

Reduction of ARNT in myeloid cells causes immune suppression and delayed wound healing

Christopher Scott,^{1,2} James Bonner,³ Danqing Min,³ Philip Boughton,^{4,5} Rebecca Stokes,¹ Kuan Minn Cha,¹ Stacey N. Walters,⁶ Kendle Maslowski,⁷ Frederic Sierro,⁸ Shane T. Grey,⁶ Stephen Twigg,^{2,3} Susan McLennan,^{2,3} and Jenny E. Gunton^{1,2,9,10}

¹Diabetes and Transcription Factors Group, Department of Immunology and Inflammation, Garvan Institute of Medical Research, Sydney, New South Wales, Australia; ²Faculty of Medicine, University of Sydney, Sydney, New South Wales, Australia; ³Department of Endocrinology, Royal Prince Alfred Hospital, Sydney, New South Wales, Australia; ⁴St. George Clinical School, St. George Hospital, Kogarah, New South Wales, Australia; ⁵Department of Biomedical Engineering, University of Sydney, Sydney, New South Wales, Australia; ⁶Department of Immunology and Inflammation, Garvan Institute of Medical Research, Sydney, New South Wales, Australia; ⁷Department of Biochemistry, University of Lausanne, Lausanne, Switzerland; ⁸Liver Immunology, Centenary Institute, Sydney, New South Wales, Australia; ⁹St. Vincent's Clinical School, University of New South Wales, Sydney, New South Wales, Australia; and ¹⁰Department of Diabetes and Endocrinology, Westmead Hospital, Sydney, New South Wales, Australia

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Scott C, Bonner J, Min D, Boughton P, Stokes R, Cha KM, Walters SN, Maslowski K, Sierro F, Grey ST, Twigg S, McLennan S, Gunton JE. Reduction of ARNT in myeloid cells causes immune suppression and delayed wound healing. *Am J Physiol Cell Physiol* 307: C349–C357, 2014. First published July 2, 2014; doi:10.1152/ajpcell.00306.2013.—Aryl hydrocarbon receptor nuclear translocator (ARNT) is a transcription factor that binds to partners to mediate responses to environmental signals. To investigate its role in the innate immune system, floxed ARNT mice were bred with lysozyme M-Cre recombinase animals to generate lysozyme M-ARNT (LAR) mice with reduced ARNT expression. Myeloid cells of LAR mice had altered mRNA expression and delayed wound healing. Interestingly, when the animals were rendered diabetic, the difference in wound healing between the LAR mice and their littermate controls was no longer present, suggesting that decreased myeloid cell ARNT function may be an important factor in impaired wound healing in diabetes. Deferoxamine (DFO) improves wound healing by increasing hypoxia-inducible factors, which require ARNT for function. DFO was not effective in wounds of LAR mice, again suggesting that myeloid cells are important for normal wound healing and for the full benefit of DFO. These findings suggest that myeloid ARNT is important for immune function and wound healing. Increasing ARNT and, more specifically, myeloid ARNT may be a therapeutic strategy to improve wound healing.

aryl hydrocarbon receptor nuclear translocator; hypoxia-inducible factor-1 α ; deferoxamine

THE INNATE IMMUNE SYSTEM functions as the first line of defense against infection. Mononuclear phagocytes and neutrophils of the myeloid lineage are key effector cells of the innate immune system that function in the elimination of pathogens, modulation of the adaptive immune system, and tissue repair. Aryl hydrocarbon receptor (AhR) nuclear translocator (ARNT) is a transcription factor of the basic helix-loop-helix Per/ARNT/Sim (bHLH-PAS) family (19). ARNT heterodimerizes with class II bHLH/PAS family members, including hypoxia-inducible factor (HIF)-1 α , HIF-2 α , and AhR, to form active tran-

scription complexes that regulate genes involved in hypoxic responses, cell survival, proliferation, glycolysis, angiogenesis, inflammation, and response to xenobiotics (13, 22, 39, 51).

HIF-1 α and HIF-2 α are regulated by cellular oxygen content; at normal oxygen concentrations, these proteins are rapidly degraded by the ubiquitin-proteasome pathway (15). HIF-1 α can be stabilized by inflammation, transforming growth factor (TGF), platelet-derived growth factor (PDGF), epidermal growth factor, insulin-like growth factor I, and interleukin (IL)-1 β (7, 19, 37, 56) and in human fibroblasts, and in diabetic animals its stability and activity are reduced by high glucose concentrations (1, 29, 43). Hypoxia-induced HIF-1 α protein stabilization leads to differential gene transcription compared with that induced by Toll-like receptor agonists, showing that the combination of stimuli leading to HIF-1 α stabilization is important in determining subsequent actions (18).

AhR is activated by ligand binding, which leads to its binding to ARNT, translocation to the nucleus, and gene regulation (19). Exogenous ligands are numerous and include the aromatic hydrocarbons, including 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD) and benzo[a]pyrene (8, 9). There is disagreement regarding the endogenous ligands for AhR: bile acids and cAMP are candidates, as are breakdown products from cruciferous vegetables (26, 35, 57). Importantly, different ligands have different effects; for example, TCDD and 6-formylindolo[3,2-b]carbazole produce opposing effects in experimental autoimmune encephalomyelitis (8, 38, 49).

The importance of HIF-1 α in myeloid cell function was shown by Cramer et al. (7), who reported decreased immune function. HIF-1 α regulates apoptosis, phagocytosis, and bacterial killing in macrophages and neutrophils (7, 36, 50). Imtiyaz et al. (16) showed that HIF-2 α also regulates immune responses in a similar direction to HIF-1 α , with myeloid cell knockout animals and isolated macrophages displaying reduced inflammatory cytokine production and decreased inflammation in models of immune function.

In contrast, deletion of AhR in macrophages leads to increased production of inflammatory cytokines after lipopolysaccharide (LPS) treatment (21), and AhR knockout animals

Address for reprint requests and other correspondence: J. E. Gunton, Rm. 2040, Clinical Sciences Corridor, Westmead Hospital, Sydney, NSW 2145, Australia (e-mail: jenny.gunton@sydney.edu.au and j.gunton@garvan.org.au).

are sensitive to LPS-induced shock (40). The role of macrophage and neutrophil ARNT in models of immune function and wound healing has not been examined. Since ARNT inactivation would block the effects of HIF-1 α , HIF-2 α , and AhR, it was not possible to predict the outcomes from the preexisting studies. To investigate its role in the innate immune system, floxed ARNT mice were bred with lysozyme M (LysM)-Cre recombinase (Cre) animals to generate LysM-ARNT (LAR) mice with reduced ARNT.

These studies demonstrate altered mRNA expression of cytokines and genes involved in wound healing in myeloid cells of LAR mice. LAR mice also had delayed wound healing. When the animals were rendered diabetic, the difference in wound healing between the LAR mice and their littermate controls disappeared.

MATERIALS AND METHODS

Animal Studies

All animals received humane care according to the criteria outlined in the "Australian Code of Practice for the Care and Use of Animals for Scientific Purposes." All procedures were approved by the Garvan Animal Ethics Committee. Floxed ARNT mice were created as previously described (13, 46) and bred with LysM-Cre mice (5) to produce LAR mice with reduced ARNT and floxed-control (FC) offspring. All mice were on an inbred C57BL/6 background for ≥ 12 generations.

Investigation of Effect of ARNT Depletion on Myeloid Cell Function

To obtain thioglycollate-stimulated macrophages, male mice were euthanized by cervical dislocation 4 days after peritoneal injection of 2 ml of 3% thioglycollate (Difco, Melbourne, Australia). Macrophages were isolated by injection of 10 ml of sterile ice-cold PBS into the peritoneum. Cells were cultured in RPMI medium (Gibco, Melbourne, Australia) with 10% FCS and 1% L-glutamine. After 2 h, the cells were washed twice with PBS, and the cells remaining were used in experiments. Cells were stimulated with LPS (LPS:B4, 100 ng/ml; Sigma, Castle Hill, Australia). After 24 h, the cells were lysed for RNA ($n = 5$ /group). For Western blot analysis, macrophages were cultured for 72 h ($n = 4$ /group).

Investigation of Effect of ARNT Depletion on Inflammatory Response

Skin irritation study. Mice were anesthetized, and hair was removed. After 3 days of rest, 5% sodium dodecyl sulfate (SDS) was painted onto the skin twice daily for 5 days, and the animals were euthanized ($n = 5$ –6/group). Skin was fixed in formalin for histology, and the inflammatory response was scored by two independent observers blinded to genotype using a semiquantitative scale of inflammation that included signs of infiltrate, edema, and epidermal proliferation. A scale of 0–4, with 0 being no skin inflammation and 4 being maximal inflammation, was used.

Skin transplant study. Mice were anesthetized and shaved. Skin grafts were collected from Balb/c mice (H-2d) and transplanted onto LAR or FC (C57BL/6, H-2b) mice. Grafts were sex-matched but full-major histocompatibility complex-mismatched. The survival curve of 16 grafts onto FC mice and 13 grafts onto LAR female mice is shown. Transplant performance was scored by observers blinded to mouse genotype until complete rejection.

Wound healing. Female mice at 20 wk of age were anesthetized and shaved, and hair was removed. On the next day, the mice were again anesthetized, and 0.5×0.5 cm wounds were created in the superficial skin layer. Wounds were covered with a sterile bandage and checked

daily. *Day 0* was taken as the day following wounding, and bandages were removed from the mice at *day 6* ($n = 20$). Wound tissue was obtained from a second cohort of mice ($n = 6$) at *day 4* for RNA and histological studies. For RNA collection, the wound center was collected using a 4-mm biopsy punch, and samples were homogenized in RLT buffer (Qiagen, Valencia, CA). For determination of tensile strength, a third cohort of LAR mice ($n = 6$) were euthanized at *day 18* postwounding, and the tissue containing the wound was excised for assessment of tensile strength and calculation of Young's modulus, as previously described (45).

Diabetes was induced by streptozotocin injection, as previously reported (41). Wound experiments were initiated once mice had random blood glucose levels of >15 mmol/l for 3 consecutive days. Because of differences in blood glucose levels between FC and LAR mice, only animals with blood glucose levels ≥ 28 mmol/l and no animals with blood glucose levels <15 mmol/l, before insulin, were included ($n = 12$ –20/group).

For DFO experiments, 10-wk-old female mice were rendered diabetic by alloxan injection (41), and wounding experiments were commenced once mice had a random blood glucose level of >15 mmol/l for 3 consecutive days ($n = 8$ –10). Twenty microliters of $0.0125 \mu\text{M}$ DFO were added drop-wise to wounds from *day 0* every 2nd day until wounds were healed.

Gene Expression

Total RNA was isolated and cDNA was synthesized as described elsewhere (3, 24). Real-time PCR was performed using specific primers and SYBR Green PCR master mix (Applied Biosystems, Melbourne, Australia), and amplification was performed using a light cycler (model 7900, Applied Biosystems). Results were corrected for expression of the housekeeping gene TATA box-binding protein (*Tbp*), which did not differ between groups (data not shown). Primers are shown in Table 1.

Western Immunoblotting

Sonicated protein (25 μg) was run on a 10% SDS-polyacrylamide gel in Western SDS running buffer using a Bio-Rad Protein 3 apparatus, as previously described (13). A 1:500 dilution of ARNT (BD Bioscience Pharmingen, Australia) or a 1:1,000 dilution of α -tubulin (Abcam, San Francisco, CA) primary antibody was used. α -Tubulin was used as a loading control, and signal intensity was quantitated by densitometry using ImageJ to calculate ARNT protein level relative to α -tubulin level.

Histology

Tissue was dissected from FC and LAR mice and fixed in 10% buffered formalin. Sections (5 μm) were stained with hematoxylin and eosin or Milligan's trichrome according to standard protocols, as previously described (3).

Human Monocyte Isolation

Informed consent was obtained from all participants. The study was approved by the Human Ethics Committee of Royal Prince Alfred Hospital. Blood (30 ml) was obtained from patients with a >10 -yr duration of diabetes attending the Diabetes Centre of Royal Prince Alfred Hospital and from nondiabetic subjects ($n = 9$). Mononuclear cells were isolated using OptiPrep (Sigma, St. Louis, MO). The yield was 4 – 726×10^4 peripheral blood mononuclear cells/ml. Flow cytometry using CD14 as a monocyte marker showed $>75\%$ purity (53). A total of 2 μg of RNA from each sample were transcribed to cDNA using oligo(dT)₁₈ (25 pmol) and SuperScript III RNase H[−] reverse transcriptase (Invitrogen). Gene expression analysis was performed as described elsewhere (13). Within the diabetic group, the majority of the patients had type 2 diabetes (T2D) and one patient had type 1 diabetes. Average hemoglobin A1c was 7.7 ± 0.2 . The age of

Table 1. Real-time PCR primers

mRNA	Forward Primer	Reverse Primer
<i>Mouse</i>		
<i>Tbp</i>	atgatgactgcagcaaatcg	tatcactcctgccacaccag
<i>Arnt</i>	tctccctcccagatgatgac	caatgttggtgcgggagatg
<i>Elastin</i>	atcctcttgctcaacctcct	gccccggataaatagactcc
<i>Mmp9</i>	gaaggcacaacctgtgtgtt	agagtactgcttgcccagga
<i>Timp-1</i>	aggtaggtctcgttgatttct	gtaaggcctgtagctgtgcc
<i>Col1a1</i>	taggccattgtgtatgcagc	acatgttcagctttgtggacc
<i>α-Sma</i>	gagaagccagccagtcg	ctcttgctctgggcttca
<i>Tnf-α</i>	ccagaccctcactagatca	cacttggtggtttgctacgac
<i>Mcp-1</i>	ggtcctctgcatgcttctgg	cctgctgctggtgacacctt
<i>Tgf-β1</i>	tgagtggctgtcttttgacg	ggttcatgtcatggatgggtg
<i>Il-6</i>	ccagagatacaaaagaaatgatgg	actccagaagaccagaggaaat
<i>Cxcl1</i>	tggctgggattcacctcgaa	tatgacttcgggttggtgcag
<i>F4/80</i>	ctttggctatgggcttcacgtc	gcaaggaggacagagtttatcgtg
<i>Human</i>		
<i>ARNT</i>	aacctcacttcgtgggtggtc	caatgttggtgcgggagatg
<i>TBP</i>	gttgagttgcagggtgtgg	ctcaaaccaacttgctcaacagc

Tbp, TATA box-binding protein; *Arnt*, aryl hydrocarbon receptor nuclear translocator; *Mmp9*, matrix metalloproteinase 9; *Timp-1*, tissue inhibitor of matrix metalloproteinase; *Col1a1*, collagen type 1α; *α-Sma*, α-smooth muscle actin; *Mcp-1*, monocyte chemoattractant protein 1; *Tgf-β1*, transforming growth factor-β1; *Cxcl1*, chemokine (C-X-C motif) ligand 1.

patients was significantly different: 52 ± 3 (control) vs. 67 ± 2 (diabetic) yr. There was no significant difference in weight (75 ± 5 vs. 82 ± 3 kg) or body mass index (27 ± 2 vs. 30 ± 1).

Human Cytokine Assays

The circulating concentrations of proinflammatory cytokines and chemokines [IL-6, IL-8, tumor necrosis factor-α (TNF-α), and monocyte chemoattractant protein-1 (MCP-1)] were measured using the BioPlex Pro Assay (Bio-Rad, Hercules, CA). Intra- and interassay coefficients of variation were 7–10% for all assays.

Statistics

Significance was calculated with Excel or Prism (version 5). Student's 2-tailed *t*-test was used unless otherwise specified. For time courses, repeated-measure ANOVA (rmANOVA) was used. Log-rank (Mantel-Cox) test was used to assess graft survival. Pearson's correlations (*r*) were calculated, and *P* values from two-tailed analysis are shown. Values are means \pm SE. *P* < 0.05 was considered significant.

RESULTS

Thioglycollate-Isolated LAR Macrophages Had Reduced Arnt and Decreased Cytokine mRNA After LPS Treatment

The number of macrophages isolated after thioglycollate stimulation from LAR mice was similar to controls: 14.7×10^6 and 14.3×10^6 from FC and LAR mice, respectively (*P* > 0.9). *Arnt* mRNA was reduced by 50% in macrophages from LAR mice after 24 h of LPS treatment (Fig. 1A). This relatively modest deletion efficiency is consistent with the percentage of enhanced green fluorescent protein (EGFP) expression in peritoneal macrophages isolated from lysozyme-EGFP mice (60%) and blood monocytes isolated from LysM-Cre \times Rosa26-stop flox EGFP mice (55–75%) in previous studies (10, 17). Macrophages from LAR mice had significantly reduced expression of chemokine (C-X-C motif) ligand 1 (*Cxcl1*), *Mcp-1*, *Tgf-β1*, and *Il-6* (*P* < 0.005 for all). There was no change in *Tnf-α* mRNA (*P* = 0.2; Fig. 1A). To further investigate the efficiency of ARNT reduction in macrophages, we extracted protein from thioglycollate-elicited macrophages after 72 h of treatment

with LPS and assessed ARNT protein by Western immunoblotting (Fig. 1B). We found after LPS treatment that ARNT protein was significantly reduced to 13.5% of FC levels (*P* = 0.0012; Fig. 1C).

Cutaneous Inflammatory Response Was Reduced in LAR Animals Following SDS Treatment

Macrophages play an important role in the inflammatory responses in the skin, including response to irritation induced by SDS, so the response of LAR mice to this model was investigated (44). As expected, FC animals showed erythema and keratosis (Fig. 1D, top). These responses were markedly reduced in LAR animals (Fig. 1D, bottom). Inflammatory changes in FC animals were marked by vasodilation, epidermal hyperproliferation, and edema and were reduced in LAR animals (Fig. 1E, top and bottom). Histological scores of inflammation averaged from results of two observers blinded to genotype are shown in Fig. 1F. Scores were lower in LAR than FC mice, indicating impaired response to skin irritants (*P* = 0.0013).

LAR Mice Had Delayed Skin Transplant Rejection

Transplant rejection is immune-mediated, and mononuclear phagocytes and neutrophils are important for this process (23, 42, 55). Skin graft survival after transplantation was studied. There was a significant delay in rejection of Balb/c skin transplants (full major histocompatibility complex mismatch, LAR and FC mice are C57BL/6) by LAR mice compared with FC mice (*P* < 0.006 by Mantel-Cox test; Fig. 1G).

LAR Mice Had Delayed Wound Healing

Mononuclear phagocytes and neutrophils are important to wound healing in nonsterile conditions (12, 31, 33). Given the observation that skin healing after graft rejection was delayed in LAR animals, wound healing was formally assessed. LAR mice had significantly delayed wound healing [*P* = 0.0017 (overall) by rmANOVA, *P* = 0.0023 at day 2, *P* < 0.001 at day 4, *P* = 0.027 at day 6, and *P* = 0.0038 at day 8; Fig. 2A].

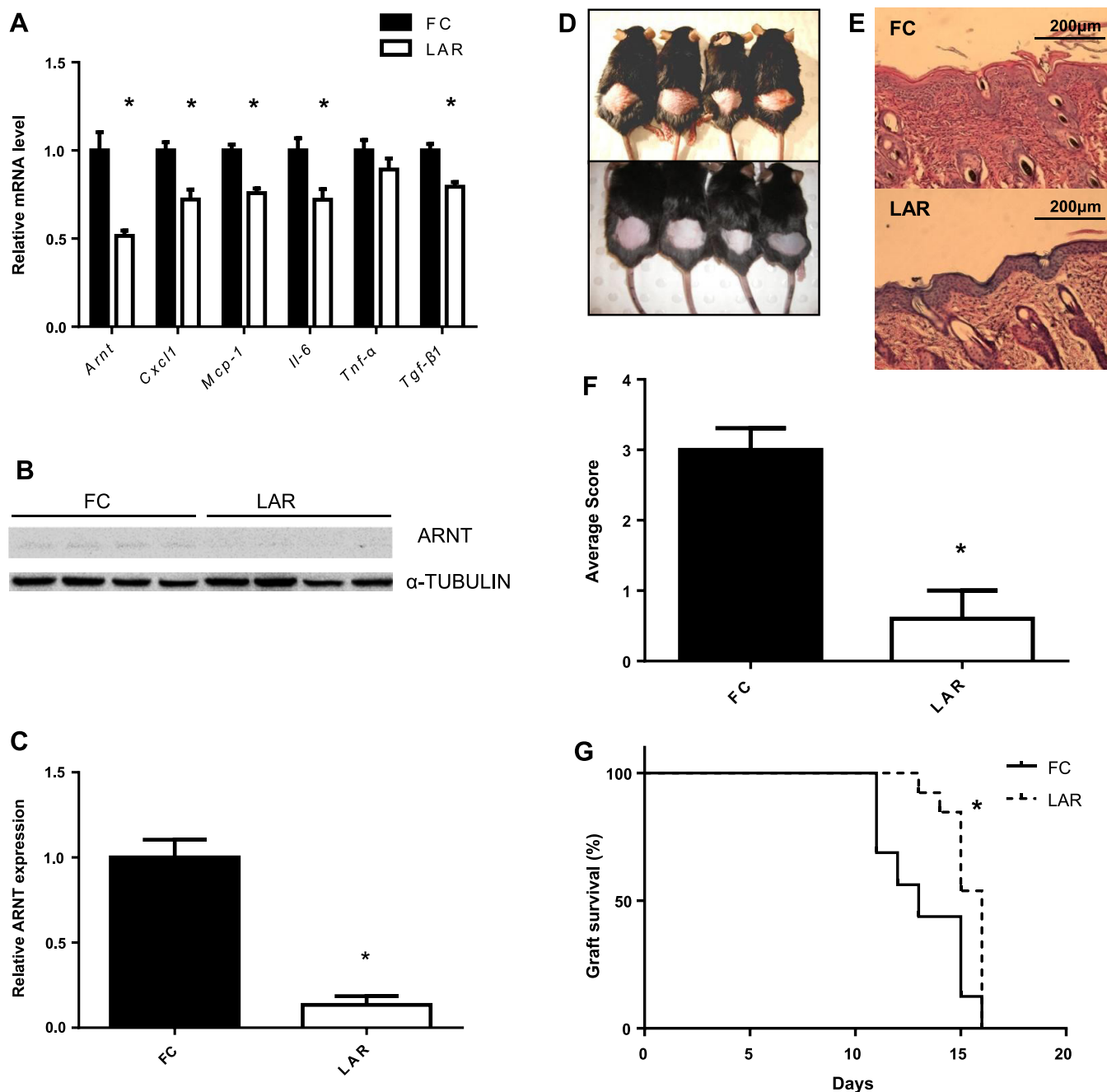


Fig. 1. Acute inflammatory response. **A**: aryl hydrocarbon receptor (AhR) nuclear translocator (ARNT) and cytokine mRNA expression in thioglycollate-elicited macrophages after 24 h of culture in RPMI medium containing 100 ng/ml LPS in floxed-control (FC) and lysozyme M-ARNT (LAR) animals. **B**: Western blot of ARNT in thioglycollate-elicited macrophages from FC and LAR mice after 72 h of LPS treatment. α -Tubulin is shown as a loading control. **C**: densitometry of ARNT protein expression in thioglycollate-elicited macrophages after 72 h of LPS treatment. **D**: FC (top) and LAR (bottom) mice treated with SDS. **E**: hematoxylin-stained section of skin from FC and LAR mice after SDS study. Magnification $\times 100$. **F**: histological inflammation score in FC and LAR mice treated with SDS. Values are means \pm SE. * $P < 0.05$ by Student's *t*-test (**A**, **C**, and **F**). **G**: survival curve of Balb/c grafts on FC and LAR. * $P < 0.05$ by log-rank (Mantel-Cox) test (**G**).

Final median tensile strength was not significantly altered at day 18 (Fig. 2B). Hematoxylin-eosin and Milligan's trichrome staining of wounds at day 4 showed no obvious histological changes (Fig. 2, C–H).

ARNT Deletion Altered Gene Expression in Wounds

In a separate cohort of mice, wound tissue was collected at day 4 of healing to measure gene expression. Cytokine expres-

sion in LAR animals was reduced. *Mcp-1* mRNA was reduced to 32% ($P < 0.0001$), *Il-6* was reduced to 14% ($P = 0.02$), and the key tissue-remodeling gene matrix metalloproteinase 9 (*Mmp9*) was reduced to 44% ($P = 0.004$). *Tnf-α* was decreased to 39% of control, and *Tgf-β1*, collagen type 1α1 (*Colla1*), α -smooth muscle actin (α -*Sma*), tissue inhibitor of metalloproteinase 1 (*Timp-1*), and *elastin* were also decreased (all non-parametrically distributed, $P < 0.05$; Fig. 3, A and B). No

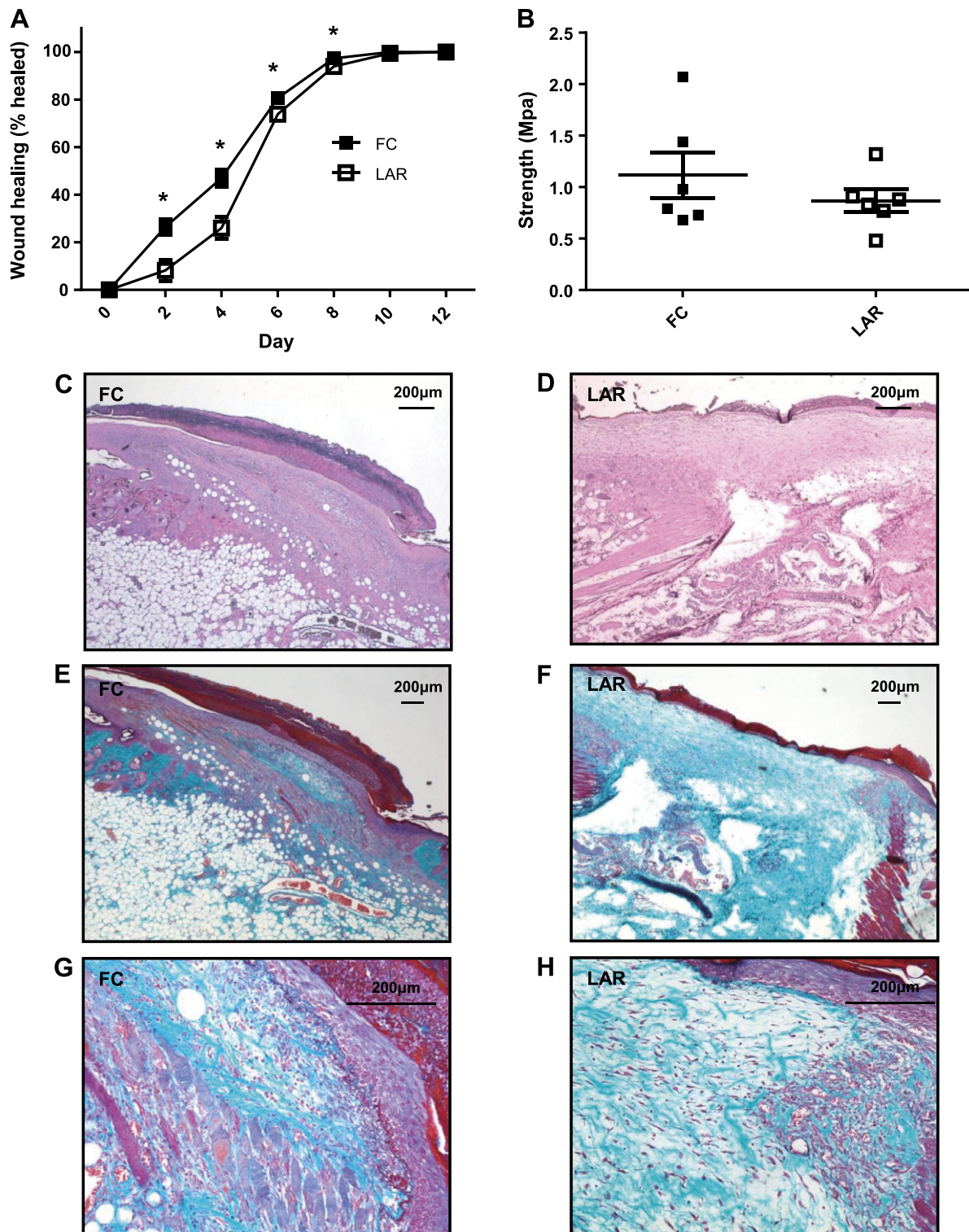


Fig. 2. *A*: wound healing in FC and LAR mice. *B*: tensile strength of FC and LAR wounds (day 18). Values are means \pm SE. * $P < 0.05$ by Student's *t*-test. *C* and *D*: hematoxylin-eosin-stained sections of FC and LAR wounds at day 4. Magnification $\times 40$. *E* and *F*: Milligan's trichrome-stained sections of FC and LAR wounds. Magnification $\times 25$. *G* and *H*: Milligan's trichrome-stained sections of FC and LAR wounds. Magnification $\times 100$ at wound edge.

significant change in macrophage staining was evident. This was confirmed by real-time PCR for *F4/80* mRNA (data not shown). Together with unchanged numbers of thioglycollate-elicited macrophages, this suggests normal tissue migration of LAR macrophages.

Diabetic LAR and FC Mice Had Equivalent Wound Healing

Because it has been reported that HIF-1 α decreases in diabetic wounds and that this contributes to impaired wound healing, wound healing was investigated in LAR and control

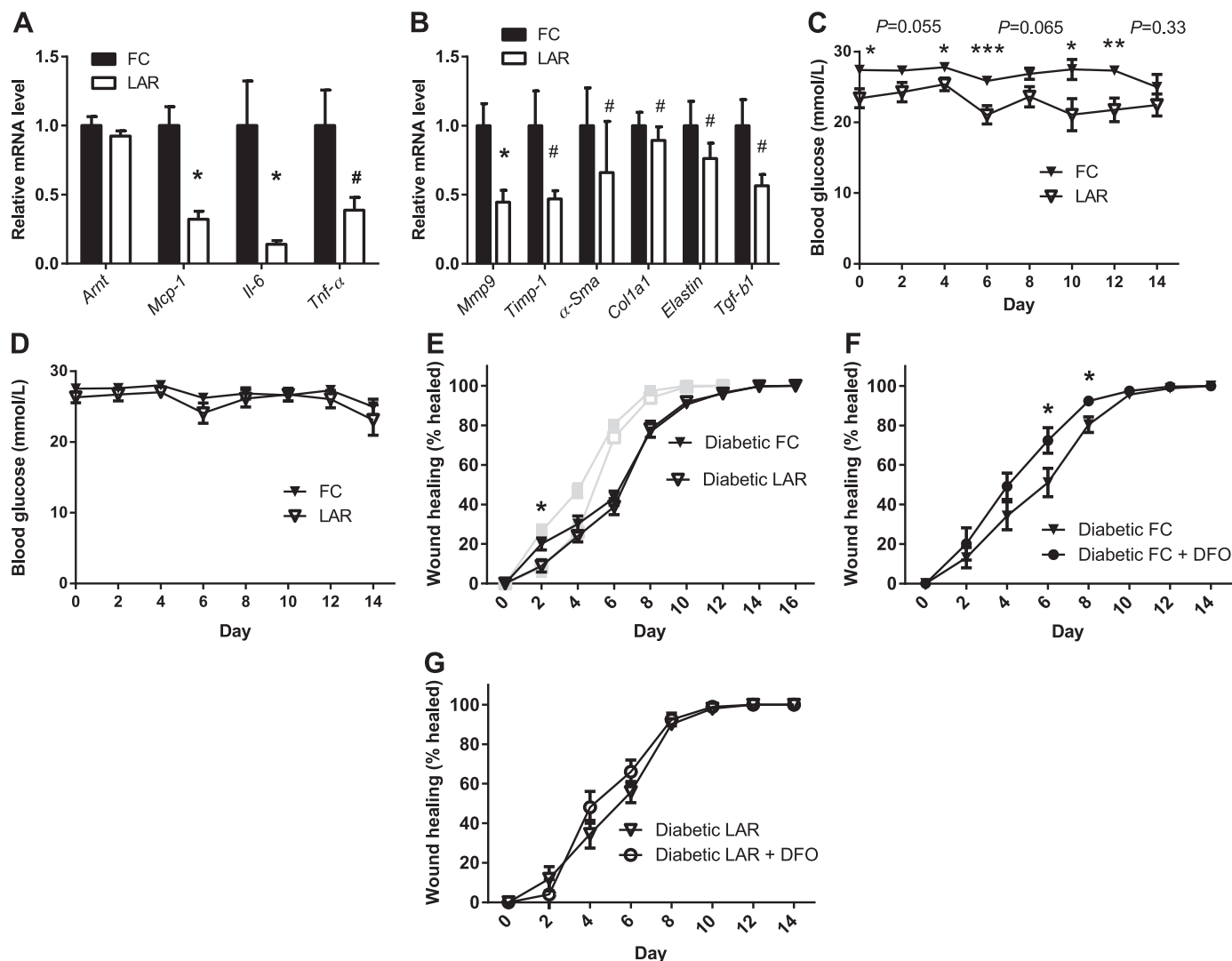


Fig. 3. Wound mRNA expression and wound healing in diabetic mice. *A*: cytokine mRNA expression in FC and LAR wounds at day 4. *B*: tissue-remodeling mRNA expression in FC and LAR wounds at day 4. *C*: blood glucose levels after streptozotocin injection in LAR and FC mice. *D*: blood glucose after streptozotocin treatment in LAR and FC mice included in the final analysis of wound healing. *E*: wound healing in diabetic FC and LAR mice. *F* and *G*: wound healing in diabetic FC and LAR mice compared with FC and LAR mice receiving DFO treatment. Values are means \pm SE. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.005$ by Student's *t*-test. # $P < 0.05$ by Wald-Wolfowitz test.

mice that were rendered diabetic (1, 2, 29, 43). Unexpectedly, overall, LAR mice had significantly reduced blood glucose levels compared with controls (Fig. 3C). For that reason, only animals with at least two blood glucose levels >28 mmol/l (506 mg/dl) and none with levels <15 mmol/l (270 mg/dl) before insulin treatment (required by the Garvan Animal Ethics Committee for animals with levels >20 mmol/l) were included in the study (Fig. 3D).

Interestingly, the differences in wound healing between LAR and control mice (Fig. 2A) were no longer significant in diabetic mice ($P = 0.3$ by rmANOVA; Fig. 3E). That is, the controls deteriorated much more than the LAR mice. This suggests that decreased function of ARNT-containing dimers in myeloid cells may contribute to the impaired wound healing observed in diabetes. We previously demonstrated decreased ARNT in the islets and liver of diabetic patients (13, 52). HIF-1 α expression has also been found to be decreased in human diabetic ulcers, while activity is reduced at high

glucose concentrations in human fibroblasts and diabetic animals (1, 2, 29).

DFO Treatment Does Not Significantly Improve Wound Healing in Diabetic LAR Mice

DFO is an iron chelator that increases HIFs (48). HIFs require ARNT for function. To delineate whether part of the mechanism of DFO in improving healing in diabetic mice (1) is increasing activity of the ARNT/HIF-1 α complex in myeloid cells, the response of diabetic FC and LAR mice to DFO was investigated (Fig. 3, F and G). DFO treatment improved (i.e., decreased) wound size of FC animals at days 6 and 8 ($P = 0.045$ and $P = 0.03$, $P = 0.046$ by rmANOVA). In contrast, DFO treatment failed to significantly improve wound healing in LAR animals ($P = 0.4735$ by rmANOVA). This indicated that activity of the ARNT/HIF dimer in myeloid cells may be required for the beneficial effects of DFO on wound healing.

Quantification of Monocyte ARNT mRNA in Control and Diabetic Patients

We previously reported that *ARNT* mRNA is reduced in the liver and islets of T2D patients and that ARNT protein is reduced in the liver of T2D patients (13, 52). To test whether *ARNT* mRNA was decreased in circulating human myeloid cells from diabetes patients, levels were measured in human blood monocytes by real-time PCR. There was very wide interindividual variability, and no difference in *ARNT* mRNA was observed ($P = 0.3$; Fig. 4A). Monocyte *ARNT* mRNA was, however, correlated with serum levels of various cytokines. Decreased *ARNT* mRNA expression correlated with increased IL-6 and IL-8 ($P = 0.0135$ and $P = 0.0232$; Fig. 4, B and C), MCP-1 ($P = 0.0403$; Fig. 4D), and TNF- α ($P = 0.0432$; Fig. 4E). No significant correlation was found for interferon- γ -induced protein 10 ($P = 0.4667$; Fig. 4F) or for interferon- γ or macrophage inflammatory protein 1 β (data not shown).

DISCUSSION

Mice with reduced ARNT in myeloid cells displayed altered cytokine transcription, skin inflammation, skin transplant rejection, and wound healing. Wound healing was similarly impaired in diabetic LAR and diabetic control mice, suggesting that decreased myeloid cell ARNT/HIF-1 α function may contribute to impaired wound healing in diabetes. DFO administration failed to significantly improve wound healing in LAR mice, suggesting that ARNT/HIF-1 α induction in myeloid cells may also be a component of the mechanism of action of DFO in improving wound healing (1, 43).

LAR granulocytes displayed altered expression of cytokine mRNA. Interestingly, ARNT deletion in Kupffer cells using the Mx-Cre promoter system has also been shown to prevent the upregulation of *Pdgf- β* , vascular endothelial growth factor (*Vegf*), *angiopoietin-1*, and *Mcp-1* in hypoxia in vitro (6). Overall, the phenotype of myeloid cells lacking ARNT in response to acute stimulation appears most consistent with the combined phenotype of HIF-1 α and HIF-2 α knockout animals, with impairment of acute inflammatory response (7, 16), rather than the phenotype of myeloid cells lacking AhR (21, 40).

Within wound biopsies we found significantly reduced *Mcp-1*, *Il-6*, and *Tnf- α* expression, suggesting that myeloid cell ARNT deletion led to an impaired acute inflammatory response to wounding in vivo. We did not assess mRNA changes at multiple time points; however, the decreased *Il-6* and *Mcp-1* are consistent with the reduced cytokine mRNA observed in isolated macrophages. Indirect effects on nonmyeloid cells may also play a role. MCP-1 and IL-6 have roles in wound healing, with those knockout animals displaying delayed healing (11, 27). Reduction of these cytokines may contribute to the delay in wound healing in our model. Although macrophages are not required for effective wound healing in sterile conditions in the absence of neutrophils (30), the importance of macrophage function in wound healing when neutrophils are present has been shown in a number of studies (12, 20, 25, 28, 33). These studies have demonstrated that macrophages play an important role in neutrophil clearance, angiogenesis, epithelialization, and collagen deposition. Reduction in the mRNAs for *Coll1 α 1*, *Mmp9*, and *Timp-1* were found and suggest altered collagen deposition and altered wound remod-

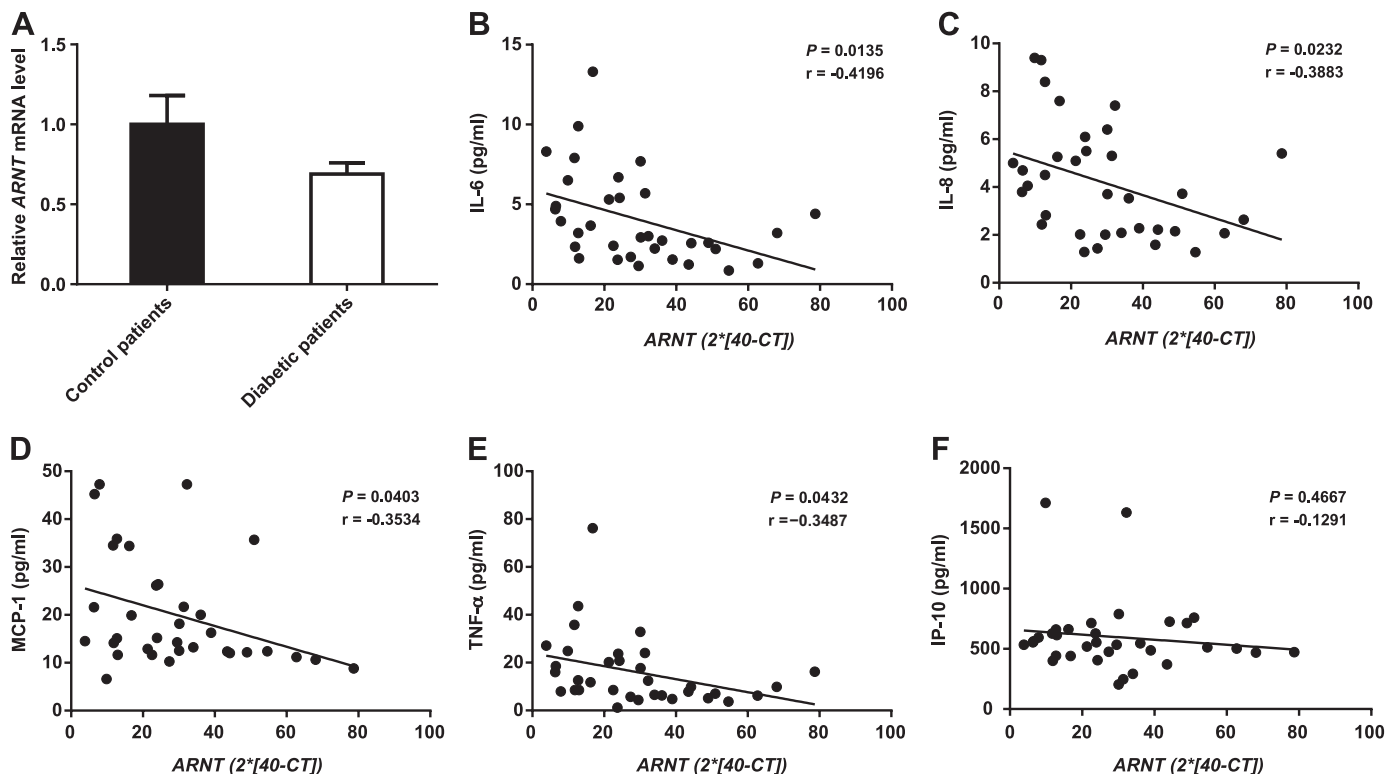


Fig. 4. *ARNT* expression in human monocytes, expressed as estimated relative fold change using the following formula: $ARNT = 2^{(40 - C_T)}$, where C_T is cycle threshold. A: average human monocyte *ARNT* mRNA expression in control subjects and diabetic patients. Values are means \pm SE. B–F: correlation of IL-6, IL-8, MCP-1, TNF- α , and interferon- γ -induced protein 10 (IP-10) to *ARNT* level. Pearson's correlations (r) were calculated using PRISM software.

eling, although tensile strength at *day 18* was equivalent. DFO was also shown to significantly decrease wound size in diabetic FC animals, but not in mice lacking myeloid ARNT, suggesting a role for myeloid cell HIF-1 α in mediating the effects of DFO on wound healing in diabetes.

Expression of ARNT in circulating human monocytes was highly variable. Ideally, future studies would also involve measurement of monocyte ARNT in wounds and blood in patients with sepsis and impaired wound healing. A novel finding was that ARNT mRNA in human monocytes correlated negatively with serum IL-6, IL-8, MCP-1, and TNF- α . Although a cause-and-effect relationship is speculative, a recent study reported increased allergic responses in mice lacking HIF-1 α in myeloid cells (47). This study suggests that myeloid ARNT/HIF-1 α may dampen some forms of excess inflammation.

Although ARNT was previously considered to be constitutively expressed, it has now been shown to be regulated in specific circumstances by reactive oxygen species, LPS, and curcumin (4, 34, 54). We recently reported that, in T2D islets, increasing HIF-1 α with DFO increased ARNT expression to near-normal levels (3). Increasing myeloid cell ARNT may be a therapeutic strategy to improve wound healing.

GRANTS

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DISCLOSURES

No conflicts of interest, financial or otherwise, are declared by the authors.

AUTHOR CONTRIBUTIONS

C.S., S.T.G., S.T., S.M., and J.E.G. are responsible for conception and design of the research; C.S., J.B., D.M., P.B., R.S., K.M.C., S.N.W., K.M., and F.S. performed the experiments; C.S., D.M., P.B., R.S., S.N.W., K.M., S.M., and J.E.G. analyzed the data; C.S., J.B., P.B., K.M., S.T., S.M., and J.E.G. interpreted the results of the experiments; C.S. and J.E.G. prepared the figures; C.S. drafted the manuscript; C.S., D.M., R.S., K.M., F.S., S.T., S.M., and J.E.G. edited and revised the manuscript; C.S., J.B., D.M., P.B., R.S., S.N.W., K.M., F.S., S.T.G., S.T., S.M., and J.E.G. approved the final version of the manuscript.

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