Pulmonary Fibrosis Induced by γ-Herpesvirus in Aged Mice Is Associated With Increased Fibroblast Responsiveness to Transforming Growth Factor-β

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Young (4 month) and aged (15–18 months) mice were given intranasal saline or γ-herpesvirus-68 infection. After 21 days, aged, but not young mice, showed significant increases in collagen content and fibrosis. There were no differences in viral clearance or inflammatory cells (including fibrocytes) between infected aged and young mice. Enzyme-linked immunosorbent assays showed increased transforming growth factor-β in whole lung homogenates of infected aged mice compared with young mice. When fibroblasts from aged and young mice were infected in vitro, aged, but not young, fibroblasts upregulated alpha-smooth muscle actin and collagen I protein. Infection with virus in vivo also demonstrates increased alpha-smooth muscle actin and collagen I protein and collagen I, collagen III, and fibronectin messenger RNA in aged fibroblasts. Furthermore, evaluation revealed that aged fibroblasts at baseline have increased transforming growth factor-β receptor 1 and 2 levels compared with young fibroblasts and are resistant to apoptosis. Increased responsiveness to transforming growth factor-β was verified by increased collagen III and fibronectin messenger RNA after treatment in vitro with transforming growth factor-β.

Key Words: Lung—Fibrosis—Herpesvirus—Aging.

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BACKGROUND

Idiopathic pulmonary fibrosis (IPF) is a chronic parenchymal disorder of the lungs associated with a progressive loss of lung function (1). It is likely secondary to dysregulated repair in response to injury leading to destruction of normal lung architecture through fibroblast accumulation, myofibroblast differentiation, and the overproduction of collagen and other extracellular matrix proteins. IPF is known to be a disease of the elderly population, with incidence, prevalence, and mortality of the disease increasing with age. The highest prevalence occurs in patients older than the age of 75 years and age is an independent predictor of having usual interstitial pneumonia (the pathological correlate of IPF) on surgical lung biopsy in comparison with other interstitial pneumonias (2,3). Patients can present with stability over several years, with a steady decline in lung function, or with a sudden accelerated decline in clinical status termed an acute exacerbation (1,4). Unfortunately, mortality remains high, with median survival times of 3 years from the time of diagnosis, and there are currently no effective therapies (5).

Although studies in the past decade have advanced our understanding of the pathogenesis and clinical presentation of the disease, the exact cause of initial injury is still unknown. However, there is accumulating data that an occult viral infection may play a role, leading to chronic lung inflammation and an abnormal healing response in certain individuals. In particular, Epstein–Barr virus (EBV), a γ-herpesvirus, has been found to have the strongest association with pulmonary fibrosis (reviewed in (6)). The virus is known to infect most people, and the initial lytic phase is followed with a latent phase and the potential for reactivation (7). Human studies have found elevated levels of EBV-specific IgA and IgG antibodies in serum, elevated EBV viral capsid–specific IgA in bronchoalveolar lavage fluid, and increased presence of EBV DNA in the lungs of IPF patients when compared with controls (8–10). In addition, evidence of actively replicating EBV was found in type 2 alveolar epithelial cells (AECs) of IPF patients and infection of type 2 alveolar cells with EBV in vitro led to increased expression of transforming growth factor (TGF)-β, a potent fibrotic mediator (11,12). Another study evaluating four human herpesviruses, cytomegalovirus, EBV, human herpesvirus-7, and human herpesvirus-8, found evidence of infection in 96% of IPF patients versus 36% of controls (13). It should be noted, however, that not all studies have found an association between γ-herpesviral infection and IPF (14,15).

Investigators have used mouse models to gain further insight into the profibrotic effects of γ-herpesviruses on the lung. γHV-68, a murine γ-herpesvirus genetically similar to...
EBV and human herpesvirus-8, infects the respiratory tract with an initial lytic phase followed by latency in lung epithelial cells (16–18) and B cells (19). Infection of Balb/c mice with \( \gamma \text{HV-68} \) before intraperitoneal bleomycin caused increased lung collagen content, increased fibrosis, and higher inflammation scores (20). Preceding latent infection also promotes fibrosis to a subthreshold dose of fluorescein isothiocyanate or bleomycin (21), suggesting that virus may alter the lung environment and predispose individuals to develop fibrosis to subsequent stimuli. However, in these studies carried out in young wild-type mice, infection with \( \gamma \text{HV-68} \) alone did not cause fibrosis (21). In contrast, infection with \( \gamma \text{HV-68} \) can cause fibrosis in Th2-biased (interferon [IFN]-\( \gamma \)-receptor–deficient [IFN\( \gamma \)-R \(-/-\)] ) mice. \( \gamma \text{HV-68} \) infection in these mice causes multiorgan fibrosis in the lungs, liver, spleen, and lymph nodes (22). Similar to patients with IPF, the viral-induced fibrosis in these mice was found to have areas of increased collagen deposition and severe peripheral fibrosis characterized by fibroblastic foci and TGF-\( \beta \) elevations (23). Later studies demonstrated that viral reactivation was key to fibrosis reactivation in this Th2-biased model (24).

It is plausible that environmental differences in an aging lung are responsible for the increased susceptibility of elderly patients to lung fibrosis, possibly potentiated by viral infections. Senescence-accelerated prone mice have increased lung fibrosis, serum fibrocyte concentrations, and TGF-\( \beta \) production in response to bleomycin when compared with senescence-resistant mice (25). There are also aged mouse studies showing a Th2 shift in cytokine profile with increased T-cell production of interleukin (IL)-4 (26–28). In addition, increased endoplasmic reticulum stress, decreased protein chaperones causing misfolded proteins, and abnormal handling of these proteins in the elderly patients drive apoptotic responses in AECs; an observation that has been implicated to promote pulmonary fibrosis (29–31). Finally, the aging lung is susceptible to changes in the immune system, predominantly in T cells important for control of viral replication (32–34).

We hypothesized that the environment of an aged lung may be sufficiently altered for \( \gamma \)-herpesvirus infection alone to cause fibrosis in wild-type mice. We compared collagen content and histology of aged (15–18 months) and young (4 month) mice 21 days after intranasal infection with \( \gamma \text{HV-68} \) and found increased fibrosis and collagen in aged mice in response to virus. This was not due to an inability to control viral replication in aged mice but rather due to an increased TGF-\( \beta \) production and a profibrotic fibroblast phenotype in aged mice. To our knowledge, this is the first demonstration of a circumstance (aging) in which viral infection alone is sufficient to cause fibrosis in wild-type mice.

**Materials and Methods**

**Mice**

Male young C57Bl/6 mice (4 months old) and aged C57Bl/6 (15–18 months old) mice were purchased from the National Institute of Aging (Bethesda, MD). All procedures were approved by the University of Michigan Committee on the Use and Care of Animals.

**\( \gamma \text{HV-68} \) Infection**

After mice were anesthetized with a low dose of ketamine and xylazine, 5 x 10^4 to 1 x 10^5 plaque-forming units (pfu) of \( \gamma \text{HV-68} \) (American Type Culture Collection, Manassas, VA) diluted in 20 \( \mu \)L sterilized phosphate-buffered saline were administered intranasally.

**Hydroxyproline Assay**

Total lung collagen was quantified via the hydroxyproline assay as described previously (35). Briefly, lung lobes were removed from perfused mice, homogenized in phosphate-buffered saline, and hydrolyzed by the addition of hydrochloric acid (Sigma, St Louis, MO). Samples were then baked at 110°C for 12 hours and aliquots were assayed by adding chloramine T solution, followed by development with Erlich’s reagent at 65°C as previously described. Absorbance was measured at 550 nm, and the amount of hydroxyproline was determined against a standard curve generated using known concentrations of hydroxyproline standard (Sigma).

**Histology**

Hematoxylin and eosin staining was performed as described previously (36). Briefly, animals were euthanized and perfused via the heart with normal saline. Lungs were inflated with 10% neutral buffered formalin, removed, and fixed overnight in formalin before being dehydrated in 70% ethanol. Lungs were processed using standard procedures and embedded in paraffin. Sections (3–5 \( \mu \)m) were cut, mounted on slides, and stained with hematoxylin and eosin.

**Virus Plaque Assay**

Lytic virus present in two lung lobes of infected mice was measured by plaque assay as described previously by plating dilutions of lung homogenates onto 3T12 cells and measuring pfu on day 5 (17).

**Semiquantitative Real-Time Reverse Transcription–Polymerase Chain Reaction**

Semiquantitative real-time reverse transcription–polymerase chain reaction (RT-PCR) was performed with ABI Prism 7000 thermocycler (Applied Biosystems, Foster City, CA) using a previously described protocol (37). Primers and probes, listed in Table 1, were made using Primer Express software (Applied Biosystems). All samples were run in triplicate.

**Analysis of Viral Genome Loads**

DNA was prepared from cells isolated from mock-infected or \( \gamma \text{HV-68} \)-infected mice using the Qiagen DNeasy
Blood and Tissue Kit (Valencia, CA), and PCR was performed to detect the gB viral coding sequence as previously described (38). Values were compared with a standard curve consisting of gB plasmid DNA diluted at known copy numbers. Reported values were normalized to 100 ng of input DNA for each reaction and represent the copy number in mock-infected samples. For gB DNA analysis, the forward primer was 5′GGCCCAAATTCAATTTGCCCT; the reverse primer, 5′CCCTGGGACAACCTCCTCAAGC; and the probe, 5′-(FAM)-ACAAGCTGAC-6-(FAM)-ACAAGCTGAC-

**Table 1. Primers and Probes Used for Analysis**

<table>
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<tr>
<th>Gene</th>
<th>Oligo</th>
<th>Sequences</th>
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<tr>
<td>Probe</td>
<td>CATGGGCAAGTGTTCATCTTAGCC</td>
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**Enzyme-Linked Immunosorbent Assay**

Cytokines and chemokines were measured in whole lung homogenate supernatants using DuoSet ELISA Development System kits (R&D Systems, Minneapolis, MN) following the manufacturer’s instructions. For analysis of TGF-β1, supernatants were first acidified to allow for measurements of total TGF-β1.

**Collagenase Digests**

Lungs were enzymatically digested using collagenase and DNase as described previously (36). Briefly, lungs were excised, minced, and enzymatically digested using digestion buffer (RPMI 1640, 5% fetal calf serum, antibiotics, 1 mg/mL collagenase [Roche, Chicago, IL], and 30 μg/mL DNase (Sigma), followed by passage through the bore of a 10-mL syringe. Contaminating erythrocytes were removed from the total cell suspension, which was then dispersed by passing through a 5-mL syringe before being filtered through a Nytex filter (Tetko, Kansas City, MO). An equal volume of 40% Percoll (Sigma–Aldrich) was added, and cells were centrifuged at 3,000 rpm (2095 × g) for 30 minutes without brake. The cell pellets were resuspended in complete medium, and leukocytes were counted on a hemocytometer in the presence of trypan blue. Recovered leukocytes were analyzed by flow cytometry.

**Flow Cytometry**

All antibodies were purchased from BD Biosciences (San Jose, CA) unless otherwise noted. Leukocytes from collagenase digests were incubated 15 minutes on ice with Fc Block before surface staining with anti-mouse CD45-PerCP-Cy5.5, followed by fixation–permeabilization with BD PharMingen Cytofix/Cytoperm kit according to the manufacturer’s instructions. Cells were then stained with the following antibodies: rat anti-mouse CD4–PE, rat anti-mouse CD8–FITC, anti-mouse CD19–APC, mouse anti-mouse NK-1.1–APC, anti-mouse CD25–PE, anti-mouse GR-1–PE, mouse anti-mouse TCR-β chain–FITC, and anti-mouse F4/80–APC (eBioscience, San Diego, CA). Cells were analyzed on a FACSCalibur (BD Biosciences) and FlowJo software (Tree Star Inc., Ashland, OR).

**AEC Isolation**

Type II AECs were isolated using dispase and DNase digestion of lungs as previously described (39). After the solution was passed successively through 100-, 40-, and 25-μm nylon mesh filters, bone marrow–derived cells were removed via anti-CD32 and anti-CD45 magnetic depletion. Mesenchymal cells were removed by overnight adherence in a Petri dish. The nonadherent cells, after this initial plating, were plated in 12-well plates coated with fibronectin. Resultant purity was approximately 94% as determined previously by intermediate filament staining. Cells were plated at 1 × 10⁶ cells per well and cultured in serum-free media for 48 hours before collection of cell supernatants.

**Alveolar Macrophage Isolation**

Alveolar macrophages (AMs) were extracted from lungs by bronchoalveolar lavage as previously described (40). Briefly, these cells were collected in lavage fluid consisting of complete medium and 5 mM ethylenediaminetetraacetic acid. These cells were enumerated by counting on a hemocytometer before plating. Cells were allowed to adhere to tissue culture plates for 1 hour and then were washed to remove nonadherent cells, resulting in an approximately 95% AM culture. The cells were then cultured in serum-free media at 3 × 10⁶ cells per well in a 96-well plate for 24 hours before collection of cell supernatants.

**Mesenchymal Cell Isolation**

Whole lung was minced and cultured for 14 days to enrich for mesenchymal cells in complete media. (See
below for the recipe for Dulbecco’s modified eagle medium.

If mice were infected in vivo prior to cell isolation, complete media were supplemented with 10 μM cidofovir (Gilead Sciences Inc., Foster City, CA) to prevent viral reactivation and cell lysis. Uninfected mesenchymal cells were cultured in serum-free media on six-well plates at 5 × 10^5 cells per well for 48 hours with or without additional treatment (2 ng/mL TGF-β or 0.1 pfu per cell γHV-68) before isolating RNA or protein lysates.

**Western Blots**

Western blots were performed on fibroblasts isolated from uninfected young and aged mice as previously described (36). Antibodies used to detect collagen I were rabbit polyclonal sera from Rockland (Gilbertsville, PA), mouse monoclonal antibodies used to detect alpha-smooth muscle actin were from Sigma, glyceraldehyde 3-phosphate dehydrogenase was identified with rabbit antibodies from Cell Signaling Technology (Danvers, MA), and β-actin was identified with mouse monoclonal antibodies from Sigma. Secondary reagents were goat anti-rabbit IgG-HRP and goat anti-mouse IgG-HRP from Pierce (Rockford, IL). Blots were scanned and densitometry was performed using the Image J program downloadable from http://imagej.nih.gov/ij/download.html.

**Apoptosis Assay**

Primary lung fibroblasts isolated from naive mice as above were used for experiments between passage 4 and 6. For assessment of apoptosis, cells were seeded into 35-mm dishes in complete media. When the cells reached 80% confluence, the media were changed to serum-free media. Apoptosis was assessed 40 hours later using the Caspase-3 Colorimetric Assay Kit (Assay Designs, Ann Arbor, MI) according to the manufacturer’s protocol.

**Reagents Used**

Complete media were Dulbecco’s modified eagle medium (Lonza, Walkersville, MD) with 10% fetal bovine serum (Fisher, Pittsburgh, PA), 1% penicillin–streptomycin (Gibco/Invitrogen, Carlsbad, CA), 1% L-glutamine (Fisher), and 0.1% amphotericin B (Lonza). Serum-free media were Dulbecco’s modified eagle medium with 1% bovine serum albumin (Sigma), 1% penicillin–streptomycin, 1% L-glutamine, and 0.1% amphotericin B.

**Statistical Analyses**

Statistical significance was measured by analysis of variance (three or more groups) or unpaired Student’s t-test (two groups) using GraphPad Prism 5 software (San Diego, CA); data represent mean ± standard error of the mean; p < .05 was considered significant.

**Results**

**Latent γ-Herpesvirus Infection in Aged Mice Is Associated With Increased Fibrosis After 21 days**

Previously, we have shown that γ-herpesvirus achieves latency in mice after 14 days of infection and that γHV-68...
infection alone in young mice does not cause fibrosis after
21 days (21). To determine whether γ-herpesvirus infection
induced pulmonary fibrosis in aged mice, we infected five
young mice and five aged mice with $1 \times 10^5$ pfu γHV-68
intranasally. For controls, the same amount of saline was
introduced intranasally into three young mice and three
aged mice. Lungs were harvested 21 days after infection or
saline administration. Hydroxyproline assay was performed
on whole lung homogenates to assess for collagen content.
There was a statistically significant increase in hydroxypro-
line content between aged mice infected with virus com-
pared with aged saline controls and young mice infected
with virus (Figure 1A). There was no difference between
young mice infected with virus and young saline controls.
This was confirmed by hematoxylin and eosin staining of
lung sections showing increased destruction of normal lung
architecture in aged mice infected with virus (Figure 1A).
Aged Mice Effectively Control Lytic γHV-68 Infection
Given the possibility of age-associated decline in viral
immunity, we next evaluated whether the increased fibrotic
response in infected aged mice was due to decreased lytic viral
clearance. We performed a viral plaque assay at 7 days follow-
ing infection with lungs of infected aged and young mice and
found significantly more lytic virus present in young infected
mice (Figure 2A). This was confirmed by increased levels in
young mice of messenger RNA encoding viral gene products,
gB and M3, 7 days following infection as evaluated by real-
time RT-PCR (Figure 2B). Thus, it appears that aged mice are
able to control initial γHV-68 viral replication and appear to do
so more effectively than younger mice.

Aged Mice Do Not Harbor Increased Latent γHV-68
We also measured total viral gB DNA in whole lung
homogenates at 21 days following infection by real-time
PCR as a measure of latent viral load and found a trend
an inability of aged mice to effectively control
latent viral infection.

Increased Production of TGF-β but Not Type 2 Cytokines
in Infected Aged Lungs
We next wanted to determine if inflammatory and profi-
brotic cytokines were elevated in aged mice infected with
γHV-68. We performed enzyme-linked immunosorbent
assays on whole lung homogenates taken from aged and
young mice 21 days after infection. In initial studies, we
found no significant differences in cytokine production
between young and aged lungs at baseline (data not shown).
Yet, after infection with γHV-68, we found increased amounts of TGF-β production at day 21 in aged
mice (Figure 4A). However, there was no difference in
type 2 cytokines (IL-4, IL-13; Figure 4B and C), type 1
inflammatory cytokines (IFNγ, IL-12), or tumor necrosis

Figure 2. Aged mice are able to effectively clear lytic virus 7 days after infection. Aged mice and young mice (n = 5 per group) were intranasally infected with $5 \times 10^5$ plaque-forming units of γHV-68. After 7 days, lungs were harvested. (A) Viral plaque assay on right lungs demonstrates that there is less preformed virus at day 7 following infection in infected aged mice versus infected young mice (p = .0005). (B) Real-time reverse transcription–polymerase chain reaction performed on left lungs demonstrates decreased lytic viral gene expression, both M3 (p = .03) and gB (p = .005), in aged mice vs young mice.

Figure 3. Latent viral load is not different in infected young and aged mice 21 days after infection. Aged and young mice (n = 3 per group) were infected with $1 \times 10^5$ plaque-forming units of γHV-68 intranasally. After 21 days, lungs were harvested and viral gB DNA content was measured by real-time polymerase chain reaction (p, nonsignificant).
factor-α (data not shown). Thus, although aged mice infected with γHV-68 do secrete more of the profibrotic cytokine, TGF-β, they do not show evidence of a fibrosis-prone Th2 environment.

Increased Production of TGF-β Is Mainly Due to Differences in Production by Infected Aged Fibroblasts

In order to understand which cells were responsible for the increased TGF-β production seen in infected aged lungs, we infected aged and young mice with γHV-68. After 21 days, we isolated AMs, fibroblasts, and AECs. Equal numbers of each cell type were plated from each group. After 24 or 48 hours (macrophages or fibroblasts, respectively), the cell supernatants were collected, and TGF-β enzyme-linked immunosorbent assay was performed. We found that infected aged fibroblasts produce twice the amount of TGF-β produced by infected young fibroblasts (Figure 5A). In addition, a threefold increase in TGF-β production was seen by infected aged AMs compared with infected young AMs, but note that the overall levels were significantly smaller amounts than produced by fibroblasts (Figure 5B). No significant differences were seen between infected aged and young AECs (data not shown). Therefore, fibroblasts appear to be the main cell type contributing to the increased TGF-β seen in infected aged mice, with some contribution also coming from macrophages.

Equal Numbers of Inflammatory Cells and Fibrocytes Are Recruited to Lungs of Aged and Young Mice After γHV-68 Infection

Since aging is sometimes associated with increased numbers of inflammatory cells (26), we hypothesized that the viral-induced fibrosis in aged mice was due to increased

Figure 4. Infection with γHV-68 induces more transforming growth factor (TGF)-β but not interleukin (IL)-4 or IL-13 production in aged mice 21 days following infection. Aged and young mice were infected with 1 × 10⁵ plaque-forming units of γHV-68 and lungs were harvested 21 days later. Enzyme-linked immunosorbent assays were run on whole lung homogenates for profibrotic and inflammatory cytokines. Significant increases were found in infected aged lungs compared with young lungs in (A) TGF-β production (p = .01, n = at least 11 per group pooled from three experiments). However, there was no difference between groups in (B) IL-4 production (p = .14, n = at least 5 per group) and (C) IL-13 production (p = .06, n = at least 5 per group).

Figure 5. Increased transforming growth factor (TGF)-β production in aged fibroblasts and macrophages infected with γHV-68 in vivo compared with young infected fibroblasts at 21 days following infection. Aged and young mice were infected with 1 × 10⁵ plaque-forming units of γHV-68 and euthanized 21 days later. Fibroblasts were isolated and 5 × 10⁵ cells were cultured in six-well plates for 48 hours in serum-free media. TGF-β enzyme-linked immunosorbent assay (ELISA) was performed on cell supernatants. (A) Fibroblast TGF-β production (n = 4 in each group, p < .0001). Alveolar macrophages were isolated by performing bronchoalveolar lavages on mice and allowing the cells to adhere to plastic after 1 hour. A total of 3 × 10⁴ cells were cultured in 96-well plates for 24 hours in serum-free media. TGF-β ELISA was performed on cell supernatants. (B) Macrophage TGF-β production (n = at least four per group, p = .004).
We performed a collagenase digest on the lungs of infected mice at days 12 and 21 following γHV-68 infection and counted the total number of inflammatory cells. There was no difference in total cell count of inflammatory cells between infected aged and young mice at either time point, although the total inflammatory cell count did drop over time as expected (Figure 6A). We next performed flow cytometry on the isolated lung leukocytes in order to determine if the composition of inflammatory cells was different between young and aged mice at day 12. The population of cell types was similar between aged and young mice (Figure 6B). Similar profiles of inflammatory cells were also noted between young and old mice at day 21 following infection (data not shown). These data suggest that differences in inflammatory cells were not responsible for the increased fibrosis found in aged mice infected with γHV-68.

Given the work by Xu and colleagues (25) implicating increased fibrocytes in the serum of senescence-accelerated prone mice in a bleomycin model of pulmonary fibrosis, we also measured fibrocyte numbers by flow cytometry in the lung leukocytes isolated as above. We found no difference in the number of fibrocytes between infected aged and young mice at day 12 (Figure 6B) or day 21 following infection (data not shown).

**Fibroblasts From Aged Mice Display a Profibrotic Phenotype in Response to Infection With γHV-68**

Because there did not appear to be differences in recruitment of cells to the lungs of our aged mice after infection...
Fibroblasts From Aged Mice Have Increased Sensitivity to TGF-β With Increased Expression of TGF-β Receptor 1 and Receptor 2

TGF-β is a known promoter of fibrosis and is a potent mediator of fibroblast differentiation to myofibroblasts (41). It has also been shown that fibroblasts from the lungs of IPF patients have increased TGF-β receptor (TGF-βR) density on their surface (42). Given the increases in whole lung TGF-β found in infected aged mice, we hypothesized that the difference in fibrotic potential between aged and young fibroblasts was TGF-β mediated. We isolated fibroblasts from untreated aged and young mice and extracted their total RNA. After analysis by real-time RT-PCR, we found that aged fibroblasts have a significant three- to fourfold increase in TGF-βR1 and TGF-βR2 messenger RNA expression compared with young fibroblasts (Figure 10A). This suggested that aged fibroblasts at baseline were more responsive to TGF-β than young fibroblasts.

To confirm increased sensitivity to TGF-β, we added 2 ng/mL of TGF-β to aged and young fibroblasts in vitro. Similar to results seen after viral infection in vivo, we found that aged fibroblasts had significantly increased messenger RNA levels of fibronectin and collagen III (Figure 10B). In addition, there was a trend toward increased collagen I production by aged fibroblasts (for baseline values between young and aged fibroblasts, see Figure 9A). Thus, aged fibroblasts are more likely to be prone to fibrosis due to their increased sensitivity to TGF-β. It is likely a combination of their increased responsiveness to TGF-β and the increased production of TGF-β found in aged lungs after viral infection that causes γHV-68-induced pulmonary fibrosis in aged but not in young mice.

Fibroblasts From Aged Mice Are Resistant to Apoptosis

To determine whether lung fibroblasts from aged mice showed altered susceptibility to apoptosis, fibroblasts from aged and young mice were seeded and grown to 80% confluence in complete media (which provides an endogenous source of TGF-β). Next fibroblasts were serum starved to induce apoptosis for 40 hours and caspase 3 activity was measured. Figure 11 demonstrates that young fibroblasts are more susceptible to apoptosis (264 ± 18.4 units/mL caspase 3) comparing uninfected aged and young fibroblast RNA production showed no significant increases in collagen I, collagen III, or fibronectin expression in aged mice (Figure 9A). After 21 days of infection, we found that collagen I, collagen III, and fibronectin messenger RNA were significantly increased in aged fibroblasts compared with young fibroblasts (Figure 9B). Thus, there are clear functional differences between aged and young fibroblasts following viral infection, with aged infected fibroblasts showing a predominantly profibrotic, myofibroblast phenotype.
than similarly cultured aged fibroblasts (193 ± 21 units/mL caspase 3, n = 4, p < .04). Thus, in total, aged fibroblasts are both more responsive to TGF-β production induced by viral infection and more resistant to apoptosis.

**Discussion**

To our knowledge, this is the first report of γHV-68 inducing pulmonary fibrosis in the absence of additional fibrotic stimuli and without genetic alterations in the lung environment. Contrary to our initial expectations, increased susceptibility to viral-induced fibrosis in the aged lung is not due to alterations in Th1 or Th2 cytokines, recruitment of inflammatory cells, or recruitment of fibrocytes. Furthermore, susceptibility in aged mice is not associated with an inability to control γHV-68 replication or latency. Rather, it appears that aged resident fibroblasts are skewed toward a profibrotic phenotype with increased responsiveness to TGF-β. This is associated with increased alpha-smooth muscle actin production over baseline levels and increased production of extracellular matrix secondary to the increased levels of TGF-β produced by infected aged fibroblasts. Additionally, aged fibroblasts are resistant to apoptosis.

Previous work by Mora and colleagues (24) in IFNγ receptor knockout mice with a Th2 cytokine bias showed that viral replication was key in that mouse model for γHV-68 to produce pulmonary fibrosis. Our results demonstrate that aged mice develop a more robust fibrotic response despite improved lytic control of virus replication when compared with young mice. We have seen a similar disconnect between viral replication and fibrotic response in mice without TLR9 signaling that have been infected with γHV-68 (43). TLR9 knockout and wild-type mice were infected with γHV-68 14 days after treatment with bleomycin and harvested 7 days later. Lungs from TLR9 knockout mice showed increased collagen expression compared with wild-type mice but no significant difference in viral replication. Thus, the degree of viral replication does not necessarily correlate with subsequent pulmonary fibrosis severity. It is also interesting that infection of fibroblasts from young mice actually decreases collagen synthesis, whereas infection of fibroblasts from aged mice does not (Figure 7). It may be the case that increased viral replication in young mice may in fact divert cellular machinery away from host proteins or may result in lysis of infected cells.

We have also seen that viral reactivation is not necessary for latent γHV-68 to augment fibrosis in response to subsequent stimuli such as bleomycin or fluorescein isothiocyanate (21). Although our experiments did not specifically address reactivation, the fact that the viral DNA load is not increased in aged mice suggests that significant viral reactivation is likely not a feature of γHV-68-induced lung fibrosis in aged mice. Our data corroborate studies recently

![Figure 8. γHV-68 infection in vivo also upregulates alpha-smooth muscle actin (α-SMA) and collagen I protein production in aged fibroblasts. Aged and young mice were infected with 1 × 10⁶ plaque-forming units of γHV-68 or equal amount of saline for controls. Twenty-one days after infection, lungs were harvested and fibroblasts were isolated (n = 2 per group). Fibroblasts were plated at 5 × 10⁵ cells in six-well plates and were cultured for 48 hours. Protein lysates were made and protein concentration was evaluated by Western blot. (A) Western blot showing increased α-SMA in aged infected fibroblasts compared with aged fibroblasts with saline controls, young infected fibroblasts and young fibroblasts with saline controls. (B) Western blot showing increased collagen I production by infected aged fibroblasts compared with other groups. (C) Densitometry showing differences in α-SMA between the four groups. (D) Densitometry showing differences in collagen I between the four groups.](image-url)
published by Yager and colleagues (44) showing no evidence of decreased viral immunity, increased viral reactivation, or increased viral latent load in mice infected with γHV-68 as they age. Finally, unlike the IFNγ receptor knockout mouse model, aged mice demonstrate a fibrotic response to viral infection in the absence of a Th2 cytokine bias. Because type 1 cytokines are largely responsible for control of viral replication, it is presumed that the Th2 cytokine bias allows viral reactivation to occur and this repetitive injury is necessary to cause virus-induced fibrosis in young mice (24). However, this Th2 bias does not appear necessary in aged mice for fibrosis to occur.

We were surprised that our aged mice did not show evidence of increased fibrocyte accumulation in their lungs in response to γHV-68 infection. Xu and colleagues (25) showed that senescence-accelerated prone mice, phenotypic equivalents to aged mice, developed worse bleomycin-induced fibrosis than did senescence-accelerated resistant mice. They also found increased TGF-β production and increased serum fibrocytes in their bleomycin-treated senescence-accelerated prone mice (25). Additionally, we have previously shown that increased numbers of fibrocytes accumulate when viral infection occurs prior to or after fibrotic stimulus (17,21). Given these prior studies, we anticipated that increased fibrosis in aged mice would be associated with increased fibrocyte accumulation. However, this does not appear to be the case in our model; both aged and young mice accumulated similar numbers of fibrocytes in response to viral infection. Given our studies in the mesenchymal cells, however, it is likely that the fibrocytes that accumulate in aged mice are more responsive to TGF-β and thus potentially skewed to a profibrotic phenotype.

Although the inflammatory cell populations appeared to be similar between the infected young and aged mice, we did not evaluate for specific differences such as the presence of alternatively activated macrophages, which have been shown to be recruited to sites of active viral-induced fibrosis in Th2-biased mice and to express fibrotic mediators (45). Because our aged mice showed no increase in IL-4 or IL-13 following infection, we think it unlikely that alternatively
activated macrophages are preferentially accumulated. It remains a formal possibility however that the increased levels of TGF-β that accumulate following infection in aged mice may serve to promote alternatively activated macrophage differentiation.

Previous work by our laboratory has shown that multiple cell types can be latently infected with γHV-68 and produce TGF-β in response to infection, but mesenchymal cells appear to produce far more TGF-β than any other cell type (21,46). Our results in aged mice display a similar trend demonstrating that infected fibroblasts in aged mice are the predominant source of TGF-β. Although AECs are well known to be a reservoir for latent γHV-68 infection (21,47), we did not find evidence of increased TGF-β release by infected aged AECs compared with infected young AECs. One possibility is that the activation of TGF-β by its main integrin, αvβ6, requires direct cell-to-cell contact, and thus, active TGF-β may be rapidly bound by epithelial cells and unavailable to be measured in the supernatants (48). If integrin expression is altered in aged mice, this could result in increased cell-associated activation of TGF-β in aged AECs that may have been missed in our studies of supernatants. Additionally, AMs are a potential reservoir for γHV-68 (46), and although we did find increases in secretion of TGF-β by infected macrophages from aged mice, the total levels were much lower than those noted in fibroblasts.

Our results demonstrate that aged fibroblasts have increased levels of TGF-βR1 and TGF-βR2 making them functionally more susceptible to TGF-β-induced fibrotic responses. It is interesting that this phenotype recapitulates the observation that TGF-βR density is increased on the surface of fibroblasts taken from lung tissue of patients with IPF (42). Like our aged mice, IPF patients have an increased sensitivity to TGF-β (42). We do not yet know whether the differences in TGF-βR1 and TGF-βR2 that characterize aged fibroblasts are also present on aged epithelial cells.

Recent work by Degryse and colleagues (49) showed that mice with a selective deficiency in TGF-βR2 on AECs had improved epithelial cell survival and decreased epithelial mesenchymal transition, resulting in markedly attenuated bleomycin-induced fibrosis. Therefore, if our aged mice have increased TGF-βR2 on their epithelial cells as they do on fibroblasts, this alteration would be predicted to decrease epithelial cell survival and promote epithelial mesenchymal transition. This may be especially relevant because epithelial cells are a known reservoir for γHV-68 (21,47) and EBV infection is associated with epithelial mesenchymal transition in human epithelial cells (12,50). Further study is needed to determine if alterations in cell types other than fibroblasts are also responsible for the viral-induced fibrosis seen in aged mice.

Finally, our results indicate that fibroblasts in aged mice are more resistant to induction of apoptosis caused by serum deprivation. This indicates that fibroblasts within dense fibrotic areas (sites that may represent areas of serum deprivation) may live longer and thus have prolonged potential to secrete extracellular matrix. Taken together, our results demonstrate that aged mice produce increased levels of TGF-β in response to γHV-68 infection largely due to increased production by infected aged fibroblasts. These same aged fibroblasts have increased sensitivity to TGF-β signaling and reduced sensitivity to apoptosis. Based on these findings, we suggest that these alterations may predispose aged individuals who experience acute or reactivated herpesvirus infections to develop a fibrotic lung response. These observations provide mechanistic insight into why fibrosis is associated with aging (3,42).

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