

Effects of Cytomegalovirus Infection on Growth Factor Production in Endothelial Cells and Fibroblasts

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ABSTRACT

To determine whether cytomegalovirus (CMV) infection alters growth factor production from endothelial cells (EC) or fibroblasts, we infected human umbilical vein EC with CMV VHL/E, a strain of CMV with affinity for human EC, and we infected human foreskin fibroblasts with CMV AD169. CMV caused cytopathic effect and positive CMV staining by immunofluorescence within 5 d, effects not seen in cells infected with UV-irradiated CMV or in uninfected (control) cells. The supernatants from the EC were assayed for platelet-derived growth factor (PDGF)-like protein using a radioreceptor inhibition assay, and EC and fibroblasts were assayed for basic fibroblast growth factor (bFGF) by Western blot analysis. There were no significant differences in PDGF production between groups of EC: CMV-infected EC, 13.5 ± 2.6 ; UV-irradiated infected EC, 12.1 ± 3.6 ; control EC, 12.9 ± 1.7 fmol/ 10^6 EC (mean \pm SD, $n = 10$, $p = \text{NS}$). There were also no significant differences in bFGF production between CMV-infected EC, UV-irradiated infected EC, and control EC as evidenced by similar intensity of migration of bFGF as a single band at approximately 18 kD ($n = 5$).

In contrast, CMV infection of fibroblasts induced a shift in production of bFGF to higher molecular weight forms migrating at 24 and 26 kD molecular mass. α -Interferon failed to alter bFGF production. We conclude that CMV VHL/E infection of EC does not directly alter PDGF or bFGF production from EC. However, CMV infection of cultured human fibroblasts qualitatively alters bFGF by inducing a shift to higher molecular weight forms. (*Pediatr Res* 38: 1003–1008, 1995)

Abbreviations

EC, endothelial cell
CMV, cytomegalovirus
PDGF, platelet-derived growth factor
bFGF, basic fibroblast growth factor
HUVEC, human umbilical vein endothelial cell
CPE, cytopathic effect
HS, heparin Sepharose
TBST, Tris-buffered saline with Tween 20

The major limiting factor to long-term survival after both pediatric and adult human heart transplantation is accelerated graft arteriosclerosis (1, 2). Although immunologic injury secondary to rejection or infection has been widely implicated as a contributing factor to its pathogenesis, the mechanisms of this process remain unknown. Pathologically, the arteriosclerosis seen after heart transplantation is quite variable, ranging from diffuse to segmental and from atheromatous to entirely composed of myointimal proliferation (3). Vascular cells im-

plicated in this process include EC, smooth muscle cells, and the so-called myofibroblasts that are internal to the internal elastic lamina (4). An association between CMV infection and an increased risk of development of graft coronary vasculopathy has been found (5–7); however, the mechanisms through which CMV may promote graft coronary vasculopathy have yet to be defined. CMV increases HLA class I antigen expression on human EC (8) and HLA class II antigen expression on rat cardiac EC (9), activates CD4⁺ T cells to induce class II antigen expression on EC (10), and increases vascular cell adhesion molecule-1 in transplanted human hearts (11). This would suggest a possible indirect method by which CMV infection causes graft coronary vasculopathy. Little is known, however, about the direct effects of CMV on growth factor expression from cells.

PDGF is a cationic polypeptide with broad mitogenic activity including stimulation of mesenchymal proliferation at sites

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of tissue injury (12). Endothelial cells synthesize and secrete a protein similar to PDGF, which can be stimulated by a variety of agents including thrombin, IL-1, and tumor necrosis factor (13–16). In human nontransplant atheroma, PDGF-like mitogen has been localized to both smooth muscle cells (17) and to macrophages within the atheromatous lesions (18). Indirect evidence for alterations in PDGF during allograft rejection comes from the finding that there is increased expression of PDGF receptors during rejection, both on vascular smooth muscle of kidney transplants (19) and in the arteries and myocardium of rat heart allografts (20).

bFGF is a mitogen for endothelial, smooth muscle, and connective tissue cells (21). bFGF is produced by vascular cells and either remains cell-associated (22) or is deposited into the subcellular matrix (23). bFGF has been found in increased amounts within the lesions of human graft coronary vasculopathy (24), suggesting that synthesis of bFGF by vascular cells may be altered during allograft rejection. However, others have found no evidence of bFGF RNA in transplanted heart biopsies, suggesting that bFGF is not increased after transplantation (25). α -Interferon is a cytokine produced *in vivo* by cells such as fibroblasts in response to viral infection (26). *In vitro*, α -interferon induces formation of endothelial cell tubules, an angiogenic-like property often ascribed to bFGF (27).

We hypothesized that CMV may either directly or indirectly (through α -interferon) cause increased production of PDGF-like protein or bFGF from endothelial cells or fibroblasts, thus providing a mechanism of altered growth factor production after CMV infection. The purpose of this study was to investigate the effects of CMV infection on PDGF and bFGF production from fibroblasts and EC.

METHODS

Cell cultures. HUVEC were cultured by methods similar to those previously described by this laboratory (28). Primary cultures through second passage cultures were used in all studies.

Swiss Albino 3T3 cells (American Type Culture Collection, Rockville, MD) were grown in 75-cm² tissue culture flasks and maintained in complete 3T3 cell medium containing 90% McCoy's 5A medium (modified with L-glutamine, without sodium bicarbonate) (Sigma Chemical Co., St. Louis, MO) and 10% FCS. Medium was changed twice weekly, and the cells were passed in subculture 1:3 upon reaching confluence. Human foreskin fibroblasts (American Type Culture Collection) were routinely passaged using trypsin and M199 with 10% FCS. Periodically (every 10–12 passages), cultures were regenerated from a stock of fibroblasts stored in liquid nitrogen.

Infection of endothelial cells and fibroblasts. CMV strain VHL/E was prepared as previously described (29). For acute infection studies of EC, HUVEC were grown to 75% confluence in fibronectin-coated 75-cm² tissue culture flasks. Cultures were inoculated with CMV VHL/E stock diluted with complete EC medium. After adsorption for 1 h, an equal amount of complete EC medium supplemented with EC growth supplement and heparin was added to each flask.

Infection of cells was allowed to continue until maximum CPE was noted, but short of cell death.

Cultured fibroblasts were infected with CMV as previously described (30). Cells were plated and grown to 80% confluence and then coincubated with CMV AD169 for 1 h. The cells were then washed and refed with complete culture medium. After this initial infection, cell cultures were incubated for 4–9 d (5.4 ± 1.8) and examined daily for CPE. Evidence of CMV infection was verified by periodic examination of cultured cells with immunohistochemical staining with a CMV culture identification test that uses MAb specific for two different human CMV antigens: an immediate-early antigen ($M_r = 72\,000$) and early antigen ($M_r = 50\,000$). Cell nuclei, which stain for the CMV nuclear antigens, display fluorescent green staining (Syva Co., Palo Alto, CA).

Sham infection was performed by incubating HUVEC or fibroblasts with CMV VHL/E or CMV AD169, respectively, which had been exposed to UV light for 60 min to inactivate the virus.

PDGF-like protein secretion from EC. EC-conditioned medium was collected from the EC monolayers and assayed for the presence of PDGF-like protein using a ¹²⁵I-PDGF (PDGF, Inc., Boston, MA) competitive radioreceptor binding assay as previously described (31). Medium used for collection and analysis of PDGF-like protein was standard EC growth medium that contained FCS at the onset of experiments and then became gradually depleted of serum by the EC during continued culture and incubation at 37°C over the ensuing 5.4 ± 1.8 d. Identical media were used in both sets of control cells (UV-irradiated virus/sham-infected EC and untreated EC), which were cultured in parallel with test plates under identical conditions. At the end of the CMV infection period, microscopic examination of the EC monolayers demonstrated viable EC without significant monolayer disruption except in the areas of CPE. Attempts at experiments without the presence of FCS in the initial EC medium resulted in a significant loss of viable cells during CMV infection.

Western blot analysis. After removal of the conditioned medium, the flasks were rinsed twice in 5 mL of ice-cold PBS with 0.05% EDTA. Cells and matrix were then taken up with scraping in three successive washings. The first washing was in 2 mL of ice-cold 2 M NaCl, 25 mM sodium acetate, pH 4, 1 mM phenylmethylsulfonyl fluoride, 0.05% sodium azide, and the second and third washings were in 5 mL of ice-cold PBS with 0.05% EDTA and 0.05% sodium azide. Cell suspensions were sonicated for 1 min on ice, pulsed at 60% with a cell disrupter (Model W-225 Heat Systems-Ultrasonics, Inc., Plainview, NY), and then centrifuged at 4°C at $13\,000 \times g$ for 30 min. Cell lysates were transferred to 12-mL centrifuge tubes (Sarstedt, Newton, NC) and either frozen at -70°C or procedures were immediately followed for preparation of a Western blot for the study of bFGF.

To recover bFGF from crude cell lysates, 40 μL of HS beads were added to each sample, incubated overnight on an end-over-end rotator at 4°C, and separated by brief centrifugation at $1000 \times g$. Using radioactively labeled recombinant bFGF, approximately 85% of bFGF was found to be retained on the HS beads under these conditions. The HS beads were then

transferred to 1.5-mL Eppendorf tubes, washed twice on ice with 1 mL of PBS/EDTA, and boiled for 10 min after the addition of 28 μ L of concentrated sample buffer consisting of 120 mM Tris, pH 6.8, 4% SDS, and 19% glycerol with 2% β -mercaptoethanol. Samples (HS beads and liquid) were loaded directly onto preprepared polyacrylamide gradient (4–20%) gels (Novex, Encinitas, CA) and electrophoresed using low molecular weight markers (Amersham Corp., Arlington Heights, IL) as standards. Proteins were then transferred to nitrocellulose for 2 h at 90 V constant voltage, washed with 20 mL of PBS, and fixed with 25 mL of PBS, 0.5% (vol/vol) glutaraldehyde for 15 min. Fixation was stopped by incubation with 25 mL of 50 mM glycine/PBS for 10 min. The nitrocellulose was next washed with 25 mL of Tris-buffered saline with TBST, blocked by overnight incubation in 5% Carnation instant nonfat dried milk, incubated with 15 mL of 100 ng/mL DE6 anti-bFGF IgG1 murine MAb (a generous gift from the DuPont Corporation, Wilmington, DE) (32) for 2 h, and incubated with 25 mL of 600 ng/mL biotinylated rabbit anti-mouse FAB2 antibody fragments for 40 min. The nitrocellulose was again washed three times with 25 mL of TBST, incubated for 25 min with 25 mL of Extravidin (Sigma Chemical Co.) diluted 1:5000 in TBST, washed an additional three times in TBST, and then washed once again with alkaline phosphatase buffer (containing 5 mM MgCl₂). Finally, the blot was developed with alkaline phosphatase substrate (nitro blue tetrazolium/5-bromo-4-chloro-3-indoyl phosphate). All incubations were performed at room temperature. Equal protein loading of lanes was verified initially with Lowry determinations and then by Coomassie Brilliant Blue staining of residual proteins in each experiment.

Data analysis. Comparisons between groups were made using a paired *t* test. All data are expressed as mean \pm SD.

RESULTS

EC monolayers could be reproducibly infected with CMV VHL/E as evidenced by the presence of CPE, whereas those incubated with UV-irradiated CMV VHL/E (sham CMV) and control EC demonstrated normal morphology without evidence of CPE (Fig. 1). Similarly, EC monolayers infected with CMV

VHL/E also demonstrated immunofluorescent staining of CMV, whereas those incubated with UV-irradiated CMV VHL/E (sham CMV) or control EC showed negative immunofluorescent staining (Fig. 2). CMV AD169 caused similar effects in fibroblasts.

CMV infection of EC monolayers did not alter PDGF-like protein secretion. In 10 experiments, PDGF-like protein secretion was similar between groups of EC monolayers: CMV-infected EC 13.5 ± 2.6 fmol/10⁶ EC; UV-irradiated infected EC 12.9 ± 1.7 fmol/10⁶ EC; control EC 12.1 ± 3.6 fmol/10⁶ EC (*p* = NS) (Fig. 3).

Western blot analysis of bFGF production from EC monolayers showed no differences in staining intensity of bFGF migrating as a single band at approximately 18 kD from cell lysates of CMV-infected EC, sham CMV-infected EC, and control ECs (Fig. 4). In contrast, Western blot analysis of cell lysates of fibroblasts showed that bFGF present in control uninfected fibroblasts (Fig. 5, lane 1) migrates as a single molecular weight band, similar to that of the recombinant bFGF standards (Fig. 5, lanes 3, 4, and 5). However, CMV-infected cell lysates demonstrated a shift to higher molecular weight forms of bFGF, migrating at 24- and 26-kD molecular mass (Fig. 5, lane 2). Additionally, diminished amounts of the smaller molecular mass 18-kD form found in control fibroblasts were present with CMV infection. Sham-infected fibroblasts showed no alteration in bFGF detection by Western blot.

Treatment of fibroblasts with α interferon failed to induce a shift to higher molecular weight forms of bFGF. Incubations as long as 72 h failed to alter bFGF production from fibroblasts. Other mitogens, such as IL-1, 12-*O*-tetradecanoylphorbol 13-acetate and thrombin, had a similar effect to that of α -interferon (Fig. 6). Finally, bFGF was not detected in conditioned medium from fibroblasts using Western blot analysis.

DISCUSSION

CMV infection of heart transplant patients is associated with arteriolar EC accumulation and with increased intimal thickness of intramyocardial vessels (33). Because EC have been implicated in the development of graft coronary vasculopathy (34–36), the question arises as to whether or not there exists a

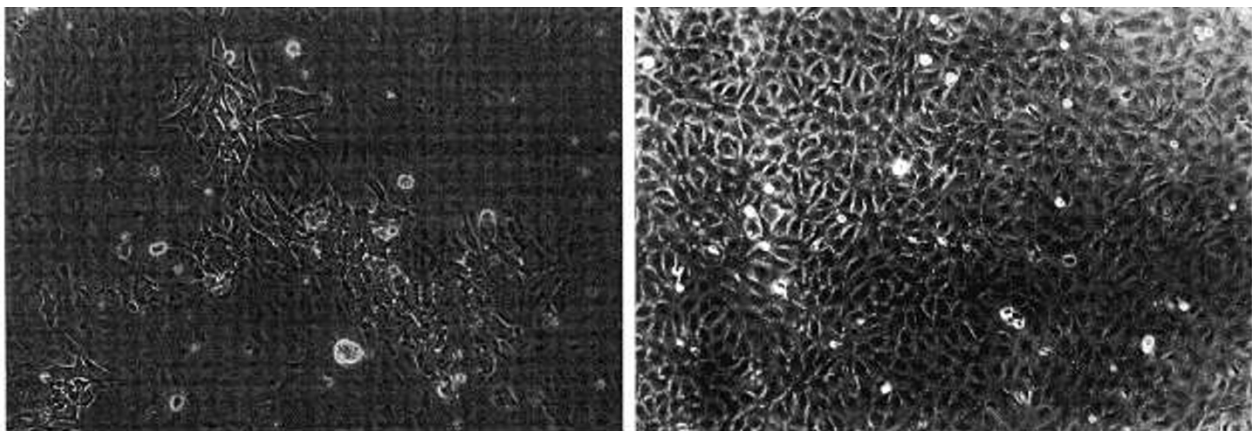


Figure 1. EC monolayer (*left*) infected with CMV VHL/E demonstrating cytopathic effect and (*right*) incubated with UV-irradiated CMV VHL/E (sham CMV) demonstrating normal morphology without evidence of cytopathic effect.

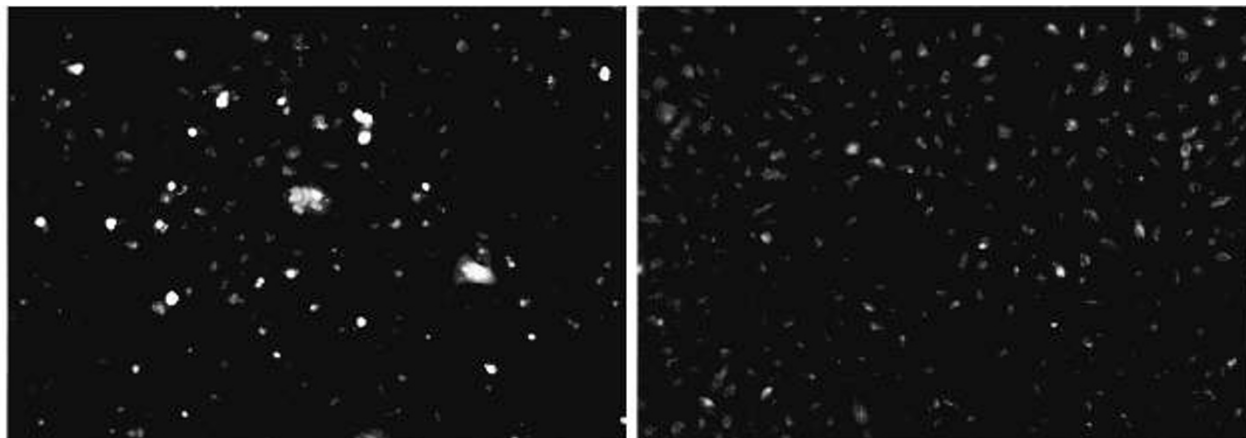


Figure 2. EC monolayer (*left*) infected with CMV VHL/E demonstrating immunofluorescent staining of CMV and (*right*) incubated with UV-irradiated CMV (sham CMV) showing negative immunofluorescent staining. The immunofluorescent staining assay employs MAb specific for two different human CMV antigens: an immediate-early antigen ($M_r = 72\ 000$) and early antigen ($M_r = 50\ 000$). Cell nuclei display fluorescent green staining.

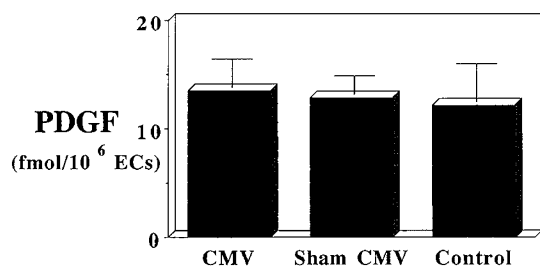


Figure 3. PDGF-like protein secretion from CMV-infected EC, sham (UV-irradiated) CMV-infected EC, and control EC ($n = 10$). No significant differences were noted between groups.

causative relationship between CMV infection, EC activation, and graft coronary vasculopathy. In this study, we attempted to determine whether *in vitro* CMV infection of EC monolayers alters the production or release of PDGF-like proteins or bFGF. Although CMV adequately infected HUVEC, there was no alteration in production of PDGF-like proteins from the EC monolayers. Residual PDGF-like protein initially present in the FCS at the time of the initiation of CMV infection could have still been present in the medium at the time of removal of the conditioned medium. This could have accounted for the somewhat elevated absolute levels of control PDGF-like protein production (12–13 fmol/10⁶ EC) compared with previously reported baseline values of 5–6 fmol/10⁶ EC in previous experiments from this laboratory, in which plasma-derived serum was used instead of FCS (31). However, it is also possible that the prolonged incubation time of the EC in this experiment (5.4 ± 1.8 d) may also account for the increased expression of PDGF-like protein. In any event, the presence of residual PDGF-like protein in the conditioned medium in these experiments cannot account for the lack of differences in PDGF-like protein secretion between CMV-infected EC and both sets of controls, because they were incubated under identical conditions.

Similar to PDGF-like protein, there were no differences in bFGF production from EC after infection with CMV. This suggests that the effects of CMV on EC does not include alterations in growth factor and thus that CMV infection of transplanted hearts does not increase the incidence of graft

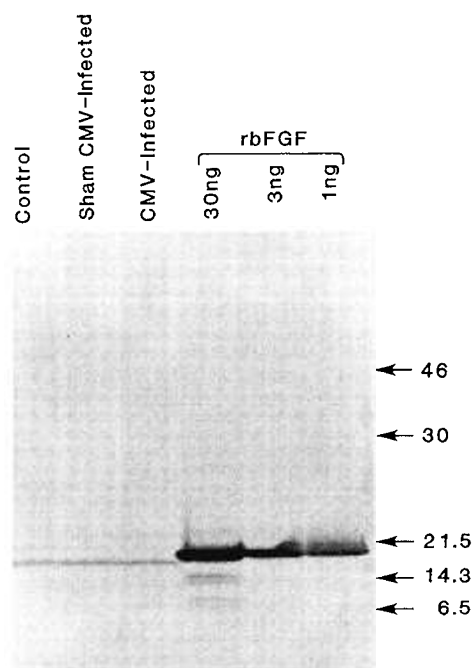


Figure 4. Western Blot analysis of bFGF production from EC monolayers showing no differences in staining intensity of bFGF migrating as a single band at 18 kD from cell lysates from control EC (lane 1), sham (UV-irradiated) CMV-infected EC (lane 2), and CMV-infected EC (lane 3). Lanes 3, 4, and 5: Recombinant bFGF (*rbFGF*) added to the gel directly, demonstrating a single band representing 30, 3, and 1 ng recombinant bFGF. Numbers to the right of the gel represent molecular weight $\times 10^3$.

coronary vasculopathy through a direct effect on PDGF or bFGF production from EC. However, because CMV alters HLA class I and vascular cell adhesion molecule-1 expression on EC and activates T cells to induce MHC class II expression on EC, this suggests that the effects of CMV on EC and graft coronary vasculopathy are indirect effects. Our data support this. Although it is certainly possible that other growth factors may be directly stimulated by CMV infection of cell cultures, PDGF and bFGF appear to be unaffected.

In contrast, CMV infection of cultured human fibroblasts does appear, under certain conditions, to qualitatively alter cellularly associated bFGF by inducing a shift to higher mo-

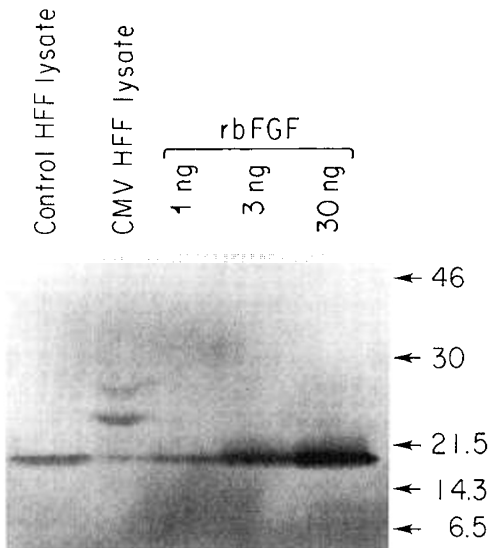


Figure 5. Western Blot analysis of bFGF demonstrating presence of higher molecular weight forms of bFGF during CMV infection. Lane 1: Uninfected (control) human foreskin fibroblast (HFF) cell lysate showing single band of bFGF. Lane 2: HFF cell lysate after 7 d of infection with CMV AD169, demonstrating three separate bands of bFGF, two of which are higher molecular weight forms. Lanes 3, 4, and 5: Recombinant bFGF added to the gel directly, demonstrating a single band representing 1, 3, and 30 ng of recombinant bFGF. Numbers to the right of the gel represent molecular weight $\times 10^3$.

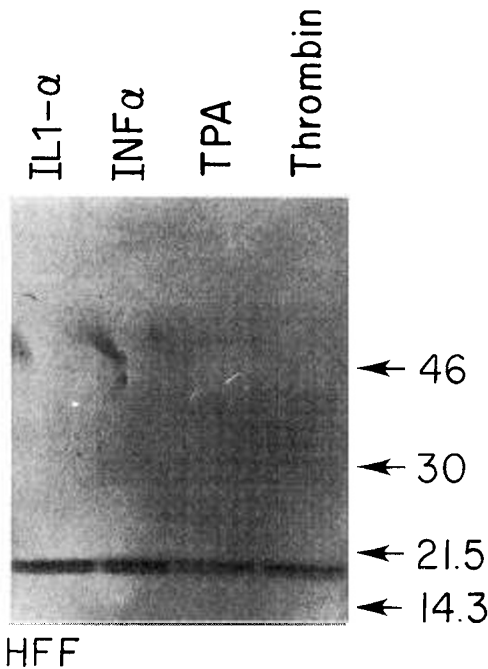


Figure 6. Western blot analysis of bFGF demonstrating no alteration in bFGF production from human foreskin fibroblast (HFF) after treatment for 24 h with IL-1 α , 2 ng/mL; α -interferon (*INF* α), 1000 U/mL; 12-*O*-tetradecanoylphorbol 13-acetate (*TPA*), 200 ng/mL; or thrombin, 10 U/mL. Numbers to the right of the gel represent molecular weight $\times 10^3$.

lecular weight forms. From the work of others (37–39), these higher molecular weight forms have been shown to represent N-terminal extensions of the protein that predominantly have a nuclear distribution. From this same work, it has been shown that, without this N-terminal extension, bFGF remains predom-

inantly cytoplasmically associated. Thus, because CMV can induce a shift to these higher molecular weight forms in human fibroblasts, it likely alters the subcellular distribution of bFGF. Both classic 18-kD bFGF and high molecular mass bFGF have similar receptor-binding activity and biologic effects on cell proliferation, chemotaxis, and angiogenesis when exogenously added to EC cultures (40). However, exogenous production of only high molecular weight bFGF in transfected cell lines has been observed to produce phenotypic differences (38). Furthermore, data from expression in NIH 3T3 fibroblasts demonstrating differences in both morphology and tumorigenicity has suggested a functional role for high molecular weight bFGF (41). Interestingly, *in vivo* expression of high molecular weight bFGF within the rat heart appears to be under both developmental and hormonal control (42), and ventricular expression of bFGF mRNA is significantly altered after cardiac transplantation (43).

It is possible that by inducing a shift to higher molecular weight forms of bFGF, CMV may alter the deposition of bFGF into the subcellular matrix and thus contribute to increased myointimal proliferation. This study also shows that α -interferon (a product of virus-infected cells) does not alter production of bFGF. The fact that fibroblasts produced, but did not actively secrete, bFGF in this study supports previous studies which state that bFGF lacks a signal peptide and thus is not normally secreted by cells. Even when higher molecular weight forms of bFGF were produced by fibroblasts, these forms were not found in the conditioned medium. In this study, the extracellular matrix was not separated from the cell layer, and thus it is not possible to know whether the bFGF was isolated to the cells, the subcellular matrix, or both.

We conclude that, although CMV VHL/E effectively infects human EC, this CMV infection does not directly alter PDGF or bFGF production from EC. However, CMV infection of fibroblasts induces a shift to production of higher molecular weight forms of bFGF. The significance of the production of higher molecular weight forms of bFGF is unknown, but we speculate that this may alter the local autocrine or paracrine function of bFGF.

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