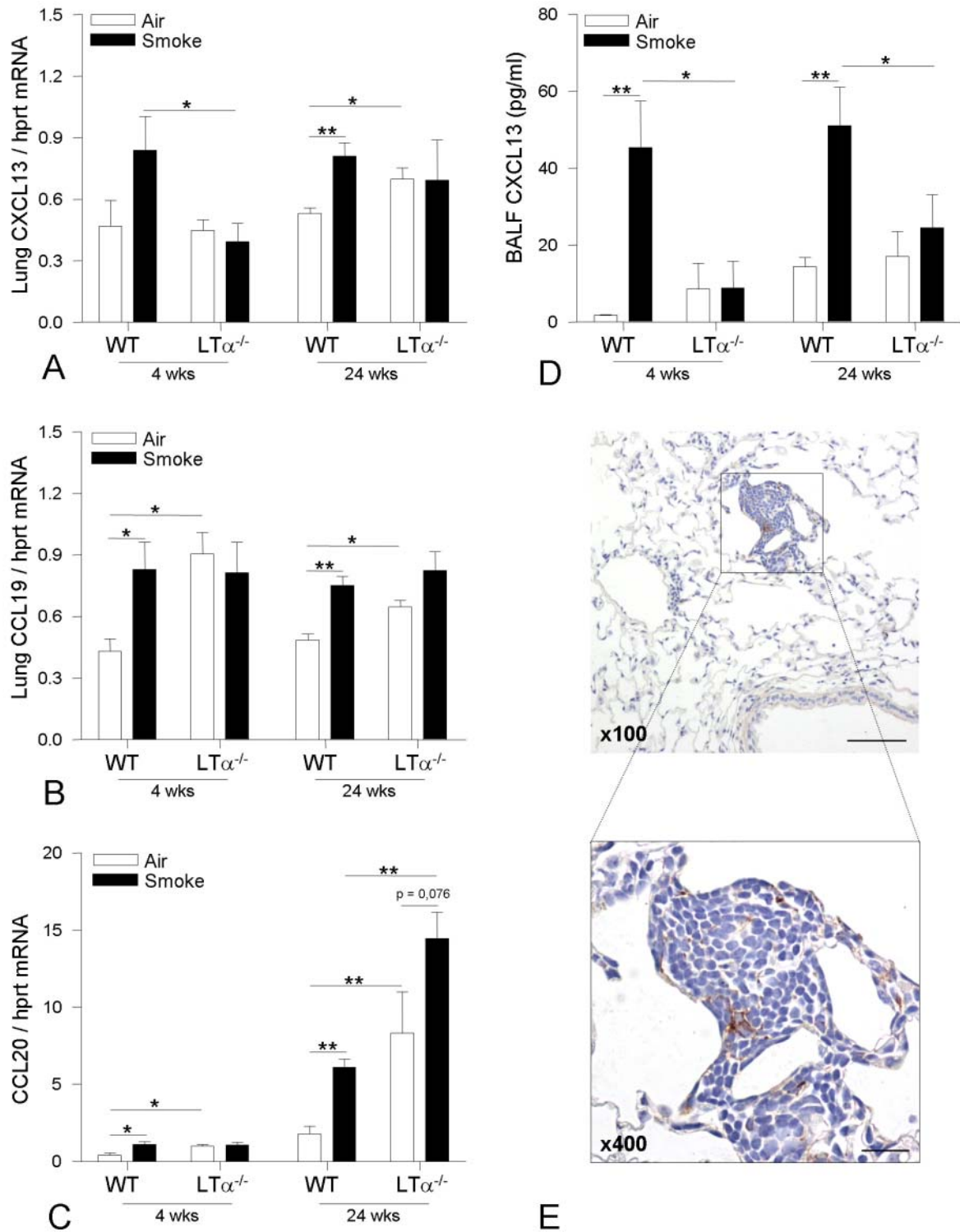


Figure 6.

FIGURE 7. Expression of lymphoid chemokines CXCL13 and CCL19 by lung fibroblasts, in response to LT β R stimulation *in vitro*. A-D, Cultured lung fibroblasts were exposed for 48 hours to DMEM, isotype control, agonistic LT β R antibody (Ab), cigarette smoke extract

(CSE) or the combination of LT β R Ab with CSE (Ab+CSE). mRNA levels of CXCL13 (*A* and *B*) and CCL19 (*C* and *D*) in WT and LT β R^{-/-} fibroblasts, as measured by RT-PCR. Results are expressed as ratio with hprt mRNA; mean of 3 experiments \pm SEM; *, $p \leq 0.05$.

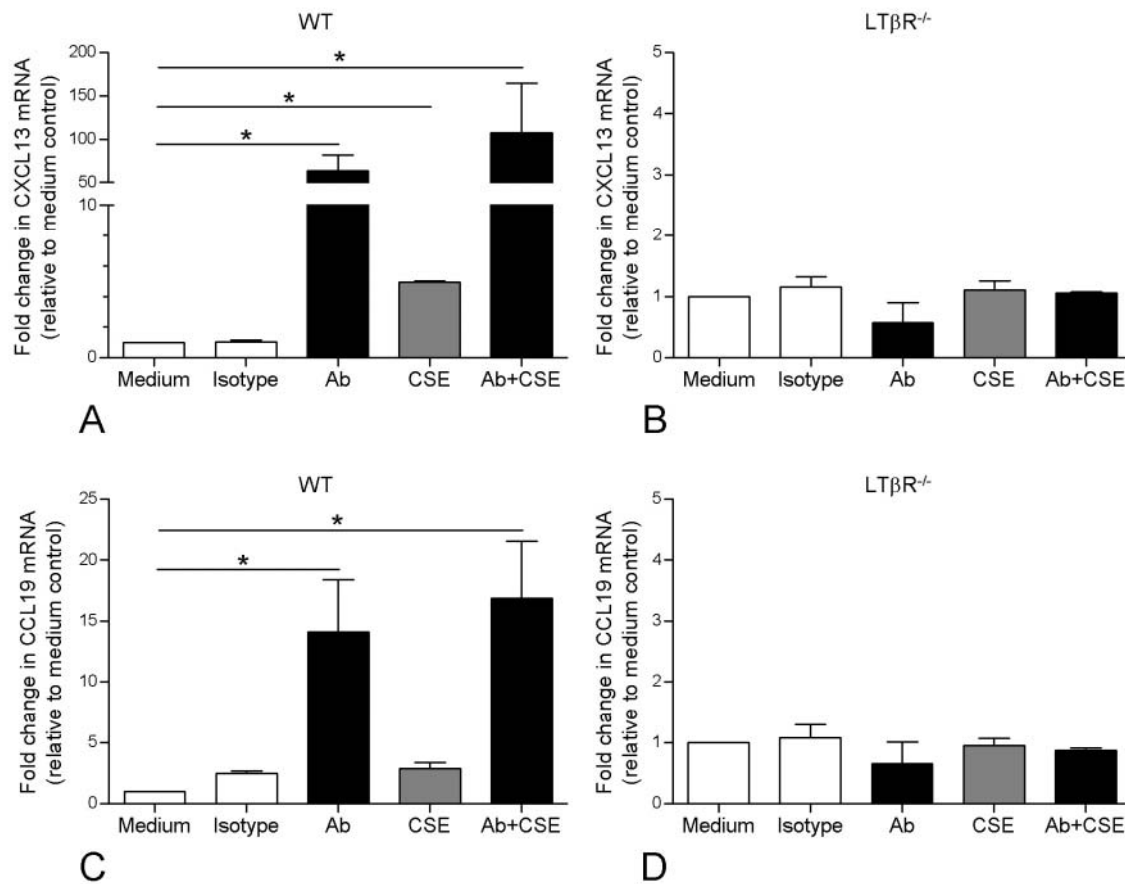


Figure 7.

FIGURE 8. Proposed mechanism for lymphoid chemokine expression in the lung upon CS exposure. CS induces pulmonary inflammation and subsequent expression of LT $\alpha\beta$ on lymphocytes, but may also directly stimulate the LT β R on stromal cells, such as lung fibroblasts. LT β R-stimulation leads to the production of CCL19 and CXCL13 which attract DCs and T-cells, and B-cells respectively. Finally, CXCL13 may induce LT $\alpha\beta$ on B-cells, creating a positive feedback loop mechanism.

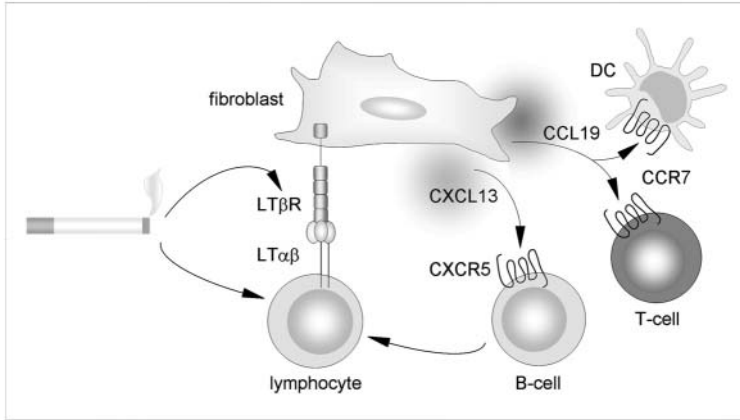


Figure 8.