Research Article

Sirtuin 2 Regulates Microvascular Inflammation during Sepsis

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Received 26 November 2016; Revised 23 February 2017; Accepted 5 March 2017; Published 19 April 2017

Objective. Sepsis and septic shock, the leading causes of death in noncoronary intensive care units, kill more than 200,000/year in the US alone. Circulating cell-endothelial cell interactions are the rate determining factor in sepsis inflammation. Sirtuin, a seven-member family of proteins (SIRT1–7), epigenetically controls inflammation. We have studied the roles of SIRTs 1, 3, and 6 in sepsis previously. In this project, we studied the role of SIRT2 on sepsis-related inflammation.

Methods. Sepsis was induced in C57Bl/6 (WT), SIRT2 knockout (SIRT2KO), and SIRT2 overexpressing (SIRT2KI) mice by cecal ligation and puncture (CLP). We studied leukocyte/platelet adhesion using intravital microscopy and E-selectin/ICAM-1 adhesion molecule expression in the small intestine with immunohistochemistry (IHC) six hours post-CLP/sham surgery. We also studied 7-day survival rates in WT, SIRT2KO, and SIRT2KI sepsis mice.

Results. Compared to WT mice, SIRT2KO mice show exaggeration while SIRT2KI mice show attenuation of cellular adhesion with sepsis in the small intestine. We also show that the small intestinal E-selectin and ICAM-1 expressions increased in SIRT2KO and decreased in SIRT2KI mice versus those in WT sepsis mice. We show that the 7-day survival rate is decreased in SIRT2KO and increased in SIRT2KI sepsis mice.

Conclusion. SIRT2 modulates microvascular inflammation in sepsis and affects survival.

1. Introduction

Sepsis and septic shock, the most expensive conditions in the US, are the leading causes of death in the noncoronary intensive care units [1, 2]. More than 30 different clinical trials for sepsis failed, leaving no molecular-based therapies currently available [3]. Sepsis leads to multiple organ failures; the number of organs involved correlates with mortality [4]. In a dysregulated acute systemic inflammatory response of sepsis, microvasculature sits at an interface between systemic circulation and local tissue. Circulating cell-endothelial cell interactions at the microvascular interface (MVI) are the rate-limiting factors in inflammation [5]. We and others have studied the role of microvascular response in early sepsis previously [6, 7]. Knowing how the MVI-cell responses are directed is critical to designing molecular-based treatments.

The sirtuin family of proteins is among the many regulators of cellular homeostasis [8, 9]. Sirtuins, first described in yeast, are the highly conserved family of proteins. The seven mammalian sirtuins (SIRT1–7) disperse through different cell compartments to guard homeostasis, sense bioenergy needs of cell, and make changes to nutrient sources [10]. Nuclear SIRT1, the most extensively studied sirtuin, epigenetically directs the pro-versus anti-inflammatory response in sepsis [12]. We reported that nuclear SIRT1and SIRT6 and mitochondrial SIRT3 inform immune and metabolic (immunometabolic) programming during acute inflammation and are dysregulated during sepsis [13, 14].

SIRT2, originally described as a tubulin deacetylase, modifies acetylation of many histone and nonhistone...
proteins including transcription factors \[15\]. While described as a cytoplasmic protein, SIRT2 translocates to nucleus during mitosis and cellular stress \[16\]. Nucleocyto-
plasmic shuttling of SIRT2 likely regulates its biological activ-
ity \[17\]. Mounting evidence suggests a critical role for SIRT2 in the control of inflammation, immunometabolism, and tumorigenesis \[15\]. We have shown recently that SIRT2, the most abundant of sirtuins in the adipose tissue, also reg-
ulates hypoinflammatory response of late sepsis in obese mice with sepsis via direct deacetylation of NFκB p65 \[18\] consistent with literature implicating the role of SIRT2 in neuroinflammation and collagen-induced arthritis among other inflammatory conditions \[19, 20\].

The effect of SIRT2 modulation on microvascular inflam-
mation during early sepsis is not very well studied. Here, we report a role for SIRT2 in inflammation during early sepsis. Using a genetic approach to modulate SIRT2 in mice, we show that in early sepsis, low SIRT2 levels enhance and high SIRT2 levels attenuate the MVI inflammatory response in vivo via modulation of adhesion molecule expression impact sepsis mortality rates in rodent sepsis.

2. Material and Methods

2.1. Mice. The studies were approved by the Institutional Animal Care and Use Committee of the Wake Forest School of Medicine; experiments were performed according to the NIH guidelines. The wild type (WT; C57Bl/6) mice were purchased from Jackson Laboratories; SIRT2 knockout (Sirt2tm1.1Fwa; SIRT2KO) and SIRT2 overexpress-
ing (SIRT2 knock in SIRT2KI) were bred in the Wake Forest Animal facility. The SIRT2KI and SIRT2KO mice were on C57Bl/6 background.

The WT mice and SIRT2KO breeding pairs were obtained from Jackson Laboratories (Bar Harbor, ME, USA), and the SIRT2KI breeding pair was a generous gift from Dr. Sinclair (Harvard Medical School, Boston, MA) and Dr. Lindsay Wu (University of New South Wales, Australia). These mice have been used in literature before \[21\]. The method for creation of these mice is described in literature. Briefly, Cre-inducible SIRT2 transgenic mice were created using transcriptional STOP element flanked with loxP sites which was then inserted between a CAGGS promoter and the murine SIRT2 cDNA. The construct was then targeted into the mouse collagen A1 locus using flp recombinase-mediated genomic integration into C10 ES cells (derived from V6.5; BL/6 × 129SvJae F1 ES cells). Mouse embryonic stem cells carrying a single copy of the SIRT2STOP construct were identified using resistance to the antibiotic marker hygromycin and Southern blotting, clones injected into blastocysts and pups genotyped for positive recombination. Mice were then backcrossed to C57Bl/6 to 10 generations. Subsequently, SIRT2STOP mice were crossed with CMV-Cre transgenic mice strains on C57Bl/6 background (Jackson Labs, Bar Harbor, ME, USA) to generate constitutive SIRT2 transgenic animals (SIRT2tg). SIRT2tg; CMV-Cre double transgenics were then backcrossed to C57Bl/6 to outcross the CMV-Cre allele \[21\]. Six eight-
week old mice were used for experiments.

SIRT2KO mice breeding pair was purchased from Jackson laboratory. Per vendor description, a targeting vector was designed to replace exon 5–6 part and part of exon 7 of the mouse SIRT2 gene with a floxed neomycin-resistance cassette. This construct was then electroporated into 129S6/SvEvTac-derived TCI embryonic stem cells. Correctly targeted embryonic stem cells were then transiently trans-
fected with a Cre expression plasmid for the purpose of removing neomycin cassette. Embryonic stem cells that had successfully undergone Cre-mediated recombination and no longer retained the cassette were injected into C57BL/6J blastocytes, and the resulting males were bred to 129Sv females. The offsprings were then backcrossed to C57Bl/6 for 8 generations to produce homozygous SIRT2KO mice. After arrival to Jackson Laboratories, these mice were then bred for at least one generation to establish colony (https://www.jax.org/strain/012772).

2.2. Cecal Ligation and Puncture (CLP). Anesthetized mice (using 1–3% isoflurane/O2 mixture via nose cone) under-
went laparotomy. A one-centimeter laparotomy incision was performed, cecum identified, ligated, perforated two times with a 22-guage needle and abdomen closed in two layers (peritoneum and skin) as described in literature \[22\]. Each mouse received subcutaneous fluid resuscitation of 1 ml normal saline. Sham-operated mice underwent laparot-
omy followed by fluid resuscitation, without cecal ligation or puncture.

2.3. Intravital Fluorescent Video Microscopy (IVM). Anes-
ethetized mice (ketamine 150 mg/kg + xylazine 7.5 mg/kg; intramuscularly) underwent carotid artery and jugular vein cannulations followed by intravital microscopy procedures 6 h post-CLP as described previously \[22\]. The laparotomy incision (from previous CLP/sham surgery) was opened to exteriorize the small intestine to study microcirculation \(n = 4–6 \text{ mice/group}\) as described previously \[22\]. Rhodame 6G (Sigma-Aldrich, St. Louis, MO, USA; 0.02% in 100 μL PBS) was injected intravenously to visualize leuko-
cytes (labeled red). Platelets from matched donors were labeled green ex vivo using carboxyfluorescein diacetate succinimidyl ester (CFSE: Sigma-Aldrich; St. Louis, MO, USA; 90 μM). This allowed leukocyte and platelet visuali-
ization simultaneously.

Platelets were obtained from donor mice of the same genetic background. Platelet donor mice underwent carotid arterial cannulation under ketamine+xyazine anesthesia (ketamine 150 mg/kg + xylazine 7.5 mg/kg; intramuscularly) to obtain blood; platelets were isolated and labeled with CFSE as described previously \[7, 23\]. Labeled platelets \(n = 100 \times 10^6\) were infused via jugular venous cannula over 5 min (yielding <5% of the total platelet count) and allowed to circulate for additional 5 min before recording on a DVD. Evidence suggests that these platelet isolation/labling pro-
duces do not affect activity/viability of isolated platelets \[24\].

Three to five postcapillary venules per mouse were recorded (4–6 mice per group) (1 min 20 seconds each venule) and cell adhesion quantified from one minute of the recording. A cell (leukocyte/platelet), stationary for at
at least 30 seconds (consecutive) of the one-minute recording analysis, was considered adherent. The mean of the average values of leukocyte adhesion determined in each mouse was then used to generate a group mean.

2.4. Western Blot Analysis. We obtained peritoneal cells from mice to confirm SIRT2 expression. WT, SIRT2KO, and SIRT2KI mice ($n = 8$ each) were injected with sodium thioglycolate medium intraperitoneally, as per literature [25]. All mice developed ascites within four days. On day, four mice were euthanized using isoflurane anesthesia and cervical dislocation (secondary method), and peritoneal lavage was performed to isolate peritoneal macrophages. Western blotting was completed as previously described [18]. The membrane was incubated with antibodies against SIRT-2 (Santa Cruz Biotechnology, sc-20966), GAPDH (Invitrogen, #AM4300), or CPA (Millipore, #07313) overnight followed by the incubation with Alexa Fluor 680-conjugated secondary antibodies (827-10080 or 827-10081; LI-COR Biosciences, Lincoln, NE), and signal was developed with a Li-COR Odyssey Infrared Imager system (LI-COR Biosciences).

2.5. Immunohistochemistry. Small intestinal tissue from CLP and sham surgery from all three strains (WT, SIRT2KO, and SIRT2KI) of mice was harvested, and fixed sections were stained using E-selectin and ICAM-1 (BD Biosciences, San Jose, CA, USA) and Von Willebrand factor (VWF, Abcam, Inc., Cambridge, MA, USA). Cy3-conjugated labeled secondary antibodies for E-selectin, ICAM-1, and FITC-conjugated secondary antibody for VWF were purchased from Jackson ImmunoResearch Laboratories, Inc. (West Grove, PA, USA). The IHC techniques and virtual image captures were as described previously [22, 23].

2.6. Experimental Protocol for Sepsis Mice. Mice were subjected to sepsis/sham surgery under isoflurane anesthesia. In IVM experiments, mice with ketamine-xylazine anesthesia underwent carotid/jugular cannulation, mean arterial blood pressure (MAP) measurement via carotid catheter followed by study of leukocyte and platelet adhesion at a 6-hour time point after surgery. A separate group of mice were euthanized with cervical dislocation under anesthesia and tissue harvested for immunohistochemistry. In a third group of mice, we observed a 7-day mortality as indicated. Mice were monitored at least two times per day. Mice were diligently followed for humane end points and euthanized per American Veterinary Medical Association guidelines.

2.7. Statistics. Leukocyte/platelet adhesion data were analyzed using Tukey-Kramer post hoc analysis (Statview, SAS; Cary, NC, USA). A $p < 0.05$ was designated as significant. Data for survival rates were analyzed using Graph Pad Prism 6.0 (Graph Pad Software, La Jolla, CA, USA). Log rank test was used to compare survival between the groups in the Kaplan-Meier survival curves and a $p < 0.05$ was designated as significant.

3. Results

3.1. Sirtuin 2 Modulates Cell Adhesion in Sepsis Microcirculation. To study microvascular inflammation, we assessed sepsis-induced leukocyte and platelet adhesion in small intestinal microcirculation of SIRT2KO and SIRT2KI mice compared to those of WT control. There were no significant differences in the mean arterial blood pressure (MAP), body weight, and total blood leukocyte counts between different groups (Table 1). As shown in Figure 1(a), the leukocyte adhesion increased significantly in sepsis versus the respective sham groups in all three strains including WT, SIRT2KO, and SIRT2KI mice. Leukocyte adhesion in the SIRT2KO sepsis group was significantly increased versus that in WT sepsis mice. In contrast, leukocyte adhesion in SIRT2KI sepsis mice was significantly decreased versus that in WT sepsis mice. Lastly, the leukocyte adhesion in SIRT2KI sepsis was significantly attenuated versus that in SIRT2KO mice. Leukocyte adhesion in the WT, SIRT2KO, and SIRT2KI sham groups were not significantly different from each other. We confirmed SIRT2 expression in the peritoneal cells from WT, SIRT2KO, and SIRT2KI. As shown in Figure 1(b), the SIRT2 expression in the peritoneal cells of SIRT2KO mice was lower while that of SIRT2KI mice was higher than that of the WT mice.

Next, we determined platelet adhesion in WT, SIRT2KO, and SIRT2KI mice with sham and sepsis (Figure 2). Similar to leukocyte adhesion, platelet adhesion in the WT, SIRT2KO, and SIRT2KI sepsis groups significantly increased versus respective sham counterparts. The platelet adhesion in the SIRT2KO sepsis group was significantly increased versus WT sepsis mice. However, unlike leukocyte adhesion, the
platelet adhesion in SIRT2KO sepsis mice was not significantly different from that in WT sepsis mice. Lastly, the platelet adhesion in the SIRT2KI sepsis group was significantly lower versus that in SIRT2KO mice. Platelet adhesion in the WT, SIRT2KO, and SIRT2KI sham groups did not differ from each other. Of note, we show total platelet adhesion here; we do not differentiate between platelet-endothelial and leukocyte-platelet-endothelial adhesion in this project.

3.2. Sirtuin 2 Regulates Adhesion Molecule Expression in Sepsis Microcirculation. Adhesion molecules are the “early responders” of inflammatory response in microcirculation. To study the mechanism of leukocyte and platelet adhesion in sepsis microcirculation, we defined adhesion molecule expression in small intestinal tissue in the WT, SIRT2KO, and SIRT2KI mice sham and sepsis groups. We focused on NFkB p65-dependent ICAM-1 and E-selectin, which are dominant endothelial cell-derived modulators of cellular adhesion [26–28].

As shown in Figures 3(a) and 3(b), there were no significant differences in E-selectin expression between WT, SIRT2KO, and SIRT2KI mice with sham surgery. However, as shown in Figures 3(c) and 3(d), the E-selectin expression in SIRT2KO sepsis mice was significantly higher compared to that in WT sepsis mice while that in SIRT2KI sepsis mice was significantly lower than that in WT sepsis mice. Figures 3(b) and 3(d) showed fluorescence quantification from small intestine IHC in each group (n = 3) while Figures 3(a) and 3(c) show representative images for E-selectin expression.

Next, we assessed the ICAM-1 expression in small intestinal tissue in WT, SIRT2KO, and SIRT2KI mice with sepsis. In Figures 4(a) and 4(b), similar to E-selectin expression, we observed no significant difference in ICAM-1 expression between WT, SIRT2KO, and SIRT2KI mice with sham surgery. However, in Figures 4(c) and 4(d), we showed increased ICAM-1 expression in SIRT2KO versus WT sepsis and decreased ICAM-1 in SIRT2KI versus SIRT2KO sepsis and WT sepsis. Figures 4(b) and 4(d) show fluorescence quantification from small intestine IHC in each group.
group \((n = 3)\) while Figures 4(a) and 4(c) show representative images for ICAM-1 expression. Together, the data support that SIRT2 regulates microvascular inflammation: low SIRT2 increases and high SIRT2 decreases leukocyte/platelet adhesion in sepsis by controlling ICAM-1 and E-selectin expression.

3.3. Sirtuin 2 Influences Survival in Sepsis Mice. To test the clinical impact of sepsis-induced inflammation on the whole animal, we determined 7-day survival rates in WT, SIRT2KO, and SIRT2KI mice with sepsis. Figure 5 depicts a 40% survival rate in WT sepsis mice, 10% in SIRT2KO sepsis, and 80% in SIRT2KI sepsis. This data suggest that the SIRT2 modulates survival in rodent sepsis.

4. Discussion

This study identifies a previously unrecognized property of SIRT2 as a regulator of microvascular inflammation. This novel function is via modulating expression of critical adhesion molecules ICAM-1 and E-selectin. Specifically, in an inverse relationship, SIRT2 deficiency is associated with increased cell adhesion and adhesion molecule expressions while SIRT2 overexpression is associated with decreased adhesion and adhesion molecule expressions. The clinical significance of this in mouse sepsis is that sepsis mortality is affected by the SIRT2 modulation as well. These phenotype shifts may represent, at least in part, the known deacetylation-dependent deactivation of NFκB p65 transcription factor as a master inflammation promoter; we have shown that SIRT2 directly deacetylates NFκB p65 [18].

This report seems to be in direct contradiction with a previous publication studying that the effect of SIRT2 inhibitor AGK-2 treatment during early sepsis improves survival [29]. There are some key differences between results from that publication and this project. Firstly, while they used a chemical compound to inhibit SIRT2, we used genetic approaches. Off target effects of a chemical
compound, namely, AGK-2, cannot be ruled out. Consistent with these results, however, our previous work using a different SIRT2 inhibitor compound during early endotoxemia showed an increased inflammatory response in mouse macrophages. We have shown previously that in fact the AK-7, a SIRT2 inhibitor treatment during the immunosuppressive phase of sepsis in obese mice improved survival. The MVI is sensitive to both local and systemic environmental stimuli such as inflammation; leukocyte adhesion is an essential link to infection outcome [5]. The growing body of evidence suggests a role for platelet activation and adhesion in sepsis related inflammation [30–33]. We have shown previously that the leukocyte and platelet adhesion increase during early sepsis [7, 34, 35]. In this project, we show the effect of SIRT2 modification on platelet adhesion along with leukocyte adhesion.

The selectins are responsible for the rolling and initial capture of leukocytes, while the engagement of leukocyte-β2-integrins with endothelial ICAM-1 mediates “firm” adhesion followed by emigration of leukocytes [36]. We previously reported that early sepsis increases adhesion molecule expression in MVI [7, 37]. Moreover, we reported that SIRT1 modulates the E-selectin and ICAM-1 expression during sepsis [22, 23]. This study adds SIRT2 as MVI regulator via E-selectin and ICAM-1 expression. Together, our findings support a role for both SIRT1 and SIRT2 in epigenetically directing inflammation. However, whether there are separate effects of the two SIRTs or redundancies will require further study.

Sirtuins are NAD+ sensors in immune and nonimmune cells. We have shown previously that SIRT2 deficiency in immune cells exaggerates inflammatory response via reacetylation of NFκB p65 [18]. SIRT2 is abundantly expressed in various tissue beds such as the brain, adipose tissue, liver, muscle, kidney, and pancreas. [15, 38–40]. In the current manuscript, we confirmed that the whole body SIRT2KO

![Figure 4: ICAM-1 expression in WT, SIRT2KO, and SIRT2KI mice. We stained small intestinal tissue for ICAM-1 expression. ICAM-1 (Cy3: red), Von Willebrand factor (VWF, FITC: green), and nuclear stain (DAPI: blue). (a) shows representative images of ICAM-1 expression in WT, SIRT2KO, and SIRT2KI mice with sham surgery while (b) shows fluorescence quantification of immunohistochemistry of three different mice in each group. There was no significant difference in ICAM-1 expression of WT, SIRT2KO, and SIRT2KI mice with sham surgery from each other. As shown in (c) (representative images) and (d) (fluorescence quantification from immunohistochemistry; n = 3 in each group), the ICAM-1 expression increased in SIRT2KO sepsis mice versus that in WT sepsis mice while it was decreased in SIRT2KI sepsis mice versus that in WT and SIRT2KO sepsis mice. *p < 0.05 versus that in WT sepsis group; #p < 0.05 versus that in SIRT2KO sepsis group; using Tukey’s post hoc analysis; error bars: sem.](image-url)
and SIRT2KO mice show changes in SIRT2 expression immune cells, namely, peritoneal macrophages as well. SIRT2 is selective for deacetylation of tubulin and contributes to HIF1-α to modulate hypoxia-dependent responses in tumor cells [41]. Clearly, SIRT2 acts in cytosol and in the nucleus during a stress response. It is possible that SIRT2 joins SIRT1 and 6 in nuclear control of inflammation but may act selectively on cytosolic tubulin. It seems likely that the sirtuin family functions in a cooperative manner for the control of inflammation. How this network is informed of stress is unknown but might involve shifts in redox signaling. Sepsis generates biphasic oxidative stress [42]. Sirtuins are known to activate an antioxidant response via the FOXO transcription family; SIRT2 modulates FOXO3a during oxidative stress [40]. Although we have not pursued this line of investigation, it is possible that SIRT2 modulates a microvascular inflammatory response via this pathway. SIRT1 and 2, in endothelial cells, might exhibit nonoverlapping activities [43]. More studies are required to separate cooperation from distinct pathway or protein contributions.

In conclusion, the study, for the first time to our knowledge, supports a critical role for SIRT2 in regulating MVI survival rate. This finding may open the way to novel therapies for adjusting MVI pathology during the inflammatory response.

Abbreviations

CLP: Cecal ligation and puncture  
ICAM: Intracellular adhesion molecule  
MVI: Microvascular interface  
NAD: Nicotinamide dinucleotide  
NFκB: Nuclear factor kappa B  
SIRT: Sirtuin  
SIRT1: Sirtuin 1  
SIRT2: Sirtuin 2  
SIRT2KI: Sirtuin 2 overexpressing  
SIRT2KO: Sirtuin 2 knockout  
WT: Wild type (lean).

Conflicts of Interest

The authors declare that they have no conflicts of interest.

Authors’ Contributions

Vidula Vachharajani conceptualized and designed the manuscript. Nancy Buechler, Xianfeng Wang, and Vidula Vachharajani contributed in data collection. The data analysis and interpretation were done by Vidula Vachharajani and Charles E. McCall. The generation of the manuscript was done by Vidula Vachharajani and Charles E. McCall.

Acknowledgments

The authors thank Dr. David Sinclair (Harvard Medical School, Boston, MA) and Dr. Lindsay Wu (University of New South Wales, Australia) for their generous gift of SIRT2KI mice. This work was supported by NIH grants: Vidula Vachharajani: R01GM099807; Charles E. McCall: (1) R01 AI065791 and (2) R01 GM102497.

References


