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Extracorporeal Membrane Oxygenation Causes Loss of Intestinal Epithelial Barrier in the Newborn Piglet

Ashish R. Kurundkar, Cheryl R. Killingsworth, R. Britt MclLwain, Joseph G. Timpa, Yolanda E. Hartman, Dongning He, Rajendra K. Karnatak, Mary Lauren Neel, John P. Clancy, G. M. Anantharamaiah, and Akhil Maheshwari

Departments of Pediatrics [A.R.K., Y.E.H., D.H., R.K.K., M.L.N., J.P.C., A.M.], Medicine [C.R.K., G.M.A.], Pathology [A.M.], and Cell Biology [A.M.], University of Alabama at Birmingham (UAB), Birmingham, AL 35294; University Hospital Services [R.B.M., J.G.T.], University of Alabama at Birmingham, Birmingham, AL35294; Clinical Perfusion Education [R.B.M.], School of Allied Health Professions, University of Nebraska Medical Center, Omaha, NE 68198

Abstract

Extracorporeal membrane oxygenation (ECMO) is an important life-support system used in neonates and young children with intractable cardiorespiratory failure. In this study, we used our porcine neonatal model of venoarterial ECMO to investigate whether ECMO causes gut barrier dysfunction. We subjected 3-week-old previously-healthy piglets to venoarterial ECMO for up to 8 hours and evaluated gut mucosal permeability, bacterial translocation, plasma levels of bacterial products, and ultrastructural changes in gut epithelium. We also measured plasma lipopolysaccharide (LPS) levels in a small cohort of human neonates receiving ECMO. In our porcine model, ECMO caused a rapid increase in gut mucosal permeability within the first 2 hours of treatment, leading to a 6–10 fold rise in circulating bacterial products. These changes in barrier function were associated with cytoskeletal condensation in epithelial cells, which was explained by phosphorylation of a myosin II regulatory light chain. In support of these findings, we also detected elevated plasma LPS levels in human neonates receiving ECMO, indicating a similar loss of gut barrier function in these infants. Based on these data, we conclude that ECMO is an independent cause of gut barrier dysfunction, and that bacterial translocation may be an important contributor to ECMO-related inflammation.

INTRODUCTION

Extracorporeal membrane oxygenation (ECMO) is an important cardiopulmonary lifesupport system used in critically-ill neonates and young children with respiratory failure, congenital heart disease, and overwhelming sepsis (1,2). While the availability of ECMO has reduced mortality in these patients by nearly 80%, concerns remain about the nearuniversal occurrence of a systemic inflammatory response syndrome (SIRS) during ECMO that is associated with considerable morbidity. ECMO-related SIRS is particularly severe in neonates (than in older children) (3,4), manifesting with the first few hours of ECMO with

Corresponding author: Akhil Maheshwari, MD, University of Alabama at Birmingham, 1670 University Boulevard, VH648C, Birmingham, AL 35294, (akhil@peds.uab.edu).

A.R.K. and C.R.K. contributed equally to this study.

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hypotension, oliguria, decreased lung compliance, anasarca, and liver dysfunction. These changes frequently persist for several days and prolong the duration of ECMO (1). Understanding ECMO-related SIRS is a critical step in the development of effective anti-inflammatory strategies.

Our understanding of the inflammatory effects of extracorporeal circulation is based primarily on studies on adult patients treated with cardiopulmonary bypass (CPB) during cardiac surgery (5). However, direct extrapolation of data from CPB to ECMO is difficult because of major pathophysiological differences between the two modalities, such as the presence of defined ischemia-reperfusion sequences in CPB (related to the placement and release of aortic cross-clamps during cardiac surgery) but not in ECMO (5). To investigate the mechanisms of ECMO-related SIRS, we have developed a porcine neonatal model of ECMO where we subject previously-healthy, 3–6 kg piglets to venoarterial ECMO in the laboratory (6). Porcine ECMO was associated with inflammatory changes similar to those reported in human studies, causing tachycardia, hypotension, and a generalized capillary leakiness manifesting as edema and ascites within the first few 2-4 hours of initiation of the procedure. ECMO caused neutrophil activation, increased expression of inflammatory cytokines in plasma and tissues, and histopathological changes of inflammation (leukocyte infiltration) and microvascular injury (focal hemorrhages, edema) in diverse organs such as lung, intestine, liver, and kidney. We showed that the initiation of ECMO was associated with mast cell degranulation, which released large amounts of pre-formed pro-inflammatory cytokines such as tumor necrosis factor (TNF/TNF- α) and mast cell-derived enzymes such as tryptase into the circulation. Since ECMO causes mast cell degranulation, which in turn, releases TNF, tryptase and chymase that can disrupt the tight junctions of the intestinal epithelial cells (IECs) (7,8), we postulate that ECMO induces gut barrier dysfunction which in turn leads to bacterial translocation and increased inflammation. In this study, we used our porcine model to test this hypothesis.

METHODS

Neonatal porcine ECMO

The ECMO procedure has been described previously (6). Briefly, mixed-breed neonatal piglets of either gender weighing 3–6 kg were subjected to venoarterial ECMO after approval by the Institutional Animal Care and Use Committee. Sham animals received anesthesia, ventilation, cannulation, and heparinization similar to ECMO animals, but were not connected to the E pump. Data in this study represent 8 animals each in sham and ECMO groups; 3 animals in each group were euthanized after 2 hours of treatment and the remaining 5 after 8 hours. ECMO-related SIRS in these animals has been described previously (6).

The piglets received general anesthesia and mechanical ventilation (volume controlled, tidal volume = 15 mL/kg, 10–15 cycles/minute, Hallowell EMC 2000 ventilator) to maintain normal blood gases. Biomedicus 8F cannulae (Medtronic, Minneapolis, MN) were inserted into the external jugular vein and the external carotid artery. Animals were heparinized to maintain activated clotting times (ACT) of 180–220 seconds. The ECMO system consisted of a Biomedicus BP-50 centrifugal pump (Medtronic, Shoreview, MN), and a Minimax hollow fiber oxygenator (Medtronic). Gas flow rates to membrane oxygenator were maintained at 0.5 L/min of 100% oxygen, and flow rates in the circuit were advanced to 250 mL/min or 1.5 L/min/m².

Measurement of bacterial products in plasma and detection of bacteria in tissues

Lipopolysaccharide (LPS) (Lonza, Walkersville, MD) and *D*-lactate (Bioassay Systems, Hayward, CA) were measured by commercially-available colorimetric assays. Lipoteichoic acids (LTA) were detected in plasma by immunoprecipitation and western blotting using our previously described protocol (9). Aerobic/anaerobic blood cultures were processed at the hospital clinical laboratory. Bacteria were identified in intestinal tissue sections using the Brown and Brenn stain (10).

Measurement of gut mucosal permeability

We used an established protocol for ex vivo measurement of gut mucosal permeability (11). Briefly, freshly-dissected jejunal or colonic mucosa was mounted in Ussing diffusion chambers (0.5 cm²; Physiologic Instruments, San Diego, CA) in Krebs-Ringer bicarbonate buffer (pH 7.4), We added fluorescein isothiocyanate (FITC)-dextran 4400 (2.2 mg/mL) tothe apical reservoirs \times 60 min, after which media were collected from apical and basal chambers and fluorescence was measured at 485/520 nm. Tissue specimens were examined histopathologically to confirm epithelial integrity.

Real-time polymerase chain reaction (PCR)

Expression of tight junction genes was measured by real-time PCR per our previouslydescribed protocol (12). Primers were designed using the Beacon Design software(Bio-Rad, Hercules, CA).

Transmission electron microscopy (TEM)

TEM analyses were performed at the UAB High Resolution Imaging Facility. Tissue samples of jejunum (about 10 inches beyond the ligament of Treitz) and the mid-transverse colon were harvested and fixed in 2% glutaraldehyde and then in 1% osmium tetroxide. Thin sections were examined with a JEOL 1200-EX/II transmission electron microscope at 80 kV. Computer-assisted morphometric measurements of intercellular spaces were made using the Metamorph software (Molecular Devices, Sunnyvale, CA).

The permeability of IEC apical junctions was evaluated using the electron-dense dye ruthenium red (13). Briefly, tissue samples were placed (luminal side up) in plastic chambers with an apical 1 cm² circular opening and the exposed to 2.5% glutaraldehyde and 0.6% ruthenium red in 0.1 M sodium cacodylate buffer, pH 7.3 × 30 min and then to 2% osmium tetroxide and ruthenium red × 30 min. Thin sections were evaluated by the presence of electron-dense RR deposits in the intercellular space between IECs. A total of 10 tight junctions were analyzed from each mucosal region.

Detection of myosin light chain (MLC) and phospho-MLC

IECs were isolated by treating intestinal tissue with HBSS containing 0.2 M ethylenediaminetetraacetic acid (EDTA), 10 mM 2-mercaptoethanol, and phosphatase and kinase inhibitors (Sigma) \times 30 min. Western blots were performed on cell lysates for phosphorylated (ser19) and total myosin II light chain (Cell Signaling Technology), using our previously-described protocol (12,14).

Plasma samples from human neonates treated with ECMO

The study was performed under appropriate oversight by the Institutional Review Board on discarded sera/plasma at the clinical laboratory from patients during the first 48 hrs of ECMO (n = 8). We included a control group (n = 8) with comparable gestational ages and birth weights who were admitted to the NICU for severe respiratory distress and received mechanical ventilation but did not require ECMO. Patients were not selected as controls if

they had perinatal depression (5 min Apgar score \leq 3) or culture-positive early-onset infection. LPS concentrations were measured using the limulus lysate assay as described above.

Statistical methods

Non-parametric tests were applied using the SigmaStat 3.1.1 software (Systat, Point Richmond, CA). The number of samples in each experiment is indicated in figure legends. Groups were compared by the Mann-Whitney U (Wilcoxon rank-sum test) or the Kruskall-Wallis H tests (with Dunn's post-test). Data are expressed as means \pm standard errors (SE). A p value of <0.05 was considered significant.

RESULTS

Bacterial translocation during ECMO

To determine whether ECMO caused important changes in gut barrier function, we first measured LPS and bacterial products in plasma samples from sham and ECMO piglets. As shown in Fig. 1A, plasma LPS increased rapidly within the first 2 hours of ECMO but not in sham animals. Similarly, plasma concentrations of *D*-lactate, a specific product of prokaryotic metabolism, also increased within the first 2 hours (Fig. 1B). We also probed for LTA, a component of the Gram-positive bacterial cell wall, in plasma samples by immunoprecipitation and western blotting and observed a similar increase in plasma LTA during ECMO (Fig. 1C). Blood cultures obtained from sham and ECMO animals remained negative, indicating that bacterial translocation remained limited at low levels during ECMO.

The rapid increase in circulating bacterial products we observed within 1–2 hours of initiation of ECMO correlated with our earlier findings of a well-established inflammatory response within the first 2 hours of ECMO (6). Therefore, to elucidate the mechanism of gut barrier dysfunction during ECMO, we focused on the 2-hour time point in subsequent experiments. To confirm that increased plasma levels of bacterial products were associated with increased bacterial translocation in the intestine, we stained intestinal tissue from sham and ECMO-treated animals for bacteria using the Brown and Brenn method. As shown in Fig. 1D, bacteria were readily detected (28 ± 3 bacteria/100 IECs) in the ECMO intestine, which was in sharp contrast with <1 bacterium seen per 100 IECs in the sham intestine (p<0.05).

ECMO increases gut mucosal permeability

In-vivo methods to assess gut mucosal permeability (15) were considered but not found suitable for our studies because of unpredictable effects of general anesthesia on bowel motility (16,17). We used an *ex vivo* method to measure the penetration of FITC-dextran, a marker of paracellular permeability, across freshly-dissected and *ex* planted specimens of jejunal and colonic mucosa (11). As seen in Fig. 2, treatment with ECMO for 2 hours increased permeability in the jejunal mucosa by nearly 8-fold (1236±281 in sham to 8226±1282 fluorescence units in ECMO group; p<0.05). Similarly, ECMO also increased permeability in the colonic samples (6987±289 units in ECMO vs. 1364±436 units in sham; p<0.05; effect not significantly different from jejunal samples).

Increased gut mucosal permeability during ECMO is not due to decreased expression of genes involved in the assembly of the tight junctions

To determine the mechanisms by which ECMO increased gut mucosal permeability, we first used a PCR array to measure the expression of 25 genes directly involved in tight junction assembly. As shown in Fig. 3, the expression of occludin, several major claudins, and the

zonula occludens protein-1 (ZO-1) remained unchanged or increased over sham after 2 hours of ECMO. In view of these findings, we hypothesized that ECMO-induced changes in gut mucosal permeability were mediated via reorganization, rather than reduced expression, of junctional proteins and therefore, next evaluated ultrastructural changes by transmission electron microscopy.

ECMO is associated with ultrastructural abnormalities in the intestinal epithelium and increased junctional permeability

ECMO was associated with loss of intercellular contact with increased intercellular spaces in IECs, which contrasted with the close apposition of cell membranes in the sham intestine (sham 6.5 ± 2 nm vs. ECMO 101.2 ± 11.7 nm; *p*<0.05; Fig. 4A). Similar changes were seen in the colon (not depicted).

To determine whether ECMO caused increased permeability of the apical junctional complex, we applied the electron-dense dye ruthenium red to the apical surface of some explants prior to fixation for electron microscopy. As shown in Fig. 4B, ruthenium red penetrated the apical junctions in the ECMO intestine and rendered those areas electron-dense (arrows), thereby indicating the functional disruption of these junctions.

ECMO causes cytoskeletal contraction in IECs

In the ECMO intestine, we noted electron-dense areas of perijunctional cytoskeletal condensation in the IECs (Fig. 5A). Consistent with previous reports (18), these changes were particularly prominent near tricellular junctions.

To investigate the intracellular signaling events by which ECMO caused cytoskeletal contraction in IECs, we considered possible mechanisms involving MLC phosphorylation, rho-rho kinase-mediated signaling, and calcium/calmodulin-activated events (7,19). Because inflammatory cytokines such as TNF can promote phosphorylation of the myosin II regulatory light chain (MLC) in IECs (7,8,20), we considered MLC phosphorylation as the most plausible mechanism for the changes we observed during ECMO-related SIRS. Using *in silico* sequence alignment, we first localized a conserved ser19 in porcine MLC that has been previously identified as a phosphorylation target in human and murine MLC. We next measured phosphorylated and total MLC in IECs by western blotting. As seen in Fig. 5B, ECMO increased MLC phosphorylation in IECs.

Human neonatal ECMO is associated with increased plasma LPS concentrations

Finally, to confirm the clinical relevance of our findings in the porcine model, we measured plasma LPS levels in a small cohort of neonates during the first day of ECMO therapy (n=8). Diagnoses among the ECMO patients included meconium aspiration syndrome (n = 7) and congenital diaphragmatic hernia (n = 2). Six neonates were on veno-venous ECMO and the other 3 on veno-arterial ECMO. The control group included neonates receiving mechanical ventilation for severe respiratory distress, who were not septic (blood culture was negative) and did not require ECMO. The gestational age and birthweight of neonates on ECMO (gestational age 39 ± 2 weeks, birthweight 3098 ± 572 grams) were similar to the control group (38.6 ± 2.7 weeks, 3289 ± 342 grams). Plasma LPS concentrations were measured using the limulus lysate assay. As seen in Fig. 6, plasma levels were higher in neonates receiving ECMO therapy than those treated with mechanical ventilation alone. While we cannot ascertain the contribution of the underlying disease to LPS elevations we observed in these samples, these data provide indirect support for our findings in the porcine model.

DISCUSSION

We present the first detailed investigation into the effects of ECMO on gut epithelial barrier and bacterial translocation. Existing evidence on ECMO-related SIRS emphasizes the role of neutrophils (activated in the ECMO circuit) (21-24), which cause diffuse microvascular injury and multi-organ dysfunction (21-25). In the present study, we detected rapid deterioration of the intestinal barrier function and a consistent rise in circulating bacterial products almost immediately after the initiation of ECMO. Because bacterial products can independently initiate/amplify all the major inflammatory pathways involved in ECMOrelated SIRS (26), strategies to protect gut barrier function and thereby prevent bacterial translocation during ECMO merit careful evaluation. Our findings in the porcine model are consistent with previous observations of Piena and colleagues, who reported abnormalities in gut mucosal permeability in human neonates receiving ECMO (27). To assess mucosal integrity before initiating enteral feedings, these investigators measured intestinal permeability by measuring urinary excretion of enterally-administered lactulose, rhamnose, D-xylose, and 3-O-methyl-D-glucose and noted abnormal mucosal permeability in 13 of their 16 patients. Similarly, Hirthler and coworkers (28) have previously reported increased plasma LPS levels in neonates who showed signs of inflammation \geq 36 hours after the initiation of ECMO. Bacterial translocation has also been noted following CPB used for cardiac surgery, where it correlated with ischemia-reperfusion injury sustained during the placement and subsequent release of aortic cross-clamps (5,29).

In the present study, we have identified cytoskeletal condensation in IECs as a mechanism for the rapid disruption of the gut epithelial barrier during ECMO. Phosphorylation of the MLC and consequent cytoskeletal contraction are emerging as a common, central event in the disruption of the epithelial barrier in diverse conditions such as inflammatory bowel disease, graft versus host disease, and infectious enterocolitis (7,19). In epithelial cells, tight junctions are affiliated with an apical perijunctional actin belt, which may contract to generate mechanical forces strong enough to retract the plasma membrane and cause internalization of various components of the apical junctional complex into subapical cytosolic compartments (30,31). Whereas inflammatory mediators such as TNF are the likely cause of MLC phosphorylation and cytoskeletal contraction in IECs during ECMO, additional procedure-related factors such as exposure to citrate in the anticoagulant solutions in banked blood also need careful evaluation. Citrate can deplete extracellular calcium (32), which, in turn, can trigger cytoskeletal condensation and disassembly of tight junctions (33– 35). Anticoagulant solutions assume great importance in neonates and young infants receiving ECMO because volumes of the ECMO circuit are often 200-300% larger than a neonate's circulating blood volume, resulting in massive exposures to these reagents per unit weight (3,4). While most anticoagulants can be eliminated to a large extent by saline prewashing of the re blood cells (RBC), the procedure may increase RBC osmotic fragility and hemolysis in centrifugal pumps (36). Rigorous clinical studies are needed to compare the effects of various anticoagulant solutions on ECMO-related SIRS.

The present study provides us with two important conclusions that might have been difficult to draw in clinical studies: (1) by subjecting previously-healthy piglets to ECMO, we established ECMO as an independent cause of gut barrier dysfunction. In clinical studies, the relative contribution of ECMO to barrier dysfunction cannot be dissected from that of the underlying disease that led to ECMO in the first place; bacterial translocation is common during hypoxemic respiratory failure, congestive cardiac failure, intractable sepsis, and shock, conditions that form the commonest indications for ECMO (1,26); (2) development of gut barrier dysfunction and bacterial translocation early during the course of ECMO suggests that barrier dysfunction is more likely to be a primary initiator/amplifier of ECMO-related SIRS and not merely an epiphenomenon reflecting loss of mucosal integrity as a part

ECMO-related SIRS cannot be prevented merely by improvements in the ECMO circuit. The physical characteristics of the ECMO circuit such as fluid dynamics, blood volume to surface area ratio, and the material's affinity for fibrinogen determine its propensity for contact activation of inflammatory pathways (37). Although smoother circuits can be designed, the structure of the membrane oxygenator calls for conflicting profiles - the necessity for the gas exchange obligates thin blood films and turbulence, factors that also favor contact activation. Thus, even though improved silicone oxygenators have shown modest benefit in reducing platelet and contact activation, these improvements face a mathematical bottleneck where reduction in turbulence and blood surface interaction compete with the gas exchange capacity of the oxygenator (38). In this context, elucidation of the inflammatory pathways involved in ECMO-related SIRS is a critical step in the development of effective anti-inflammatory therapies. Since potent inhibitors of MLC kinase are now available (39), the identification of MLC phosphorylation as a likely mechanism of gut barrier dysfunction during ECMO indicates a need for further investigation of this pathway in preclinical and clinical settings.

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List of abbreviations

CPB	cardiopulmonary bypass
ECMO	extracorporeal membrane oxygenation
IEC	intestinal epithelial cell
LPS	lipopolysaccharide
LTA	lipoteichoic acids
MLC	myosin light chain
SIRS	systemic inflammatory response syndrome

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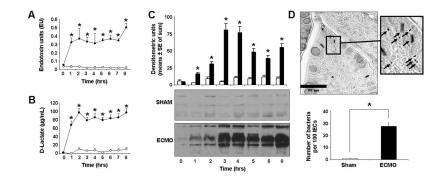


Fig.1. ECMO-mediated gut barrier dysfunction results in bacterial translocation

A. B. Plasma LPS (*panel A*) and D-lactate (*panel B*) levels increased rapidly during ECMO (n=5 animals/group). Line diagrams show means \pm SEM values as a function of time. Black circles indicate data points from the ECMO group, while open circles depict the sham animals. * indicates p<0.05 when compared to sham at corresponding data points. C. Representative western blots show bacterial LTA in plasma. Bar graph of plasma lipoteichoic acid represents the sum of densities at each point (time after initiation of ECMO) for the 3 ECMO and 3 sham control animals. * indicates p<0.05 when compared to sham at corresponding data points. C is an at corresponding data points. D. Brown and Brenn stain on intestinal sections (magnification 400x) reveals bacteria in the ECMO intestine. Inset (1500x) shows rods and cocci (arrows). Bar diagram shows means \pm SE of the number of bacteria counted per 100 IECs in all 3 ECMO animals (2 hrs after initiation of ECMO) and sham controls. * indicates p<0.001 when compared to sham at corresponding data points.

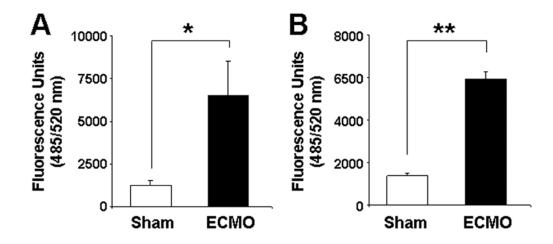


Fig.2. ECMO increases gut mucosal permeability

ECMO increased the mucosal permeability in *ex* planted jejunal (*panel A*) and colonic (*panel B*) samples, measured as mucosal-to-basal penetration of FITC-dextran 4400 across tissue specimens in 60 min (n=3–4 tissue replicates per animal. N=3 animals per group; tissue replicates from each animal were averaged and used as a single data point. * indicates p<0.001, ** indicates p<0.001.

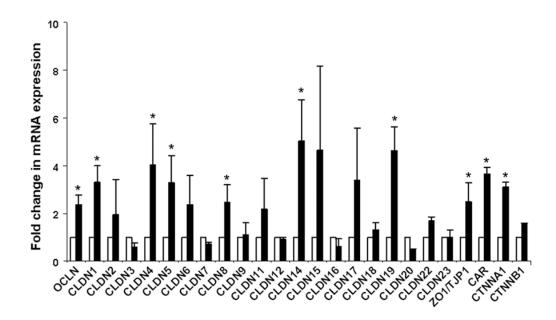


Fig.3. Increased gut mucosal permeability during ECMO is not due to decreased expression of genes involved in the assembly of the tight junctions

Bar diagram (means \pm SEM) depicts the expression of 25 porcine genes involved in tight junctions after 2 hours of ECMO (black bars), shown as fold change in mRNA expression above sham (grey bars marked at fold change = 1); n= 3 animals/group. The expression of 20 of these 25 genes was increased (80%). * indicates *p*<0.05 when compared to sham. Analysis for reduced gene expression by calculating 95% confidence intervals did not show significant reductions. Claudin 10 mRNA was not detected in either sham or ECMO intestine. Abbreviations: OCLN = occludin, CLDN1-23 = claudin 1–23, ZO1/TJP1 = zonula occludens protein-1/tight junction protein-1, CAR = coxsackie-adenovirus-receptor homolog, CTNNA1 = catenin (cadherin-associated protein), alpha 1, CTNNB1 = catenin (cadherin-associated protein), beta 1.

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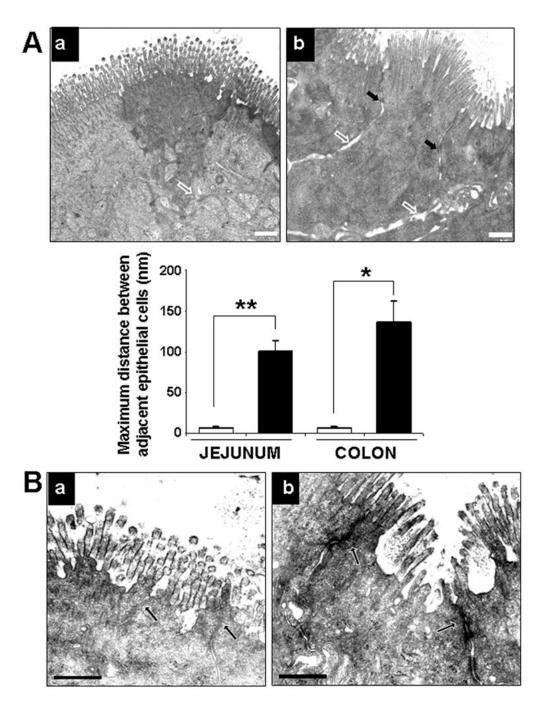


Fig.4. Ultrastructural changes in IECs in the ECMO intestine

A. Loss of intercellular apposition in IECs in the ECMO intestine: Transmission electron micrographs (TEMs) of (a) sham and (b) ECMO intestine showing increased separation of IECs after 2 hours of ECMO (open arrows), except at desmosome sites (black arrows). Bar diagram (means \pm standard error) on the lower left summarizes the distances between adjacent epithelial cells in sham and ECMO intestine. Distance bar represents 500 nm. Data include junctions in 2 villi (or colonic crypts) in 3 randomly chosen areas. * indicates p < 0.001, ** indicates p < 0.001. B. Increased junctional permeability in the ECMO intestine: TEMs of (a) sham and (b) ECMO intestine show increased penetration of the

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electron-dense dye ruthenium red in the ECMO intestine (arrows). Distance bar represents 500 nm.

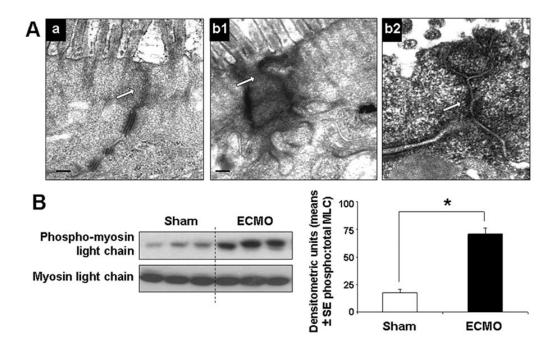


Fig.5. ECMO causes cytoskeletal contraction in IECs

A. TEMs of (a) sham and (b1) ECMO intestine (magnification 15000x). Unlike the uniform appearance of the perijunctional IEC cytoskeleton in the sham intestine, IECs in the ECMO intestine show electron-dense areas of cytoskeletal condensation (arrows). High-magnification image b2 shows prominent cytoskeletal condensation near a tricellular junction. **B. ECMO increases the phosphorylation of myosin II light chain (MLC) in IECs:** Western blots show increased phosphorylation of MLC in IECs. Bar diagrams shows densitometric analysis (means \pm SE densitometric units) of the three bands (each representing a different animal) from sham and ECMO groups. Data were confirmed by repeating the blots. * indicates p < 0.001, ** indicates p < 0.001.

