Alveolar Macrophage Dysregulation in Hermansky-Pudlak Syndrome Type 1

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Rationale: Individuals with Hermansky-Pudlak syndrome type 1 (HPS-1), an autosomal recessive disorder characterized by defective biogenesis of lysosome-related organelles, develop an accelerated form of progressive fibrotic lung disease. The etiology of pulmonary fibrosis associated with HPS-1 is unknown.

Objectives: To investigate the potential pathogenesis of pulmonary fibrosis in HPS-1, lung cells and proteins from individuals with HPS-1 were studied.

Methods: Forty-one subjects with HPS-1 with and without pulmonary fibrosis were evaluated with pulmonary function tests, high-resolution computed tomography scan, and bronchoscopy. Bronchoalveolar lavage cells and analytes were analyzed.

Measurements and Main Results: Concentrations of total bronchoalveolar lavage cells and alveolar macrophages were significantly higher in epithelial lining fluid from subjects with HPS-1 with and without pulmonary fibrosis compared with healthy research volunteers. Concentrations of cytokines and chemokines (i.e., monocyte chemoattractant protein-1, macrophage inflammatory protein-1 α , and granulocyte-macrophage colony-stimulating factor) in alveolar epithelial lining fluid were significantly higher in subjects with HPS-1 with and without pulmonary fibrosis compared with healthy research volunteers (P < 0.001). In vitro, HPS-1 pulmonary fibrosis alveolar macrophages, which did not express HPS1 mRNA, secreted significantly higher concentrations of monocyte chemoattractant protein-1, macrophage inflammatory protein-1 α , and regulated upon activation, normal T cell expressed and secreted (RANTES) protein compared with normal cells (P = 0.001, P = 0.014, and P = 0.011, respectively). Pirfenidone suppressed HPS-1 alveolar macrophage cytokine and chemokine secretion in vitro in a dose-dependent manner.

Conclusions: In HPS-1, alveolar inflammation predominantly involves macrophages and is associated with high lung concentrations of cytokines and chemokines. HPS-1 alveolar macrophages provide a model system in which to study the pathogenesis and treatment of HPS pulmonary fibrosis.

Keywords: inflammation; cytokines; chemokines; bronchoalveolar lavage; pirfenidone

Hermansky-Pudlak syndrome type 1 (HPS-1), a rare autosomal recessive disease, occurs commonly in northwest Puerto Rico but can also affect individuals without Puerto Rican heritage

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AT A GLANCE COMMENTARY

Scientific Knowledge on the Subject

Limited information is available regarding the pathogenesis of Hermansky-Pudlak syndrome pulmonary fibrosis, a progressive and often fatal disease.

What This Study Adds to the Field

This study demonstrates that high concentrations of activated alveolar macrophages, cytokines, and chemokines are found in individuals with Hermansky-Pudlak syndrome type 1. Dysregulation of alveolar macrophages may contribute to the pathogenesis of Hermansky-Pudlak syndrome pulmonary fibrosis.

(1-3). Clinical manifestations include oculocutaneous albinism, platelet dysfunction due to a storage pool defect, granulomatous colitis, and fibrotic lung disease, which typically leads to death in the fourth or fifth decade of life (1-3). The pulmonary fibrosis of HPS-1 is largely unresponsive to conventional medical treatment, although patients with mild to moderate disease may respond to treatment with pirfenidone (4). In patients with HPS-1 and advanced pulmonary fibrosis, lung transplantation has been successfully performed (5, 6).

To date, eight genetic subtypes of HPS with various phenotypes have been identified in humans, and six subtypes are not associated with fibrotic lung disease (3). Although the exact function of most HPS proteins remains unknown, they can interact to form complexes that regulate biogenesis of lysosome-related organelles. Individuals with HPS-2 have mutations in *AP3B1*, which encodes the β 3A subunit of the adaptor protein-3 complex (7, 8). Patients with HPS-3, HPS-5, and HPS-6 have mutations resulting in abnormalities in biogenesis of lysosome-related organelles complex-2 (BLOC-2), and those with HPS-7 and HPS-8 have mutations affecting BLOC-1 (3, 9–11).

Individuals with HPS-1 or HPS-4 can develop pulmonary fibrosis (3, 12). Patients with HPS-1 have mutations in the *HPS1* gene, located on chromosome 10q23.1–23.3 (13). Virtually all Puerto Rican individuals with HPS-1 are homozygous for a 16-base pair duplication in exon 15 of *HPS1* [c.1472_1487dup16; p.H497QfsX90 (ex 15)], but other mutations in *HPS1* have been reported, and many are associated with progressive pulmonary fibrosis (1, 2, 14–18). The *HPS1* gene encodes HPS1, a 79.3-kD cytosolic and membrane-bound protein involved in biogenesis of lysosome-related organelles (19). The *HPS4* gene encodes HPS4, a 76.9-kD protein; HPS1 associates with HPS4 to form BLOC-3, a protein complex that regulates the intracellular localization of lysosomes and endosomes (20, 21).

Although basic mechanisms causing pulmonary fibrosis in HPS-1 are unknown, findings in lung pathology specimens and murine models of HPS have provided insights into the disease

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pathogenesis. Distinctive histopathologic features of the HPS pulmonary fibrosis include accumulation of ceroid lipofuscin within alveolar macrophages and proliferation of type II alveolar epithelial cells that appear foamy due to the formation of giant lamellar bodies (22, 23). These characteristics have also been described in the lungs of murine models of HPS. Pale ear mice, which have mutations in Hps1, and pearl mice, which have mutations in Ap3b1, serve as models for HPS-1 and HPS-2, respectively (24-29). Pale ear mice develop giant lamellar bodies in type II alveolar epithelial cells, interstitial inflammation, and foamy alveolar macrophages (26, 29). Furthermore, constitutive activation of macrophages in the lung, but not in the blood or peritoneum, occurs in pale ear and pearl mice (28). Notably, pale ear mice do not develop spontaneous fibrosis, but exposure to silica induces persistent alveolar accumulation of activated macrophages and collagen fibers (26, 27). In addition, intratracheal administration of bleomycin to pale ear or pearl mice causes diffuse pulmonary fibrosis, deposition of extracellular matrix protein, and apoptosis of type II cells (29). Taken together, these findings suggest that dysregulation of lung immune and epithelial cells, in combination with certain environmental exposures, may contribute to the pulmonary fibrosis in HPS.

To investigate the potential pathogenesis of lung disease in HPS-1, we studied cells and proteins in the bronchoalveolar lavage fluid of affected individuals. We report that individuals with HPS-1 accumulate increased numbers of total bronchoalveolar lavage cells and macrophages in alveolar fluid, with accompanying elevations in cytokine and chemokine concentrations. We also show that alveolar macrophages cultured from individuals with HPS-1 secrete high concentrations of cytokines, and that pirfenidone down-regulates this cytokine secretion *in vitro*.

METHODS

Subjects

Forty-one adult subjects with HPS-1, evaluated at the National Institutes of Health (NIH) Clinical Center (Bethesda, MD), were enrolled in protocols 95-HG-0193, 97-HG-0085, and 04-H-0211, which were approved by the Institutional Review Boards of the National Institute of Child Health and Human Development, National Human Genome Research Institute, or National Heart, Lung, and Blood Institute. HPS was diagnosed by the presence of oculocutaneous albinism and the absence of platelet dense bodies; the diagnosis of HPS-1 was confirmed by molecular analysis of the HPS1 gene (GenBank NM_000195) (1, 19). Eighty-five healthy research volunteers without lung disease were enrolled in protocols 96-H-0100 and 99-H-0068, which were approved by the National Heart, Lung, and Blood Institute Institutional Review Board. Written informed consent was obtained from all subjects. For patients with HPS, high-resolution computed tomography scan, pulmonary function tests, and bronchoscopy were performed during the same visit.

High-resolution Computed Tomography

High-resolution computed tomography (HRCT) scans of the chest were performed without intravenous contrast during end-inspiration in the prone position, using a 1-mm collimation at 30-mm intervals and a high-spatial-frequency reconstruction algorithm (General Electric Medical Systems, Milwaukee, WI) (2). Pulmonary fibrosis was diagnosed by characteristic findings on HRCT scans of the chest, as previously described (2). Briefly, HRCT scores were based on the following scheme: 0 = normal CT; 1 = mild disease (1 to 15 thickened interlobular septa per segment, 1 to 5 patches of reticulation, subpleural cysts, and pleural/parenchymal scars); 2 = moderate disease (moderate reticulation, peribronchovascular thickening, traction bronchiectasis, tracheal retraction); 3 = severe disease (areas of parenchymal consolidation, diffuse areas of peribronchovascular thickening, traction bronchiectasis, and patches of reticulation).

Pulmonary Function Testing

Measurements were made with standard equipment according to American Thoracic Society recommendations (Collins Medical, Inc., Braintree, MA; SensorMedics, Yorba Linda, CA) (30). FEV₁, FVC, total lung capacity (TLC), and diffusion capacity (DL_{CO}) were expressed as percentages of predicted values.

Fiberoptic Bronchoscopy with Bronchoalveolar Lavage

Bronchoscopy with lavage as well as isolation of alveolar macrophages and bronchoalveolar lavage (BAL) fluid were performed as described (31-33). Briefly, subjects were sedated and received 1% lidocaine for topical anesthesia. A fiberoptic bronchoscope was inserted into the proximal airways, and up to four 25- to 30-ml aliquots of sterile 0.9% normal saline were instilled into the right middle lobe medial or lateral segments and/or the lingula. The BAL fluid was collected by suction into sterile specimen containers, filtered through gauze, and centrifuged at 1,000 \times g for 10 minutes at 4°C to pellet the BAL cells. Supernatant was aliquoted, stored at -80° C, and thawed immediately before analysis. Pelleted cells were washed twice in RPMI 1640, centrifuged at 1,000 \times g for 10 minutes at 4°C, dispersed in RPMI 1640, transferred to sterile polystyrene tissue culture dishes, and incubated for 2 hours at 37°C in a humidified atmosphere containing 5% CO₂. Adherent cells (about 95% alveolar macrophages) were analyzed further in the experiments described later. A blood sample was obtained after bronchoscopy, and urea concentrations in BAL fluids and plasma were measured to calculate dilution of BAL fluids (Sigma Diagnostics, St. Louis, MO). Differential cell counts of BAL cells were performed on Cytospin slides stained with Diff-Quik, using a hemocytometer, and concentrations of BAL cells were calculated (Thermo Fisher Scientific, Inc., Waltham, MA).

Northern Blot Analysis and Reverse Transcriptase-Polymerase Chain Reaction

Total RNA was isolated from alveolar macrophages of a subject with HPS-1 and a healthy research volunteer as well as from normal fibroblasts. Northern blots containing 20 µg of mRNA were analyzed with a ³²P-labeled human HPS1 cDNA as previously described (16). Reverse transcriptase-polymerase chain reaction (RT-PCR) amplification was performed with total RNA isolated from alveolar macrophages and peripheral blood mononuclear cells (PBMCs) from three different healthy research volunteers as well as from PBMCs from three different subjects with HPS-1. cDNA was generated with the SuperScript first strand synthesis system for RT-PCR in accordance with the manufacturer's instructions (Invitrogen, Carlsbad, CA). A 414-bp fragment of HPS1 was amplified in a 25-µl reaction mixture containing cDNA and primers (5'-CCTGTTTGGAGAATGCCTGTTC-3' and 5'-CCAAGCTGCTGGCATTCTACTC-3'), using standard PCR methods. A 353-bp RT-PCR product of *β-actin* was generated in accordance with the manufacturer's instructions (Invitrogen). GeneRuler 100-bp DNA ladder (Fermentas, Inc., Glen Burnie, MD) and PCR products were subjected to electrophoresis on a 1.5% agarose gel and stained with ethidium bromide.

Quantification of Cytokines and Growth Factors

Concentrations of monocyte chemoattractant protein (MCP)-1, macrophage inflammatory protein (MIP)-1a, granulocyte-macrophage colony-stimulating factor (GM-CSF), macrophage colony-stimulating factor (M-CSF), epidermal growth factor (EGF), granulocyte colonystimulating factor (G-CSF), regulated upon activation, normal T cell expressed and secreted (RANTES), transforming growth factor (TGF)-B₁, and platelet-derived growth factor (PDGF)-AB were measured in BAL fluid specimens that were concentrated 10-fold by lyophilization. ELISA kits were used in accordance with manufacturers' instructions (Amersham Biosciences, Piscataway, NJ; Biosource International, Camarillo, CA; R&D Systems, Minneapolis, MN). Concentrations of IL-4, IL-6, IL-10, IL-12p40, MIP-4, IFN-y, tumor necrosis factor (TNF)-a, PDGF-AA, and PDGF-BB were measured by multiplex ELISA (SearchLight; Pierce Biotechnology, Inc., Woburn, MA). Final epithelial lining fluid (ELF) concentrations were determined by the urea method (31-33).

After bronchoscopy, alveolar macrophages (AMs, 1×10^{6} /ml) were cultured in cell culture medium (RPMI, $1 \times$ glutamine, $1 \times$ penicillin–streptomycin). Plates were incubated in 5% CO₂ at 37°C. After 2 hours, AMs were washed twice with fresh medium, and then cultured with medium with or without pirfenidone. Concentrations of pirfenidone were 200, 600, 1,200, and 2,400 µg/ml. Conditioned medium was isolated after 24 hours, and immediately stored in a liquid nitrogen cryotank. Samples were thawed once, and concentrations of MCP-1, MIP-1 α , RANTES, M-CSF, MIP-4, GM-CSF, IL-12p40, and IFN- γ were measured by multiplex ELISA (SearchLight; Pierce Biotechnology, Inc.).

Statistical Analysis

The Student *t* test was used to determine significant differences (P < 0.05) between the mean values of age, pulmonary function test measurements, BAL cell concentrations, and BAL analyte concentrations. Data are reported as mean values \pm standard error of the mean.

RESULTS

Subject Characteristics

Forty-one subjects with HPS-1 and 85 healthy research volunteers participated in this study. PCR amplification of genomic DNA demonstrated that 35 of 41 subjects with HPS-1 were homozygous for a 16-base pair duplication in exon 15 of *HPS1* [c.1472_1487dup16; p.H497QfsX90 (ex 15)]. Six of 41 subjects had 5 different truncating *HPS1* mutations, including [c.1744-2A > C (IVS16)], [c.1375delA; c.1388C > A; p.S459VfsX16 (ex 14)], [c.97_100delTCAG; p.S33RfsX18 (ex 3)], [c.972dupC; p.M325HfsX128 (ex 11)], and [c.1189delC; p.Q397SfsX2 (ex 13)]. In 34 of 41 subjects with HPS-1, pulmonary fibrosis was diagnosed by characteristic findings on HRCT scan of the chest, including ground glass opacification in 20 (59%) (Figure 1). Seven subjects with HPS-1 did not have interstitial infiltrates on their HRCT scan images.

The mean age of subjects with HPS-1 pulmonary fibrosis, 38.6 ± 1.5 years, did not differ significantly from that of subjects with HPS-1 without pulmonary fibrosis, 31.0 ± 4.2 years (P = 0.054) or our healthy research volunteers, 38.8 ± 1.4 years (P = 0.947) (Table 1). The most common symptoms of lung disease included dyspnea on exertion and cough. Six (86%) of 7 subjects with HPS-1 and 7 (21%) of 34 with HPS-1 pulmonary fibrosis were asymptomatic. One subject with HPS-1 without pulmonary

fibrosis experienced dyspnea on exertion at age 43 years, and 27 (79%) of 34 subjects with HPS-1 with pulmonary fibrosis developed symptoms at approximately 34.9 ± 1.77 years of age. A few subjects in this cohort were being treated with inhaled steroids with or without a β -agonist, including one subject with HPS-1 and four subjects with HPS-1 pulmonary fibrosis. Eleven (32%) of 34 subjects with HPS-1 pulmonary fibrosis, 5 (71%) of 7 with HPS-1, and 55 (65%) of 85 healthy research volunteers were male. No subjects with HPS-1 with or without pulmonary fibrosis and no healthy research volunteers were current smokers, and 3 (8.8%) of 34 subjects with HPS-1 pulmonary fibrosis were former smokers who had stopped smoking for at least 2.5 years and had a mean history of smoking for 17.7 \pm 1.20 pack-years. Many subjects with HPS-1 were unemployed or were receiving disability benefits, largely due to their visual impairment, including 4 of 7 with HPS-1 and 13 of 34 with HPS-1 pulmonary fibrosis. Among the subjects who were employed, one individual with HPS-1 pulmonary fibrosis was exposed to wood dust in the construction industry.

Analysis of pulmonary function tests revealed that measurements of FEV₁, FVC, TLC, and D_{LCO} were significantly lower in subjects with HPS-1 with and without pulmonary fibrosis than in healthy research volunteers. Although no statistically significant differences were found between subjects with HPS-1 with and without pulmonary fibrosis, pulmonary function tests indicated that subjects with HPS-1 pulmonary fibrosis had mild restriction and a mild defect in gas exchange, and subjects with HPS-1 without pulmonary fibrosis had normal lung function.

High Numbers of Total Alveolar Cells and Alveolar Macrophages in HPS-1 with and without Pulmonary Fibrosis

On fiberoptic bronchoscopy, subjects with HPS-1 without pulmonary fibrosis had significantly higher concentrations of total cells in BAL fluid (4.47 [± 0.52] × 10⁵ cells/ml) and ELF (25.2 [± 3.29] × 10⁶ cells/ml) compared with healthy research volunteers (1.57 [± 0.08] × 10⁵ and 14.1 [± 0.77] × 10⁶ cells/ml, respectively) (P = 0.002 and P < 0.001, respectively) (Figures 2A and 2B). To a lesser degree, subjects with HPS-1 pulmonary fibrosis had significantly higher concentrations of total cells in BAL fluid (3.13 [± 0.34] × 10⁵ cells/ml) and ELF (17.5 [± 1.12] × 10⁶ cells/ml) compared with healthy research volunteers



Figure 1. (*A*–*D*) High-resolution computed tomography (HRCT) scan findings in subjects with Hermansky-Pudlak syndrome type 1 (HPS-1) pulmonary fibrosis. Representative HRCT scan images from four individual subjects with HPS-1 pulmonary fibrosis are shown. Reticulation (*arrow*-*heads*), honeycombing (*arrows*), and ground glass opacification (*open arrows*) are found in subjects with HPS-1 pulmonary fibrosis.

TABLE 1. SUBJECT CHARACTERISTICS

	HPS-1 (<i>n</i> = 7)	HPS-1 PF $(n = 34)$	NV (n = 85)	P Value		
				HPS-1 vs. HPS-1 PF	HPS-1 vs. NV	HPS-1 PF vs. NV
Age, yr	31.0 ± 4.2	38.6 ± 1.5	38.8 ± 1.4	0.054	0.134	0.947
Sex, M/F	5 / 2	11 / 23	55 / 30			
Current smokers	0	0	0			
Former smokers	0	3	6			
Dyspnea or cough	1	27	0			
FEV ₁ , % of predicted	87.5 ± 4.3	85.6 ± 3.4	106.7 ± 1.9	0.803	0.004	< 0.001
FVC, % of predicted	82.1 ± 5.1	79.5 ± 3.4	101.5 ± 1.6	0.738	< 0.001	< 0.001
TLC, % of predicted	$84.9~\pm~5.6$	78.8 ± 3.3	99.5 ± 1.5	0.478	0.006	< 0.001
DLCO, % of predicted	$85.8~\pm~7.7$	75.0 ± 3.3	102.8 ± 1.8	0.186	0.009	<0.001

Definition of abbreviations: DL_{CO} = diffusion capacity for carbon monoxide; F = female; HPS-1 = Hermansky-Pudlak syndrome type 1; HPS-1 PF = Hermansky-Pudlak syndrome type 1 with pulmonary fibrosis; M = male; NV = healthy research volunteer; TLC = total lung capacity.

(P < 0.001 and P = 0.018, respectively). In addition, differential cell counts revealed that concentrations of alveolar macrophages, but not lymphocytes or polymorphonuclear leukocytes, in ELF were significantly higher in subjects with HPS-1 without and with pulmonary fibrosis (23.4 [± 2.87] and 15.5 [± 0.96] $\times 10^{6}$ cells/ml, respectively) than in healthy research volunteers (12.3 $[\pm 0.70] \times 10^6$ cells/ml) (P < 0.001 and P = 0.013, respectively) (Figure 2C). Compared with subjects with HPS-1 without pulmonary fibrosis, subjects with HPS-1 pulmonary fibrosis had significantly lower concentrations of total cells and alveolar macrophages in ELF (P = 0.010 and P = 0.003, respectively). In contrast, total white blood cells and percentage of monocytes in peripheral blood did not differ significantly in subjects with HPS-1 with and without pulmonary fibrosis (Figure 2D), and there were no significant differences in percentages of peripheral blood polymorphonuclear leukocytes, lymphocytes, or eosinophils between these two groups (data not shown).

High Alveolar Concentrations of Cytokines and Chemokines in HPS-1 with and without Pulmonary Fibrosis

In subjects with HPS-1 pulmonary fibrosis, concentrations of MCP-1, MIP-1 α , GM-CSF, and M-CSF in ELF were significantly higher (9,116 ± 1,122, 4,641 ± 498.0, 673 ± 77.0, and 19,190 ± 3,629 pg/ml, respectively) compared with the concentrations in healthy research volunteers (261 ± 29.0, 839 ± 108, 53 ± 7.0, and 5,318 ± 795.0 pg/ml, respectively) (P < 0.001) (Figure 3). Concentrations of MCP-1, MIP-1 α , and GM-CSF

were also significantly higher in subjects with HPS-1 without pulmonary fibrosis (3,668 ± 1,519, 2,663 ± 843.3, and 316.7 ± 104.9 pg/ml, respectively) than in healthy research volunteers (P < 0.001). Furthermore, concentrations of MCP-1 α , GM-CSF, and M-CSF were significantly higher in subjects with HPS-1 pulmonary fibrosis compared with those in subjects with HPS-1 without pulmonary fibrosis (P = 0.041, P = 0.049, and P <0.001, respectively). No significant differences in concentrations of IL-4, IL-6, IL-10, IL-12p40, MIP-4, IFN- γ , TNF- α , EGF, G-CSF, RANTES, TGF- β_1 , PDGF-AA, PDGF-AB, or PDGF-BB in ELF were found between subjects with HPS-1 with and without pulmonary fibrosis and healthy research volunteers (data not shown).

To determine whether alveolar concentrations of MCP-1, MIP-1 α , GM-CSF, and/or M-CSF correlate with severity of HPS-1 pulmonary fibrosis, data from subjects with HPS-1 pulmonary fibrosis were segregated into those with mild or moderate/severe disease and compared with subjects with HPS-1 without pulmonary fibrosis. We found that concentrations of MCP-1 correlated with HPS-1 pulmonary fibrosis disease severity (*see* Figure E1 in the online supplement). In contrast, concentrations of MIP-1 α and GM-CSF were significantly higher in subjects with HPS-1 with mild, and not moderate/severe, pulmonary fibrosis compared with those without lung disease (P = 0.041 and P = 0.033, respectively). Although concentrations of M-CSF were significantly higher in subjects with HPS-1 with some degree of pulmonary fibrosis compared



Figure 2. Increased alveolar concentrations of total cells and alveolar macrophages (AMs) in subjects with Hermansky-Pudlak syndrome type 1 (HPS-1). (A-C) Concentrations of total cells in bronchoalveolar lavage (BAL) fluid $(\times 10^3 \text{ cells/ml})$ and alveolar epithelial lining fluid (ELF) $(\times 10^6 \text{ cells/ml})$ were significantly higher in subjects with HPS-1 (HPS) and HPS-1 pulmonary fibrosis (HPSPF) compared with healthy research volunteers (NV). In ELF, concentrations (\times 10⁶ cells/ml) of AMs were significantly higher in subjects with HPS-1 and HPSPF compared with NV subjects. Subjects with HPS-1 had significantly higher concentrations of BAL cells and AMs in ELF compared with subjects with HPSPF. (D) No significant differences were found in peripheral blood concentrations of total leukocytes (WBC) or percentage of monocytes (mono) in subjects with HPS-1 (solid columns) and those with HPSPF (open columns).



Figure 3. High alveolar concentrations of cytokines and chemokines in subjects with Hermansky-Pudlak syndrome type 1 (HPS-1) with and without pulmonary fibrosis. Concentrations (pg/ml) of monocyte chemoattractant protein (MCP)-1, macrophage inflammatory protein-1 α (MIP-1a), granulocyte-macrophage colony-stimulating factor (GM-CSF), and macrophage colony-stimulating factor (M-CSF) were significantly higher in alveolar epithelial lining fluid (ELF) from subjects with HPS-1 pulmonary fibrosis (HPSPF) compared with healthy research volunteers (NV), and concentrations of MCP-1, MIP-1 α , and GM-CSF were significantly higher from subjects with HPS-1 (HPS) compared with NV subjects. Subjects with HPSPF had significantly higher concentrations of MCP-1, GM-CSF, and M-CSF compared with subjects with HPS-1.

with subjects with HPS-1 without lung disease (P = 0.002 and P = 0.035, respectively), levels of M-CSF did not correlate with disease severity.

Cellular Expression of HPS1 in Healthy Research Volunteers, But Not in Subjects with HPS-1

Northern blot analysis demonstrated that alveolar macrophages from subjects with HPS-1 do not express *HPS1* mRNA, whereas the *HPS1* transcript is expressed in both alveolar macrophages and fibroblasts from healthy research volunteers (Figure 4A). In agreement with these data, amplification of cDNA by RT-PCR demonstrated that an *HPS1* PCR product is amplified in alveolar macrophages and PBMCs from healthy research volunteers, but not in PBMCs from subjects with HPS-1 (Figure 4B).

Pirfenidone-induced Suppression of Cytokine Secretion by Alveolar Macrophages from Subjects with HPS-1

To determine whether absence of *HPS1* transcript in alveolar macrophages is associated with cellular dysregulation, we cultured these cells and measured the concentrations of analytes secreted *in vitro* (Figure 5A). Concentrations of MCP-1, MIP-1 α , and RANTES were significantly higher in conditioned medium from alveolar macrophages cultured for 24 hours from subjects with HPS-1 pulmonary fibrosis (548.1 ± 114.3, 787.3 ± 93.2, and 203.4 ± 69.6 pg/ml, respectively) compared with healthy research volunteers (96.5 ± 38.2, 318.7 ± 111.4, and 3.62 ± 2.17 pg/ml, respectively) (P = 0.001, P = 0.014, and P = 0.011, respectively).

To determine the effect of the antiinflammatory drug pirfenidone on alveolar macrophage cytokine and chemokine secretion, alveolar macrophages from subjects with HPS-1 pulmonary fibrosis were cultured with increasing concentrations of pirfenidone for 24 hours. The alveolar macrophages incubated with pirfenidone secreted lower amounts of MIP-1 α , MCP-1, RANTES, M-CSF, MIP-4, and IFN- γ in conditioned medium in a dose-dependent manner, whereas concentrations of GM-CSF and IL-12p40 remained essentially unchanged (Figure 5B).

DISCUSSION

Alveolar inflammation may be associated with high concentrations of lung cytokines and chemokines. Indeed, we found this in patients with HPS, who also had higher concentrations of total BAL cells and alveolar macrophages in ELF. We also found that pirfenidone suppresses chemokine secretion by cultured HPS-1 alveolar macrophages in a dose-dependent manner. Finally, HPS-1 alveolar macrophages and PBMCs do not express *HPS1* mRNA, whereas normal cells do.

Several attributes of HPS-1 make this a uniquely suitable disease in which to study pulmonary fibrosis. Individuals with HPS-1 comprise a relatively homogeneous cohort, because most are homozygous for the same 16-bp duplication in the *HPS1* gene. This mutation is associated with absence of HPS1 protein and, thus, dysregulation of biogenesis of lysosome-related organelles is found in the cells of individuals with HPS-1. In addition, patients with HPS-1 are distinguishable from the general pop-



Figure 4. Cellular expression of *HPS1* in healthy research volunteers, but not in subjects with Hermansky-Pudlak syndrome type 1 (HPS-1). (*A*) Northern blot analysis revealed that alveolar macrophages (AM) and fibroblasts from healthy research volunteers (NV), and not alveolar macrophages from subjects with HPS-1, express *HPS1* mRNA. (*B*) Amplification of cDNA by reverse transcriptase-polymerase chain reaction (RT-PCR) using primers for *HPS1* demonstrated that a 414-base pair (bp) *HPS1* PCR product is expressed in peripheral blood mononuclear cells (PBMCs) and alveolar macrophages (AM) from two representative healthy volunteers (NV1 and NV2), but not in PBMCs from a representative subject with HPS-1. A 353-bp RT-PCR product was generated using primers for human β -actin.



Figure 5. Cytokine secretion by Hermansky-Pudlak syndrome type 1 (HPS-1) and normal cultured alveolar macrophages and suppression by pirfenidone in HPS-1 cells. (*A*) Concentrations (pg/ml) of monocyte chemoattractant protein (MCP)-1, macrophage inflammatory protein- 1α (MIP-1a), and regulated upon activation, normal T cell expressed and secreted (RANTES) were significantly higher in conditioned medium from alveolar macrophages cultured for 24 hours from subjects with HPS-1 pulmonary fibrosis (HPSPF) (n = 18) compared with healthy research volunteers (NV) (n = 6). (*B*) Concentrations of MIP- 1α , MCP-1, RANTES, macrophage colony-stimulating factor (M-CSF), macrophage inflammatory protein (MIP)-4, and IFN- γ (IFNg), but not granulocyte-macrophage colony-stimulating factor (GM-CSF) or IL12p40, in conditioned medium from alveolar macrophages from subjects with HPS-1 pulmonary fibrosis (n = 6) cultured for 24 hours were lower with increasing concentrations of pirfenidone.

ulation because of their oculocutaneous albinism, and there is nearly universal development of progressive pulmonary fibrosis in these individuals. To our knowledge, there are no other populations that share such characteristics, except possibly for the few people with HPS-4 (12, 34).

Despite progress in the understanding of genetic, molecular, and cellular aspects of HPS, the etiology of pulmonary fibrosis in HPS-1 remains unknown. Our findings of high concentrations of activated alveolar macrophages, cytokines, and chemokines in individuals with HPS-1 provide evidence that immune cells and proinflammatory proteins contribute to the pathogenesis of pulmonary fibrosis. We demonstrate that concentrations of MCP-1 correlate with severity of pulmonary fibrosis in HPS-1. We also found that concentrations of MIP-1a, GM-CSF, and M-CSF are high in subjects with mild HPS-1 pulmonary fibrosis, and it is possible that these proteins may contribute to the pathogenesis of early, and not advanced, HPS-1 pulmonary fibrosis. Overall, these findings are consistent with studies focusing on lung disease in murine models of HPS, which demonstrate lung inflammation and constitutive activation of alveolar macrophages (26, 28).

Although the role of inflammation in the pathogenesis of pulmonary fibrosis is controversial, several reports indicate that dysregulation of immune cells contributes to the development of lung fibrosis. Specifically, high alveolar concentrations of activated lymphocytes were found in first-degree relatives of probands with familial pulmonary fibrosis with asymptomatic lung disease, suggesting that early stages of fibrotic lung disease are associated with accumulation of abnormal immune cells in the lung (32). In addition, impairment of general transcription, associated with decreased concentrations of transcription factor II-H, has been reported in alveolar macrophages of patients with idiopathic pulmonary fibrosis (33). High concentrations of MCP-1, MIP-1a, and M-CSF were found in BAL fluid from patients with idiopathic pulmonary fibrosis, and concentrations of MCP-1 correlated negatively with pulmonary function measurements (35, 36). Furthermore, mice deficient in MCP-1 or M-CSF had decreased lung fibrosis and inflammatory cell accumulation after receiving bleomycin, and anti-MCP-1 gene therapy attenuated lung disease in a murine model of pulmonary fibrosis (26, 37).

In agreement with these reports, several publications demonstrate a key etiologic role for macrophages in extrapulmonary fibrosis. For example, a decrement of macrophages or absence of macrophage-derived galectin-3 ameliorated renal fibrosis in a murine model of progressive renal fibrosis (38). Also, selective depletion of macrophages in an animal model of hepatic inflammation significantly attenuated liver fibrosis (39). Furthermore, in vivo and in vitro studies of wound healing demonstrated that macrophage-derived PDGF contributed to expression of fibroblast osteopontin, which may be partially responsible for the development of wound fibrosis (40). Taken together, these data confirm the importance of macrophages in the pathogenesis of fibrosis. Our findings in this cohort of individuals with HPS-1 are consistent with these reports, and further substantiate the hypothesis that inflammation and macrophages contribute to the pathogenesis of pulmonary fibrosis.

Although treatment options for HPS-1 are limited, oral pirfenidone has been shown to slow the rate of lung function decline in some individuals with HPS-1 (4). Other studies have demonstrated that pirfenidone reduced the incidence of acute exacerbations and stabilized lung function in patients with idiopathic pulmonary fibrosis (41, 42). In animal models of pulmonary fibrosis, pirfenidone ameliorated lung disease (43). The mechanism of action of pirfenidone is undefined, but it inhibits TGF- β , PDGF, and TNF- α and has antioxidant effects (44-47). Our results indicate that pirfenidone reduced alveolar macrophage cytokine secretion in vitro. It is unlikely that cytotoxic effects of pirfenidone contributed to these findings, because stable concentrations of GM-CSF and IL-12p40 indicate that the alveolar macrophages remained viable at these levels of pirfenidone. Notably, a limitation of this study is that plasma concentrations of pirfenidone achieved with therapeutic doses are lower than concentrations of pirfenidone used for these experiments (4, 48). Although lung epithelial lining fluid concentrations of pirfenidone are unknown, it is likely that the concentrations of pirfenidone used for these in vitro experiments are higher than those achieved locally in the alveolar space. These data suggest, however, that down-regulation of alveolar macrophage activation is a possible mechanism contributing to the clinical response to pirfenidone in HPS-1 pulmonary fibrosis, and this supports results of animal models indicating that pirfenidone has both antiinflammatory and antifibrotic properties (43-47, 49).

Our data indicate that lung inflammation and alveolar macrophage activation in HPS-1 are associated with high lung concentrations of cytokines and chemokines, including MCP-1, MIP-1 α , and GM-CSF. Although alveolar fluid concentrations of TGF- β_1 were not significantly different in HPS-1, it is unclear whether TGF- β_1 activity is altered in this disease. Our findings also indicate that alveolar macrophages may be a potential source of cytokines and chemokines and, thus, may contribute to a self-perpetuating proinflammatory cycle in the HPS-1 lung. Because alveolar macrophages derive from the bone marrow, additional research using murine models of HPS is indicated to further explore the potential utility of bone marrow transplantation for treatment of pulmonary fibrosis in HPS-1 (50, 51).

Our findings provide a basis for future investigations into the causes and treatment of the pulmonary fibrosis of HPS. Can down-regulation of *HPS1* expression in normal alveolar macrophages recapitulate the HPS phenotype, that is, increased cytokine secretion? Can transfection of HPS-1 alveolar macrophages with normal *HPS1* reverse the abnormal cellular phenotype? Can HPS-1 alveolar macrophages be used to test new drug therapies, using reduced cytokine secretion as an outcome parameter? BAL cells and fluid hold promise in answering these questions.

Conflict of Interest Statement: None of the authors has a financial relationship with a commercial entity that has an interest in the subject of this manuscript.

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