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Article Angiogenesis

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# Pathogenesis of persistent lymphatic vessel hyperplasia in chronic airway inflammation

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**Edema occurs in asthma and other inflammatory diseases when the rate of plasma leakage from blood vessels exceeds the drainage through lymphatic vessels and other routes. It is unclear to what extent lymphatic vessels grow to compensate for increased leakage during inflammation and what drives the lymphangiogenesis that does occur. We addressed these issues in mouse models of (a) chronic respiratory tract infection with *Mycoplasma pulmonis* and (b) adenoviral transduction of airway epithelium with VEGF family growth factors. Blood vessel remodeling and lymphangiogenesis were both robust in infected airways. Inhibition of VEGFR-3 signaling completely prevented the growth of lymphatic vessels but not blood vessels. Lack of lymphatic growth exaggerated mucosal edema and reduced the hypertrophy of draining lymph nodes. Airway dendritic cells, macrophages, neutrophils, and epithelial cells expressed the VEGFR-3 ligands VEGF-C or VEGF-D. Adenoviral delivery of either VEGF-C or VEGF-D evoked lymphangiogenesis without angiogenesis, whereas adenoviral VEGF had the opposite effect. After antibiotic treatment of the infection, inflammation and remodeling of blood vessels quickly subsided, but lymphatic vessels persisted. Together, these findings suggest that when lymphangiogenesis is impaired, airway inflammation may lead to bronchial lymphedema and exaggerated airflow obstruction. Correction of defective lymphangiogenesis may benefit the treatment of asthma and other inflammatory airway diseases.**

## Introduction

Lymphatic vessels provide key routes for drainage of interstitial fluid from tissues. Edema, a cardinal sign of inflammation and clinically significant feature of inflammatory disease, results when the amount of leakage from inflamed blood vessels exceeds the capacity of lymphatic vessels for drainage. Despite the recognition that angiogenesis and mucosal swelling can increase airflow resistance in obstructive lung disease (1, 2), remarkably little is known about the status of lymphatic vessels and factors that influence fluid drainage from the respiratory tract. Lymphatic vessels may proliferate under inflammatory conditions (3), yet the time course, extent, molecular mechanisms, and functional consequences of lymphangiogenesis are largely unknown.

Vascular endothelial growth factors VEGF-C and VEGF-D drive lymphangiogenesis via VEGFR-3 signaling (4–7). The exclusivity of this effect is unclear, because VEGF-C and VEGF-D may also induce angiogenesis (6, 8, 9). Added to the list of potential growth factors, VEGF (VEGF-A, also known as vascular permeability factor [VPF]), well known for its potent angiogenic actions, and PDGF appear to drive lymphatic growth under certain conditions (10, 11).

Impairment of lymphatic drainage, most commonly caused by genetic mutation, filarial infection, surgery, or trauma, results in lymphedema with sometimes marked swelling of limbs or other regions (12). It follows that defects in lymphangiogenesis in the respiratory tract, especially when plasma leakage is exaggerated by inflammation, may produce edema of the airway mucosa (bronchial lymphedema). Because of the strategic role of mucosal thickness in determining the effect of bronchial constriction on airway caliber and airflow resistance (1), bronchial lymphedema could exacerbate airway obstruction in inflammatory disease.

Blood vessels that proliferate and undergo remodeling in inflammatory airway disease rapidly return to normal as the inflammation resolves during treatment (13, 14). However, lymphatic growth may not be so readily reversible (15, 16). Persistence of new lymphatic networks after resolution of inflammation would leave in place a drainage system for fluid and immune cells that may accelerate responses in subsequent inflammatory episodes.

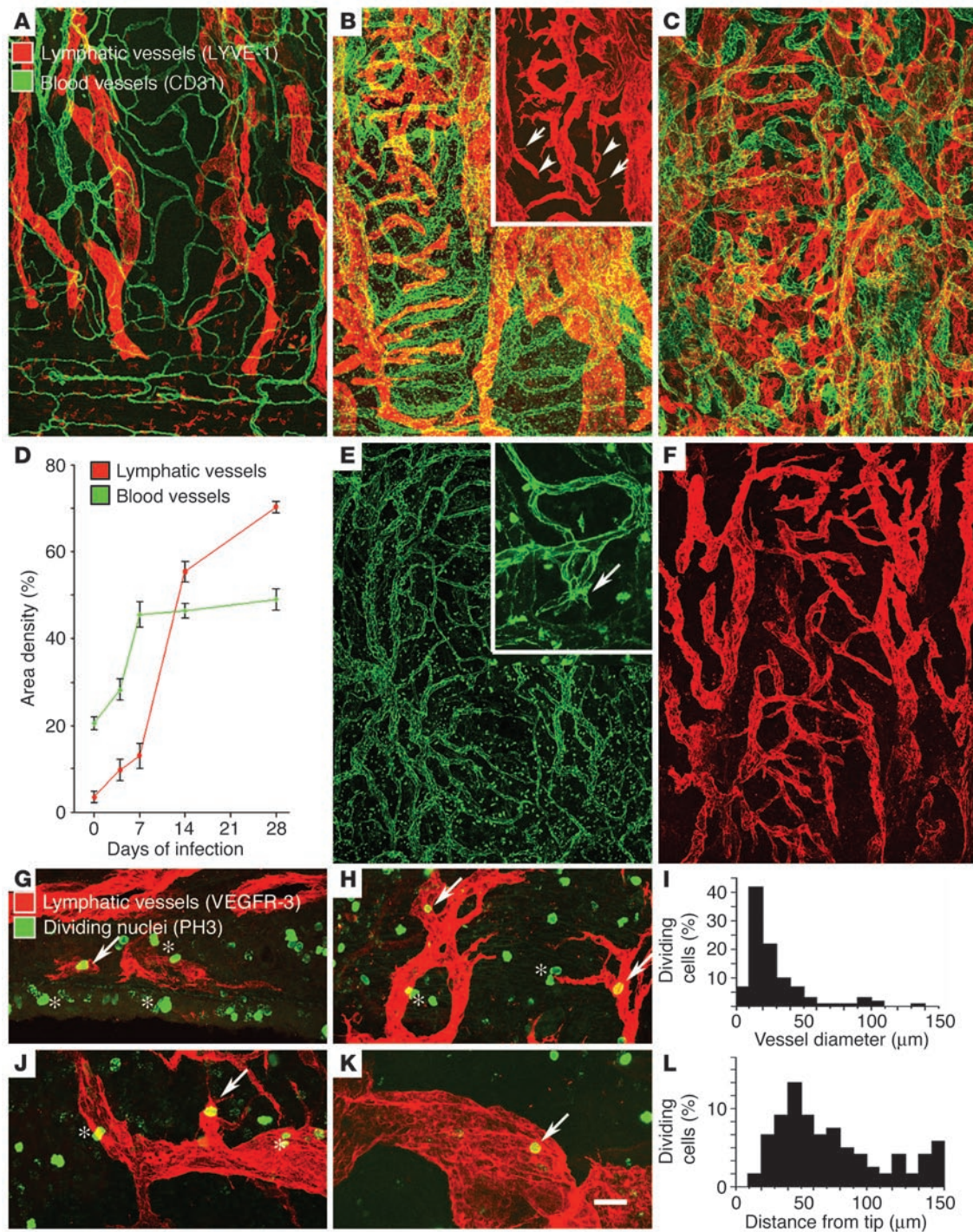
The present study exploited 2 novel and versatile mouse models of lymphangiogenesis to develop a better understanding of airway lymphatic vessels in health and disease. Here, we show that *Mycoplasma pulmonis* infection, which causes inflammatory cell influx, angiogenesis, mucosal edema, epithelial changes, and fibrosis typical of chronic airway inflammation (17), also induces robust lymphangiogenesis. Inhibition of VEGFR-3 signaling entirely blocked sprouting of lymphatic vessels but had no effect on angiogenesis. Similarly, adenoviral transduction of airway epithelium with VEGFR-3 ligands VEGF-C or VEGF-D induced lymphangiogenesis similar to that in *M. pulmonis* infection but no angiogenesis. Conversely, adenoviral delivery

**Nonstandard abbreviations used:** VEGF-DANAC, processed VEGF-D truncated at N and C termini.

**Conflict of interest:** D.J. Hicklin and B. Pytowski are employees of ImClone Systems Inc. and hold stock or stock options in the company. M.G. Achen owns shares in Lymphatix Ltd., which has an interest in the field of research described in this work.

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**Figure 1**

Lymphangiogenesis and angiogenesis after *M. pulmonis* infection. (A–C, E, and F) Confocal micrographs of tracheal whole mounts stained for lymphatic vessels (red) and blood vessels (green). (A) Pathogen-free C3H mouse. (B) C3H mouse infected for 14 days; inset shows lymphatic sprouts (arrowheads) and filopodia (arrows). (C) C3H mouse infected for 28 days. (D) Proliferation of lymphatic vessels (red) and blood vessels (green) in tracheas of C3H mice over 28 days of infection. (E) In a C57BL/6 mouse infected for 14 days, blood vessels (CD31) exhibit sprouting (arrow) and enlargement. (F) Same region as shown in E. Lymphatic vessels (LYVE-1) have a growth pattern similar to that of C3H mice. (G, H, J, and K) Dividing endothelial cells stained for phosphohistone H3 (PH3, green) in tracheal lymphatic vessels (red) of C3H mice infected for 14 days. (G) Section of trachea showing dividing cells, which are sparse in lymphatic vessels (arrow) and numerous in epithelial cells and leukocytes (asterisks). (H, J, and K) Tracheal whole mounts. (H) Dividing lymphatic endothelial cells (arrows) in stalks of medium-sized sprouts. (I) Size distribution of 100 dividing lymphatic endothelial cells. Most dividing cells are near sprout tips (J) or in larger lymphatic vessels (K). (L) Distribution of distances of dividing lymphatic endothelial cells from sprout tips. Scale bar in K applies to all figures: 100  $\mu$ m in A–C, E and F, and 20  $\mu$ m in G, H, J, and K.





of VEGF induced angiogenesis but not lymphangiogenesis. Inflammatory cells and some resident cells were identified as cellular sources of VEGF-C or VEGF-D in infected airways. Studies of the functional consequences of blocking lymphangiogenesis revealed that inhibition of VEGFR-3 signaling in *M. pulmonis*-infected mice exaggerated mucosal edema and reduced lymph node hypertrophy. Comparison of the reversibility of angiogenesis and lymphangiogenesis after antibiotic treatment showed that inflammation and blood vessel remodeling were largely resolved within 2 weeks, but an extensive network of new lymphatic vessels remained in the airway mucosa even after 12 weeks of treatment.

## Results

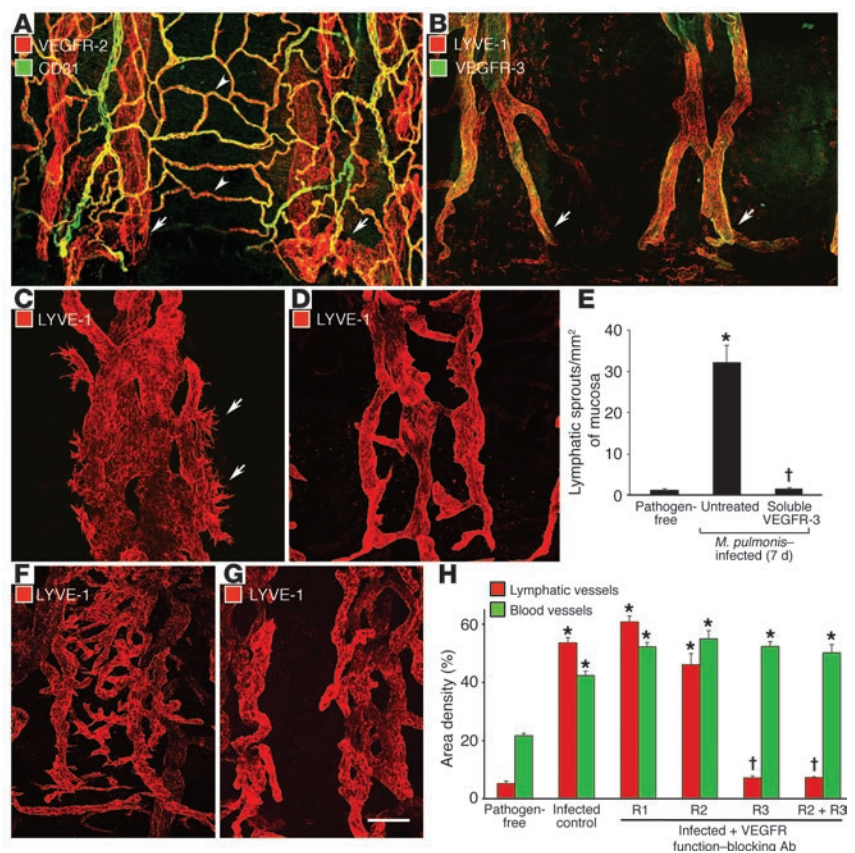
**Blood and lymphatic vasculature of normal airways.** In the airways of pathogen-free mice, blood vessels and lymphatic vessels formed discrete, segmented networks aligned with the framework of cartilage rings (Figure 1A). Most arterioles and venules, identified by CD31 immunoreactivity, were located in the tissue between cartilage rings, whereas most blood capillaries were in regions of mucosa overlying rings. Lymphatic vessels, identified as blind-ended, LYVE-1-immunoreactive vessels approximately 70  $\mu\text{m}$  in diameter on average, were restricted to the mucosa between rings. Lymphatic vessels also showed CD31 immunoreactivity, but this was much weaker than in blood vessels.

**Both blood vessels and lymphatic vessels grow in airway inflammation.** During the first few days after *M. pulmonis* infection, mucosal blood vessels enlarged conspicuously after endothelial cell proliferation (18), but no changes were evident in lymphatic vessels. Blood vessel growth reached a plateau at 14 days (Figure 1B). The earliest signs of lymphatic growth were detected at 7 days, when numerous delicate filopodia and larger tubular sprouts were visible along lymphatic vessels (Figure 1B, inset). Lymphatic sprouts were most abundant on the surface facing cartilage rings and oriented toward the overlying airway epithelium. During the second week, lymphatic sprouts enlarged and fused to form a 3-dimensional network (Figure 1, B and C). Although blood vascular and lymphatic networks were intimately intertwined, no connections between the 2 systems were detected. Measurements confirmed that blood vessel growth was greatest during the first week, but the rate of lymphatic growth peaked in the second week, and thereafter lymphatic vessels were more abundant than blood vessels (Figure 1D).

Growth of airway lymphatic vessels after infection was similar in C3H mice and C57BL/6 mice. However, remodeling of mucosal blood vessels differed in the

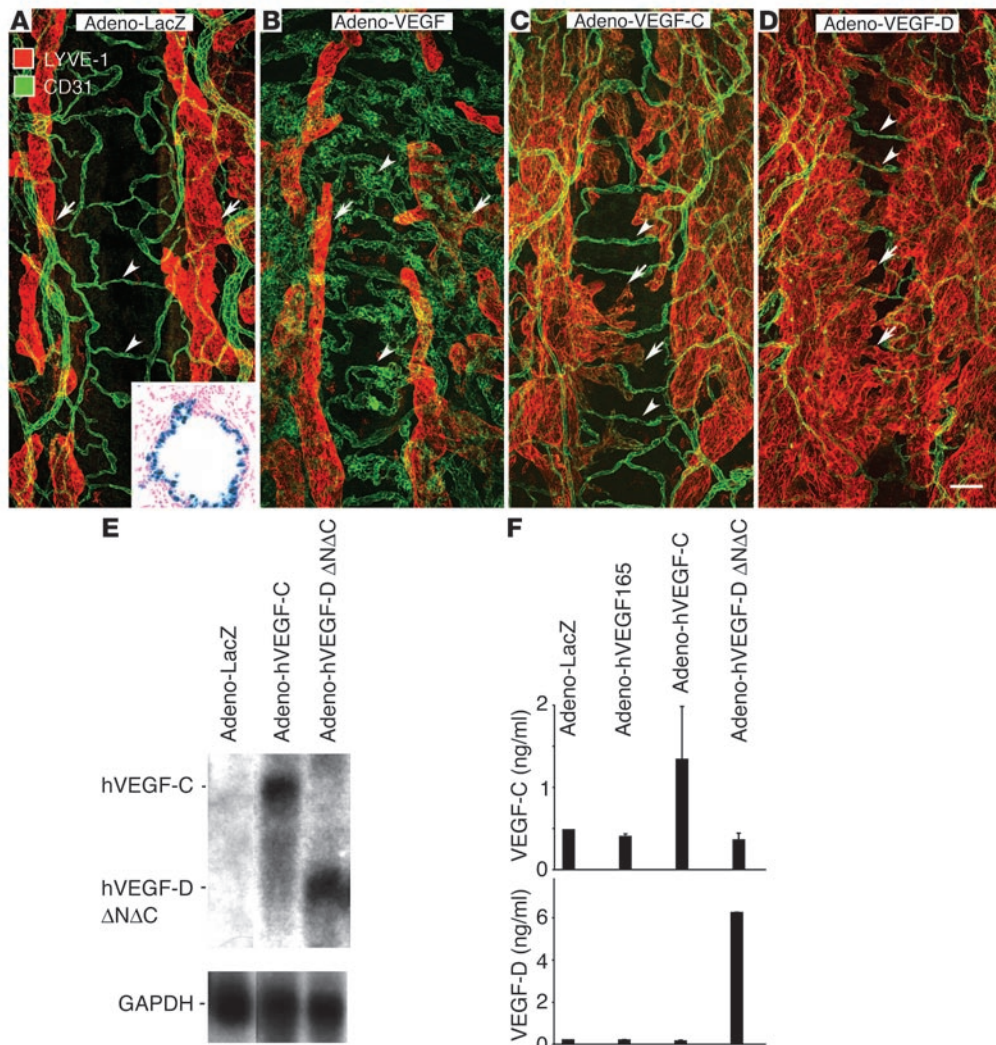
2 strains (19). Vascular enlargement was most conspicuous in C3H mice, but enlargement was accompanied by extensive proliferation in C57BL/6 mice (Figure 1, E and F). In both strains, vascular remodeling resulted in the transformation of capillaries into venules that supported abundant leukocyte adhesion.

Endothelial cells of growing blood vessels divide at the base of sprouts, advancing the filopodia-rich but nondividing tip cells (20). To determine the location of dividing endothelial cells in growing lymphatic vessels, we stained tracheal whole mounts for phosphohistone H3, a marker of M-phase of cell division (20). Most dividing LYVE-1-positive endothelial cells were located in small lymphatic vessels (with a median diameter 22  $\mu\text{m}$ ), not at sprout tips (Figure 1, G-I). The separation between dividing lymphatic endothelial cells and sprout tips was on average 63  $\mu\text{m}$  (Figure 1L). Only 10% of



**Figure 2**

VEGF receptor distribution and inhibition in tracheal vessels. (A) Strong VEGFR-2 immunoreactivity on blood vessels (arrowheads) and weaker staining of lymphatic vessels (arrows) in pathogen-free mouse. (B) Immunoreactivities for VEGFR-3 and LYVE-1 colocalize on lymphatic vessels (arrows) in pathogen-free mouse. (C and D) Lymphatic vessels in mice infected with *M. pulmonis* for 7 days, either untreated (C) or pretreated with soluble VEGFR-3-Ig (D). Sprouts (arrows) emerge from enlarged lymphatic vessels after infection (C) but not if pretreated with soluble VEGFR-3-Ig (D). (E) Number of lymphatic sprouts in tracheas from pathogen-free mice, or from mice after 7 days of infection, with or without soluble VEGFR-3-Ig pretreatment. (F and G) After 14 days of infection, lymphangiogenesis is not affected by concurrent treatment with antibody against VEGFR-1 (F) but is blocked by antibody against VEGFR-3 (G). (H) Area density of lymphatic vessels (red) and blood vessels (green) showing inhibition of infection-induced lymphangiogenesis by anti-VEGFR-3 (R3) antibody but not by anti-VEGFR-1 (R1) or anti-VEGFR-2 (R2). Anti-VEGFR-2 did not augment the effect of anti-VEGFR-3 (R2 + R3). None of the antibodies reduced angiogenesis. \* $P < 0.05$  vs. pathogen-free group; † $P < 0.05$  vs. infected control group. Scale bar in G applies to all figures: 100  $\mu\text{m}$  in A and B and 50  $\mu\text{m}$  in C, D, F, and G.



**Figure 3** Overexpression of VEGF family growth factors by adenoviral vectors. Tracheal lymphatic vessels (red) and blood vessels (green) 10 days after intranasal inoculation of nude mice with adenoviral vectors. (A) Transduction of airway epithelial cells by adenovirus encoding LacZ (Adeno-LacZ), shown by X-gal staining (inset, blue), had no apparent effect on lymphatic vessels (arrows) or blood vessels (arrowheads). (B) Adenoviral VEGF induced angiogenesis (arrowheads) but no detectable lymphangiogenesis (arrows). (C) Adenoviral VEGF-C induced extensive lymphangiogenesis (arrows) but no angiogenesis (arrowheads). (D) Adenoviral VEGF-D induced widespread lymphangiogenesis but no apparent change in blood vessels (arrowheads); lymphatic vessels appear to coalesce (arrows). Scale bar in D (100 μm) applies to all figures. (E) Northern blot showing human VEGF-C (hVEGF-C) and VEGF-D mRNA in extracts of lungs 10 days after inoculation with adenoviral LacZ, human VEGF-C, or human VEGF-D ΔNΔC. (F) Results of ELISA showing VEGF-C and VEGF-D protein in lungs 10 days after inoculation with adenoviral LacZ, human VEGF165, VEGF-C, or VEGF-D ΔNΔC.

dividing endothelial cells were within 20 μm of the tip of lymphatic sprouts (Figure 1J). Another 10% were in large collecting lymphatic vessels (mean diameter 130 μm; Figure 1K).

*VEGFR-3 signaling is required for lymphangiogenesis in inflamed airways.* We used multiple strategies to identify the growth factor signaling systems that drive angiogenesis and lymphangiogenesis in chronically inflamed airways. In the first step, we determined whether receptors for VEGF family members were expressed on airway blood vessels or lymphatic vessels. VEGFR-2 immunoreactivity was strong on blood capillaries, moderate on lymphatic vessels,

and weak on arterioles and venules (Figure 2A). VEGFR-3 immunoreactivity was absent on blood vessels but strong on lymphatic vessels (Figure 2B). These results suggest that VEGFR-2 signaling might drive angiogenesis but either VEGFR-2 or VEGFR-3 could mediate lymphangiogenesis.

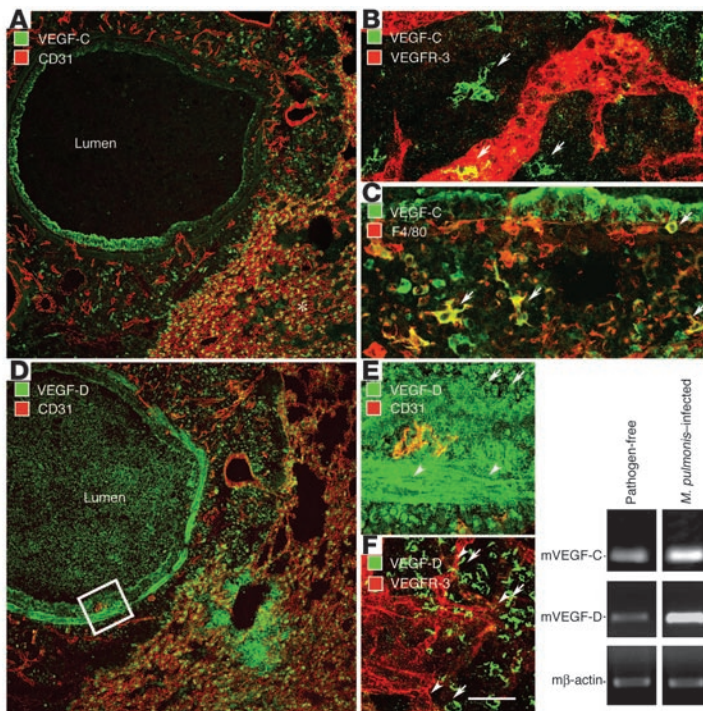
Because VEGFR-3 signaling is essential for lymphangiogenesis in some conditions (4, 21), we sought to determine whether the pathway is important for lymphangiogenesis in *M. pulmonis*-infected mice. Treatment with soluble VEGFR-3-Ig, which blocks VEGFR-3 signaling by binding VEGF-C and VEGF-D, almost completely prevented lymphatic growth during the first week of infection (Figure 2, C and D). Measurements showed a 96% reduction in the number of lymphatic sprouts (Figure 2E). No impairment in blood vessel remodeling or growth was evident after treatment with soluble VEGFR-3-Ig.

In another strategy of blocking VEGF receptor signaling, mice were treated with function-blocking monoclonal antibodies against VEGFR-1, VEGFR-2, or VEGFR-3 at doses reported to be effective in other mouse models (22–24). Treatment with antibodies against VEGFR-1 (Figure 2F) or VEGFR-2 had no effect on growth of lymphatic vessels after *M. pulmonis* infection, but the anti-VEGFR-3 antibody almost totally blocked lymphangiogenesis during 2 weeks of infection (Figure 2G).

Measurements showed an 87% reduction in lymphatic growth after VEGFR-3 antibody treatment but no significant change in lymphatic vessels after VEGFR-1 or VEGFR-2 antibody treatment (Figure 1H). A combination of antibodies against VEGFR-2 and VEGFR-3 had an effect no greater than anti-VEGFR-3 alone (Figure 1H). None of these antibodies reduced growth or enlargement of airway blood vessels after infection (Figure 2H).

*VEGF-C and VEGF-D drive lymphangiogenesis in airways.* To address the role of VEGFR-3 ligands in lymphangiogenesis in the respiratory tract, we used an adenoviral delivery system to generate VEGF-C or



**Figure 4**

VEGF-C and VEGF-D in *M. pulmonis*-infected airways. Immunohistochemical staining of VEGF-C (A–C) and VEGF-D (D–F) in mouse airways and lung 14 days after infection. (A) VEGF-C (green) in epithelium, peribronchial inflammatory cells, and type II alveolar epithelial cells (asterisks) of lung. Blood vessels stained for CD31 (red). (B) VEGF-C (green) in inflammatory cells (arrows) near sprouting lymphatic vessels (red) in tracheal whole mount. (C) Colocalization of VEGF-C immunoreactivity (green) and F4/80 immunoreactivity (red) in macrophages (yellow, arrows). (D) Strong VEGF-D staining (green) of neutrophils in airway lumen. (E) VEGF-D-positive neutrophils (boxed area in D) shown at higher magnification in airway lumen (arrows, green) and airway smooth muscle cells (arrowheads). Blood vessels stained for CD31 (red). (F) VEGF-D immunoreactivity in neutrophils (arrows, green) in tracheal mucosa near sprouting lymphatic vessels (arrowheads) stained for VEGFR-3 (red). Scale bar in F applies to all figures: 200  $\mu\text{m}$  in A and D and 50  $\mu\text{m}$  in B, C, E, and F. (G) Results of RT-PCR analysis showing higher expression of mouse mRNA for VEGF-C (mVEGF-C) and VEGF-D in tracheas after 14 days of infection.

VEGF-D in airway epithelial cells. VEGF was generated in the same way for comparison. All 3 of these growth factors have been reported to induce angiogenesis and/or lymphangiogenesis (6, 8–10).

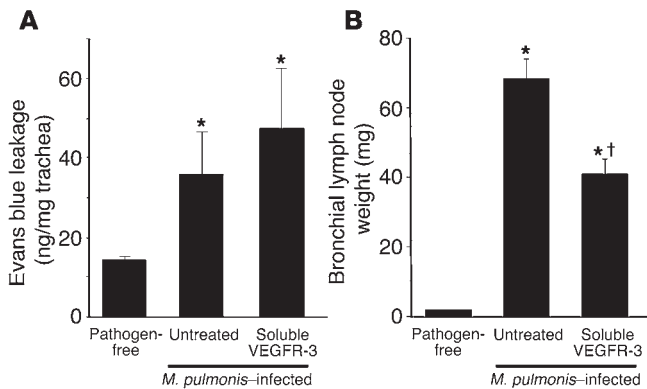
Mice were inoculated intranasally with LacZ-expressing adenovirus to determine whether transduction of airway epithelial cells by itself had effects on blood vessels or lymphatic vessels. Airway epithelial cells of these mice had strong  $\beta$ -gal expression (Figure 3A, inset) but no change in blood or lymphatic vasculature (Figure 3A). In contrast, inoculation with VEGF adenovirus induced prolific sprouting and growth of blood vessels but no lymphangiogenesis (Figure 3B). Conversely, inoculation with VEGF-C adenovirus induced growth of lymphatic filopodia and sprouts similar to those seen in *M. pulmonis*-infected mice but did not cause detectable changes in blood vessels (Figure 3C). Adenoviral processed VEGF-D truncated at N and C termini (VEGF-D  $\Delta\text{N}\Delta\text{C}$ ), the mature form of VEGF-D (25), induced the formation of giant sheet-like lymphatic vessels in the trachea (Figure 3D) and twice as many lymphatic sprouts as did VEGF-C ( $30 \pm 4$  vs.  $15 \pm 2$  sprouts/ $\text{mm}^2$  of mucosa;  $P < 0.05$ ). However, VEGF-D  $\Delta\text{N}\Delta\text{C}$ , which stimulates angiogenesis in skeletal muscle (6), had no apparent effect on airway blood vessels (Figure 3D).

Inoculation of mice with VEGF-C or VEGF-D adenovirus conspicuously increased the corresponding mRNA in Northern blots of lung (Figure 3E). VEGF-C protein was increased approximately 3-fold, and VEGF-D was increased 30-fold above baseline amounts, as determined by ELISA (Figure 3F).

**Cellular sources of VEGF-C and VEGF-D in inflamed airways.** Inflammatory cells were rare in the airways of pathogen-free mice but were abundant after *M. pulmonis* infection. Neutrophils were not detected in the airway mucosa of pathogen-free C3H mice but were abundant after infection, reaching a peak at 7 days ( $1,581 \pm 332$  neutrophils/ $\text{mm}^2$ ). Neutrophils were also abundant in the airway lumen. The massive influx of neutrophils was followed by a smaller, more gradual influx of macrophages and lymphocytes into the mucosa but not the lumen.

In the respiratory tract of pathogen-free mice, most VEGF-C immunoreactivity was in airway epithelial cells and surfactant-producing type II alveolar epithelial cells of the lung (data not shown). However, after infection, VEGF-C immunoreactivity was strong in mucosal inflammatory cells as well as in airway epithelial cells (Figure 4, A and B). In addition, some F4/80-positive macrophages had VEGF-C immunoreactivity (Figure 4C). VEGF-D immunoreactivity was minimal in the airways of pathogen-free mice but, after *M. pulmonis* infection, was strong in neutrophils in the mucosa and lumen (Figure 4, D–F). VEGF-D staining was also strong in macrophages, airway smooth muscle cells, and focal regions of some lymphatic vessels. In addition, staining for both VEGF-C and VEGF-D was diffuse and weak in basement membrane (Figure 4E). No staining was present when the primary antibody was omitted or replaced by preimmune serum (data not shown). Consistent with these findings, mRNAs for VEGF-C and VEGF-D increased in the airways after *M. pulmonis* infection, according to semiquantitative RT-PCR (Figure 4G).

**Blocking lymphangiogenesis impairs lymphatic function and promotes mucosal edema.** The amount of plasma leakage in the airway mucosa increases after *M. pulmonis* infection because the new and remodeled blood vessels are leaky and have increased surface area (17). The blood vessels are also enlarged and unusually sensitive to inflammatory mediators (17). Expansion of the lymphatic network after infection should accommodate the increased flux of tissue fluid. We therefore asked whether failure of lymphangiogenesis after infection impaired fluid drainage and immune cell transport to local lymph nodes and promoted bronchial lymphedema. Using Evans blue dye accumulation as a measure of leakage, we found that the amount of extravasated dye increased significantly after infection and increased further when lymphangiogenesis was inhibited more than 90% by systemic treatment with soluble VEGFR-3-Ig delivered by adenovirus prior to infection (Figure 5A).

**Figure 5**

Consequences of blocking lymphangiogenesis in inflamed airways. Comparison of 3 groups of C3H mice: pathogen-free, infected with *M. pulmonis* for 14 days, and infected with lymphangiogenesis blocked by concurrent treatment with soluble VEGFR-3-Ig delivered by adenovirus. (A) Evans blue accumulation in trachea over 30 minutes showing significantly increased dye accumulation after infection ( $*P < 0.05$ ) but even greater accumulation when lymphangiogenesis was blocked by soluble VEGFR-3-Ig. (B) Lymph nodes were enlarged after infection ( $*P < 0.05$ ), but significantly less so when lymphangiogenesis was inhibited by soluble VEGFR-3-Ig ( $†P < 0.05$ ).

Bronchial lymph nodes that receive drainage from the airways and lungs increase in weight with time after *M. pulmonis* infection (18), a finding consistent with the proliferation of lymphocytes triggered by activated antigen-presenting cells and superantigens or mitogens from the organisms (26). Soluble VEGFR-3-Ig treatment prior to infection diminished this response, as reflected by an approximately 40% reduction in bronchial lymph node enlargement from the usual 20- to 30-fold increase after 2 weeks of infection (Figure 5B).

The weights of bronchial lymph nodes and lungs turned out to be useful indicators of disease severity. Lung weight increased 2- or 3-fold during the infection, as leukocytes and fluid accumulated, but was significantly lower after 2 weeks of antibiotic treatment and essentially normal by 6 weeks of treatment (Figure 6A). Bronchial lymph nodes decreased in size more slowly during treatment. There was no change after 2 weeks of treatment, but after 12 weeks, bronchial lymph nodes lost 88% of the enlargement present at the onset of treatment (Figure 6A).

*Persistence of hyperplastic lymphatic vessels during resolution of inflammation.* Treatment of *M. pulmonis* infection with antibiotics or steroids rapidly and fully reverses the remodeling and proliferation of blood vessels in the airway mucosa (13, 14). Consistent with those reports, the present experiments confirmed that remodeled blood vessels returned to their normal state within 4 weeks of antibiotic treatment (Figure 6, B–D). Lymphatic sprouting stopped during treatment, but the extensive network of new lymphatic vessels did not show a corresponding decrease even after 12 weeks of treatment.

Some normalization of lymphatic vessels was evident after 2 and 4 weeks of treatment with antibiotic. Amounts of lymphatic filopodia were reduced by 70% and 95% at 2 and 4 weeks of treatment with antibiotic, respectively, compared with the number present at the onset of treatment at 2 weeks of infection (Figure 6B). With this change, lymphatic vessels acquired smoother contours. In addition, some lymphatic vessels were thinner or had constrictions along their length. Seemingly isolated segments of LYVE-1-

positive lymphatic vessel were also present. The overall intensity of LYVE-1 immunoreactivity of lymphatic sprouts was reduced after treatment but varied in individual lymphatic endothelial cells from intense to almost absent (Figure 6, C and D). Immunoreactivity to LYVE-1 was weaker than immunoreactivity to CD31 in some lymphatic vessels (Figure 6D).

Compared to the large reduction in sprouts, measurements of the overall extent of the lymphatic network changed relatively little over 12 weeks of treatment (Figure 6B). The pattern suggested a small decrease during the first 6 weeks of treatment followed by a plateau with little change thereafter, but the changes were not statistically significant (Figure 6B).

## Discussion

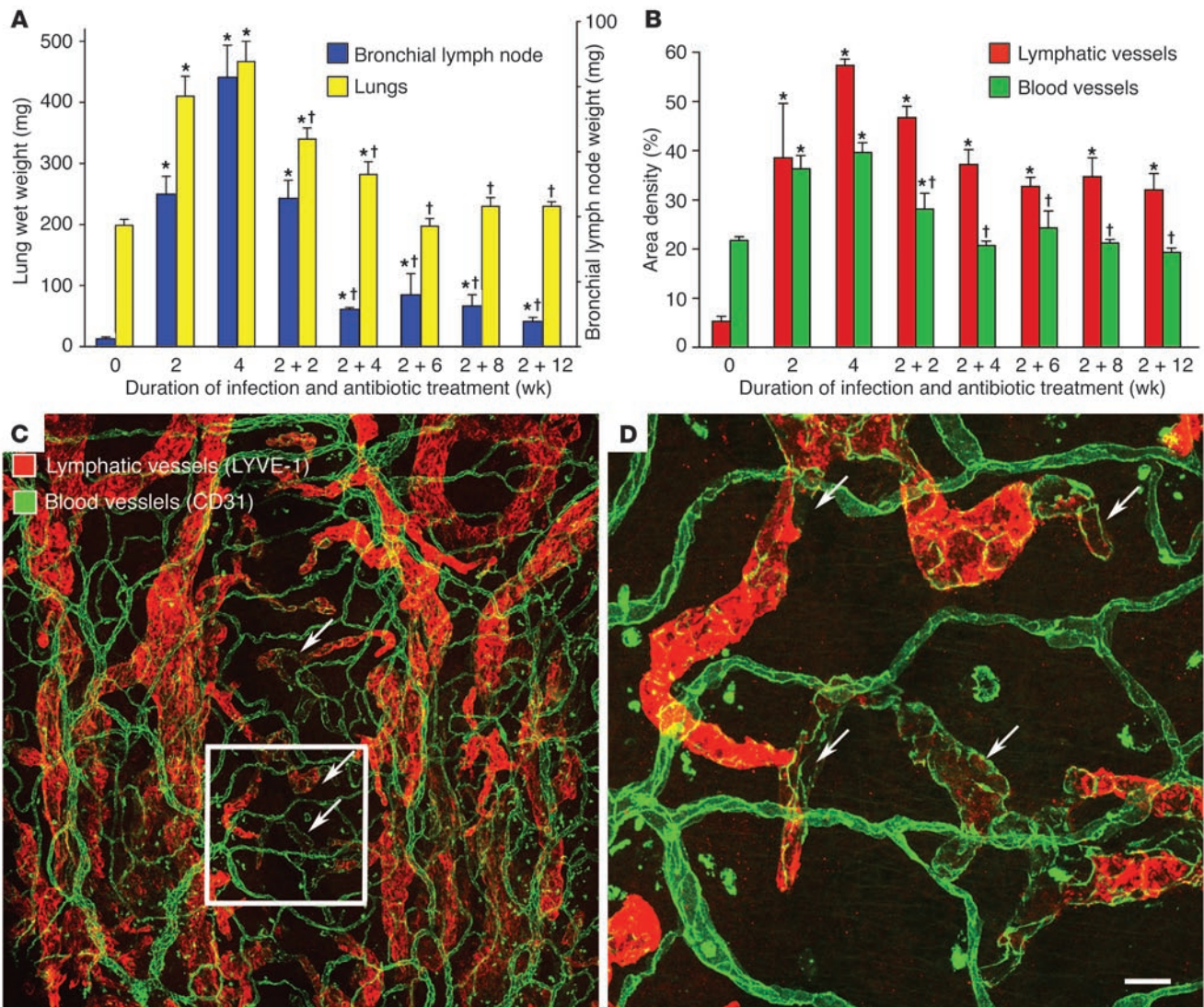
The present study revealed that airway inflammation is accompanied not only by angiogenesis but also by extensive growth of lymphatic vessels. In mice with *M. pulmonis* infection, lymphangiogenesis appears to be driven by VEGF-C and VEGF-D from inflammatory cells that migrate into the airways. Adenoviral delivery of either of these growth factors induced lymphangiogenesis but not angiogenesis in the airways. Even the mature form of VEGF-D, which in humans binds to VEGFR-2 as well as VEGFR-3 (6, 25), induced abundant lymphangiogenesis without apparent angiogenic activity. After infection, expression of VEGF-C and VEGF-D was greatest in airway leukocytes. Inhibition of growth factor signaling through VEGFR-3 entirely blocked lymphatic sprouting after infection but had no apparent effect on blood vessel remodeling. Inhibition of lymphangiogenesis predisposed to bronchial lymphedema and reduced the infection-induced hypertrophy of draining lymph nodes. Newly formed lymphatic vessels — unlike the remodeled blood vessels — did not regress after the infection was treated.

*New tools for studying lymphangiogenesis.* With the advent of new markers for and reagents that can be used to manipulate lymphatic vessels, the findings of many classical studies are beginning to be understood on a molecular level (21, 27, 28). Earlier studies showed that lymphatic vessels proliferate in wounds and inflammation of skin (3, 16, 29) and cornea (30). Some experiments also suggested that lymphangiogenesis occurs more slowly than angiogenesis (15, 16, 31). More recent studies show that lymphangiogenesis can be stimulated or inhibited by agonists or antagonists of VEGFR-3 signaling independent of angiogenesis (32, 33). Lymphatic vessels proliferate in inflammatory bowel disease, renal fibrosis, and transplant rejection (34, 35), but few models have been available for mechanistic studies of lymphangiogenesis under pathological conditions.

The present study revealed attributes of *M. pulmonis* infection of the respiratory tract in mice as a model for studying lymphangiogenesis in chronic inflammation. These features complement the model's attributes for studying vascular remodeling and angiogenesis (18, 19). The simple, orderly arrangement of blood vessels and lymphatic vessels in mouse airways coupled with the experimental conditions for examining the step-by-step progression toward chronic disease make it possible to dissect the cellular and molecular mechanisms of angiogenesis and lymphangiogenesis and their reversibility. Because the anatomical nature of the mouse trachea lends itself to the preparation of tissue whole mounts, blood vessels, lymphatic vessels, and inflammatory cells can be viewed simultaneously in three dimensions by fluorescence and confocal microscopy.

In lymphatic vessels that grow after *M. pulmonis* infection, sites of endothelial cell proliferation and vessel enlargement fit with a recent model of blood vessel growth (20). Delicate filopodia on





**Figure 6** Persistence of new lymphatic vessels after treatment. (A and B) Weight of lungs and bronchial lymph nodes (A) and abundance of tracheal lymphatic vessels and blood vessels (B) in pathogen-free mice (0), mice infected with *M. pulmonis* for 2 or 4 weeks (2 and 4), and mice infected for 2 weeks and then treated with oxytetracycline for 2 to 12 weeks (2 + 2 through 2 + 12). The weight of both organs increased after infection and decreased toward normal after treatment. Airway blood vessels (green) showed a similar pattern, whereas lymphatic vessels (red) proliferated after infection but regressed little, even after 12 weeks of treatment. \* $P < 0.05$  vs. pathogen-free group; † $P < 0.05$  vs. 2-week-infected group without treatment. (C and D) Confocal micrographs showing tracheal lymphatic vessels (red) and blood vessels (green) after infection for 2 weeks and then oxytetracycline treatment for 8 weeks. Blood vessels regressed almost to the pathogen-free state. Lymphatic vessels showed some changes but regressed only slightly; lymphatic vessels with constrictions and no LYVE-1 immunoreactivity in some cells are indicated by arrows. (D) Enlargement of the boxed area in C. Scale bar in D applies to both figures: 100  $\mu\text{m}$  in C, 25  $\mu\text{m}$  in D.

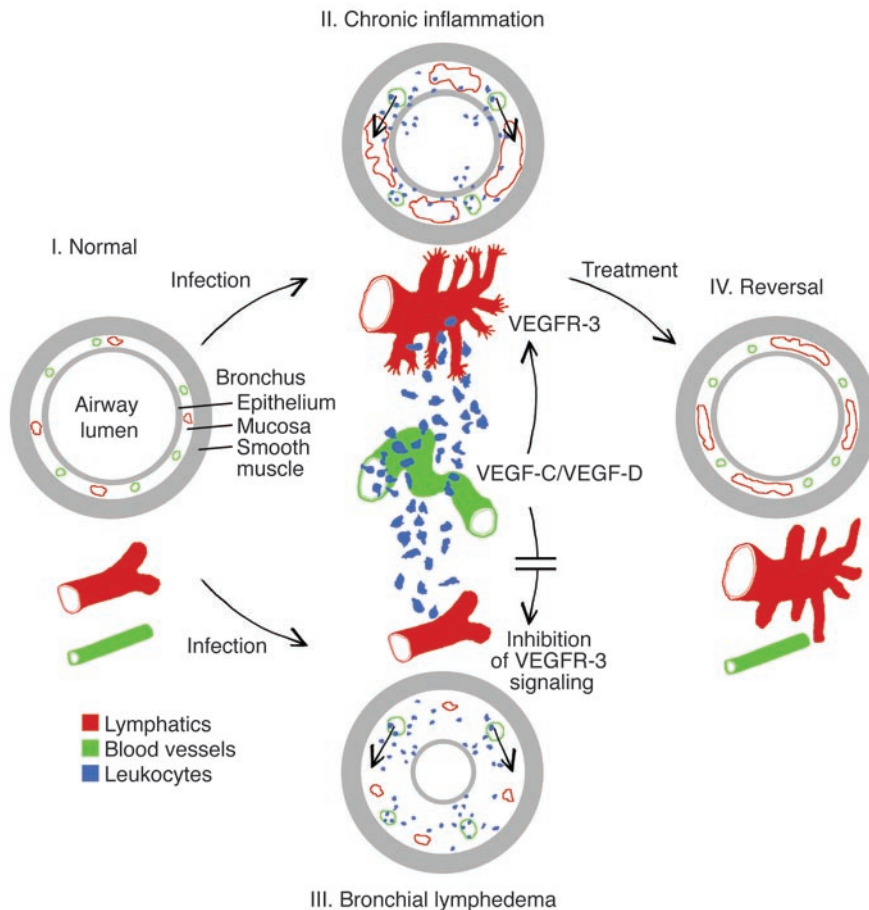
lymphatic endothelial cells at the tips of sprouts marked sites of vessel growth, but cell division was most common in the stalk, one or two cells behind the tip. Enlargement of large collecting lymphatic vessels apparently resulted from proliferation of endothelial cells combined with distention by lymph.

**Role of VEGFR-3 signaling.** In our experiments, we used 2 approaches to block VEGFR-3 signaling, with the goal of determining whether lymphangiogenesis in the *M. pulmonis* model of chronic inflammation is driven by VEGFR-3 signaling, as it is in some other systems (4, 21). Both approaches — using soluble VEGFR-3-Ig and using function-blocking antibody — almost completely blocked lymphatic

growth. By comparison, inhibition of VEGFR-1 or VEGFR-2 had no apparent effect on lymphatic growth, and inhibition of VEGFR-2 did not augment the action of VEGFR-3 blockade.

Next we found that either VEGF-C or VEGF-D triggered intense lymphangiogenesis in the airways when delivered by adenoviral vector. VEGF stimulated angiogenesis but not lymphangiogenesis, despite some expression of VEGFR-2 on endothelial cells of lymphatic vessels. Together, these findings are consistent with the identification of VEGFR-3 signaling as the principal driving force for lymphatic growth in this model. VEGF and VEGFR-2 signaling play little or no apparent role.





**Figure 7**

Role of lymphangiogenesis in airway inflammation. The diagram shows relationships of changes in lymphatic vessels (red), blood vessels (green), inflammatory cells (blue), and plasma leakage (short arrows) in inflamed airway mucosa. Cross-sections of bronchi and vessel schematics compare 4 conditions. (I) In normal airways, blood vessels have little or no leukocyte traffic; baseline leakage drains via lymphatic vessels. (II) After infection, activated antigen-presenting cells traffic from airways to local lymph nodes, evoking an immune response. Mucosal capillaries remodel into activated venules that mediate leukocyte influx. These cells release VEGF-C and VEGF-D, which drive lymphangiogenesis via VEGFR-3 signaling in lymphatic endothelial cells. The lymphatic network expands by sprouting from existing lymphatic vessels to accommodate the increased leakage from venules and increased trafficking of immune cells to lymph nodes. (III) If lymphangiogenesis does not occur, leakage exceeds drainage, bronchial lymphedema develops, trafficking of immune cells to lymph nodes decreases, and the immune response is reduced. (IV) Treatment decreases the inflammatory stimulus, allowing blood vessels to return to the baseline state. Leukocyte influx decreases, and the stimulus for lymphangiogenesis diminishes, but lymphatic vessels that formed during the infection persist — ready for the next infection.

The signaling system that drives angiogenesis after *M. pulmonis* infection is unknown. Antibodies that block the function of VEGFR-1 or VEGFR-2 did not significantly reduce blood vessel remodeling, which indicates that VEGF, VEGF-B, and placental growth factor (PlGF) are not responsible for changes in these vessels. Similarly, neither VEGF-C nor VEGF-D stimulated angiogenesis in the model, yet both can stimulate angiogenesis under certain conditions (5, 6, 8, 9). Local concentration and not overall dose of growth factors determine the exact pattern of vessel remodeling (36). Even the mature form of VEGF-D, which can induce tyrosine phosphorylation of VEGFR-2 (6, 9), produced only lymphangiogenesis in our model.

**Sources of VEGF-C and VEGF-D in inflamed airways.** Cells in the airways of infected mice had significantly greater expression of VEGF-C and VEGF-D, as reflected by mRNA analysis and immunohistochemistry. Localization studies revealed that VEGF-C was expressed by mononuclear inflammatory cells that were not present in pathogen-free mice. Some of these cells, likely macrophages or dendritic cells, were close to growing lymphatic vessels. Mononuclear cells are a known source of VEGF-C (33, 35, 37). VEGF-C was also expressed in airway epithelial cells, both with and without infection. VEGF-D was expressed in neutrophils in the mucosa and lumen of inflamed airways and in other inflammatory cells. Little VEGF-D expression was detected in airways or lung of pathogen-free mice, consistent with previous findings (25).

Dendritic cells, as antigen-presenting cells activated by bacterial antigens, migrate via lymphatic vessels to local lymph nodes within 24 hours of infection to initiate an immune response, with the

proliferation of lymphocytes causing hypertrophy of the lymph nodes (38). Activated dendritic cells, which express VEGF-C (37), may help to drive the development of the lymphatic network they use as a route to lymph nodes. In inflammation, dendritic cells are believed to migrate toward lymphatic vessels along a chemotactic gradient of mediators, including chemokines CCL19 and CCL21 from lymphatic endothelial cells (39, 40). Dendritic cells express CCR7 receptors for these ligands (39, 40). Thus, dendritic cells and lymphatic vessels express ligands and receptors for one another that may act in a reciprocal fashion.

The full range of factors that activate dendritic cells is unclear, but IL-1 $\beta$  and TNF- $\alpha$  are both present after *M. pulmonis* infection (41) and can increase VEGF-C expression (42). Although many of the details are still missing, it appears that key cells, mediators, and feedback loops necessary to induce lymphangiogenesis are present in the airways after *M. pulmonis* infection (Figure 7).

**Functional consequences of impaired lymphangiogenesis in airways.** Results of the present study suggest that inhibition of VEGFR-3 signaling reduces the transport of fluid and antigen-presenting cells from airways to lymph nodes via lymphatic vessels, thereby exaggerating mucosal edema and dampening the afferent limb of the immune response. Consistent with these consequences of impaired lymphangiogenesis is the finding that blocking lymphatic transport from inflamed rabbit ears may exacerbate local edema and eventually lead to necrosis (43). Similarly, inhibition of VEGFR-3 signaling suppresses dendritic cell trafficking from the cornea to draining lymph nodes and corneal transplant rejection (44).



Additional experiments will be needed to develop a more complete understanding of the net effect of these actions on the severity of inflammatory responses.

Angiogenesis and mucosal swelling are part of the pathophysiology of chronic inflammatory airway diseases such as asthma and chronic bronchitis (2, 45). Like other causes of sustained airway inflammation, mycoplasmal infection in rodents is accompanied by inflammatory cell influx, angiogenesis, mucosal edema, epithelial changes, and fibrosis (17). *Mycoplasma pneumoniae* infection also results in bronchial hyperreactivity but not necessarily all the features of asthma (46, 47).

To our knowledge, the involvement of lymphatic abnormalities in bronchial hyperreactivity or airflow obstruction has not been examined. Impaired drainage of edema fluid would be expected to lead to airway wall thickening, reduced wall compliance, and narrowing of the airway lumen (Figure 7). If lymphangiogenesis is impaired in inflamed airways, restoration of lymphangiogenesis could complement conventional therapies for inflammatory airway disease.

**Persistence of lymphatic vessels after treatment.** Although additional growth of lymphatic vessels induced by *M. pulmonis* infection was readily prevented by treatment with antibiotic, once formed, the new lymphatic vessels persisted after the inflammation resolved. The relative stability of lymphatic vessels under these conditions clearly differs from the corresponding changes in blood vessels, which rapidly regressed after antibiotic treatment. Because lymphatic vessels lack pericytes and basement membrane (16, 30), their resilience is at odds with current dogma that such features protect endothelial cells from regression (48). Even though most lymphatic vessels did not regress, some had markedly reduced LYVE-1 expression. Downregulation of LYVE-1 is also observed in lymphatic vessels of some tumors and acute skin inflammation (L. Johnson and D.G. Jackson, unpublished data), which suggests that expression is responsive to tissue microenvironment.

In conclusion, lymphangiogenesis begins slowly in airway inflammation but, after a few weeks, overtakes the remodeling and proliferation of blood vessels and persists after the inflammation resolves. Lymphangiogenesis in inflamed airways is mediated by VEGF-C and VEGF-D derived mainly from inflammatory cells via VEGFR-3 signaling in lymphatic endothelial cells. The expanded lymphatic network provides conduits for extravasated fluid from leaky blood vessels and activated antigen-presenting cells to local lymph nodes. Impaired lymphangiogenesis in airway inflammation may interfere with fluid clearance and lead to bronchial lymphedema. A better understanding of the factors that stimulate or inhibit lymphatic growth or accelerate the regression of new lymphatic vessels may offer new targets for intervention in the treatment of asthma and other inflammatory airway diseases.

## Methods

**Animals.** Specific pathogen-free C3H or C57BL/6 mice (Charles River Laboratories Inc.) at 8 weeks of age were used for *M. pulmonis* infection experiments, and immunodeficient NMRI nu/nu mice (Harlan) were used to study adenoviral transduction of airway epithelial cells in vivo. Mice were housed under barrier conditions, and sentinels were routinely tested serologically for common pathogens. Before being subject to experimental procedures, mice were anesthetized with ketamine (100 mg/kg) and xylazine (10 mg/kg) injected intramuscularly. The Institutional Animal Care and Use Committees of the UCSF and the University of Helsinki approved all procedures.

***M. pulmonis* infection and antibiotic treatment.** C3H or C57BL/6 mice were inoculated intranasally on day 0 with 50  $\mu$ l of broth containing  $10^5$  or  $10^6$  CFU, respectively, of *M. pulmonis* organisms (strain CT7; University of Alabama at Birmingham) (19). In reversal experiments, infected C3H mice were injected subcutaneously once daily with oxytetracycline (20 mg/kg; Phoenix Pharmaceuticals Inc.) starting on day 14 of infection and continuing for 2–12 weeks (13). At the end of the experiment, mice were anesthetized and perfused with fixative; trachea, lungs, and bronchial lymph nodes were removed.

**Inhibition of VEGF receptor signaling.** Soluble VEGFR-3 was used to block VEGFR-3 signaling. This decoy receptor was produced systemically by adenoviral delivery of soluble chimeric VEGFR-3 coupled to a mouse Ig fragment (49). Adenovirus ( $1 \times 10^9$  PFU in 100  $\mu$ l) was injected i.v. into C3H mice on the day of infection. Lymphangiogenesis and angiogenesis were assessed 7 or 14 days later. To prepare the vector, we produced cDNA encoding the first 3 Ig loops of mouse VEGFR-3 (GenBank accession number NM008029.1, nucleotides 117–1053) by RT-PCR using the primers 5'-GCGCAAGCTTGTACTCCATGACCCCTCCAAC-3' and 5'-GAGCGATATCCAGGACAGGTCCTTTGAGCCA-3' and total mouse liver RNA as a template. The PCR product and mouse Fc fragment from the plasmid mFc/plasmid Bluescript were cloned into pSecTag expression vector (Invitrogen Corp.), and a fragment containing both mVEGFR-3 and Fc domains was further subcloned into pAdCMV vector for production of recombinant adenoviruses (50).

Alternatively, signaling by VEGFR-1, VEGFR-2, or VEGFR-3 was inhibited using selective function-blocking rat monoclonal antibodies from clones MF1, DC101, and 31C1, respectively, at doses known to be effective (ImClone Systems Inc.; refs. 22–24). Rat IgG was used as a control. Antibodies were injected i.p. in C3H mice at an initial dose of 2 mg/mouse on the day of infection followed every other day with doses of 0.8 mg/mouse for 14 days.

**Adenoviral transduction of airway epithelium with growth factors.** Airway epithelial cells were transduced with genes encoding VEGF, VEGF-C, VEGF-D, or LacZ by intranasal inoculation of nude mice with adenovirus at 5 to approximately  $10 \times 10^8$  PFU in 50  $\mu$ l. Adenovirus expressing full-length cDNA for human VEGF165, VEGF-C, or VEGF-D  $\Delta$ NAC (6), or a nuclear-targeted LacZ gene as an inactive control, was prepared as previously described (8, 51). Each adenovirus was analyzed to ensure absence of replication-competent virus, lipopolysaccharide, mycoplasma, or other contaminants. Lymphangiogenesis and angiogenesis were assessed 10 days after inoculation. Transduction of epithelial cells was verified by X-gal staining (7) or  $\beta$ -gal immunoreactivity after inoculation with the LacZ adenovirus. Expression of VEGF, VEGF-C, and VEGF-D was verified by Northern blotting of RNA extracted from lungs and tracheas.

**Immunohistochemistry.** The vasculature of anesthetized mice was perfused for 2 minutes with 1% paraformaldehyde in PBS, pH 7.4 (52). Trachea, lung, and liver were removed and immersed in the fixative for 1 hour. Tracheas were stained as whole mounts and lung and liver as 20- $\mu$ m cryostat sections with 1 or more primary antibodies (52): (a) lymphatic vessels: rabbit polyclonal antibody against LYVE-1, 1:1000 (D.G. Jackson laboratory) (53); (b) blood vessels: monoclonal antibodies against mouse CD31, hamster clone 2H8, 1:500, Chemicon International Inc. or rat clone MEC 13.3, 1:500 (BD Biosciences – Pharmingen); (c) VEGF-C: rabbit polyclonal antibody no. 6 to full-length human VEGF-C, 1:500; (d) VEGF-D: rabbit polyclonal antibody no. 490 to mouse VEGF-D, 1:500 (Herbert Weich, ReliaTech, Braunschweig, Germany); (e) VEGFR-2: rabbit polyclonal antibody, 1:2000 (Rolf Brekken, University of Texas Southwestern, Dallas, Texas, USA); (f) VEGFR-3: goat polyclonal antibody, 1:500 (R&D Systems); (g) leukocytes: rat monoclonal antibody against CD45, 1:250 (BD Biosciences – Pharmingen); (h) macrophages: rat monoclonal antibody against F4/80, 1:500 (BD Biosciences – Pharmingen); (i) dividing cells: rabbit polyclonal antibody against phosphohistone H3, 1:500 (Upstate





Biotechnology Inc.); (j) LacZ: rabbit polyclonal antibody against  $\beta$ -gal, 1:500 (Chemicon International Inc.). Species-specific secondary antibodies were labeled with Cy3, Cy5, or FITC (1:400; Jackson ImmunoResearch Laboratories Inc.). For control experiments, the primary antibody was omitted or the preimmune serum substituted. Specimens were examined with a Zeiss Axiophot microscope or Zeiss LSM-510 confocal microscope (52).

**Dividing lymphatic endothelial cells.** Dividing lymphatic endothelial cells were identified by colocalization of staining for phosphohistone H3 and VEGFR-3. Tracheal whole mounts were examined by fine focusing with an oil-immersion lens ( $\times 40$ ; numerical aperture, 1.0; final screen magnification  $\times 1,200$ ) or in optical sections (0.7  $\mu$ m thick) obtained by confocal microscopy. The diameter of lymphatic vessels and distance of 20 labeled nuclei to the tip of lymphatic sprouts were measured in tracheas from 5 mice per group.

**RT-PCR analysis.** Tracheas and lungs of pathogen-free mice and mice infected with *M. pulmonis* for 14 days were removed and immersed in RNAlater (QIAGEN Inc.). Total RNA was extracted from tissues using the RNeasy kit (QIAGEN Inc.). Genomic DNA was eradicated by treatment with DNase (QIAGEN Inc.), and cDNA was generated using Omniscript reverse transcriptase (QIAGEN Inc.), oligo-dT primers, and 150 ng of total RNA. Equal amounts of cDNA were subjected to PCR with primers specific for mouse *Vegfc* (7) or *Vegfd* (GenBank accession number NM010216, base pairs 903–1348) cDNA, and  $\beta$ -actin was used as a loading control.

**Northern blots.** RNA extracted as above was electrophoresed, transferred to a nylon membrane, and hybridized to radioactive probes for human *VEGF-C* (GenBank accession number NM005429.2, base pairs 80–2076) and *VEGF-D*  $\Delta$ NAC (GenBank accession number NM004469.2, base pairs 509–1572) (5). A GAPDH probe served as a loading control.

**ELISA.** VEGF-C and VEGF-D in the lungs of nude mice treated intranasally with adenoviruses encoding LacZ, VEGF165, VEGF-C, or VEGF-D  $\Delta$ NAC 10 days previously were measured by ELISA (human VEGF-C ELISA [no. BMS297; Bender MedSystems], human VEGF-D ELISA [no. DVED00; R&D Systems]). Full-length recombinant human VEGF-C and supernatant from HeLa cells transfected with adenoviral VEGF-D  $\Delta$ NAC were used as positive controls. The mean sensitivities of the ELISA tests were 11.4 pg/ml for VEGF-C and 12.8 pg/ml for VEGF-D.

**Extravasation of Evans blue dye.** Anesthetized mice received an i.v. injection of Evans blue dye (30 mg/kg; Sigma-Aldrich) and 30 minutes later were perfused via the left ventricle with 1% paraformaldehyde in citrate buffer, pH 3.5 (18). Tracheas were weighed, and the dye was extracted in 1 ml formamide overnight at 60°C and measured by spectrophotometry at 610 nm.

**Morphometric measurements and statistics.** Area densities (percentage of total tissue area) of LYVE-1-positive lymphatic vessels and CD31-positive blood vessels viewed in real-time fluorescent images of tracheal whole mounts were measured by stereological point counting of 10 regions per trachea, each 1.7 mm<sup>2</sup> in area. Lymphatic sprouts, defined as tapered projections visible at a screen

magnification of  $\times 180$ , were counted in 5 regions per trachea, each 1.5 mm<sup>2</sup> in area. Filopodia arising from lymphatic sprouts were counted at a screen magnification of  $\times 600$  in 5 regions per trachea, each 0.13 mm<sup>2</sup> in area. Values were expressed per square millimeter of tracheal mucosa. Neutrophils were counted in sections stained with H&E. Values presented are mean  $\pm$  SEM;  $n = 4–8$  mice per group. Significance of differences between means was assessed by analysis of variance followed by the Dunn-Bonferroni test for multiple comparisons, with the level of significance set at  $P < 0.05$ .

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