Serotonin Mechanisms in Heart Valve Disease I

Serotonin-Induced Up-Regulation of Transforming Growth Factor- β 1 via G-Protein Signal Transduction in Aortic Valve Interstitial Cells

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Clinical disorders associated with increased serotonin [5-hydroxytryptamine (5-HT)] levels, such as carcinoid syndrome, and the use of serotonin agonists, such as fenfluramine have been associated with a valvulopathy characterized by hyperplastic valvular and endocardial lesions with increased extracellular matrix. Furthermore, 5-HT has been demonstrated to up-regulate transforming growth factor (TGF)- β in mesangial cells via G-protein signal transduction. We investigated the hypothesis that increased exposure of heart valve interstitial cells to 5-HT may result in increased TGF- β 1 expression and activity because of serotonin receptor-mediated signal transduction with activation of G α q, and subsequently up-regulation of phospholipase C. Thus, in the present study we performed a clinical-pathological investigation of retrieved carcinoid and normal valve cusps using immunohistochemical techniques to detect the presence of TGF- β 1 and other proteins associated with TGF- β expression, including TGF- β receptors I and II, latent TGF- β -associated peptide (LAP), and α -smooth muscle actin. Carcinoid valve cusps demonstrated the unusual finding of widespread smooth muscle actin involving the interstitial cells in the periphery of carcinoid nodules; these same cells were also positive for LAP. Normal valve cusps were only focally positive for smooth muscle actin and LAP. In sheep aortic valve interstitial cell cultures 5-HT induced TGF- β 1 mRNA production and increased TGF- β 1 activity. 5-HT also increased collagen biosynthesis at the dosages studied. Furthermore, TGF- β 1 added to SAVIC cultures increased the production of sulfated glycan and hyaluronic acid. In addition, overexpression of G α q using an adenoviral expression vector for a constitutively

active G α q mutant (Q209L-G α q) resulted in increased phospholipase C activity as well as up-regulation of TGF- β expression and activity. These results strongly support the view that G-protein-related signal transduction is involved in 5-HT up-regulation of TGF- β 1. In conclusion, 5-HT-associated valve disease may be, in part, because of TGF- β 1 mechanisms. (*Am J Pathol* 2002, 161:2111–2121)

Heart valve disease associated with increased serotonin [5 hydroxytryptamine (5-HT)] levels has been observed with carcinoid tumors,¹ and a 5-HT mechanism has been hypothesized to be responsible for the valvulopathy that has been described after prolonged administration of the diet drug combination of fenfluramine-phentermine.² Cuspal lesions associated with increased 5-HT levels have characteristically demonstrated subendothelial plaques with cuspal interstitial cell hyperplasia, and increased amounts of extracellular matrix constituents.² The presence of distinct subsets of 5-HT receptors in human heart valves^{3,4} suggests that valvular interstitial cells have the potential to respond to serotonin.

A previous study of a clinical series of carcinoid valves removed at cardiac surgery investigated transforming growth factor (TGF)- β mechanisms and demonstrated that increased amounts of latent TGF- β -associated peptide and latent TGF- β -binding protein were present in the cuspal interstitial cells and extracellular matrix, respectively.⁵ Furthermore, related research investigating mesangial cells has shown that 5-HT induces the up-regulation of TGF- β 1 in this cell type.⁶ Thus, serotonin could hypothetically mediate comparable effects by up-regulating TGF- β 1 within heart valve interstitial cells.

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Furthermore, because others using reverse transcriptase-polymerase chain reaction (RT-PCR) techniques have demonstrated the 5-HT type 2 receptor to be present in aortic valve interstitial cells (AVICs),³ we further hypothesized that 5-HT control of TGF- β 1 occurs through a G-protein signal transduction pathway. Thus, we hypothesized that 5-HT-associated heart valve disease is, in part, because of 5-HT-receptor-mediated up-regulation of TGF- β 1 with resulting increased deposition of extracellular matrix components.

Materials and Methods

Human Pathology Specimens and Immunohistochemical Studies

Human aortic valves were obtained either at the time of cardiac surgery or at autopsy, under exemptions granted by the Institutional Review Boards of the Hospital of the University of Pennsylvania and the Children's Hospital of Philadelphia. Carcinoid human tricuspid and pulmonary valve cusps were obtained from three female patients (age range, 47 to 75 years) with severe carcinoid heart disease undergoing valve removal and valve replacement. Normal human tricuspid valves were obtained at autopsy from human patients ranging in age from 65 to 74 years, including three males and two females. All valve specimens were fixed in neutral buffered formalin. Paraffin thin sections were prepared and routine microscopy hematoxylin and eosin staining was performed. Immunohistochemical studies were performed using an avidin-biotin complex technique (Vector Laboratories, Burlingame, CA), with immunoperoxidase methodology using diaminobenzidine tetrahydrochloride (Vector Laboratories) as a final chromogen. The primary antibodies, anti-TGF- β 1 (1:400), anti-TGF- β RI (1:100), and anti-TGF- β RII (1:200), were obtained from Santa Cruz Biotechnology (Santa Cruz, CA). Antibody to human latent associated peptide-TGF- β 1 (LAP-TGF- β 1, AB-246-NA) was obtained from R&D Systems (Minneapolis, MN). Antigen was retrieved by heating the slides at 95°C in 10 mmol/L of sodium citrate (pH 6.0) for 20 minutes. Paraffin sections of human high-grade osteosarcoma and salivary gland were used as positive controls for TGF- β staining.^{7,8} Slides incubated with nonspecific mouse or rabbit IgG (DAKO, Carpinteria, CA) were used as a negative control. Antibodies used for cell characterization studies included the following: a mouse monoclonal antibody against human (α / τ)-smooth muscle actin (SMA) was purchased from DAKO. A Serotonin 2A (5-HT_{2A}) receptor polyclonal antibody was obtained from Oncogene Research Products (Boston, MA). No other anti-5-HT receptor antibodies were commercially or privately available for these studies. Other antibodies used in the study include: von Willebrand factor (DAKO), anti-desmin, anti-vimentin, and anti-calponin (all from Sigma, St. Louis, MO).

Cell Culture

Aortic valve leaflets were isolated from mature female sheep (Western Cross from Thomas Morris, Reisterstown, MD) as approved by the Institutional Animal Care and Use Committee (IACUC) of the Children's Hospital of Philadelphia. Fresh excised cusps were rinsed in Medium 199 (Life Technologies, Inc., Gaithersburg, MD). Leaflets were then scraped lightly on both surfaces with a scalpel blade to remove endothelial cells. To obtain sheep aortic valve interstitial cells (SAVICs), 1- to 2-mm pieces of leaflets were cut by gentle rolling of the scalpel blade, and were cultivated in Medium 199 supplemented with 10% fetal bovine serum (FBS) (HyClone, Logan, UT) and penicillin/streptomycin (Life Technologies, Inc.) in a humidified atmosphere of 95% air-5% CO₂ at 37°C. In most of the studies, cells were cultivated in six-well plates coated with bovine type I collagen (Cohesion Technologies, Inc., Palo Alto, CA) as previously described.⁹ Cells from passages 3 to 10 were used in all experiments. Quiescence was induced by plating the cells in Medium 199 with 0.5% FBS and penicillin/streptomycin for 24 hours before the treatment with reagents of interest.

For cell characterization studies, SAVICs were plated on collagen gels at a density of 20 cells/mm². After 24 to 72 hours of culture in Medium 199 (10% FBS), cells were fixed in 10% neutral buffered formalin. Cultures used for detection of 5-HT_{2A} receptor were fixed in 4% paraformaldehyde for 30 minutes. Immunohistochemistry studies were performed with specific antibodies for α / β -SMA, the serotonin 5-HT_{2A} receptor, von Willebrand factor, desmin, vimentin, and calponin. Immunostaining via immunoperoxidase used biotin-labeled peroxidase-conjugated secondary antibody incubations with diaminobenzidine tetrahydrochloride as a final chromogen as described above.

Transient Expression of Constitutively Active Mutant G α q in SAVICs

A replication defective adenovirus (type V, E1, E3 deleted) containing the gene sequence for the constitutively active GTPase-deficient G α q (Q209L) under the control of the human cytomegalovirus promoter (CMV) termed AdCMV-G α q, was kindly provided by Dr. Morris Birnbaum (Department of Medicine, University of Pennsylvania, Philadelphia, PA).¹⁰ SAVICs were plated on six-well plates overnight in Medium 199 containing 0.5% FBS. After washing, adenovirus suspensions were added at a titer of 10⁷ or 10⁸ plaque-forming units (PFU) per well (~25 to 250 PFU/cell). A replication defective adenoviral vector encoding for the green fluorescent protein (AdCMV-GFP), a control vector, was obtained from the Institute for Human Gene Therapy of the University of Pennsylvania (Philadelphia, PA).

Real-Time RT-PCR of TGF- β 1 Transcripts

Real-time RT-PCR was performed to measure TGF- β 1 expression semiquantitatively as described.¹¹ Total

RNAs were isolated (Trizol reagents, Life Technologies, Inc.) from SAVICs cultivated on bovine type I collagen stimulated with various concentrations of serotonin in Medium 199 with 0.5% fetal calf serum. Typically, 1 μ g of total RNA was treated with amplification grade deoxyribonuclease I (Life Technologies, Inc.) and reverse-transcribed using the TaqMan reverse transcription kit (Perkin-Elmer, Foster City, CA). TGF- β 1 primers and probe, based on the sheep TGF- β 1 sequence, used for real time RT-PCR were: primers, TGTTTCGTCAGCTCTACATTGACTTC and GGCCCCAGGCAGAAATTG; probe: TGGATTACGAACCCAAGGGCTACC. The probe was labeled at the 5' end with the reporter dye 6-FAM and on the 3' end with the dye TAMRA (Perkin-Elmer, Foster City, CA). TaqMan ribosomal RNA was used as an internal control. TaqMan runs were performed in triplicate.

Phospholipase C (PLC) Assay

PLC activity was evaluated via the following procedures:¹² SAVICs cultivated on type I collagen-coated plates were labeled with [3 H]myo-inositol (5 μ Ci/ml; NEN Life Science Products, Inc., Boston, MA) for 24 hours in Dulbecco's modified Eagle's medium (without inositol) (Life Technologies, Inc.) supplemented with 0.5% FBS. After washing, serotonin in different concentrations was diluted in 15 mmol/L of LiCl (Sigma) containing Dulbecco's modified Eagle's medium and incubated with [3 H]inositol-labeled cells for various time points. Inositol-containing fractions were extracted from cells using chloroform/methanol/HCL (1:2:0.05). Inositol 1-phosphate (InsP₁), inositol 1,4-bisphosphate [Ins(1,4)P₂], and inositol 1,4,5-phosphate [Ins(1,4,5)P₃] were eluted sequentially with 100 mmol/L formic acid and 200 mmol/L ammonium formate, 100 mmol/L formic acid and 600 mmol/L ammonium formate, and 100 mmol/L formic acid and 1 mol/L ammonium formate, respectively, on an anion-exchange columns (Agx8 resin, formate form). The columns were calibrated with each inositol phosphate standard to confirm complete separation of InsP₁, Ins(1,4)P₂, and Ins(1,4,5)P₃.

Luciferase Assay for TGF- β Activity

Equal numbers of quiescent cells ($\sim 4 \times 10^5$ cells) on collagen-coated six-well plates were treated with 10 μ mol/L of 5-HT for 24 to 72 hours. After incubation, medium (test sample) was collected, centrifuged, and assayed for TGF- β 1 activity by the plasminogen activator inhibitor 1 luciferase (PAI/L) assay, as described by Abe and colleagues.¹³ Briefly, 1.6×10^4 mink lung epithelial cells, which were stably transfected with a portion of the plasminogen activator inhibitor 1 promoter (a generous gift from Dr. DB Rifkin, New York University Medical Center, New York, NY), were plated in 96-well tissue culture plates and allowed to attach for at least 3 hours at 37°C in a 5% CO₂ incubator. The medium was replaced with the test sample for 14 hours at 37°C to assay the active TGF- β activity. For total TGF- β 1 (active and latent), test samples were heated for 5 minutes at 80°C and diluted 4-

to 10-fold before adding to mink lung epithelial cells. Cell extracts were prepared and assayed for luciferase activity using a luciferase assay kit (Tropix, Inc., Bedford, MA). TGF- β 1-neutralizing antibody (AB-100-NA; R & D Systems, Minneapolis, MN) was also used in control studies to document the specificity of the assay.

Collagen and Total Protein Synthesis

Collagen synthesis was assessed by measuring the cellular uptake of 3 H-proline as described by Hafizi and colleagues¹⁴ with minor modifications. Briefly, SAVICs were seeded on type I collagen-coated 24-well plates at a density of 3×10^4 cells per well in growth medium overnight. To make the cells quiescent, cells were incubated with M199/0.5% FBS for at least 24 hours before adding various concentrations of 5-HT, TGF- β 1, or anti-TGF- β 1 antibody (R & D Systems). L-[2,3,4,5- 3 H]proline (NEN Life Science Products, Inc.) was added to each well at a final concentration of 10 μ Ci/ml and the cells were incubated for 48 hours before trichloroacetic acid precipitation. For total protein synthesis, cells were incubated with test samples for 24 hours and L-[3 H]leucine (1 μ Ci/ml, NEN Life Science Products, Inc.) incorporation throughout the last 5 hours was measured.¹⁵ The specificity of 5-HT stimulation of 3 H-proline by a TGF- β 1 mechanism was assessed by adding anti-TGF- β 1 antibody (R & D Systems) to the 3 H-proline samples just described.

Cell Proliferation Assay

A colorimetric assay was used according to manufacturer's direction (Boehringer Mannheim, Indianapolis, IN) for the quantification of cell proliferation and cell viability, based on the cleavage of the tetrazolium salt WST-1 by mitochondrial dehydrogenases in viable cells. SAVICs were cultured on type I collagen-coated 96-well plates in a density of 1×10^4 cells/well. After 48 hours of incubation in reduced serum culture medium (M199/0.5% FBS), without or with test compounds, or a control positive stimulus (10% FBS), 10 μ l of the cell proliferation reagent WST-1 was added to each well and incubated for 2 hours. The absorbance was measured using a microtiter plate reader at a wavelength of 418 nm.

Quantitative Assay for Hyaluronic Acid

Hyaluronic acid (HA) synthesis was measured by an enzyme-linked immunosorbent-like assay as described previously.¹⁶ Conditioned medium and standards were first preincubated with protease (bacterial type XIV, Sigma) to remove interfering proteins, then preincubated with biotinylated HA-binding protein (bHABP; Seikagaku, Japan), and finally applied to microtiter plates coated with HA (ICN, Irvine, CA). The plates were washed and the bound bHABP was quantitated using an avidin-peroxidase system (Vectastain ABC kit, Vector Laboratories) with o-phenylenediamine (Sigma) and 0.015% hydrogen peroxide. The intensity of the color was thus inversely related to the amount of HA available for the bHABP.

Table 1. Distribution of α/γ SMA, TGF- β 1, LAP-TGF- β 1, TGF- β Receptor (RI), TGF- β RII on Normal Autopsied Tricuspid Valves and Valves Retrieved from Patients Undergoing Aortic Valve Replacement because of Carcinoid Valvular Disease

	SMA	TGF- β 1	LAP-TGF- β	TGF- β RI	TGF- β RII
Control tricuspid valves from autopsy					
1	++*†	—	+	+	—
2	++*†	—	—	—	—
3	++*	+	—	++	+
4	++*	++	++	++	+
5	++*	+	+	++	+++
Carcinoid valves from patients underwent valve replacement surgery					
6	+++	—	+	++	+
7	+++	—	++	—	++
8	+++	—	++	—	++
9 (pulmonary)	+++	—	+	+	+++

*Subendothelial staining.

†Focal spindle cells.

‡Endothelial staining.

Glycosaminoglycan Assay

Glycosaminoglycans in the conditioned medium were determined by the Blyscan assay (Accurate Chemical and Scientific Corporation, Westbury, NY) following the manufacturer's instructions. Briefly, cells were cultured on bovine type I collagen (Vitrogen) as described above. Before adding 5-HT, cells were incubated in M199 containing 0.5% FBS for 48 hours to make the cells quiescent. Serum-free medium that did not contain phenol red was then used for the experiments. Conditioned media samples were collected at various time points, freeze dried, and kept at -70°C until the assays were performed.

Statistical Methods

All studies were performed in triplicate and data are the means of at least three independent experiments. Error bars indicate standard errors of the means. Differences between means were tested with one-way analysis of variance followed by Tukey-Kramer multiple comparisons post test (InStat; GraphPad Software, Inc., San Diego, CA). A value of $P < 0.05$ was considered statistically significant.

Results

Immunohistochemical Studies of Carcinoid Heart Valves and Autopsied Control Valves

We examined three tricuspid valve cusps and one pulmonary valve cusp taken from patients undergoing valve replacement surgery for severe carcinoid heart disease (Table 1). Five clinically normal tricuspid valve cusps were included in the study as controls (Table 1). Carcinoid valve cusps demonstrated numerous dense nodules, enriched in extracellular matrix, that were surrounded by hyperplasia of the valvular interstitial cells. These carcinoid perinodular interstitial cells showed a distinct spindle-like morphology (Figure 1A). Immunostaining of the sectioned valves for SMA showed strong differences between the carcinoid valve cusps and con-

trol valve cusps (Figure 1, Table 1). In carcinoid valve cusps, most of the interstitial cells surrounding the carcinoid nodules were immunopositive for SMA (Figure 1A). In contrast, control valve cusps only showed focal subendothelial SMA staining (Figure 1B). In one of the controls, a small focus of spindle-shaped cells was observed that was also immunopositive for SMA (data not shown). Nonspecific IgG exposure resulted in no apparent immunopositive staining (Figure 1E).

We also performed immunostaining of the valve cusps for active TGF- β 1, LAP-TGF- β 1, and TGF- β receptors type I and II (Table 1). Both receptors were present in carcinoid valve cusps and control valve cusps in variable amounts. Focal LAP-TGF- β 1 staining was found in all carcinoid valve cusps and three of the control valves with important differences in immunostaining pattern. In carcinoid valve cusps, positive staining was observed mainly in the spindle-shaped cells around carcinoid nodules (Figure 1C). In control cusps, positive immunostaining was found mainly in endothelial or subendothelial cells (Figure 1D). However, TGF- β 1 staining was not observed in carcinoid valve cusps and was detected in sparse amounts in normal cusps (Table 1).

Characterization of SAVICs

Virtually all SAVICs in culture were immunopositive for the 5-HT_{2A} receptor, compared to nonspecific IgG controls (Figure 2, A and B). Unfortunately, no other antibodies were commercially available to screen for other 5-HT₂ receptor subtypes. Other cell markers used in the characterization of SAVICs included von Willebrand factor, vimentin, desmin, and calponin. SAVICs were negative for vimentin and von Willebrand factor, and weakly positive for desmin and calponin (data not shown). When SAVICs were cultured in the presence of 10 ng/ml of TGF- β 1, significantly more cells were immunopositive for α/τ -SMA than cells not exposed to TGF- β 1 (Figure 2, C and D). In control cultures (no added TGF- β 1), $6.5 \pm 1.2\%$ of cells were α/τ -SMA-positive. However, after TGF- β 1 treatment, $56.9 \pm 10.7\%$ of cells showed α/τ -SMA positivity ($P < 0.05$). Nonspecific IgG control showed no immunopositive staining (Figure 2E).

Figure 1. **A:** Representative micrographs of a carcinoid tricuspid valve cusp with prominent SMA staining by peroxidase immunohistochemistry (brown staining) with a hematoxylin counterstain. **B:** Normal control tricuspid valve cusp shows sparse subendothelial SMA staining (arrows indicate SMA-positive cells). **C:** A carcinoid valve cusp with strongly positive LAP-TGF- β 1 immunostaining (arrows) localized in hyperplastic areas around carcinoid nodules. **D:** Focal subendothelial LAP-TGF- β 1 immunostaining (arrows) in control valves. **E:** Negative IgG control shows an absence of peroxidase activity. Asterisk denotes carcinoid nodule. All micrographs were prepared with immunoperoxidase and hematoxylin staining techniques. Original magnifications, $\times 400$.

TGF- β 1 Is Up-Regulated by Serotonin (5-HT) in Aortic Valve Interstitial Cells

SAVICs were cultivated in the presence of 5-HT, and were monitored for TGF- β 1 expression in terms of mRNA levels and TGF- β 1 activity, using the PAI/luciferase assay. TGF- β 1 mRNA, evaluated by real-time-RT-PCR, was increased significantly after 6 hours of 10 μ mol/L 5-HT exposure and remained relatively constant up to 72 hours (Figure 3A, $P < 0.05$). For calibration, increasing concentrations of TGF- β 1 were directly added to mink lung ep-

ithelial cells, and this resulted in a dose-dependent rise in luciferase activity (Figure 3B). Furthermore, treatment of SAVICs with 10 μ mol/L of 5-HT for 72 hours resulted in an increase in both direct and total TGF- β activity (Figure 3, C and D). However, these increased luciferase activities were blocked completely by a TGF- β 1 function-blocking antibody at a concentration of 10 μ g/ml (Figure 3, C and D). These studies conclusively demonstrate that 5-HT causes SAVICs to up-regulate TGF- β 1, resulting in increased amounts of active (Figure 3C) and latent TGF- β 1 (Figure 3D).

Figure 2. Immunostaining results demonstrating that SAVICs are immunopositive (brown) for the 5-HT_{2A} receptor (A), compared to B, which demonstrates an absence of immunopositive cells using a nonspecific antibody. TGF- β 1 exposure results in increased α /t-SMA-immunopositive cells (brown) with an elongated cytoskeleton (C), compared to D. Cells not exposed to TGF- β 1 are predominant by immunonegative for α /t-SMA. Nonspecific IgG results were negative for α /t-SMA (E). Peroxidase immunohistochemistry. Original magnifications: $\times 200$ (A, B); $\times 400$ (C-E).

5-HT and TGF- β 1 Increase the Synthesis of Both Collagen and Glycosaminoglycans

Both 5-HT and TGF- β 1 (Figure 4) increased collagen synthesis by SAVICs, as measured by ^3H -proline incorporation. The addition of TGF- β 1 to SAVIC cultures resulted in significantly increased collagen synthesis at concentrations of 0.01 ng/ml to 10 ng/ml of TGF- β 1 (Figure 4A) with results comparable to those seen with 5-HT. Total protein synthesis was assessed using a [^3H]leucine incorporation assay. TGF- β 1 has no effect on total protein synthesis at a concentration of 1 ng/ml or lower, but total protein synthesis was inhibited ($P < 0.05$) at 10 ng/ml of TGF- β 1 (Figure 4B). A stimulatory effect of 5-HT on collagen synthesis was observed at doses ranging from 1 $\mu\text{mol/L}$ to 100 $\mu\text{mol/L}$ of 5-HT (Figure 4C). However, collagen synthesis stimulated by 5-HT was significantly reduced by an anti-TGF- β antibody (Figure 4D), thus indicating the specificity of 5-HT stimulation of TGF- β 1.

The secretion of glycosaminoglycan by SAVICs after 5-HT or TGF- β 1 treatment was also investigated. The

secretion of sulfated glycosaminoglycan was increased significantly after TGF- β 1 treatment (Figure 5A), but no significant increase was observed after 5-HT treatment (data not shown). Hyaluronic acid (HA), a nonsulfated glycosaminoglycan, was also studied. HA is the most abundant acidic glycosaminoglycan (AGAG) in aortic valve cusps, constituting almost 50% of the total AGAG.¹⁷ It is hypothesized that the presence of HA is a marker for an initial phase of the extracellular matrix remodeling.¹⁸ These experiments demonstrated that TGF- β 1 increases SAVIC HA synthesis in a dose-dependent manner, and a significant cumulative effect was observed at a concentration of 10 ng/ml or greater of TGF- β 1 after 24 hours of incubation (Figure 5B). The effect of TGF- β 1 on HA synthesis was also cumulative throughout time (Figure 5C). We also examined the effect of direct addition of 5-HT on HA secretion. At a concentration of 10 $\mu\text{mol/L}$ of 5-HT, no significant change in HA secretion was observed after 24 hours incubation (Figure 5B). The effects of 5-HT and TGF- β 1 on cell proliferation were also evaluated by a colorimetric assay¹⁹ for the

Figure 3. Serotonin up-regulates TGF-β1 expression and activity in AVICs. **A:** Time-dependent TGF-β1 mRNA induction by 10 μmol/L of 5-HT treatment evaluated by real-time RT-PCR. Results are expressed as mean ± SEM of four to six independent experiments (**, $P < 0.01$; *, $P < 0.05$; analysis of variance followed by Tukey-Kramer multiple comparisons post test). **B:** A standard calibration curve for the PAI/L construct with directly added TGF-β1 demonstrating a dose response. **C:** Active TGF-β stimulated by 10 μmol/L of 5-HT after 72 hours, per PAI/luciferase assays, and blocked by a TGF-β1 function-blocking antibody. **D:** Total TGF-β (active and latent), examined by PAI/luciferase assays, increased because of 72 hours exposure to 5-HT, and blocked by TGF-β1 function-blocking antibody. For **B**, **C**, and **D**, the results are expressed as mean ± SEM of triplicates, which are typical of four independent experiments (*, $P < 0.05$, significant from control; **, $P < 0.01$, significant from 5-HT; analysis of variance followed by Tukey-Kramer multiple comparisons post test).

quantification of cell proliferation (WST-1, Boehringer Mannheim; see Materials and Methods). The concentration ranges of 5-HT and TGF-β1 used in these studies had little effect on proliferation: for serotonin, 0 to 10 μmol/L, cell counts ranged from 5.0×10^4 to 6.0×10^4 ; for TGF-β1, 0 to 10 ng/ml, cell counts ranged from 4.5×10^4 to 5.6×10^4 , whereas a positive stimulus (10% FBS) produced a major increase (6×10^3 increasing to 3×10^4) in cell number.

Overexpression of Gαq via a Replication-Defective Adenovirus Increases PLC Activity, TGF-β1 Expression, and TGF-β1 Activity

It was hypothesized that 5-HT-induced TGF-β1 production by SAVICs was because of receptor-mediated activation of the G-protein signal transduction pathway. Related research by our group established that 5-HT administration to SAVICs in the same dose range used in the present TGF-β1 studies, results in a dose-dependent

increase in phospholipase C (PLC) activity.²⁰ Thus, using an adenoviral vector construct with constitutively active, GTPase-deficient mutant Gαq (AdCMV-G29), we investigated the effects of overexpression of Gαq on both PLC and TGF-β1 activity. Initial studies investigated the effects of increasing dosages of Gαq-adenovirus, compared to AdCMV-GFP on PLC activity. It was observed that a 40-fold (10^7 PFU Gαq) to 70-fold (10^8 PFU Gαq) increase in PLC activity occurred over the dosage range studied (Figure 6A).

The effects of Gαq on TGF-β1 mRNA level were not significantly increased until 72 hours (Figure 6B). However, active TGF-β1 content, as assessed with the PAI/luciferase system using a dose of 10^7 PFU Gαq, was increased more than sixfold as early as 24 hours and remained constant for 72 hours (Figure 6C). However, total TGF-β1 activity did not increase significantly in response to Gαq overexpression until 72 hours after the addition of adenoviral vector, in association with the in-

Figure 4. Both 5-HT and TGF-β1 stimulate collagen synthesis by SAVICs. **A:** Stimulation of [³H]proline incorporation into cells by TGF-β1, significant stimulation was observed at 0.01 to 10 ng/ml of TGF-β1 (**, $P < 0.01$). **B:** The effects of TGF-β1 on total protein synthesis measured by [³H]leucine incorporation, demonstrating significant inhibition of protein synthesis at 10 ng/ml of TGF-β1 (*, $P < 0.05$). **C:** Stimulation of [³H]proline incorporation into cells by 5-HT, significant stimulation was observed from 1.0 to 100 μmol/L of 5-HT (*, $P < 0.05$; **, $P < 0.01$). **D:** Anti-TGF-β antibody blocked [³H]proline incorporation into cells stimulated by 5-HT. A significant blocking effect was observed at concentrations as low as 5 μg/ml of anti-TGF-β antibody (*, $P < 0.05$; **, $P < 0.01$). Results are expressed as mean ± SEM of four independent experiments (*, $P < 0.05$; **, $P < 0.01$, analysis of variance followed by Tukey-Kramer multiple comparisons post test).

crease of TGF-β1 mRNA levels (Figure 6B). AdCMV-GFP had no effect on either PLC activity or on TGF-β expression and activity (Figure 6; A, C, and D).

Discussion

The present studies have demonstrated that 5-HT-related heart valve disease may be, in part, because of TGF-β mechanisms. These results are in agreement with those reported previously^{21–23} and provide insights concerning the mechanisms responsible for serotonin-mediated heart valve disease. Carcinoid heart valve cusps have been demonstrated to accumulate TGF-β latency associated peptide and latent binding protein.⁵ Furthermore, serotonin has been demonstrated by our study and others to stimulate increased collagen synthesis by heart valve interstitial cells.¹⁴ Serotonin up-regulation of TGF-β has also been observed in mesangial cells with associated increased PLC activity indicating G-protein signal transduction.⁶ In addition, 5-HT₂ receptors (A and B)

have been demonstrated by RT-PCR studies to be present in human and porcine aortic valve cells.³ We demonstrated 5-HT_{2A} receptors with immunostaining in all SAVICs. Our results are consistent with these previous studies. Thus, the present results and related previous research are consistent with the view that heart valve interstitial cells are serotonin responsive, and thus it is likely that with an excessive serotonin exposure, as in the carcinoid syndrome, a TGF-β-related mechanism may explain at least in part the pathophysiological events.

Our clinical pathology study was comparable to the results of others⁵ revealing relatively greater amounts of LAP in carcinoid cusps compared to normals, very likely reflecting up-regulation of TGF-β because of serotonin. However, relatively lower levels of the active form of TGF-β were detected compared to normal cusps. The mechanisms responsible for this observation are not completely understood, but may be, in part, because of rapid turnover of the active form of TGF-β with relatively little accumulation in the metabolically dynamic carcinoid

Figure 5. TGF- β 1 stimulates both total glycosaminoglycan and HA production by SAVICs in a dose- and time-dependent manner. **A:** Time- and dose-dependent stimulation of total sulfated glycan by TGF- β 1. **B:** Dose-dependent stimulation of HA production by TGF- β 1, with a significant increase in HA production observed at 10 ng/ml or greater of TGF- β 1 for 24 hours treatment. 5-HT at 10 μ mol/L showed no significant effect on HA production. **C:** Time-dependent stimulation of HA production by TGF- β 1, with the maximum HA production observed after 48 hours of incubation with TGF- β 1. Results are expressed as mean $\pm$ SEM of triplicates, which are typical of three or four independent experiments (*, $P < 0.05$; **, $P < 0.01$, followed by Tukey-Kramer multiple comparisons post test).

cusps. Furthermore, hyperplastic interstitial cells within carcinoid cusps were for the most part positive for SMA, which was present to a markedly lesser extent in normal cusps. This may reflect up-regulation of SMA in cardiac valve interstitial cells because of TGF- β (Figure 2). This observation is also comparable to the up-regulation of SMA by TGF- β in fibroblasts observed by others.²⁴ Thus, in carcinoid valves increased SMA could be hypothetically because of serotonin-mediated up-regulation of TGF- β .

Serotonin-related heart valve disease is a chronic process. Thus, the relatively rapid signaling events reported in this study can only explain the potential initiating mechanism that could be hypothetically associated with a cumulative effect throughout time, via changes in the extracellular matrix including accumulation of TGF- β . As discussed above, latent TGF- β -binding protein accumulates in the extracellular matrix, via covalent attachment,^{25,26} and thus carcinoid heart valve disease provides an experiment in nature that supports our hypothesis. Furthermore, other studies from our group have shown that calcified human heart valves, but not normals, have high amounts of TGF- β accumulating in the extracellular matrix.²⁷ Although this observation is unlikely to be related to serotonin mechanisms, it nevertheless supports the view that chronic accumulation of

TGF- β in human heart valve disease is a common pathological event that could have multiple pathophysiological impacts on heart valve disease in general. Therefore, the cell culture studies reported in this study with serotonin-induced TGF- β expression and activity, and increased biosynthesis of both collagen and glycosaminoglycans, may reflect incremental events of a chronic cumulative process.

The use of an adenoviral vector that overexpresses G α q provided a unique opportunity to establish a strong association between G-protein signal transduction and TGF- β expression in heart valve interstitial cells. Our present studies demonstrate that there were significant increases in both PLC activity and TGF- β expression and activity because of overexpression of G α q after adenoviral vector transduction. Interestingly, these results demonstrate that G-protein signal transduction appears to act at three levels to enable TGF- β 1 expression and activity. As described above, an increase in the active TGF- β occurs after only 24 hours. Furthermore, TGF- β 1 mRNA synthesis increases throughout a somewhat longer time course of 72 hours. In addition, 72 hours after G α q exposure, both active and total TGF- β activity are significantly elevated (Figure 6). Taken together, these data indicate a profound effect of G-protein signal transduction on TGF- β 1 expression and processing in heart valve

Figure 6. Overexpression of Gαq via transduction of SAVICs with a replication defective adenovirus stimulates PLC activity and TGF-β1 expression, where S1 is the M199 medium (control); S2 is the AdCMV-GFP, 10⁷ PFU; S3 is the AdCMV-Gαq, 10⁷ PFU. Overexpression of Gαq significantly increases the accumulation of total inositol phosphates (**A**), stimulates TGF-β1 mRNA expression (**B**); increases the production of active TGF-β1 evaluated by PAI/L assay (**C**); and increases the production of total (active and latent) TGF-β1 evaluated by PAI/L assay (**D**). Results are expressed as mean ± SEM of triplicates, which are typical of three independent experiments (*, *P* < 0.05; **, *P* < 0.01, analysis of variance followed by Tukey-Kramer multiple comparisons post test).

interstitial cells via Gαq, that may be related in part to the pathophysiological events in serotonin-associated heart valve disease.

The absence of any effect of 5-HT on glycosaminoglycan synthesis by SAVICs in culture in our studies (Figure 5) and a modest effect on proline incorporation (reflecting increased collagen synthesis) could be because of the fact that a far longer duration of exposure may be needed for a cumulative effect in a cell culture model system. Furthermore, our results also show that the TGF-β activity resulting from 5-HT stimulation is primarily in the latent form (thus requiring activation for physiological effects to ensue), in contrast to our studies involving direct addition of active TGF-β1 that had potent effects on both glycosaminoglycan and collagen production. These results concerning latent TGF-β are in agreement with our clinical pathology data. Because of the potential for serotonin exacerbation of heart valve disease, pharmaceuticals that have the capacity to increase serotonin levels could also contribute to the progression of heart valve disease in general. Thus, new and existing drugs could be screened in AVIC culture systems to determine whether the potential for TGF-β-related adverse effects exists. Further, it is conceivable that a novel therapeutic approach could arise based on the serotonin TGF-β pathway investigated in these studies. Numerous serotonin

receptor antagonists have already been described,^{28–30} and thus could be screened for their effects on cell cultures of AVICs. A similar approach could also be taken with the various PLC inhibitors that have been reported.^{31,32} Therefore, because our results show that heart valve interstitial cells are unusually sensitive to serotonin, a serotonin-based therapeutic strategy is worthy of investigation.

Conclusions

Serotonin may contribute to the development and progression of cardiac valve disease through up-regulation of TGF-β1, and related increases in extracellular matrix components, including collagen and glycosaminoglycans. Serotonin signaling in heart valve interstitial cells leading to up-regulation of TGF-β1 likely takes place via a G-protein signal transduction mechanism, thus indicating potential therapeutic strategies that could be investigated.

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