Age-Associated Remodeling of the Intestinal Epithelial Barrier

Lee Tran^{[1](#page-0-0)} and Beverley Greenwood-Van Meerveld^{1-[3](#page-0-1)}

1 Oklahoma Center for Neuroscience, ²Department of Physiology, and ²Department of Physiology, and
³VA Medical Center, University of Oklahoma Health Science Center, Oklahoma City.

Address correspondence to Beverley Greenwood-Van Meerveld, PhD, FACG, AGAF, VA Medical Center, Research Admin. Rm. 151G, 921 N.E. 13th Street, Oklahoma City, OK 73104. Email: Beverley-Greenwood@ouhsc.edu

Disorders of the gastrointestinal tract are common in the elderly people; however, the precise trait(s) of aging that contribute to the vulnerability of the gastrointestinal tract are poorly understood. Recent evidence suggests that patients with gastrointestinal disorders have increased intestinal permeability. Here, we address the hypothesis that disruption of the intestinal barrier is associated with aging. Our results demonstrated that permeability was significantly higher in colonic biopsies collected from old baboons compared with young baboons. Additionally, colonic tissue from the older animals had decreased zonula occluden-1, occludin, and junctional adhesion molecule-A tight junction protein expression and increased claudin-2 expression. Upregulation of miR-29a and inflammatory cytokines IFN-γ, IL-6, and IL-1β was also found in colonic biopsies from old baboons relative to young baboons. These results show for the first time that a pivotal contributing factor to geriatric vulnerability to gastrointestinal dysfunction may be increased colonic permeability via age-associated remodeling of intestinal epithelial tight junction proteins.

Key Words: Aging—Permeability—Tight junctions—MicroRNA—Cytokines.

Received November 28, 2012; Accepted June 5, 2013

Decision Editor: Rafael de Cabo, PhD

AGING significantly increases the vulnerability to gastro-
intestinal (GI) disorders with approximately 35%–40% of geriatric patients reporting at least one GI complaint during routine physical examination [\(1\)](#page-9-0). Despite the need to further understand age-associated factors that increase the susceptibility to GI dysfunction, there is a paucity of studies investigating the key factors in aging that affect the GI tract. Thus, far studies in rodents have demonstrated that aging alters intestinal smooth muscle contractility [\(2](#page-9-1)), as well as the neural innervations of the GI tract musculature ([3\)](#page-9-2). Several studies in rodents have also reported an increase in intestinal permeability to macromolecules with age [\(4–7\)](#page-9-3). Specifically, advancing age was shown to correlate with an enhanced transepithelial permeability of D-mannitol, indicating that there may be an age-associated decline in barrier function [\(8\)](#page-9-4).

The gut barrier protects the body from potentially harmful compounds and microorganisms while permitting the absorption of nutrients, electrolytes, and water. A loss of gut barrier integrity can lead to increased intestinal permeability allowing for the passage of noxious substances across the intestinal mucosa $(9-11)$. A key factor contributing to abnormalities in permeability is aberrant structuring of tight junction proteins that support gut barrier function [\(12\)](#page-9-6). Tight junction proteins link adjacent epithelial cells, regulating the paracellular movement of substances across the small and large intestinal lumen. However, the efficiency of tight junctions in controlling paracellular transport is dependent on the composition of the tight junction proteins associated in the junction, which include various combinations of zonula occludens (ZOs), occludins, claudins, and junctional adhesion molecules (JAMs). Substrates that are known to induce aberrant structuring of tight junction proteins include enhanced expression of microRNAs (miRNAs) [\(13](#page-9-7),[14](#page-10-0)), decreased circulating concentrations of glutamine ([15](#page-10-1)), and induction of inflammatory cytokines [\(16](#page-10-2),[17](#page-10-3)).

The overall goal of the current study was to investigate the role of aging on colonic permeability. The colon was specifically selected as the region of interest due to the technical feasibility and because many GI disorders associated with aging are predominantly colonic, that is, constipation, chronic diarrhea, and incontinence. To this end, we hypothesized that there is an increase in permeability with age that can be attributed, at least in part, to a structural decline in tight junction proteins. To test our hypothesis, we designed a comparative study to investigate the differences in permeability between colonic mucosal biopsies from young and old nonhuman primates, which have virtually identical rates and patterns of aging to humans ([18\)](#page-10-4).

Materials and Methods

Animals

Colonic mucosal biopsies were isolated from baboons (*Papio anubis*) housed at the University of Oklahoma Health Science Center's Department of Comparative Medicine Annex, Oklahoma City, Oklahoma, which is an Association for Assessment and Accreditation of Laboratory Animal Care accredited facility. The animal studies were approved by the University of Oklahoma Health Science Center Institutional Animal Care and Use Committees and were performed in accordance with the *Guide for the Care and Use of Laboratory Animals* and National Research Council guidelines. All baboons were free from GI pathologies, and a full report of the pathologies of the baboon has been reviewed ([19\)](#page-10-5). Baboons aged 4–10 years were considered "young" baboons, whereas baboons aged 18 and older were considered "old" in this study. A total of 10 old and 10 young baboons were used for this study.

Collection of Mucosal Biopsies

Baboons were sedated with an intermuscular injection of ketamine 10 mg/kg, intubated and maintained under anesthesia with 1.25%–1.5% isoflurane in O_2 . During the biopsy collection, physiological parameters were monitored using a noninvasive blood pressure cuff, pulse oximeter, and visual observation of respiration. The depth of the anesthesia was confirmed by the absence of corneal reflex. To collect the biopsies, a colonoscope was inserted into the rectum of the baboon and mucosal biopsies (8mm) were obtained from the transverse colon at a distance of 60–70 cm from the rectum using standard endoscopic forceps. Biopsy specimens were immediately placed in an iced oxygenated Krebs solution (pH 7.2–7.4; 120mM NaCl, 6mM KCl, $1.2 \,\text{mM} \,\text{MgCl}_2$, $1.2 \,\text{mM} \,\text{NaH}_2\text{PO}_4$, $2.5 \,\text{mM} \,\text{CaCl}_2$, $14.4 \,\text{mM}$ NaHCO₃, and 11.5 mM of glucose), paraformaldehyde (4%), or flash frozen on dry ice. The colonoscope was removed at the end of the procedure, and the baboons were allowed to recover from anesthesia while being continually monitored by a veterinarian.

From each baboon, a total of eight colonic biopsies were collected. Immediately following collection, four out of the eight biopsies were mounted in individual Ussing chambers for permeability experiments. From the remaining four colonic biopsies, one was processed for routine histology and another for the assessment of myeloperoxidase (MPO) activity. Out of the remaining two biopsies, one was placed in 4% formaldehyde, paraffin embedded, and sectioned for tight junction histology, and the other was flash frozen in dry ice and stored at −80°C for extraction of RNA and protein.

Histology and MPO Activity

Formalin-fixed biopsies were paraffin embedded and sectioned at 10 μm. Hematoxylin and eosin staining was performed by Precision Histology (Oklahoma City, OK). A pathologist blind to the experiment performed examination of the histology for morphological characterization, using the validated Chiu score system for gut pathology ([20\)](#page-10-6). MPO activity was assessed using EnzChek MPO Activity Assay Kit (Life Technologies, Carlsbad, CA). Colonic biopsies were homogenized in five volumes of phosphatebuffered saline followed by three cycles of freeze and thawing and by sonication for 10 seconds. After 15 minutes of centrifugation at 14,000*g* at 4°C, the supernatant was collected and Amplex UltraRed reagent (phosphate-buffered saline, 2% dimethyl sulfoxide, and 0.005% of H_2O_2) was added for peroxidation detection. The fluorescent intensity of each sample was detected using an excitation of 530 nm and emission at 590 nm using a fluorescent plate reader. Each value was converted into MPO values by using curves obtained from a standard sample of human MPO provided in the kit.

Assessment of Permeability in Colonic Biopsies

Four biopsies were mounted in a modified Ussing chambers with an exposed tissue area of 2.0 mm² and bathed with 10 mL of Krebs buffer (pH 7.2–7.4; 120mM NaCl, 6mM KCl, $1.2 \text{ mM } MgCl_2$, $1.2 \text{ mM } NaH_2PO_4$, $2.5 \text{ mM } CaCl_2$, 14.4 mM NaHCO₃, and 11.5 mM of glucose). The Krebs solution was continuously stirred and oxygenated with a steady gas flow of 95% O_2 and 5% CO_2 and maintained at 37°C with a heating block system. Two pairs of agar–salt bridge electrodes were connected to an automatic voltage clamp apparatus (ECV4000, World Precision Instruments, Sarasota, FL) to monitor transmucosal potential difference and short-circuit current $(I_{\rm sc})$. Transmucosal resistance was calculated using Ohm's law from the open-circuit potential difference and the $I_{\rm sc}$. The bathing solution on the mucosal side was then replaced with Krebs containing horseradish peroxidase (HRP type II; ~44 kDa protein antigen). Two samples of 250 µL were withdrawn from the submucosal side of the chambers every 20 minutes for 100 minutes $(t_0 - t_5)$ and replaced with fresh Krebs buffer to maintain equal hydrostatic pressure. HRP concentration from the submucosal samples was quantified by spectrophotometer analysis of the reaction of HRP with 3,3′,5,5′-tetramethylbenzidine liquid substrate. After 10-minute reaction period, a stop solution $(0.1 \text{ mM } H_2\text{SO}_4)$ was added to the sample extracted from the submucosal chamber. The absorbance of the colorimetric reaction product was measured at wavelength 450 nm and concentration was determined using a standard curve. HRP flux was expressed as quantity of HRP/20 min/2mm2 (Ussing chamber aperture) and recorded as pgHRP/min/mm2 .

Expression of Tight Junction Proteins

A colonic biopsy from each baboon was placed in 4% formaldehyde, paraffin embedded, and sectioned. Sections were then deparaffinized in xylene and rehydrated with gradient ethanol dips. Antigen retrieval was performed using proteinase K digestion (20 μ g/mL) in Tris– ethylenediaminetetraacetic acid buffer (50nM Tris, 1mM ethylenediaminetetraacetic acid, 0.5% Triton X-100, pH 8.0) for 10 minutes at 37°C and allowed to cool to room temperature and rinsed in Tris-buffered saline (TBS). Sections were then blocked with ImageIT signal enhancer (Life Technologies) for 30 minutes at room temperature. Sections were washed with TBS and incubated overnight with dual antibodies of either mouse anti-ZO-1 antibodies conjugated to Alexa Fluor 488 and rabbit anti-occludin or mouse anti-JAM-A and rabbit anti-claudin-2 (Life Technologies) at room temperature. Slides with no primary antibodies were used as negative controls. The following day, slides were rinsed in TBS and incubated in Alexa Fluor 546–conjugated anti-rabbit antibodies and Alexa Fluor 488–conjugated anti-mouse antibodies (Life Technologies) for 2 hours at room temperature. The slides were then rinsed again and cover slipped with ProLong Gold Anti-Fade solution (Life Technologies). Slides were imaged using a Zeiss LSM 510 confocal microscope at the Oklahoma Medical Research Foundation Imaging Core (Oklahoma City, OK). The negative control slides were used to calibrate the confocal microscope. Two randomly selected sections of 480,000 pixels at 40× magnification out of a total of six sections processed from each biopsy were analyzed with ImageJ software (NIH) for optical density [\(21](#page-10-7)). The densities were then averaged into a single *n* value used for statistical calculation.

Western Blot Analysis

Colonic tissue samples were homogenized in 20 volumes of lysis buffer (50mM Tris–HCl, pH 7.4; 1% NP-40; 0.25% sodium deoxycholate; 150mM NaCl; 1 mM ethylenediaminetetraacetic acid; 1mM phenylmethylsulfonyl fluoride; $1 \text{ mM } \text{Na}_3\text{VO}_4$; $1 \text{ mM } \text{Na}$ F) containing a protease inhibitor cocktail (104mM 4-(2-aminoethyl)benzenesulfonyl fluoride, 0.08mM aprotinin, 2.1mM leupeptin, 3.6mM bestatin, 1.5mM pepstatin A, and 1.4mM E-64; Sigma, St Louis, MO). Samples were then set on ice for 30 minutes and centrifuged at 14,000*g* for 20 minutes. The supernatants were collected and the protein concentration was determined by Bradford assay (Millipore, Billerica, MA). Approximately 30 μg of protein was separated on a 4%– 20% gradient polyacrylamide gel (Bio-Rad, Hercules, CA)

and transferred to a nitrocellulose membrane (Millipore). The membranes were blocked with 3% blocking solution in TBS-T for 1 hour. Blots were then incubated for 2 hours with the same antibodies used for immunohistochemistry in addition to anti-glyceraldehyde 3-phosphate dehydrogenase and anti-β-actin (Life Technologies) for normalization. Following incubation, the blots were washed in three changes of TBS-T and incubated for 1 hour with secondary anti-rabbit or anti-mouse antibodies (Millipore). Following three more washes in TBS-T, bands were visualized with ECL Western Blot Detection Kit (Amersham, Piscataway, NJ) and imaged using an Omega 12iC chemiluminescent imager (UltraLum, Claremont, CA).

Quantitative Reverse Transcription–Polymerase Chain Reaction

Whole tissue RNA and miRNA were extracted using RNeasy Mini Kit and miRNeasy Kit (Qiagen, Valencia, CA) following the manufacturer's instructions. Concentration of total RNA was determined by spectrophotometry (λ = 260nm). Extraction was followed by cDNA synthesis using RT2 First Strand cDNA Kit for mRNA and miScript II RT Kit for miRNA and qPCR using SYBR Green qPCR Mastermix or miScript SYBR Green in a total reaction volume of 25 μL (Qiagen). Samples were run in triplicates and "no template" conditions served as a negative control. The housekeeping gene ubiquitin-C was used to normalize mRNA samples and RNU-6B was used to normalize miRNA samples. Additionally, mRNA samples were normalized to the housekeeping gene 18S to validate results. The housekeeping gene β-actin was also used but found to be inconsistent between old and young baboon samples. The miRNA samples were also normalized to housekeeping genes SNORD-25 and SCARN-A7 to verify the results. The validated cytokine primers, 18S primers, and β-actin primers summarized in [Table 1](#page-2-0) were adapted from conserved sequences from multiple species ([22](#page-10-8)[,23](#page-10-9)) and acquired from Invitrogen (Carlsbad, CA). Human glutamine synthase (GLUL), ubiquitin-C, and miRNA Primer Assays were purchased from Qiagen. The reaction was performed on an Applied Biosystems StepOnePlus Real-Time PCR System Thermal Cycling Block (Life Technologies) with the initial

Table 1. Baboon Primers

Gene	Sense Primer	Antisense Primer	Product Size
IL-1 β	TGTACGATCACTGAACTGCA	GAAGTCAGTTATATCCTGGC	423
$IL-2$	AACTGGAGCATTTACTGCTG	GTGTTGAGATGATGCTTCGAC	357
$IL-6a$	ATGAAGTTCCTCTCTGCAAGAG	CACTAGGTTTGCCGAGTAGAT	421
$IL-10$	AAACCACAAGACAGACTT	GATTTTGGAGACCTCTA ATTTA	427
$IL-12a$	AGCAGCTGGTCATCTCTTGGTT	CCAGCATCTCCAAACTCTTTGA	559
TNF- α	TCTTCACTGGAAAGGACACCA	GAAGGAGAAGAGGCTGAGGAA	540
IFN- γ	TGCAGAGCCAAATTGTCTCCTTTTAC	GGGATGCTCTTCGACCTCGAAA	453
18S	TCAAGACAAGTCGGAGG	GGACATCTA AGGGCATCACA	489
β -Actin	CTACAATGAGCTGCGTGTGG	AAGGAAGGCTGGAAGAGTGC	528

denaturation at 95°C 15 minutes. Subsequent denaturations were at 94°C for 10 seconds, annealing at 60.0°C for 30 seconds, and extension at 72°C for 30 seconds, for a total of 40 cycles, and a final extension at 72°C for 10 minutes. Melting curves were performed at the end of each run from 72°C to 95°C for 90-second intervals and showed only a single peak for each gene analyzed. Calculation of the gene expression was performed as previously described [\(24](#page-10-10)). Relative quantity of mRNA from each sample was calculated as the difference in $C_{(T)}$ for target minus $C_{(T)}$ for housekeeping rRNA [$\Delta C_{(T)}$], and calibrated to $C_{(T)}$ of young control sample $[\Delta \Delta C_{T}]$. Fold change in transcription is expressed as 2^[−ΔΔC(T)].

Statistical Analysis

Data are represented as the mean ± *SEM*. Data from all Ussing chambers were averaged for each animal into a single *n* value. Means from old and young baboons were compared using a Student's unpaired *t* test using GraphPad Prism Software (version 5.2; La Jolla, CA), and significance was defined as *p* < .05.

Results

The Effect of Aging on Colonic Mucosal Histology and MPO Activity

The histological analysis revealed no marked differences between the mucosal biopsies from young and old baboons [\(Figure 1A](#page-4-0) and [B](#page-4-0)). Specifically, a review by a pathologist blind to the study indicated that there was no evidence of increased fibrosis, crypt atrophy, or acute inflammation in the aged samples when compared with the young samples. To further substantiate the histological observations, total MPO activity was measured to biochemically quantify levels of neutrophil infiltration [\(Figure 1C](#page-4-0)). Our results showed no difference in MPO activity between young and old baboons ($p < .05$).

The Effects of Aging on Colonic Permeability

Transepithelial electrical resistance, an electrophysiological measure of the ionic gradient across the colonic mucosal biopsy, was recorded and summarized in [Figure 2A](#page-5-0). Overall, there was a lower transepithelial electrical resistance in old baboon colonic biopsies versus young baboon colonic biopsies ($p < .01$), suggesting an abnormality in the ability of the tissue from aged animals to maintain an ionic gradient. To further corroborate the electrophysiological finding, we assessed epithelial barrier function by quantifying the amount of HRP that traversed the epithelial barrier from the mucosal to the submucosal chamber. As illustrated in [Figure 2B,](#page-5-0) we found that there was a greater HRP flux in colonic biopsies from old baboons compared with young baboons $(p < .05)$ indicating an increase in colonic permeability to macromolecules in the tissue from aged animals.

The Effects of Aging on Tight Junction Protein Expression in Colonic Biopsies

In the current study, four tight junction proteins were assessed: ZO-1, occludin, JAM-A, and claudin-2, all of which have been shown to play an important role in intestinal barrier function. Immunofluorescence was used to visualize the structure of the tight junctions in the epithelium of the colonic biopsies, and the fluorescent optical density was quantified (Figure 3). We found that aging decreased ZO-1, occludin, and JAM-A tight junction protein expression in the epithelium ($p < .05$). Although there was a trend toward an increase in claudin-2 protein expression, the changes were not statistically significant $(p = .09)$. Western blot analysis was used to quantify the total expression levels in the colonic biopsies for each tight junction protein [\(Figure 4\)](#page-7-0). In comparison with young baboon colonic biopsies, the old baboon biopsies exhibited significant decreases in ZO-1, occludin, and JAM-A $(p < .05)$, in contrast to a significant increases in claudin-2 expression ($p < .05$).

The Effect of Aging on miRNAs and GLUL Expression in Colonic Biopsies

Previous studies have shown that tight junction proteins can be regulated by miRNAs $(13-15)$ and glutamine ([15](#page-10-1)[,25](#page-10-11)). In the current study, miRNAs were measured by quantitative reverse transcription–polymerase chain reaction (qRT–PCR) and included miR-29a, miR-212, and miR-122a ([Figure 5A\)](#page-8-0). Overall, miR-29a was significantly increased in old baboon colonic tissue ($p < .05$), but no changes in miR-212 or miR-122a were observed with age $(p > .05)$. GLUL mRNA expression was also measured by qRT–PCR and shown in [Figure 5B.](#page-8-0) Our results showed no differences in GLUL expression between old and young colonic samples ($p > .05$).

The Effect of Aging on Cytokine Production in the Colonic Biopsies

Inflammatory cytokines modulate intestinal permeability through regulation of tight junction protein expression and trafficking $(16,17)$ $(16,17)$ $(16,17)$; thus, we measured cytokine expression by qRT–PCR in the colonic mucosal biopsies ([Figure 6](#page-8-1)), which is the optimal method of detecting cytokine levels in isolated tissue ([26\)](#page-10-12). Because mRNA levels are more stable than protein, quantifying mRNA also eliminates confounding factors such as the short half-life of cytokines. Previous studies have also shown excellent correlation between cytokine mRNA levels quantified by qRT–PCR and protein levels quantified by enzyme-linked immunosorbant assays ([27](#page-10-13)[,28](#page-10-14)). Compared with young baboons, we found IFN- γ ,

Figure 1. Hematoxylin and eosin staining revealed no histological differences between colonic biopsies from young and old baboons (**A**). Review by a pathologist blind to the study confirmed that there was no evidence of increased fibrosis, crypt atrophy, or acute inflammation in the aged samples when compared with the young samples. Quantification of myeloperoxidase (MPO) activity within the colonic biopsies showed no difference between young or old baboons (**B**). Statistical significance was **p* > .05 by Student's unpaired *t* test.

IL-6, and IL-1β to be significantly increased (*p* < .05) in old baboon colonic biopsies without changes in IL-12 or TNF- α ($p > .05$). Anti-inflammatory cytokines IL-10 and IL-2 were undetectable by our methods.

Discussion

The overarching goal of the current study was to delineate the effects of aging on intestinal barrier function. We found that epithelial permeability was greater in colonic biopsies isolated from older baboons. Supporting this observation, we discovered that there is significant tight junction remodeling including a decrease in ZO-1, occludin, and JAM-A proteins, and an increase in claudin-2 expression in old baboon colon compared with young baboon colon. Upon investigation of the potential mechanisms that may be responsible for ageassociated changes in tight junction protein expression, we found an increase in miR-29a but no observable differences

Figure 2. Aging increases colonic permeability. A reduction in transepithelial electrical resistance (TEER; **A**) and an increase in horseradish peroxidase (HRP) flux (**B**) was apparent in biopsies from old baboons compared with young baboons. Statistical significance was **p* < .05 by Student's unpaired *t* test.

in GLUL expression. We also measured an elevated level of proinflammatory cytokines, in the absence of overt inflammation as assessed via routine histology and MPO activity, supporting the claim that inflammatory cytokines modulate intestinal permeability through regulation of tight junction protein expression and trafficking [\(16](#page-10-2)[,17](#page-10-3)).

A growing body of evidence suggests that digestive disorders such as constipation, diarrhea, diverticular disease, and celiac disease may be the result of abnormal intestinal permeability due to compromised gut barrier function [\(9–11](#page-9-5)). Although elderly people often report abnormalities in bowel function, the etiology and pathophysiology are not well understood. To our knowledge, there is limited information on the effects of aging on the epithelial barrier; however, ischemic changes and an increase in the use of nonsteroidal anti-inflammatory drugs are thought to contribute to impaired epithelial barrier in the duodenum of elderly patients ([29\)](#page-10-15). Several studies in rodents have indicated an increase in overall absorption of macromolecules with age $(4–7)$ $(4–7)$, as well as specifically in the colon (8) (8) , confirming that there may be an age-associated increase in intestinal permeability, at least in a rodent model. In the current study, we advanced these earlier observations by demonstrating in a nonhuman primate model that aging significantly increases colonic permeability. Specifically, in colonic biopsies, we found a decrease in transepithelial electrical resistance across the colonic epithelium in a nonhuman primate model implying that the intestinal barrier from aged animals is less effective at separating ionic charge, most likely due to increased mucosal permeability. Further substantiating these electrophysiological findings, we then went on to show that there was an increase in HRP flux across the epithelial membrane in old baboon colonic mucosal tissue compared with young baboon colonic mucosal tissue. Our experiments focused primarily on paracellular permeability of the colonic tissue, but it is important to note that transcellular movement also influences the overall permeability of the colon. The colonic epithelium contains ion channels, pumps, and carriers that actively transport vitamins, minerals, and electrolytes, and their functional capacity varies

with ontogeny ([30](#page-10-16)[,31](#page-10-17)). Although there are limited studies evaluating transcellular mechanisms in the aging colon, investigations in the small intestine indicate that there is an overall decline in active absorption with age including carrier-mediated transport [\(32](#page-10-18),[33\)](#page-10-19), and sodium-dependent transport of amino acids [\(34](#page-10-20)) and sugars [\(35](#page-10-21)).

Many studies have reported that the permeability of the intestinal epithelial barrier is dependent on a network of protein strands made up of ZOs, occludins, claudins, and JAMs. In the current study, we quantified tight junction proteins expression and, despite the lower resolution in the colonic biopsies when compared with whole intestinal mounts or cultured cells, we were able to visualize ageassociated restructuring of the junctions. Specifically, our results indicated a significant decrease in ZO-1, occludin, and JAM-A in colonic biopsies from old baboons, indicating a disruption of tight junction barrier function. Whereas ZO-1, occludin, and JAM-A function in adhering the tight junctions together, claudin-2 creates "leaky" pores through the tight junctions (36) (36) . In the current study, although we found via Western blot analysis that claudin-2 levels were increased suggestive of enhanced gut permeability [\(36](#page-10-22)), this increase in claudin-2 expression was not confirmed via immunofluorescence. The discrepancy in the data remains to be resolved, but we postulate that the further activation of age-independent mechanisms may be required to induce insertion of claudin-2 proteins into the epithelial membrane. In summary, our results show that a pivotal factor contributing to the enhanced permeability in colonic biopsies from old baboons is senescence-induced remodeling of tight junction proteins, and taken together, these findings extend the previous studies in rodents, which demonstrated an increased intestinal permeability with age.

The mechanisms underlying these potential age-associated remodeling of tight junctions remain to be determined. However, recent evidence suggests that miRNAs are involved in regulating gut permeability via a decrease in the expression of tight junction proteins in response to specific environmental factors. For example, alcohol exposure increases gut "leakiness" by downregulating

Figure 3. Aging restructures colonic epithelial tight junctions. Confocal micrographs of cross-sections imaged from colonic biopsies are shown for young (left) and old (middle) baboons in addition to a quantification of the fluorescent optical densities of the micrographs (right). Overall, there was a significant decrease in zonula occluden-1 (ZO-1; **A**), occludin (**B**), and junctional adhesion molecule (JAM)-A immunofluorescence (**C**). Although there was a trend toward increased claudin-2 expression, the change was not statistically significant (**D**). Statistical significance was **p* < .05 by Student's unpaired *t* test.

Figure 4. Aging alters expression of tight junctions. Western blot was used to quantify total expression levels of tight junction proteins. Quantification of the optical densities is shown on the left panel with representative blots on the right. Detection by chemiluminescence revealed positive bands for the tight junction proteins zonula occluden-1 (ZO-1) at ~225kDa (**A**), occludin at ~65kDa (**B**), junctional adhesion molecule (JAM)-A at ~36kDa (**C**), and claudin-2 at ~22kDa (**D**). Glyceraldehyde 3-phosphate dehydrogenase (~36kDa) was used to normalize ZO-1, occludin, and claudin-2 optical densities, and β-actin (~42kDa) was used to normalize JAM-A. There was a significant decrease in ZO-1, occludin, and JAM-A expression in contrast to a significant increase in claudin-2 protein expression. Statistical significance is **p* < .05 by Student's unpaired *t* test.

 $_{\rm old}$

 0.0

Young

Figure 5. MicroRNA and glutamine synthase expression in colonic biopsies. With age there was a significant increase in miR-29a but not miR-122a or miR-212 expression within the colonic biopsies (**A**). There was no significant difference in GLUL expression in the same samples (**B**). Statistical significance was **p* < .05 by Student's unpaired *t* test.

Figure 6. From the cytokine panel, there was a significant increase in IFN-γ, IL-6, and IL-1β expression in aged colonic tissue compared with young colonic tissue. There was also a trend toward increased IL-12 production that was not statistically significant. Statistical significance was **p* < .05 by Student's unpaired *t* test.

ZO-1 proteins though a mechanism involving increased miR-212 expression [\(37\)](#page-10-23). Moreover, TNF- α has been shown to enhance intestinal permeability by inducing miR-122-mediated degradation of occludin mRNA ([14\)](#page-10-0). Our study showed a significant upregulation of miR-29a in colonic mucosal tissue from aged animals with enhanced gut permeability, which is supported by a previous study that found an intrinsic upregulation in miR-29a in normal aged mice when compared with young mice ([38\)](#page-10-24) and another study showing that miR-29a was increased in irritable bowel syndrome patients with enhanced gut permeability (15) (15) .

There is also evidence that suggests a role for glutamine in the regulation of gut permeability. Glutamine supplementation reverses abnormal paracellular permeability induced by nutrient deprivation in caco-2 cells ([39](#page-10-25)) and in clinical experiments involving elderly patients [\(40](#page-10-26)). GLUL catalyzes the endogenous production of glutamine, and therefore changes in permeability may result from dysfunction in GLUL expression, as reported in irritable bowel syndrome patients with enhanced gut permeability [\(15](#page-10-1)). We therefore examined the expression of GLUL in our colonic biopsies taken from young and old baboons. In our study, there was no difference in intestinal GLUL expression in tissue taken from aged baboons compared with young, suggesting normal glutamine production.

Immune system–gut interactions, in particular cytokine expression in the epithelium, were examined as a potential mechanism for the enhanced permeability of the mucosal barrier observed in aged animals. Specific cytokines produced by the intraepithelial leukocytes, such as eosinophils, mast cells, macrophages, and TCRγδ-positive T cells ([41](#page-10-27)[,42](#page-10-28)), have been documented to regulate tight junction proteins [\(16](#page-10-2),[17\)](#page-10-3). In support of neuroimmune-mediated mechanism accounting for the increase in gut permeability observed in response to aging, we found significant increases in IFN-γ, IL-6, and IL-β in colonic biopsies from aged animals. In other studies, IFN-γ exposure has been shown to

(i) decrease epithelial resistance while increasing epithelial permeability ([43\)](#page-10-29) and (ii) induce endocytosis of tight junction proteins including occludin and JAM-A ([44,](#page-10-30)[45\)](#page-10-31). Systemic IFN-γ has also been shown to increase with age in human plasma without the presence of inflammation or infection [\(46](#page-10-32)), supporting our data and implicating IFN-γ as an important factor in age-related changes in gut permeability. IL-6 has been shown to decrease the expression of ZO-1 in the intestine of mice [\(47\)](#page-10-33) and increase claudin-2 expression ([48\)](#page-10-34). Furthermore, previous studies demonstrate that there is a significant increase in human and rhesus monkey plasma IL-6 with age without inflammation or infection, which supports the data in this study identifying IL-6 as an important mediator of age-associated increase in intestinal permeability (49) (49) . IL-1 β is also a proinflammatory cytokine that causes an increase in intestinal epithelial permeability ([50\)](#page-10-36) and has been shown to increase with age in human plasma [\(51](#page-10-37)). Remodeling of tight junction proteins in response to IL-1β includes the downregulation of occludin [\(52](#page-10-38)), increased claudin-2 expression and hyperphosphorylation of occludin [\(53](#page-11-0),[54\)](#page-11-1). Interestingly, not only does our data show decreased occludin and increased claudin-2 but we also show a second occludin band in our Western blot experiments in old baboon colonic tissue, which may represent differential phosphorylation states of occludin. Endogenous changes in cytokine production as a function of age can be caused by a number of factors including remodeling of the gut microbiome. In the colon, studies have shown that the microenvironment changes drastically with age ([55,](#page-11-2)[56\)](#page-11-3), which can significantly affect local immunity [\(57](#page-11-4)). Although primary studies in humans have implicated a remodeling of the gut microbiota in facilitating a proinflammatory state in the elderly people [\(58\)](#page-11-5), future studies in nonhuman primate models, beyond the scope of the current study, could more definitively determine the relationship between immunosenescence in the gut and the microbiome.

Other cytokines examined in the current study include IL-12 and TNF-α, which were previously demonstrated to affect tight junction proteins but were unchanged in our aging baboon model. Furthermore, we were unable to detect the anti-inflammatory cytokines IL-10 and IL-2, which function to decrease permeability. The inability to detect IL-10 and IL-2 may be because these cytokines are not constitutively present and expressed only upon activation such as in the case of IL-10 ([59\)](#page-11-6) or alternatively may be expressed in cells located in Peyer's patches, which are deeper in the lamina propria and may not have been captured by the biopsies. It is important to note in the interpretation of our data that the increases in cytokines expression measured in the current study occurred in the absence of any inflammation within the colonic biopsies, which had normal histological appearance and levels of MPO activity resembling that in biopsies from younger animals. Moreover, because the cytokine levels measured were much lower than profiles from inflamed tissue (60) (60) , we can assume that the changes in cytokine expression are due to intrinsic changes in cytokine expression with age and not due to extrinsic induction of inflammation and immune infiltration.

In conclusion, our data in a nonhuman primate model suggest that normal aging increases intestinal permeability in colonic biopsies through the restructuring of tight junction proteins. Moreover, we have shown that via a mechanism involving increased miR-29a expression and/or enhanced inflammatory cytokine expression with age, there is increased intestinal permeability that may enhance the susceptibility to age-related GI disorders.

FUNDING

This work was supported by the National Center for Research Resources/ National Institute of Health (P40RR012317).

Acknowledgments

The authors would like to acknowledge Gary L. White, DVM, MMS, for providing funding and Roman F. Wolf, DVM, for assistance in collecting the colonic biopsies. The authors would also like to acknowledge the Oklahoma Medical Research Foundation Imaging Core for their assistance and use of their confocal microscope, and Stan Lightfoot, MD, and Megan Lerner for the preparation and examination of the colonic histology.

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