Peroxisome Proliferator-Activated Receptor γ Regulates the V-Ets Avian Erythroblastosis Virus E26 Oncogene Homolog 1/microRNA-27a Axis to Reduce Endothelin-1 and Endothelial Dysfunction in the Sickle Cell Mouse Lung

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Abstract

Pulmonary hypertension (PH), a serious complication of sickle cell disease (SCD), causes significant morbidity and mortality. Although a recent study determined that hemin release during hemolysis triggers endothelial dysfunction in SCD, the pathogenesis of SCD-PH remains incompletely defined. This study examines peroxisome proliferatoractivated receptor γ (PPAR γ) regulation in SCD-PH and endothelial dysfunction. PH and right ventricular hypertrophy were studied in Townes humanized sickle cell (SS) and littermate control (AA) mice. In parallel studies, SS or AA mice were gavaged with the PPAR_Y agonist, rosiglitazone (RSG), 10 mg/kg/day, or vehicle for 10 days.In vitro, human pulmonary artery endothelial cells (HPAECs) were treated with vehicle or hemin for 72 hours, and selected HPAECs were treated with RSG. SS mice developed PH and right ventricular hypertrophy associated with reduced lung levels of PPARg and increased levels of microRNA-27a (miR-27a), v-ets avian erythroblastosis virus E26 oncogene homolog 1

(ETS1), endothelin-1 (ET-1), and markers of endothelial dysfunction (platelet/endothelial cell adhesion molecule 1 and E selectin). HPAECs treated with hemin had increased ETS1, miR-27a, ET-1, and endothelial dysfunction and decreased PPARg levels. These derangements were attenuated by ETS1 knockdown, inhibition of miR-27a, or PPAR γ overexpression. In SS mouse lung or in hemin-treated HPAECs, activation of PPAR_y with RSG attenuated reductions in PPAR_y and increases in miR-27a, ET-1, and markers of endothelial dysfunction. In SCD-PH pathogenesis, ETS1 stimulates increases in miR-27a levels that reduce PPARg and increase ET-1 and endothelial dysfunction. PPARg activation attenuated SCD-associated signaling derangements, suggesting a novel therapeutic approach to attenuate SCD-PH pathogenesis.

Keywords: peroxisome proliferator-activated receptor γ ; sickle cell disease; microRNA-27a; v-ets avian erythroblastosis virus E26 oncogene homolog 1; pulmonary hypertension

Sickle cell disease (SCD), an autosomal recessive disorder caused by a mutant b-hemoglobin (HbS), is associated with a large spectrum of pathologic complications caused by the aggregation and polymerization of HbS within red blood cells (RBCs) (1, 2). The sickling of abnormal RBCs results in anemia and

vasoocclusion, causing acute chest syndrome (3, 4) and multiorgan dysfunction. SCD affects millions of people worldwide. In the United States, one of

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Clinical Relevance

The current study provides novel evidence that microRNA-27a (miR-27a) is up-regulated in mice with sickle cell disease and in hemintreated human pulmonary artery endothelial cells through v-ets avian erythroblastosis virus E26 oncogene homolog 1 activation. Increases in miR-27a down-regulate peroxisome proliferator-activated receptor γ (PPAR γ) and contribute to endothelial dysfunction. Activation of PPARg stimulates reciprocal inhibition of v-ets avian erythroblastosis virus E26 oncogene homolog 1, miR-27a and pulmonary endothelial dysfunction. We believe these findings provide the first indication that targeting PPARg activation may represent a novel therapeutic approach to attenuate sickle cell disease–pulmonary hypertension pathogenesis through modulation of microRNA levels.

every 365 African Americans and one of every 36,000 Hispanic Americans are born with SCD, and \sim 100,000 people are living with SCD (5).

Pulmonary hypertension (PH), defined as an elevation of the mean pulmonary artery pressure to >25 mm Hg (6, 7) is an increasingly recognized complication of SCD that is associated with high morbidity and mortality (8–10). Two-year mortality rates among patients with SCD and PH have been reported to be as high as 40–50% (10, 11), and \sim 6–10.4% of patients with SCD ultimately develop PH (12–14). Despite the magnitude of this problem, the pathogenesis of SCD-PH remains incompletely defined. Current evidence suggests that SCD-PH, similar to other forms of PH, involves endothelial dysfunction with increased production of vasoconstrictors (e.g., endothelin-1 [ET-1]) (15), and reduced production of vasodilators (e.g., prostacyclin [16] and nitric oxide [17, 18]). Interactions between RBCs and endothelium may contribute to the pathogenesis of vasoocclusive crisis in SCD (19). The release of hemoglobin and hemin during hemolysis triggers endothelial dysfunction and contributes to PH pathogenesis (20). Recent studies in sickle cell (SS) patients demonstrate that EC adhesion molecules, including selectins and vascular endothelial cadherin, play an important role in the recruitment and binding of inflammatory cells to vascular endothelium (21). Because SCD-PH morbidity and mortality remain unacceptably high, there is an urgent need for novel therapeutic strategies.

Mouse models of hemolysis, including the Berkeley SS mouse (18, 22), the spherocytosis mouse (23), and the alloimmune hemolysis mouse, develop spontaneous PH and right heart failure (18). To further examine the pathogenesis of SCD-PH and explore novel therapeutic targets, the current study capitalized on an existing mouse model of SCD (24–26). In this study, we used a humanized knockin mouse model of SCD (27) (the Townes model) in which mouse HbS genes are replaced with human γ -and β^S -hemoglobin
genes and mouse α -hemoglobin genes are genes and mouse α -hemoglobin genes are replaced with human α -hemoglobin genes. Because our laboratory and others have shown that stimulating peroxisome proliferator-activated receptor γ (PPAR γ) with thiazolidinedione (TZD) ligands attenuates PH in experimental models $(28-32)$, whereas reduced PPAR γ expression is associated with PH pathogenesis (29, 33, 34), we compared SS mice with littermate control (AA) mice to determine if PPARg plays an important role in SCD-associated pulmonary vascular dysfunction and PH pathogenesis.

PPAR_y is a ligand-activated transcription factor that belongs to the nuclear hormone receptor superfamily. The PPAR γ receptor is activated by a diverse spectrum of ligands that includes naturally occurring fatty acids and their metabolites, as well as synthetic TZD medications, which are currently used in this country to treat patients with type 2 diabetes. Once activated, PPARg forms a heterodimer with the retinoid X receptor and binds to PPAR response elements in the promoter region of responsive genes to increase their expression. PPAR γ activation can also reduce the expression of selected genes though transrepression mechanisms (35). For example, PPAR γ inhibits transcription factors, such as early growth response 1 (36, 37), Sp1 transcription factor (37, 38), and v-ets avian erythroblastosis virus E26 oncogene homolog (ETS)1 (ETS1) (39, 40). ETS1, a member of the ETS family of transcription factors (41), plays an important role in proliferation, migration,

vascular remodeling, apoptosis, and senescence (42). Recent studies demonstrate that PPAR_y reduction increases ETS1 expression and vascular smooth muscle cell migration and proliferation (39), whereas PPARg activation inhibits ETS1 expression and attenuates migration by inhibiting ETS1 expression (40, 43). Furthermore, PPAR γ attenuates hypoxic increases in proliferative mediators (e.g., ET-1 and nicotinamide adenine dinucleotide phosphate oxidase 4), in part through inhibition of nuclear factor-kB p65 (22, 32, 44, 45). Interestingly, mounting evidence demonstrates that pharmacological activation of PPARg attenuates pulmonary vascular cell proliferation and PH (46), whereas loss of PPAR_Y expression or function is associated with PH (47). Taken together, these findings led us to postulate in the current study that: (1) chronic hemolysis in SCD stimulates loss of PPARg and increased expression of the proliferative mediator, ET-1, which contributes to PH pathogenesis; and (2) targeting PPAR γ with existing pharmacological ligands may represent a novel therapeutic approach in SCD to attenuate increases in proliferative mediators and SCD-PH pathogenesis.

Pathways leading to reductions in PPAR γ in PH continue to be defined (29, 33, 34). We reported recently that microRNA-27a (miR-27a) plays an important role in hypoxic reductions in PPAR_y (37). Evolving evidence indicates that dysregulation of microRNAs (miRNA or miR) contribute to PH pathogenesis (48–52). miRNAs are endogenous, noncoding, single-stranded RNAs of \sim 22 nucleotides that constitute a novel class of post-transcriptional gene regulators (53, 54). miRNAs negatively regulate the expression of their target genes through translational repression or mRNA degradation (54). Recent studies have provided compelling evidence that miRNAs regulate pulmonary vascular remodeling by controlling endothelial cell (EC) and smooth muscle cell differentiation and proliferation (48, 49, 51, 52, 55–57). The potential role of miRNAs as master regulators of cell differentiation in physiological and pathological processes in the lung has been reviewed recently (58–60). However, the precise posttranscriptional mechanisms by which PPAR_y activation regulates SCD-PH have not been defined. In the current study, we

explored the hypothesis that hemolysis increases the levels of miR-27a to reduce PPAR_y expression through posttranscriptional mechanisms, leading to increased ET-1 expression, endothelial dysfunction, and PH. The current study demonstrates that SCD-related hemolysis stimulates miR-27a expression, reduces PPAR_y expression, and increases ET-1 in human pulmonary artery endothelial cells (HPAECs) in vitro and in the SS mouse lung in vivo. Pharmacological activation of $PPAR\gamma$ attenuates these derangements, suggesting that therapeutic targeting of this receptor provides a novel strategy for reducing SCD-associated pulmonary endothelial dysfunction and PH.

Materials and Methods

Reagents

HPAECs were obtained from Invitrogen (Carlsbad, CA). ET-1, PPARg, ETS1, and glyceraldehyde phosphate dehydrogenase antibodies were obtained from Santa Cruz Biotechnology (Santa Cruz, CA). The following reagents were obtained from Sigma-Aldrich (St. Louis, MO): hemin, fetal bovine serum (FBS), dimethyl sulfoxide (DMSO), and methyl cellulose. 3-(4,5-Dimethylthiazol-2-yl)-2,5 diphenyltetrazolium bromide (MTT) was purchased from ATCC (Manassas, VA). The PPAR_Y ligand, rosiglitazone (RSG), was obtained from Cayman Chemical (Ann Arbor, MI).

Measurement of PH in SS Mice

The recently described Townes humanized mouse model of SCD was used in the current study (27), and male and female SS and AA mice were examined at age 15–17 weeks, as reported (32). All animals were given unrestricted access to water and standard mouse chow. To assess PH, SS and AA mice were subjected to measurements of right ventricular systolic pressure (RVSP) and right ventricular hypertrophy (RVH), as reported (32). Briefly, after being anesthetized with isoflurane, RVSP was measured by introducing a 1.2-F microtip pressure transducer catheter (Transonic, Ithaca, NY) into the right internal jugular vein and advancing it to the right ventricle (RV), and RV pressures were monitored continuously for 10 minutes. RVH was assessed by

dissecting the heart and calculating the ratio of the weight of the RV to the weight of the left ventricle (LV) and septum (S) [Fulton's Index, $RV:(LV + S)$ weight ratio]. Selected animals were gavaged daily with ¹⁰⁰ ml RSG (10 mg/kg/d in 100 ml 0.5% methyl cellulose) or with an equivalent volume of vehicle for 10 days, as we reported previously (32). All experiments using mice were approved by the Institutional Animal Care and Use Committee of the Atlanta Veterans Affairs Medical Center.

Hemin-Treated HPAECs and Cell Proliferation Assay

To examine the effect of RSG on hemininduced cell proliferation, HPAECs were treated with DMSO as a control, or hemin (HEM, 5 μ M), for 72 hours, as reported (61). Hemin concentrations in this range have been measured in patients with SCD (62). Selected HPAECs were treated with RSG (10 μ M) or an equivalent volume of vehicle added directly to the culture media during the last 24 hours of hemin treatment, and cell proliferation was measured using MTT assays, as we have reported (46).

miR-27a Down-Regulation

To confirm the role of miR-27a in alterations in PPARg expression, HPAECs (passages 3–6) were transfected with anti–miR-27a (50 nM) or an equivalent amount of scrambled anti-miR negative control using lipofectamine RNAiMax (QIAGEN, Valencia, CA). After transfection for 6 hours, media were replaced with endothelial growth medium (EGM) containing 10% FBS. HPAECs were treated with DMSO or HEM $(5 \mu M)$ and then cultured for 72 hours. Alterations in miR-27a, ET-1, PPARg, platelet/endothelial cell adhesion molecule 1 (PECAM1), and E-selectin or endothelial adhesion molecule 1 (E-SEL) levels were examined using real-time quantitative reverse transcription-polymerase chain reaction (qRT-PCR).

PPAR_y Overexpression in HPAECs In Vitro

To overexpress PPARg, HPAECs were transfected with adenovirus containing a PPAR_y plasmid (AdPPAR_y, 25 multiplicity of infection) or control green fluorescent protein (GFP) plasmid, as we reported previously (63). Six hours after transfection,

adenovirus-containing medium was replaced with fresh 10% FBS EGM, and HPAECs were then treated with DMSO or HEM (5 μ M) for 72 hours. Levels of miR-
273. FTS1. FT-1. PPAR α . PECAM-1. and 27a, ETS1, ET-1, PPARγ, PECAM-1, and
E-SEL were determined by aRT-PCR E-SEL were determined by qRT-PCR.

HPAEC Transfection and RNA Interference Studies

HPAECs were transfected with scrambled or ETS1 (5'-UCAGUAGAGAGAAGUGUA CGGCU TT-3', 3'-UAAGUCAUCUC UCUUCACAUGCCGAA A-5') RNA interference (RNAi) duplex (10 nM) using RNAiMax transfection reagent (Invitrogen) according to the manufacturer's instructions. After transfection for 6 hours, the transfection media were replaced with EGM containing 10% FBS. HPAECs were then treated with DMSO or HEM $(5 \mu M)$ for 72 hours. HPAEC lysates were then harvested and examined for ETS1, miR-27a, ET-1, and PPAR γ levels using qRT-PCR and Western blot assays.

miRNA and mRNA qRT-PCR Analysis

To measure miR-27a, PPARg, ETS1, ET-1, E-SEL, and PECAM1 levels in HPAECs or SS mouse lungs, small RNAs $(<200$ nt) and large RNAs $(>200$ nt) were isolated using the mirVana kit (Invitrogen). The levels of miR-27a expression were analyzed by qRT-PCR using the QIAGEN miRNA primer assay according to the manufacturer's instructions. RNU6B (miRNA) was used as an endogenous control. PPARg, ETS1, ET-1, PECAM1, and E-SEL mRNA levels in the same sample were determined and quantified using specific mRNA primers, as described previously (46). Levels of 9S mRNA were used as an endogenous control.

Western Blot Analysis

After treatment with RSG or vehicle, protein homogenates from mouse lungs or hemintreated HPAECs were subjected to Western blot analysis, as reported (46). Primary antibodies were purchased from Santa Cruz Biotechnology and included ET-1 rabbit polyclonal antibody (1:250 dilution, Cat. No. SC-98727, 24 kD), PPARg rabbit polyclonal antibody (1:500 dilution, Cat. No. SC-7196, 54 kD), ETS1 antibody (1:500 dilution, Cat. No. SC-350, 55 kD), PECAM1 (1:500 dilution, Cat. No. SC-8306, 130 kD), E-SEL (1:500 dilution, Cat. No. SC-14011, 115 kD), and CDK4 antibody (1:500 dilution, Cat. No. SC-260,

32 kD). Proteins were visualized using a peroxidase-coupled antigoat, antirabbit, or antimouse IgG in the presence of LumiGlo reagent (Thermo Scientific, Waltham, MA). Bands were identified by chemiluminescence, quantified by laser densitometry (Bio-Rad, Hercules, CA), and normalized to CDK4 levels within the same lane.

Statistical Analysis

For all measurements, data are presented as mean \pm SE. All data were analyzed using analysis of variance. Post hoc analysis used the Student-Newman-Keuls test to detect differences between specific groups. In studies comparing only two experimental

groups, data were analyzed with Student's t test to determine the significance of treatment effects. The level of statistical significance was taken as $P < 0.05$. Statistical analyses were performed using GraphPad Prism, Version 5.0 software (LaJolla, CA).

Results

Townes SS Mice Develop PH and RVH Spontaneously

Because PH is an important complication of SCD (64, 65) and is associated with an increased mortality risk (12–14), we measured RVSP and RVH in SS and AA

mice. As illustrated in Figure 1, Townes SS mice of either sex at 15–17 weeks of age spontaneously developed PH (Figure 1A) and RVH (Figure 1B). As reported previously (27), Townes SS mice showed decreased hematocrit values compared with AA mice (Figure 1C), whereas no differences in heart rate were observed between AA and SS mice (Figure 1D). To evaluate pulmonary vascular remodeling in SS and AA mice, α -smooth muscle actin–stained lung sections were used, and the number of muscularized vessels per square millimeter was counted. As illustrated in Figure 1E, pulmonary vascular remodeling was increased in SS compared with AA lungs.

Figure 1. Compared with littermate control (AA) mice, sickle cell (SS) mice had increases in right ventricular systolic pressure (RVSP) and right ventricular hypertrophy and reduced hematocrit. (A and D) AA and SS mice were anesthetized, and the right internal jugular vein of each mouse was surgically exposed and cannulated with a pressure transducer that was advanced to record RVSP and heart rate. Each bar represents the mean RVSP in mm Hg \pm SE. $P < 0.05$ versus AA (n = 6–13). (B) The right ventricular (RV) to left ventricular (LV) + septum (S) weight ratio [RV:(LV + S)] is presented as an index of right ventricular hypertrophy. *P < 0.05 versus AA (n = 6). (C) Hematocrits were measured as the volume of red blood cells in AA and SS mice whole blood. $*P < 0.05$ versus AA (n = 6-13). (E) The number of α smooth muscle actin (α SMA)–positive vessels (0-20, 21-30, 31-50, and 51-100 µm in luminal diameter) per square millimeter was determined. Each bar represents the mean \pm SE α SMA-positive vessels per square millimeter (n = 6). *P < 0.05 versus AA. Scale bar, 25 µM.

Alterations in PPAR γ and ET-1 in SS Mice and Hemin-Treated HPAECs

Because mounting evidence demonstrates that PPAR γ activation attenuates PH (46), whereas loss of PPAR_Y expression is associated with PH (47), we examined PPAR_y levels in SCD-PH. As illustrated in Figure 2A, $PPAR\gamma$ mRNA and protein levels were reduced in SS mouse lungs. Because ET-1 levels are increased in the plasma of patients with SCD (66–68) and because PPAR γ negatively regulates ET-1 (46), we measured ET-1 mRNA and protein levels in these same models. As expected, ET-1 expression was increased in SS compared with AA lungs (see Figure E1A in the online supplement). Circulating hemin levels are increased in SCD and they contribute to vascular endothelial dysfunction (69, 70), and plasma hemin concentrations in patients with SCD were \sim 4.2 µM compared with 0.2 µM in healthy control subjects (62). Hemin, released into the circulation during hemolysis, causes inflammation, oxidative damage, and endothelial dysfunction and contributes to SCD-PH pathogenesis (71). Furthermore, plasma cell free hemin levels are elevated in Townes SS mice (72) and they contribute to endothelial barrier dysfunction (73). Thus, we examined the dose-dependent $(0-10 \mu M)$ effects of hemin on HPAECs (Figure E2).

Treatment with hemin (5 μ M) was sufficient to reduce PPAR_y mRNA and protein levels (Figure 2B) and increase HPAEC ET-1 mRNA and protein levels (Figure E1B). Hemin treatment was therefore selected to further model SCDinduced endothelial signaling pathways in vitro.

To examine the role of endothelial dysfunction in SCD-PH pathogenesis, we examined levels of markers of endothelial dysfunction, including E-SEL (Figure E3A) and PECAM1 (Figure E3B). Each of these markers of endothelial dysfunction was increased in the lungs of SS mice compared with those of AA mice. In parallel, hemin treatment was sufficient to increase HPAEC mRNA and protein levels of these same markers of endothelial dysfunction in vitro (Figures E3C and E3D).

miR-27a Is Increased in SS Mice and Hemin-Treated HPAECs

To examine potential miRNAs that negatively regulate PPARg, a bioinformatics approach using multiple prediction algorithms (miRBase, PicTar, and TargetScan v6.1) was used to identify binding sites for miRNAs in the 3'UTR of PPAR_y. This analysis indicated miR-27a/b, miR-130a/b, miR-301a/b, and miR-454 as potential regulators of PPARg. Samples from SS and AA mouse lungs or hemintreated HPAECs were used to screen for alterations in these miRNAs. As illustrated in Figure 3A, of the miRNAs that were examined and predicted to regulate PPARg, miR-27a was increased in SS mouse lung and miR-130b, miR-301a/b, and miR-454 were selectively increased in SS mouse lung (Figure E4A). As illustrated in Figure 3B, compared with control conditions, hemin increased the levels of miR-27a and miR-27b only in HPAECs in vitro (Figure E4B). Because increased levels of miR-27a are predicted to reduce PPARg, and because miR-27a reduced lung PPARg levels in a hypoxia-induced model of PH (37), subsequent studies focused on miR-27a in SS mouse lung and hemin-treated HPAECs.

miR-27a Inhibition Reduces ET-1 Expression and Endothelial Dysfunction through PPAR γ Up-Regulation in Hemin-Treated HPAECs

We have reported previously that 50 nM anti–miR-27a reduced miR-27a levels by 50% and increased PPARg levels in HPAECs (37). To confirm the role of miR-27a in hemin-induced reductions in PPAR_y and increases in ET-1 and endothelial dysfunction, loss-of-miR-27a function was accomplished by transfecting HPAECs with an miR-27a inhibitor. Anti–miR-27a increased PPARg levels in control HPAECs and attenuated reductions in PPAR_y levels

Figure 2. Levels of peroxisome proliferator-activated receptor γ (PPAR γ) are reduced in SS mice and in hemin-treated human pulmonary artery endothelial cells (HPAECs). Whole lung homogenates were collected from SS and AA mice. (A) Lung PPARy levels were measured with quantitative RT-PCR (qRT-PCR) and/or Western blotting and are expressed relative to lung mRNA (9S mRNA) or CDK4. * $P < 0.05$ versus AA (n = 6). HPAECs were treated with dimethyl sulfoxide vehicle (CON) or hemin (HEM, 5 µM) for 72 hours. (B) Mean HPAEC PPAR_Y levels were measured with qRT-PCR and/or Western blotting. Each bar represents the mean PPARy level \pm SE expressed as fold change versus CON. *P < 0.05 versus CON (n = 6).

Figure 3. MicroRNA-27a (miR-27a) regulates PPARg in SS mice and in hemin-treated HPAECs. Anti–miR-27a reduces endothelin-1 (ET-1) and markers of endothelial dysfunction and increases PPAR_Y expression in hemin-treated HPAECs. (A) Whole lung homogenates were collected from SS and AA mice. qRT-PCR was performed on lung tissue for miR-27a predicted to regulate PPARg expression. Each bar represents the mean miR-27a level \pm SE relative to RNA, U6B small nuclear (RNU6B) and normalized to control values. * $P < 0.05$ versus AA ($n = 3-6$). (B) HPAECs were treated with dimethyl sulfoxide vehicle (CON) or hemin (HEM, 5 μ M) for 72 hours. *P < 0.05 versus CON (n = 3–6). HPAEC miR-27a was isolated and subjected to qRT-PCR analysis together with RNU6B. Each bar represents the mean miR-27a level \pm SE relative to RNU6B and normalized to control values. $*P < 0.05$ versus CON (n = 3–6). (C–F) HPAECs were treated with either scrambled or 50 nM anti–miR-27a for 6 hours and then treated with dimethyl sulfoxide vehicle (CON) or hemin (HEM, 5 µM) for 72 hours. HPAECs were then collected, and mRNAs were isolated. qRT-PCR was performed for (C) PPARg, (D) ET-1, (E) endothelial adhesion molecule 1 (E-SEL), and (F) platelet/endothelial cell adhesion molecule 1 (PECAM1) levels. Each bar represents the mean mRNA level \pm SE relative to ribosomal S9 (9S RNA) expressed as fold change versus CON. $P < 0.05$ versus CON; $P < 0.05$ versus HEM $(n = 6-9)$.

in hemin-treated HPAECs (Figure 3C). Anti–miR-27a also lowered HPAEC ET-1 mRNA levels in control cells and attenuated hemin-stimulated increases in ET-1 mRNA (Figure 3D). In addition, inhibition of miR-27a attenuated hemininduced increases in E-SEL and PECAM1 mRNA levels (Figures 3E and 3F).

ETS1 Knockdown Attenuates Increases in miR-27a and ET-1

To further examine the mechanisms underlying increases in miR-27a in SS mouse lungs and hemin-treated HPAECs, in silico analysis of the miR-27a promoter using PuTmiR software was performed (74). The upstream promoter region of miR-27a (chromosome X: 13,808,373–13,809,373 bp) contains a putative ETS binding site (5'-ACTTCCT-3'), suggesting that ETS may directly regulate transcriptional expression of miR-27. As shown in Figures 4A and 4B, ETS1 mRNA levels were increased in SS mouse lungs and hemin-treated HPAECs. In preliminary studies, we screened two different targeted regions, and subsequent studies focused on small interfering RNA targeting ETS1 transcript 1 (V1) that significantly decreased ETS1 levels (Figure E5). To further examine the role of ETS1 in hemin-induced miR-27a expression, RNA interference duplexes to ETS1 (10 nM) were used to directly reduce HPAEC ETS1 levels. Figures 4C and 4D illustrate that knockdown of HPAEC ETS1 decreased ETS1 mRNA and protein levels, but increased HPAEC PPARg mRNA and protein levels. The findings in Figures 4E–4G illustrate that depletion of ETS1 was sufficient to attenuate hemin-induced (1) increases in HPAEC miR-27a, (2) decreases in PPAR γ expression, and (3) increases in ET-1.

PPAR_y Regulates miR-27a and ET-1 Expression and Endothelial **Dysfunction**

We demonstrated previously that miR-27a and PPARg were mutually repressive in the hypoxic pulmonary vasculature (37). To further determine if alterations in PPAR γ expression are sufficient to regulate miR-27a, ET-1, and endothelial dysfunction, AdPPARg expression construct was used to examine PPARg gain of function in hemin-induced signaling. As shown in Figure E6, AdPPAR γ caused large and

dose-dependent increases in HPAEC PPAR_y protein levels. As shown in Figure 5, AdPPAR γ at 25 multiplicity of infection increased PPAR γ (Figure 5A) and decreased miR-27a (Figure 5B) and ETS1 (Figure 5C) mRNA levels in control cells. PPAR_y overexpression also prevented hemin-induced reductions in HPAEC PPAR_y and increases in miR-27a and ETS1 levels. Similarly, PPARg overexpression reduced ET-1, E-SEL, and PECAM1 mRNA levels in control HPAECs and attenuated hemin-induced increases in mRNA levels of these markers of endothelial dysfunction (Figures 5D–5F).

PPAR_y Activation Attenuates Increases in miR-27a and ETS1 Levels and Restores PPAR γ in SS Mouse Lungs and Hemin-Treated HPAECs

Because PPARg levels were reduced in the lungs of SS mice, and because, as we have reported previously, pharmacological PPAR_y activation attenuates increases in miR-27a levels in the lung and attenuates hypoxia-induced PH (37), we sought to examine the impact of PPAR_Y activation on miR-27a levels in SS mouse lungs and hemin-treated HPAECs. As shown in Figure 6, consistent with Figures 2 and 3, miR-27a was significantly increased in SS mouse lungs, whereas $PPAR\gamma$ levels were significantly reduced. Treating mice with the PPAR γ ligand, RSG, using a regimen that attenuated hypoxia-induced PH (32), attenuated increases in miR-27a and ETS1 mRNA levels and restored PPARg levels in SS mouse lungs (Figures 6A–6C). Similarly, in HPAECs treated with hemin for 72 hours, treatment with RSG during the last 24 hours attenuated hemin-induced increases in miR-27a and ETS1 and decreases in PPARg (Figures 6D–6F). To further examine phenotypic changes in hemin-treated HPAECs and the effect of RSG on hemin-induced cell proliferation, HPAEC proliferation was measured with MTT assays. Figure E7 shows that hemin significantly increased HPAEC proliferation and that RSG treatment attenuated hemin-induced HPAEC proliferation. Taken together, the findings in Figures 3–5 provide compelling evidence that miR-27a down-regulates $PPAR\gamma$ to stimulate ET-1 expression and endothelial dysfunction, and that activation of PPARg can negatively regulate miR-27a in these conditions.

Discussion

In the current study, we demonstrate, we believe for the first time, that (1) the Townes humanized SS mouse model spontaneously develops PH, RVH, and pulmonary vascular remodeling, (2) PPAR γ levels are reduced in lungs from SS mice and in hemin-treated HPAECs, and (3) miR-27a levels are increased in SS mouse lung and in hemin-treated HPAECs through pathways involving the ETS1 transcription factor. Furthermore, reductions in PPARg promote increased expression of ET-1 and markers of endothelial dysfunction in vivo and in vitro. Activation of PPAR_Y with the pharmacological ligand RSG or with adenoviral-mediated PPARg overexpression attenuates these derangements, suggesting that therapeutically targeting PPARg provides a novel strategy for reducing SCD-associated pulmonary endothelial dysfunction and PH (Figure 7).

ET-1 is elevated in both patients with SCD and those with PH and plays a key role in the pathophysiology of SCD and PH. Although several potential therapeutic strategies have been investigated in SCD-PH, none have significantly affected SCD-PH outcomes (11, 75–79). The importance of ET-1 in the pathogenesis of SCD-PH is well established (80–83). ET-1 causes endothelial dysfunction and stimulates vasoconstriction and migration and proliferation of pulmonary vascular wall cells in PH (81, 84). However, in clinical trials in SCD-PH, the ET-1 receptor antagonist, bosentan, failed to significantly reduce pulmonary vascular resistance (85). Thus, despite existing evidence that ET-1 plays an important pathophysiological role in SCD-PH pathogenesis, new therapeutic strategies that regulate ET-1 as well as other pathways involved in SCD-PH may lead to more effective and widely applicable therapeutic approaches for SCD-PH. We demonstrated previously that PPARg ligands favorably modulate several components of the ET-1 signaling pathway (46) and also attenuate other proliferative signaling pathways including nicotinamide adenine dinucleotide phosphate oxidase 4 (32), platelet-derived growth factor (32), and thrombospondin-1 (86) in hypoxiainduced PH pathogenesis. These findings indicate that the therapeutic potential of targeting $PPAR\gamma$ in SCD-PH merits

additional evaluation and led us to the current study.

Chronic hemolysis in SCD stimulates steady-state inflammation (87, 88) through subclinical vascular endothelial injury and transient vasoocclusive events (70, 89). The vascular endothelium in the lung and other organs senses these SCD-related stimuli including hypoxia, ischemia-reperfusion, hemoglobin, and hemin. Studies in SS mice demonstrate a role for adhesion molecules in the interaction of erythrocytes with the endothelium (22, 90–93). Furthermore, adhesion molecules are increased in SCD and are associated with PH and mortality (21). Current evidence suggests that SCDrelated hemolysis and its byproducts, hemoglobin and hemin, induce endothelial dysfunction and PH in SCD. We postulated that these hemolysis-induced derangements in endothelial function play an important role in overall pulmonary vascular dysfunction, transducing signals from the blood to other cellular compartments of the vascular wall (72, 73). Our findings are consistent with these concepts and demonstrate that endothelial adhesion molecules, including E-SEL and PECAM1, are increased significantly in the lungs of SS mice and in hemin-treated HPAECs (Figure E3). Hemin triggers vasoocclusion through increases in and activation of these adhesion molecules (90). The current studies demonstrate that PPAR_y activation attenuates endothelial dysfunction in the SS mouse lung and in hemin-treated HPAECs (Figures 5 and 6), supporting the concept that therapeutic targeting of this receptor provides a novel strategy for reducing SCD-PH. In addition, unpublished studies by our group indicate that PPARg activation may also exert direct and beneficial effects on the myocardium in experimental models of PH.

We reported recently that hypoxiainduced PH was associated with reductions in PPAR γ , and that activating the PPAR γ receptor attenuates hypoxia-induced reductions in PPAR γ in mouse lungs and HPAECs (32, 46). The current study extends these observations by providing novel evidence that PPARg is reduced in the lungs of SS but not AA mice. Because treatment with pathobiologically relevant concentrations of hemin was sufficient to reduce PPARγ in HPAECs in vitro (Figure 1), these findings suggest that products of intravascular hemolysis trigger these reductions in PPARg. Furthermore,

Figure 4. ETS1 knockdown attenuates increases in miR-27a and ET-1 levels and decreases in PPAR_Y expression in hemin-treated HPAECs. (A) Whole lung homogenates were collected from SS and AA mice. Lung ETS1 levels were measured with qRT-PCR and expressed relative to lung 9S mRNA. * P < 0.05 versus AA ($n = 6$). (B) HPAECs were treated with dimethyl sulfoxide vehicle (CON) or hemin (HEM, 5 μ M) for 72 hours. Mean HPAEC ETS1 levels were measured with qRT-PCR. Each bar represents the mean ETS1 level \pm SE relative to 9S expressed as fold change versus CON. *P < 0.05 versus CON (n = 5). (C–E) HPAECs were treated with 10 nM scrambled (CON) or small interfering RNA duplexes to ETS1 for 72 hours. HPAEC mRNA or protein was then isolated. Each bar represents the mean \pm SE (C) ETS1 or (D) PPAR_Y relative to 9S in the same sample expressed as fold change versus CON. $*P < 0.05$ versus CON ($n = 3$). Western blotting was used to detect (C) ETS1 and (D) PPAR γ protein levels. Representative blots depicting ETS1 or PPAR γ protein in siETS1-treated HPAECs are shown. (E-G) HPAECs were treated with 10 nM scrambled (CON) or small interfering RNA duplexes to ETS1 for 6 hours and then treated with dimethyl sulfoxide vehicle (CON) or hemin (HEM, 5 µM) for 72 hours. MicroRNAs (miRNA) or mRNA were isolated and subjected to qRT-PCR analysis for miR-27a, PPAR γ , ET-1, and RNU6B or 9S. Each bar represents the mean \pm SE (E) miR-27a, (F) PPAR γ , or (G) ET-1 relative to RNU6B or 9S expressed as fold change versus CON. * $P < 0.05$ versus CON; $^+P < 0.05$ versus HEM (n = 3–6). ETS1, v-ets avian erythroblastosis virus E26 oncogene homolog 1; SCR, scrambled small interfering RNA; siETS1, ETS1 small interfering RNA.

Figure 5. PPAR_Y gain-of-function attenuates miR-27a, ETS1, and ET-1 expression and endothelial dysfunction in hemin-treated HPAECs. HPAECs were transfected with adenovirus containing a PPAR_Y plasmid (AdPPAR_Y, 25 multiplicity of infection) or green fluorescent protein (AdPPAR_Y-) constructs for 6 hours and were then treated with dimethyl sulfoxide vehicle (CON) or hemin (HEM, 5 µM) for 72 hours. miRNA or mRNA were isolated and subjected to qRT-PCR analysis ($n = 3-9$). (A) HPAEC PPAR γ mRNA levels are expressed relative to 9S in the same sample as fold change versus CON. (B) HPAEC miR-27a levels are expressed relative to RNU6B as fold change versus CON. HPAEC (C) ETS1, (D) ET-1, (E) E-SEL, and (F) PECAM1 mRNA levels are expressed relative to 9S as fold change versus CON/AdPPAR γ -. All bars represent mean \pm SE. *P < 0.05 versus CON; ^+P < 0.05 versus HEM.

our findings indicate that $PPAR\gamma$ activation restores $PPAR\gamma$ in SS mouse lung and in hemin-treated HPAECs (Figures 5 and 6), supporting the concept that stimulating PPAR_y may provide a novel therapeutic strategy in SCD-PH.

We demonstrated previously that hypoxic exposure sufficient to induce PH in vivo and pulmonary artery EC proliferation in vitro increased miR-27a and reduced PPAR γ (37). Furthermore, PPAR_y activation attenuated hypoxiainduced reductions in PPAR γ by attenuating hypoxia-induced increases in miR-27a levels. Emerging evidence supports the role of miRNAs in PH pathogenesis (94). Both miR-648 (95) and miR-199a2 (96) were reported to contribute to SCD pathogenesis in the Berkeley SS mouse. Our findings extend these observations to suggest that miR-27a also contributes to SCD-PH. Levels of miR-27a were increased in the lungs of SS mice and hemin-treated HPAECs (Figure 3). Anti–miR-mediated inhibition of miR-27a (Figure 3), activation of PPAR γ with RSG (Figure 6), or overexpression of PPARg (Figure 5) attenuated increases in miR-27a levels in SS mouse lung and HPAECs. These results provide novel evidence that PPAR_Y activation suppresses increases in miR-27a in SCD, which is consistent with previous reports of the repressive effect of PPAR_Y activation on miR-27a under hypoxic conditions (37).

To further explore the mechanisms for increases in miR-27a levels in SCD, we

performed an in silico analysis of the upstream promoter region of miR-27a and identified putative ETS1 binding sites. Recent studies reported that hemin increases ETS1 mRNA levels in human erythroleukemic cell lines (97), that dominant-negative loss of $PPAR\gamma$ function increases ETS1 expression (39), and that PPAR_y activation attenuates ETS1 activity (36). Therefore, to further investigate whether ETS1 activates the miR-27a promoter in SCD and whether it might also be regulated by PPARg, we examined the effects of PPARg activation on ETS1 in SS mouse lung and in hemin-treated HPAECs. The importance of ETS1 was supported by evidence that small interfering RNAmediated ETS1 depletion attenuated

Figure 6. The PPAR_Y ligand rosiglitazone (RSG) attenuates increases in miR-27a and ETS1 levels and reductions in PPAR_Y in SS mouse lung and hemin-treated HPAECs. (A) Whole lung homogenates were collected from AA and SS mice. Selected AA and SS mice were gavaged with RSG (10 mg/kg/d) or vehicle for 10 days. qRT-PCR was performed on AA and SS lung tissues. Lung miR-27a levels are expressed relative to RNU6B and normalized to control values. Each bar represents the mean \pm SE. *P < 0.05 versus AA/-RSG; *P < 0.05 versus SS/-RSG, (n = 3-6). Lung (B) PPAR γ and (C) ETS1 mRNA levels were measured with qRT-PCR and expressed relative to lung 9S mRNA levels. Each bar represents the mean PPARy and ETS1 mRNA level \pm SE expressed as fold change versus AA/-RSG. $*P$ < 0.05 versus AA/-RSG; $*P$ < 0.05 versus SS/-RSG (n = 3–6). HPAECs were treated with dimethyl sulfoxide vehicle (CON) or hemin (HEM, 5 μ M) for 72 hours. During the final 24 hours of hemin exposure, selected HPAECs were treated with or without RSG (10 µM). HPAECs were then collected, and miRNAs or total RNAs were isolated. qRT-PCR was performed for (D) miR-27a, (E) PPARy, and (F) ETS1 levels. Each bar represents the mean \pm SE miR-27a, PPARy, or ET-1 level relative to RNU6B or 9S as indicated. $*P < 0.05$ versus CON/-RSG; $+P < 0.05$ versus HEM/-RSG (n = 3–6).

hemin-induced (1) increases in HPAEC miR-27a and ET-1 and (2) decreases in PPAR_Y expression (Figure 4). Taken together, these novel findings indicate that hemin and perhaps other stimuli in SCD can activate ETS transcription factors to increase miR-27a levels. As reported previously (37), miR-27a reduces PPARg, which attenuates ET-1 levels and prevents endothelial dysfunction. Because of the mutually repressive relationship between miR-27a and PPAR γ (37), our findings suggest that stimulating the PPAR γ receptor with exogenous $PPAR\gamma$ ligands can reverse pathogenic programs of gene expression in SCD. These findings highlight

a previously unrecognized pathobiological cascade in SCD that may be amenable to therapeutic intervention.

This study has several important limitations. First, although the current study demonstrates that hemin increases ETS1 and miR-27a levels, it does not directly address how hemin increases ETS1 levels. ETS1 knockdown attenuated hemininduced increases in miR-27a levels and increased PPARg mRNA and protein levels in HPAECs. These findings indicate that ETS1 stimulates miR-27a expression in HPAECs and suggest that PPAR γ activation could attenuate ETS1 and miR-27a promoter activity through

transrepression. Therefore, additional studies will be required to further define how hemin directly stimulates ETS1. Second, although discordant patterns of miRNA expression have been reported in experimental rodent models and in human PH (98), reductions in PPAR_y have been reported in several experimental models of PH (28, 30, 32, 33, 46, 99), in human PAH lung tissue (33), and in pulmonary artery ECs isolated from patients with idiopathic pulmonary arterial hypertension (37). These findings suggest that common mechanisms that regulate PPAR γ during PH pathogenesis in multiple models and species merit additional investigation and

Figure 7. Schema of putative pathways increasing miR-27a and reducing PPAR_Y that increase ET-1 and endothelial dysfunction in sickle cell disease (SCD). The current findings suggest that products of hemolysis in SCD, such as hemin, activate the transcription factor ETS1 to stimulate the expression of miR-27a. Increases in miR-27a reduce PPARg, leading to increased ET-1, endothelial dysfunction, and pulmonary hypertension (PH). Stimulation of PPAR_Y with pharmacological ligands such as RSG attenuates increases in miR-27a and downstream ET-1 levels and endothelial dysfunction, suggesting that this pathway may provide an opportunity for new therapeutic targeting and intervention in SCD-PH.

may provide new insights into the pathobiology and therapy of PH. Finally, the current study used RSG as a pharmacological agonist to stimulate PPAR γ activity. RSG is a TZD PPAR γ ligand currently used in the clinical management of patients with type 2 diabetes. However, RSG, as well as other drugs in this class, has been recognized to have significant adverse effects in patients with diabetes (100). Cardiovascular responses elicited by different TZD PPARg ligands in human subjects are distinct. Furthermore, it is difficult to extrapolate the therapeutic effects of drugs across different groups of PH (101). Thus, although the current study, together with other reports, suggests therapeutic benefits of pharmacological PPARg activation in PH, the potential of this therapeutic strategy will require additional study and more refined pharmacological tools that activate PPARg without causing adverse effects.

Conclusions

To our knowledge, the current study of SCD-PH pathogenesis provides the first evidence that miR-27a reduces $PPARv$ expression and increases ET-1 expression and endothelial dysfunction through posttranscriptional mechanisms in vivo and in vitro. Furthermore, this study provides novel evidence that ETS1 mRNA levels were increased in SS mouse lungs and in hemin-treated HPAECs and that ETS1 loss of function was sufficient to attenuate hemin-induced increases in HPAEC miR-27a and ET-1 and decreases in PPARg expression. These results suggest that targeting PPARg activation may represent a novel therapeutic approach in SCD-PH pathogenesis. \blacksquare

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