



Morphomolecular motifs of pulmonary neoangiogenesis in interstitial lung diseases

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Pulmonary injury patterns are the common denominator in progression of fibrosing lung diseases. This study identified distinct manifestations of neoangiogenesis as drivers of pulmonary remodelling in human ILDs, modulated by bone marrow-derived monocytes. <http://bit.ly/34F7KNS>

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ABSTRACT The pathogenetic role of angiogenesis in interstitial lung diseases (ILDs) is controversial. This study represents the first investigation of the spatial complexity and molecular motifs of microvascular architecture in important subsets of human ILD. The aim of our study was to identify specific variants of neoangiogenesis in three common pulmonary injury patterns in human ILD.

We performed comprehensive and compartment-specific analysis of 24 human lung explants with usual interstitial pneumonia (UIP), nonspecific interstitial pneumonia (NSIP) and alveolar fibroelastosis (AFE) using histopathology, microvascular corrosion casting, micro-computed tomography based volumetry and gene expression analysis using Nanostring as well as immunohistochemistry to assess remodelling-associated angiogenesis.

Morphometrical assessment of vessel diameters and intervascular distances showed significant differences in neoangiogenesis in characteristically remodelled areas of UIP, NSIP and AFE lungs. Likewise, gene expression analysis revealed distinct and specific angiogenic profiles in UIP, NSIP and AFE lungs.

Whereas UIP lungs showed a higher density of upstream vascularity and lower density in perifocal blood vessels, NSIP and AFE lungs revealed densely packed alveolar septal blood vessels. Vascular remodelling in NSIP and AFE is characterised by a prominent intussusceptive neoangiogenesis, in contrast to UIP, in which sprouting of new vessels into the fibrotic areas is characteristic. The molecular analyses of the gene expression provide a foundation for understanding these fundamental differences between AFE and UIP and give insight into the cellular functions involved.

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Introduction

The term interstitial lung disease (ILD) refers to a complex group of hundreds of non-neoplastic pulmonary diseases with variable morphologies, clinical presentations and progression tendencies. ILD is characterised by an aberrant response to injury, generally leading to compartment-specific disruptions of lung architecture [1]. In wound healing, the impact of angiogenesis and physiological vascular architecture has long been recognised as a crucial element, particularly in the interaction of granulation tissue and extracellular matrix [2]. Although blood vessels make up >70% of normal lung parenchyma volume [3, 4], the influence of pulmonary microvascular changes on fibrotic remodelling in humans and especially on the progress of ILD remains poorly understood.

The pivotal role of neoangiogenesis in idiopathic pulmonary fibrosis (IPF), an aggressive variant of the idiopathic interstitial pneumonias, was revealed by EBINA *et al.* [5], who could demonstrate an increased capillary density with increased endothelial proliferation in areas of minimal fibrosis, but decreased density in the adjacent, most extensively remodelled fibrotic lesions in afflicted patients.

Microvascular corrosion casting and high-resolution computed tomography (micro-computed tomography) are well suited to visualise and analyse the vascular structure of lung tissue [6]. Using these methods, the heterogeneous complexity of the vasculature in neoplasias was characterised [7–9]. Here, highly vascularised, capsule-like structures around (myo)fibroblastic foci [10] were described. In ILD, the only previous demonstration of prominent vascular obstruction was in lungs of patients with IPF [11].

Aberrant expression of a multitude of angiogenesis-related molecular mediators has been described in the lungs and blood of IPF patients, including vascular endothelial growth factor (VEGF), basic fibroblast growth factor, CXC chemokines with ELR motif and transforming growth factor (TGF)- β . In addition, the highly angiostatic molecule pigment epithelium-derived factor was found to be overexpressed within the fibroblastic foci, typical (but not specific) for the usual interstitial pneumonia (UIP) pattern commonly seen in IPF [10, 12]. In a bleomycin-induced murine fibrosis model, CXCL11 reduced pulmonary fibrosis and the absolute number of endothelial cells found in the remodelled compartments [13]. Our own studies on the antiangiogenic multikinase inhibitor nintedanib [14, 15] revealed an attenuation of aberrant microvascular architecture. In detailed studies, we and others found an overall decreased vascular density with normalisation of vessel diameters [16] and a reduction of CCL18 secretion and macrophage polarisation in treated lungs [17, 18].

In a computed-tomography analysis of the degrees of fibrotic remodelling in IPF patients, JACOB *et al.* [19] reported on the impact of radiological variables for predicting progression and mortality. They demonstrated that pulmonary vascular volumes had the strongest correlation with mortality [19]. We and others have repeatedly hypothesised that microvascular alterations represent (one of) the first step(s) in the pathogenesis of IPF [20]. The loss of vascular integrity and the destruction of the alveolar–capillary membrane drives the continuous remodelling of the pulmonary architecture by pro-angiogenic and pro-fibrotic stimuli [16]. Recent observations also demonstrated that pulmonary capillaries can undergo an endothelial–mesenchymal transition in bleomycin-induced fibrosis [21].

Given the significant role of angiogenesis in repair reactions, one can reasonably assume that varying microvascular alterations may be reflections of both the differing response-to-injury patterns and the subsequent remodelling of the pulmonary architecture observed in ILD [22, 23]. Prior studies have focused predominantly on epithelial–mesenchymal pathogenesis of ILD, while possible vascular mechanisms remain poorly characterised or understood. Therefore, the aim of our present study was to characterise the morphological and molecular differences of vascularisation and angiogenesis in ILD, focusing on the

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predominant and clinically important patterns/entities of UIP, nonspecific interstitial pneumonia (NSIP) and alveolar fibroelastosis (AFE).

Methods

Patient selection, specimens and study groups

Fresh lung explants of 24 patients with different subtypes of ILD were included in this study. All diagnoses were made by experienced pulmonary pathologists following the European Respiratory Society and American Thoracic Society guidelines. Patient groups consisted of patients with characteristic radiological and histological patterns: UIP (n=6, 0 females, mean±SD age at transplantation 61.4±6.8 years), idiopathic NSIP (n=6, 4 females, age at transplantation 52.3±9.2 years) and AFE (n=6, 5 females, age at transplantation 40.7±27.8 years) in the context of chronic graft *versus* host disease, following radio-/chemotherapy, as the histological hallmark of chronic lung allograft dysfunction and idiopathically (idiopathic pleuroparenchymal fibroelastosis). As controls, we used surgically resected lung allograft samples which had been obtained immediately prior to transplantation for donor–recipient size matching (n=6, 1 female, age at transplantation 29.0±22.8 years). Despite careful specimen selection, a significant difference in age between UIP and controls remains (p-value 0.046 (<0.05), Holm–Bonferroni method). All IPF/UIP patients showed varying degrees of mild pulmonary hypertension consistent with sclerosis of the elastic and muscular pulmonary arteries. Tissue specimens and samples were handled anonymously according to the principles expressed in the Declaration of Helsinki. The study was designed and performed following the requirements of the local ethics committee at Hannover Medical School (ethics vote no. 2702-2015). Sampling and histology, microvascular corrosion casting, micro-computed tomography and gene expression are described in the supplementary methods.

Gene expression analysis of ILD explants

All mRNA expression data was obtained *via* the nCounter® Analysis System (NanoString Technologies, Seattle, WA, USA) using the PanCancer Progression Panel (770 genes including 30 reference genes). Normalisation of raw counts was performed using the nSolver™ analysis software version 3.0 (NanoString Technologies) and a modified version of the nCounter® advanced analysis module (version 1.1.5). The normalisation process included positive normalisation (geometric mean), negative normalisation (arithmetic mean) and reference normalisation (geometric mean) using the five most suitable reference genes from the total of 30 available reference genes selected by the geNorm algorithm [24].

Protein expression analysis of ILD explants

Detection of expressed proteins was done using standard immunohistochemistry and multiplex labelling using the OPAL 7 system. Additional information can be found in the supplementary methods.

Functional analysis

In order to gain insight on the regulation of physiological functions conveyed by the mRNA expression in different ILD entities, we made use of the Ingenuity Pathway Analysis tool (IPA; Qiagen, Hilden, Germany) [25]. We considered p-values significant according to the following levels of confidence: p<0.05, p<0.01 and p<0.001. Additional information can be found in the supplementary methods.

Statistical analysis of vascular morphometry

Analysis of vascular morphometry was based on measurements in ≥24 lung samples (intervascular distances: 2566 measuring points; vessel diameters: 2259 measuring points). Intervascular distances and vessel diameters were analysed by a one-way ANOVA. If the ANOVA indicated significant differences (p<0.05) between samples within a group, an all-pairwise analysis was performed (Tukey). Commercially available statistical analysis software (JMP® 8.0.2.2; SAS Institute Inc., Cary, NC, USA) was used to perform the statistical tests.

Results

UIP, NSIP and AFE present a specific vascular morphogenetic pattern

In contrast to regular lung morphology (figure 1a–c), histopathological subtypes of ILDs reveal specific morphologic alterations. A characteristic feature of the UIP pattern is a transition between a spatially and temporally heterogeneous interstitial fibrosis, driven by aggregates of myofibroblasts (fibroblast foci), and morphologically normal lung tissue (figure 1f). UIP shows patchy and dense fibrosing remodelling predominantly in the subpleural spaces (figure 1d and e). NSIP shows homogeneous and mild interstitial fibrosis with a uniform and diffuse thickening of alveolar septa as well as some degree of interstitial inflammation (figure 1g–i). The development of AFE involves three stages [26, 27]. The AFE pattern (figure 1j–l) is characterised by preserved elastic fibres outlining the former alveolar septa and a

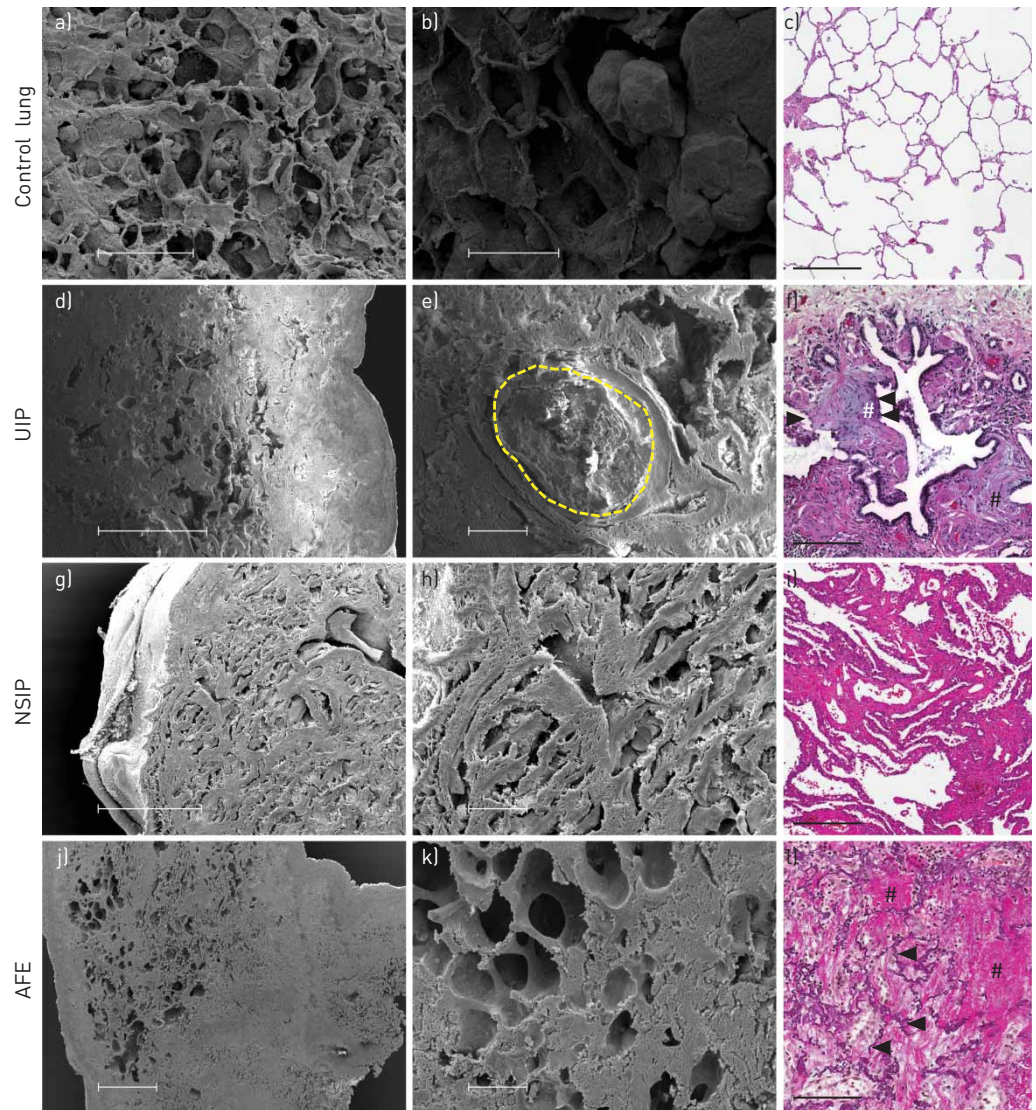


FIGURE 1 Different morphological patterns in usual interstitial pneumonia (UIP), nonspecific interstitial pneumonia (NSIP) and alveolar fibroelastosis (AFE) lungs. Scanning electron micrographs and histological sections demonstrate the injury-specific morphological pattern. a–c) In healthy control lungs, lung architecture includes thin-walled alveolar ducts and alveoli, with sparse bronchial mucous remnants. d–f) The cardinal features of UIP represent a loss of architecture with d) heterogeneous, patchy parenchymal, mostly subpleural fibrotic involvement and fibroblastic foci which are indicative of the progression of the fibrosis; e) adjoining the fibrotic foci, organised thrombotic occlusion (yellow dotted line) is frequently observed in the draining veins; f) adjacent to fibroblastic foci (#), perifocal alveolar type I cells are replaced by hyperplastic alveolar type II cells (arrowheads). g–i) NSIP is characterised by a diffuse and uniform injury pattern with a moderate fibrotic remodelling of alveolar septa with an interstitial infiltration by lymphocytes, histiocytes and plasma cells (i). j–l) In contrast to conventional fibrotic remodelling, the development of alveolar fibroelastosis (AFE) involves three stages: 1) the “fibrin stage” with fibrin filled alveoli in response to an initial alveolar epithelial damage; 2) the “histiocytic stage” with degradation of alveolar fibrin exudates by local macrophages and accumulation of fibroblasts followed by deposition of collagen within the alveolar space; and 3) the “full developed stage” of AFE presents a concomitant elastosis of alveolar septa (l) (arrowheads), whereas alveolar spaces are consolidated with collagenous fibres (l) (#). Scanning electron microscopy was performed using a Philips XL30 microscope (Philips, Eindhoven, the Netherlands) at 15 keV and 21 μ A. Hematoxylin and eosin staining (c, f and i), and Elastica van Gieson staining (l). Scale bars=200 μ m).

continuously increased deposition of elastin in the alveolar interstitium, whereas the airspaces are completely filled up by extracellular matrix.

The immunohistochemical analysis allows a morphological characterisation of fibrotic areas where an expression of the endothelial marker CD31, the bronchoepithelial marker CK7 and the lymphatic marker

podoplanin can all be frequently observed in close spatial proximity (figure 2). However, immunohistochemical staining of the fibrotic areas cannot reflect the three-dimensional architectural aspects of fibrotic and vascular remodelling.

Our analysis of microvascular corrosion casts of UIP, NSIP and AFE lungs show the underlying morphological patterns (supplementary material 1). The microvascular architecture of UIP lungs reveals a tumour-like and chaotic aberrant distribution of blind-ending, partly occluded sinusoid-like vessels with a lack of hierarchy and wide alterations of vessel diameters with reduced vessel density (figure 3b). The NSIP pattern is characterised by densely packed, dilated tufts of vessel formation with abnormal tortuosity within the alveolar septa (figure 3c). The AFE pattern exhibits a certain resemblance with the NSIP pattern whereby the microvasculature in the preserved alveolar septa suggests some degree of compression (figure 3d).

Vascular morphometry reveals significant alterations in vascular remodelling in UIP, NSIP and AFE

We assessed the microvascular unit of fibrotic areas by three-dimensional morphometry of the microvascular corrosion casts. The individual vessel diameters in the fibrotic foci, in alveolar vessels of UIP, NSIP, AFE and healthy control lung revealed highly significant differences (figure 4a). The highest vessel diameters were seen in UIP lungs (mean \pm SD 16.5 \pm 7.7 μ m) compared with NSIP lungs (12.7 \pm 5.1 μ m; p <0.001), AFE lungs (13.1 \pm 5.1 μ m; p <0.001, ANOVA) or healthy control lungs (mean 5.9 \pm 1.9 μ m; p <0.001, ANOVA). The mean intervascular distances (distance between adjacent individual vessel segments) yielded similar significant results. The highest intervascular distances were observed in UIP lungs (59.4 \pm 24.1 μ m) in comparison to NSIP lungs (18.8 \pm 9.0 μ m; p <0.001, ANOVA), AFE lungs (11.6 \pm 4.1 μ m; p <0.001) or the healthy controls (12.0 \pm 3.7 μ m; p <0.001). Vascular volumetry was conducted on micro-computed tomography of microvascular corrosion casts (supplementary material 2a–c) to assess the vascular changes of upstream vasculature with a diameter >50 μ m (figure 4c). UIP lungs showed a higher, albeit significant, vascular volume (36.1 \pm 5.9%; p =0.013) and vascular surface–volume ratio (0.43 \pm 0.7; p =0.055, ANOVA) compared with NSIP lungs (18.9 \pm 2.0% and 0.54 \pm 0.6, respectively), AFE lungs (mean 16.6 \pm 3.7% and 0.39 \pm 0.04, respectively) and controls (6.24 \pm 0.8% and 0.58 \pm 0.11, respectively) (data not shown).

Intussusceptive and sprouting angiogenesis accelerate fibrotic neovascularisation

We identified intussusceptive and sprouting angiogenesis within the microvascular corrosion casts of all analysed lungs. The formation of intussusceptive pillars was progressed by initial vasodilation followed by incorporation of CD34/31-positive progenitor-like [28, 29] as well as the activation of mesenchymal stem cells which span a transluminal bridge, the “intussusceptive pillar” (figure 5a and b) (supplementary material 5; quantification of proliferation of CD34/CD31 positive cells by Ki-67). The occurrence of

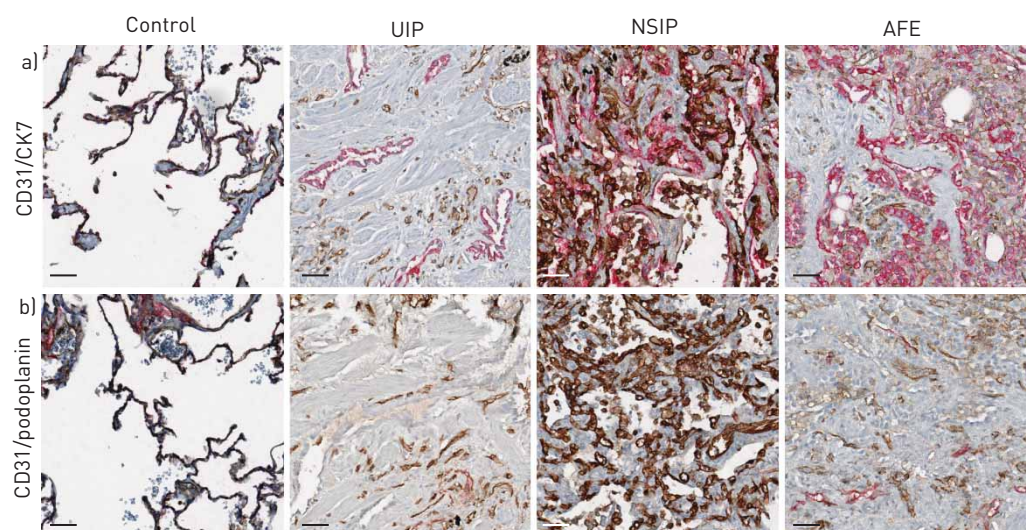


FIGURE 2 Immunohistochemical detection of CD31, CK7 and podoplanin expression in alveolar fibroelastosis (AFE), nonspecific interstitial pneumonia (NSIP) and usual interstitial pneumonia (UIP). Double-staining against the endothelial marker CD31 (brown) and a) the bronchoepithelial marker cytokeratin CK7 (red) and b) the lymphatic marker podoplanin (red). Staining results demonstrate the spatial heterogeneity of vascularisation in the fibrotic areas in AFE, NSIP and UIP, as compared to controls. Scale bars=50 μ m.

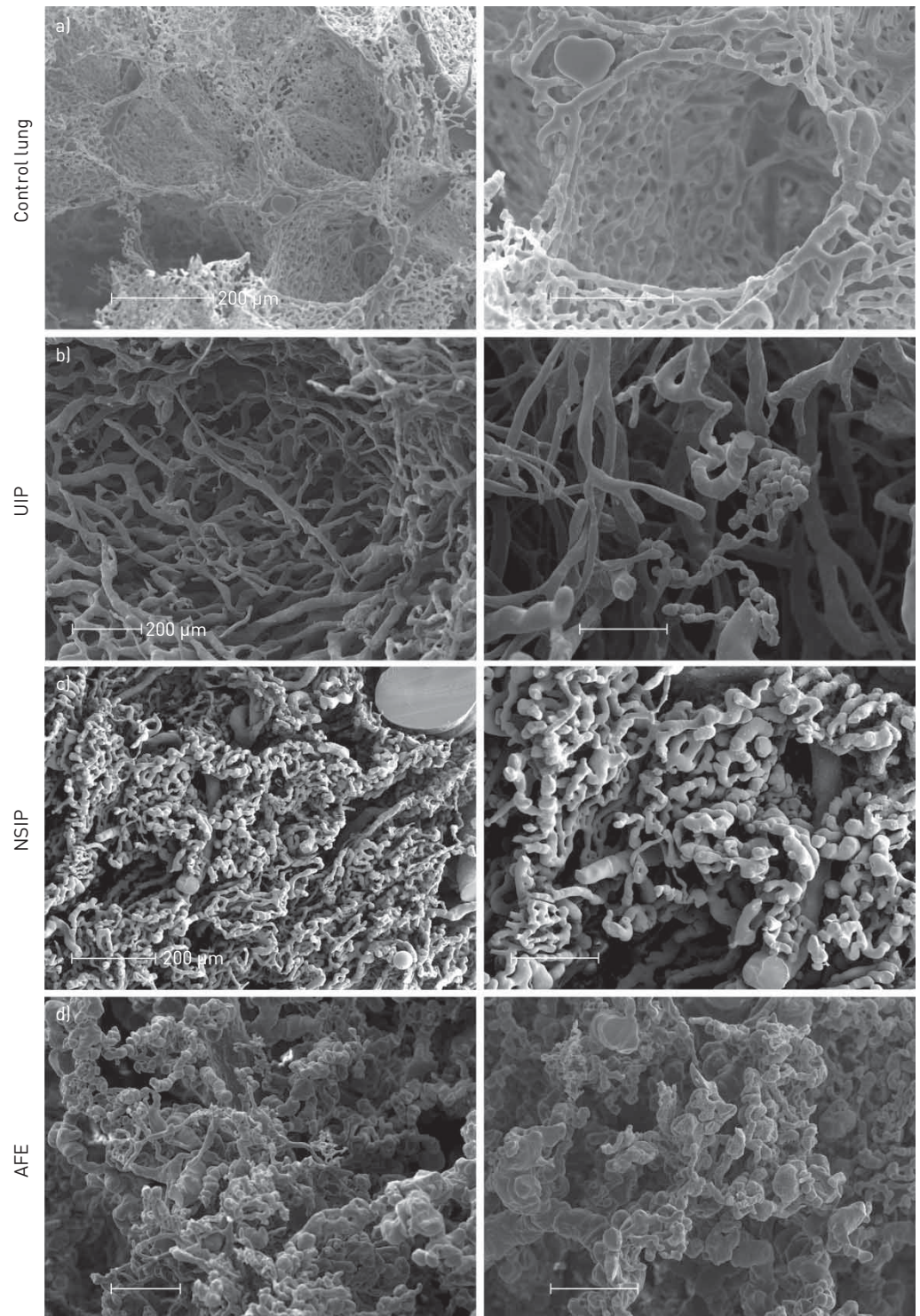


FIGURE 3 Microvascular architectural alterations in usual interstitial pneumonia (UIP), nonspecific interstitial pneumonia (NSIP) and alveolar fibroelastosis (AFE). Scanning electron micrographs of microvascular corrosion casts illustrate the substantial architectural differences between the different injury patterns. a) Healthy control lung vasculature is characterised by thin-walled alveolar capillary plexuses aligned along the alveolar duct; b) UIP lungs demonstrate an aberrant vasculature with blunt, sinusoid-like vessels, without a clear vessel hierarchy, but instead high variability of vessel diameters and small vessel sprouts; c) NSIP lungs present with dense, tortuous dilated tufts of vessel formations in the alveolar septa and frequent intussusceptive pillars; d) AFE lungs resemble the appearance of the NSIP lungs with pronounced vascular alterations in the alveolar septa. Scale bars=100μm, unless otherwise stated.

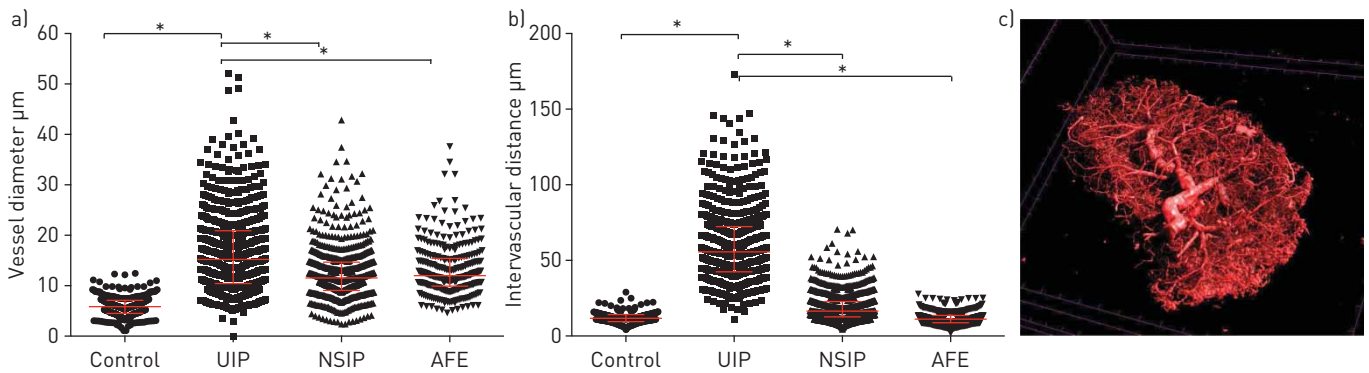


FIGURE 4 Vascular morphometry of usual interstitial pneumonia (UIP), nonspecific interstitial pneumonia (NSIP) and alveolar fibroelastosis (AFE). The morphometrical assessment was carried out in the alveolar plexus of fibrotic areas in UIP, NSIP and AFE, and in healthy control lungs. a) Vessel diameters and b) intervascular distances showed significant differences between the different injury patterns. Scatter plots show the respective median and the 25th and 75th percentile in red. ***: $p < 0.001$ (ANOVA). c) Micro-computed tomography on corrosion casts were performed to assess the volumetric changes of upstream vasculature. Only vascular changes of $> 50 \mu\text{m}$ diameter were evaluated. Three-dimensional visualisation of microvascular corrosion casts shows exemplarily the upstream vascular hierarchy of a NSIP lung.

sprouting and intussusceptive angiogenesis was observed in all fibrotic lungs, whereas NSIP lungs showed significantly higher frequencies of pillar formations in comparison to UIP and AFE lungs (figure 5c; $p < 0.001$, ANOVA). In contrast to intussusception, the highest frequencies (> 10 -fold increase) of sprout formation were seen in UIP lungs compared to NSIP and AFE lungs ($p < 0.001$, ANOVA) (figure 5c). The high occurrence of intussusceptive angiogenesis in NSIP lungs was predominantly observed in the

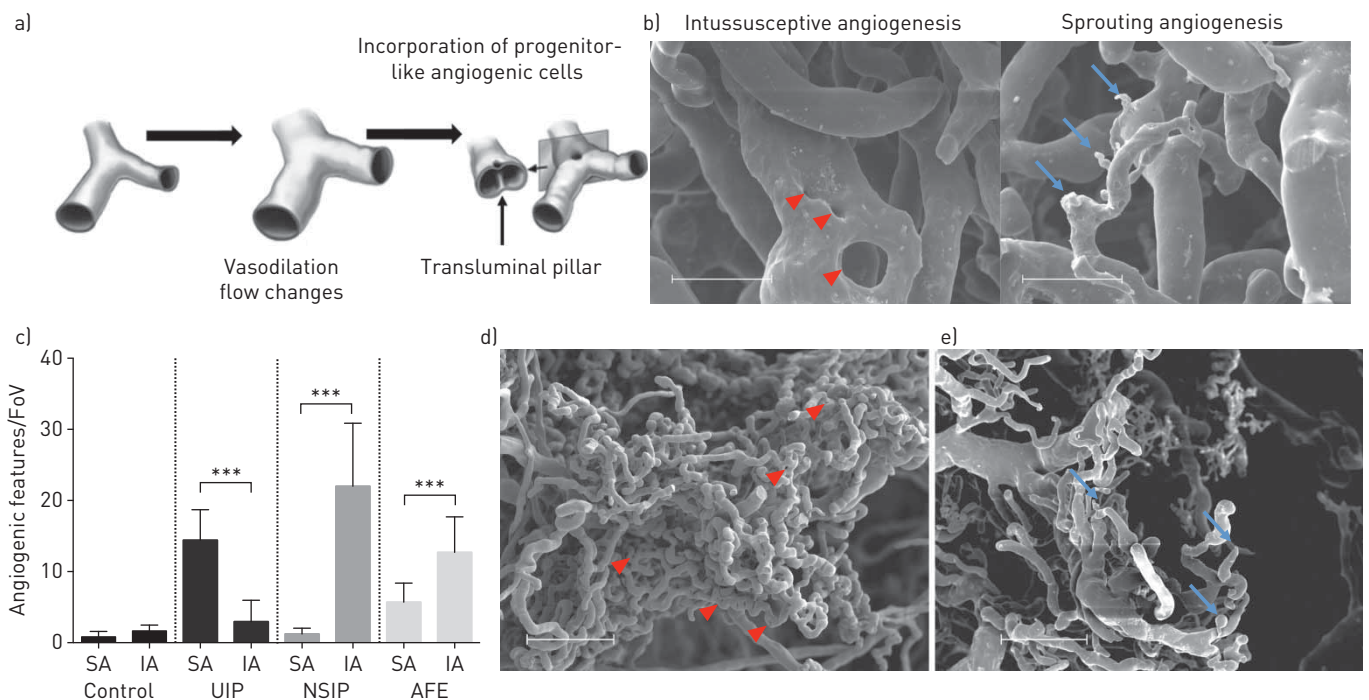


FIGURE 5 Neoangiogenic remodelling of fibrotic lung tissue by intussusceptive and sprouting angiogenesis. a) Schematic representation of intussusceptive angiogenesis. Blood flow and velocity are reduced by marked vasodilation. After the dilation of the vessel, intraluminal pillars form at vessel bifurcations by an intraluminal intussusception of myofibroblasts and incorporation of tissue resident or blood-borne progenitor-like angiogenic cells. b) The intussusceptive transluminal pillar, a tissue bridge that leads to multiple septations of the vessel lumina, appears as a "hole" in microvascular corrosion cast (arrowheads) whereas sprouting angiogenesis forms little extraluminal "sprouts" (arrows) from pre-existing blood vessels; scale bars = $100 \mu\text{m}$. c) The numerical density newly formed sprouts (SA) and intussusceptive pillars (IA) per field of view (FoV) assessed on microvascular corrosion casts revealed the highest frequency of intussusceptive pillars in nonspecific interstitial pneumonia (NSIP) lungs, whereby the highest occurrence of sprouts was observed in usual interstitial pneumonia (UIP) lungs. ***: $p < 0.001$ (ANOVA). d) The septal thickening in NSIP lungs is mainly characterised by a dense vascular network which expands by numerous intussusceptive pillars (arrowheads); scale bar = $100 \mu\text{m}$. e) Microvascular architecture of UIP-injured lungs outlines perifocally aberrant blind-ending small calibre vessels with vessel sprouts which are aligned towards the fibroblastic foci (arrows); scale bar: $200 \mu\text{m}$. AFE: alveolar fibroelastosis.

expanded septa, in which a prominent expansion of tuft-like blood vessels was observed (figure 5d). In the UIP lungs, blunt, blind-ending vessels with dilated parent venules were located adjacent to fibroblastic foci. At the tip of these blunt vessels, sprouts were mostly aligning centrally towards the fibroblastic foci (figure 5e).

Differential regulation of mRNA expression and biological functions in UIP, NSIP and AFE

We performed a digital multiplexed gene expression analysis *via* NanoString nCounter® technology (supplementary material 3a–d). Transcript analysis showed significant differences in the expression of 20 genes among the three lung injury patterns compared to healthy control lungs (figure 6a). For the sake of validation, the protein expression of selected genes was also analysed in a compartment specific manner *via* immunohistochemistry. Here, collagens (COL I, III, IV) and collagen remodelling matrix metalloproteinases (MMP2, MMP9, MMP14) correlated well with the mRNA expression in the corresponding sites of neoangiogenesis (supplementary material 4). Comparative gene analysis demonstrated an overlap of statistically significant differentially expressed genes between UIP, NSIP and AFE (figure 6b). 16 genes were significantly upregulated in AFE lungs, whereas only four genes were differentially expressed in UIP lungs and one gene in NSIP lungs. One gene (*IGF1*) was significantly upregulated ($p < 0.001$) as a common denominator of AFE and UIP, whereas *MEG3* was significantly ($p < 0.01$) expressed in both injury patterns of UIP and NSIP. *COL1A1* and *THY1* are highly significantly ($p < 0.001$) expressed in NSIP and AFE lungs (supplementary material 3). In addition to canonical pathways, differentially expressed genes were also categorised to related functional pathways using IPA (figure 6c). By this approach, AFE lungs exhibited an upregulation of different function pathways compared to NSIP and UIP. The analysis argued for significant ($p < 0.01$) and highly significant ($p < 0.001$) activation of epithelial cell and fibroblast migration, as well as macrophage differentiation in AFE and NSIP lungs. Moreover, activation of epithelial–mesenchymal transition was observed in AFE lungs. Neoangiogenesis, migration of endothelial cells and fibrogenesis were also significantly activated in AFE compared to UIP lungs ($p < 0.01$). In contrast to AFE, the lungs of NSIP and UIP showed that the pathways are less activated (NSIP > UIP). In the analysis, signalling pathways of inflammation are only upregulated in UIP lungs in comparison to AFE and NSIP.

Discussion

In this study, we assessed the morphogenetic and molecular aspects of angiogenesis in the interstitial lung injury patterns of UIP, NSIP and AFE. For this we assembled patient cohorts which showed no significant differences with regard to age or sex, with the exception of the UIP group (older, in general). However, since the statistical analysis of the gene expression data of the healthy control group revealed no age-dependent effects, it seems plausible that NSIP, AFE and UIP could be analysed in comparison to the healthy control group.

First, our data demonstrate that the histopathological morphological features widely depend on the underlying vascular architectural alterations. NSIP and AFE reveal a similar vascular injury pattern with pronounced remodelling in the thickened alveolar septa, whereas UIP lungs are characterised by upregulation of vascular density in the upstream vessels and a reduction of vessel density in the perifocal and alveolar perifocal capillary plexus. Furthermore, the difference in vessel diameters and intervascular distances reflect these individual changes. Second, our data provide evidence for a role of sprouting and intussusceptive angiogenesis in the expansion of perifocal and alveolar blood vessels. Third, our findings on differences and similarities of angiogenesis-related gene expression underline the morphological features of vascular remodelling in UIP, NSIP and AFE (figure 7).

The impact of angiogenesis on the pathogenesis and progression of ILD has become a subject of controversy [12, 20, 25]. While several studies proposed that stimulated aberrant angiogenesis is a core characteristic of ILD, other authors have reported on reduced vascular densities in lung biopsies of IPF patients, in particular [5, 20, 30]. The legitimate question “angiogenesis in pulmonary fibrosis – too much or not enough?” raised by HANUMEGOWDA *et al.* [12] cannot be easily addressed due to the architectural complexity of the vascular system and its density on two-dimensional histological slides. Therefore, in this study, using three-dimensional analysis, we put effort into unravelling the complexity of both spatial and temporal heterogeneity of microvascular changes present in UIP, NSIP and AFE lungs. It seems contradictory that the overall vascular volume in UIP is upregulated, whereas the perifocal vascular density is reduced and thinned out while the exact opposite was evident with NSIP lungs. The predictive clinical value of vascular volume as a marker for mortality was recently discussed in depth by JACOB *et al.* [19], using a computer-based algorithm to assess pulmonary vascular volumes measured by computed tomography. However, this technique only captures the alterations of upstream blood vessels from the alveolar and perifocal capillary plexus, but does not address the heterogeneous vascular morphological aspects of newly formed blood vessels that probably contribute to the progression of the disease. The

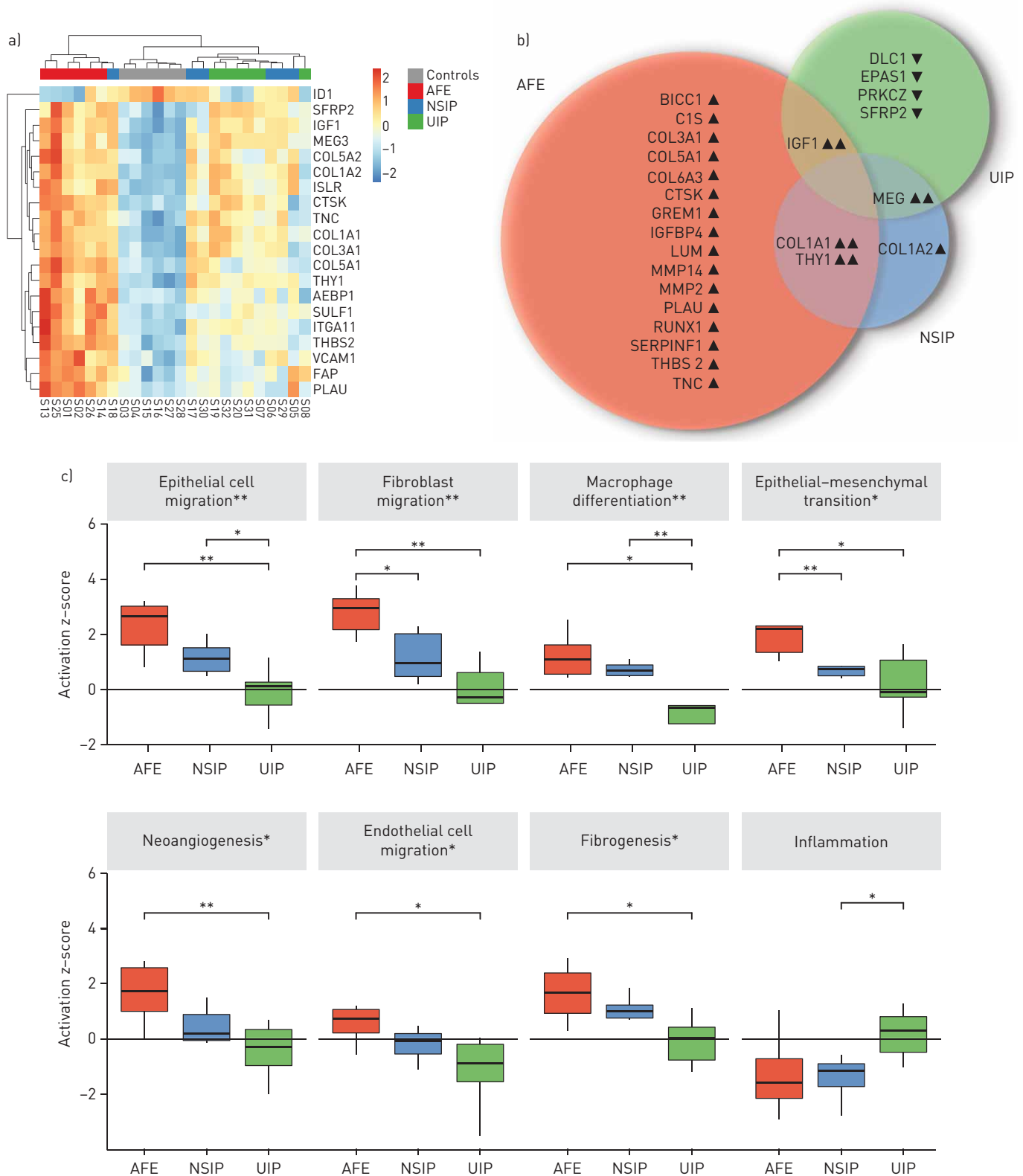
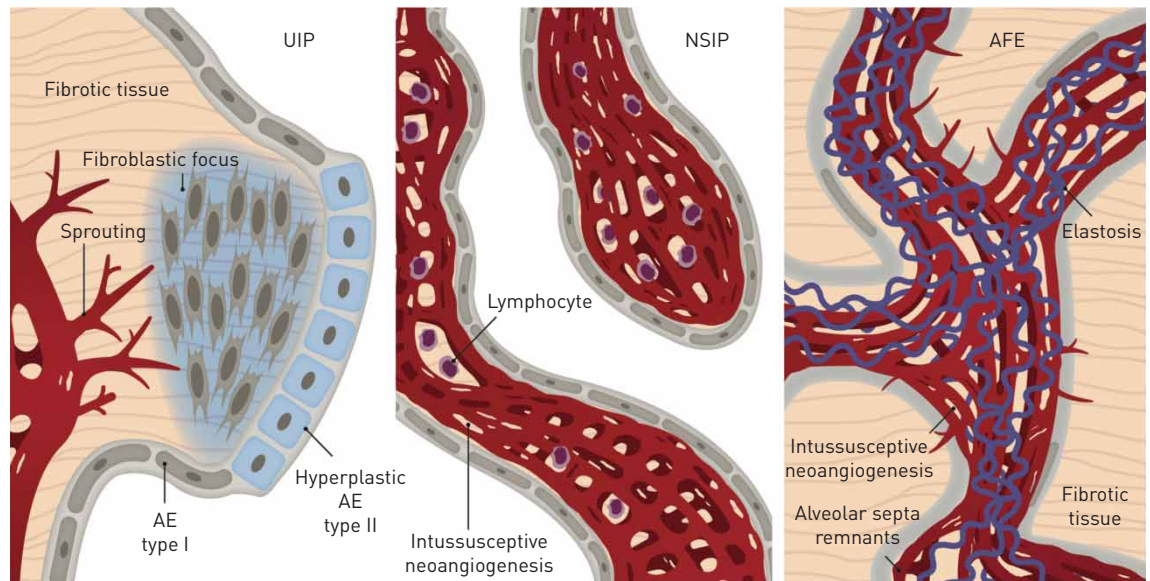


FIGURE 6 Differential regulation of messenger (m)RNA expression and biological functions in usual interstitial pneumonia (UIP), nonspecific interstitial pneumonia (NSIP) and alveolar fibroelastosis (AFE). **a)** A heatmap of hierarchically clustered mRNA expression data shows distinct clusters for AFE and control samples. The displayed gene set is limited to 20 significantly differentially expressed genes in a multigroup comparison of AFE, NSIP, UIP and control samples. **b)** Venn diagram of genes that are expressed differentially, with a p-value of ≥ 0.05 , compared to control samples. Arrows indicate up- or downregulation of genes compared to controls. While a large number of genes in AFE samples is significantly upregulated, no gene is commonly regulated in all three entities. **c)** Significant differences in the presumed regulation of biological functions. Significance level in a multigroup test of all three entities: *: $p < 0.05$; **: $p < 0.01$. AFE and UIP frequently exhibit significant differences.



	UIP	NSIP	AFE
Vascular volume	↑↑	↑	↗
Intervascular distances	↑↑	↘	↔
Vessel diameter	↑↑	↗	↑
Sprouting	↑↑	↔	↑
Intussusception	↔	↑↑	↑
Gene expression	DLC1, EPAS1, PRKCZ, SFRP2, IGF1, MEG3	COL1A1, COL1A2, THY1	BICC1, C1S, COL1A1, COL3A1, COL5A1, COL6A3, CTSK, GREM1, IGF1, IGFB4, LUM, MMP14, MMP2, PLA1, RUNX1, SERPINF1, THBS2, THY1, TNC

FIGURE 7 Schematic illustration of morphomolecular motifs in histopathological subtypes usual interstitial pneumonia (UIP), nonspecific interstitial pneumonia (NSIP) and alveolar fibroelastosis (AFE). UIP is characterised by prominent basophilic fibroblastic foci with adjacent hyperplastic type 2 alveolar epithelial (AE) cells. Here, we frequently find newly formed blood vessels *via* sprouting reaching out towards the fibroblastic foci. In NSIP thickened septa are composed of blood vessels with intussusceptive pillars and increased amounts of lymphocytes. In AFE former alveolar air spaces are filled with collagen, while former alveolar walls are remodelled, showing elastosis and increased amounts of intussusceptive blood vessels.

formation of fibrotic areas leads inevitably to haemodynamic changes with an opening of anastomoses between the pulmonary and bronchial circulation. These haemodynamic and micromechanical forces are known to enhance the expansion of vascular plexuses by intussusceptive angiogenesis [26, 31]. Thus, this pillar formation and branch remodelling may represent a pivotal adaptive response to the continually increasing blood flow and blood pressure during inflammation and regeneration. Previously, it has been shown by our group that chronic inflammation triggers the migration and recruitment of progenitor-like angiogenic cells to the site of remodelling [25–27]. In the inflammatory ILD patterns of NSIP and AFE [32, 33], we observed a frequent occurrence of intussusceptive pillars.

In contrast, angiogenesis in UIP lungs was driven by hypoxia with perifocal sprouting angiogenesis, which, among other factors, was indicated by the upregulation of insulin-like growth factor (IGF)-1 [34]. Comparable to malignant tumours, the UIP-specific vascularity showed a chaotic vessel-like arrangement with numerous blunt ends. This hypoxia-induced vascular remodelling is associated with the modification of the extracellular matrix by MMPs, and as we showed earlier, inhibition of IGF-1 can impede sprouting angiogenesis significantly [35]. In addition, localised hypoxia results in metaplasia and hyperplasia of alveolar type II cells by Notch signalling and Krt5^{pos} basal-like cell expansion in alveolar repair and remodelling [36].

Nintedanib, a multikinase inhibitor with anti-angiogenic properties, has been shown to have efficacy in reducing the decline of lung function in IPF patients [37]. Besides anti-inflammatory and anti-fibrotic effects,

the question remains whether an anti-angiogenic therapy may play a pivotal role in improving the functional outcome in IPF patients. Our own studies in a bleomycin-induced mouse fibrosis model demonstrated that the altered vascular architecture appears to be “normalised” after nintedanib treatment [16], a phenomenon also found in tumours treated with an anti-angiogenic therapy [35, 38]. Moreover, intervascular distances and vessel diameters showed significant differences in nintedanib-treated animals, whereas lung function improved under the therapy. Therefore, the structure–function dynamics of microvascular architecture are likely to underlie the injury and progression patterns of ILD. Otherwise, FARKAS *et al.* [39] could show in an IPF rat model with transient overexpression of active TGF- β 1 that a decreased vascularity is associated with an extensive remodelling and a secondary pulmonary hypertension. In this model, the administration of VEGF showed an amelioration of pulmonary hypertension *via* inhibition of endothelial apoptosis, yet also an aggravation of pulmonary fibrosis.

Some authors suggest that vascular alterations might represent an initial trigger in the pathogenesis and the progression of ILD [20, 40] and conclude that patients with pulmonary fibrosis show aspects of pulmonary venous occlusive disease (PVOD) with aberrant postcapillary expansion by haemangiomatosis. Similar observations were made in non-fibrotic areas of patients with IPF [40], in which 65% of cases demonstrated occlusive venopathy. Our own recent work [41] on PVOD highlights multifocal intussusceptive neoangiogenesis and vascular sprouting as representing the three-dimensional correlate of progressive pulmonary haemangiomatosis. In addition, we described comparable morphogenetic phenomena in chronic thromboembolic pulmonary hypertension in the form of plexiform vasculopathy with pronounced intussusceptive angiogenesis [42].

NSIP and AFE revealed a shared upregulation of COL1A1 and Thy1/CD90. Thy1/CD90 is expressed on endothelial cells at sites of chronic inflammation and adjacent to malignant neoplasms, as well as on adventitial mesenchymal progenitor cells in adult human arteries, where both are involved in the microvascular expansion by intussusceptive neoangiogenesis [43]. Other studies reported that the intussusceptive capillary growth is the main driver in post-injury angiogenesis in a Thy1.1-nephritis model [43], which is consistent with the recovery of septal blood vessels in human NSIP and AFE lungs. We observed a higher frequency of intussusceptive features in NSIP and AFE fibrotic lungs. The expansion of the microvascular network by intussusception is associated with the recruitment of progenitor-like angiogenic cells forming transluminal pillars [26, 31]. These circulating progenitor-like angiogenic cells are recruited as tissue-resident or bone-marrow derived monocytes which can transdifferentiate among other types into proangiogenic M2-polarised alveolar macrophages [29, 44, 45] or into endothelial cells which then incorporate into newly formed intussusceptive blood vessels [42, 45]. ALVAREZ *et al.* [44] characterised endothelial cells isolated from the lung microcirculation as highly vasculogenic and capable of renewing the entire hierarchy of endothelial cell growth.

Recent evidence [46] confirmed that bone marrow derived mononuclear cells are major constituents in the vascular expansion by intussusceptive angiogenesis, controlled by SDF-1/CXCR4 signalling. Our group observed similar effects on the migration of CD11b⁺ monocytes during lung regeneration [29]. In summary, we provide the first evidence that the histopathological injury patterns of life-threatening ILDs are highly dependent on the underlying vascular architectural alterations and vessel densities in UIP, NSIP and AFE lungs. Whereas UIP lungs showed a higher density of upstream vascularity and lower density in perifocal blood vessels, NSIP and AFE lungs revealed densely packed alveolar and septal blood vessels. Microvascular alterations by sprouting (UIP) and intussusceptive angiogenesis (NSIP, AFE) may contribute as one of the pivotal pathogenetic mechanisms in the progression of disease in the capillary–alveolar interaction. Our data can serve to identify novel therapeutic targets with the potential to favourably alter the course of ILD by influencing aberrant angiogenesis in the different subtypes of ILDs.

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