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Intestinal Lymphatic Dysfunction in Kidney Disease

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ABSTRACT: Kidney disease is associated with adverse consequences in many organs beyond the kidney, including the heart, lungs, brain, and intestines. The kidney-intestinal cross talk involves intestinal epithelial damage, dysbiosis, and generation of uremic toxins. Recent studies reveal that kidney injury expands the intestinal lymphatics, increases lymphatic flow, and alters the composition of mesenteric lymph. The intestinal lymphatics, like blood vessels, are a route for transporting potentially harmful substances generated by the intestines. The lymphatic architecture and actions are uniquely suited to take up and transport large macromolecules, functions that differentiate them from blood vessels, allowing them to play a distinct role in a variety of physiological and pathological processes. Here, we focus on the mechanisms by which kidney diseases result in deleterious changes in intestinal lymphatics and consider a novel paradigm of a vicious cycle of detrimental organ cross talk. This concept involves kidney injury–induced modulation of intestinal lymphatics that promotes production and distribution of harmful factors, which in turn contributes to disease progression in distant organ systems.

Key Words: intestines ■ kidney ■ lymphatic ■ renal insufficiency, chronic

K idney disease is a growing global epidemic with significant health care and economic consequences.
Even modest kidney impairment dramatically increases extrarenal morbidity, most notably as it relates idney disease is a growing global epidemic with significant health care and economic consequences. Even modest kidney impairment dramatically to cardiovascular disease (CVD) but also infection, bone disease, cognitive dysfunction, and cancer.¹ For example, impaired kidney function is a more important predictor of CVD than traditional cardiovascular risks such as elevated blood pressure, cholesterol, and diabetes.² Nonetheless, questions remain regarding the mechanistic basis of how kidney disease affects distant organs.

Recent studies have demonstrated a central role for the gut in mediating kidney disease–related complications.3 Kidney disease disrupts the intestinal barrier and modulates the composition and metabolism of the intestinal microbiome, producing bioactive metabolites and toxins.4 Traditionally, blood vessels and nerves were thought to be the primary pipelines by which diseaseassociated factors could spread from the gut to distant organs. Only recently have researchers turned their attention to the lymphatics, which route fluids, macromolecules, and cells from the interstitial compartments of virtually every organ for return into the systemic circulation. The intestinal lymphatics are unique within the lymphatic vascular network because, in addition to the usual functions, they are responsible for absorption, remodeling, and transport of dietary lipids and intestinally generated lipoproteins, making them a potentially significant player in the dissemination of gut-borne disease-associated factors. Thus, disruptions in lymph transport and lymphatic vessel integrity are powerful potentiators of disease, including CVD, inflammatory bowel disease, bone disease, cognitive dysfunction, and chronic kidney disease.

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This review examines the novel paradigm of a vicious cycle of detrimental organ cross talk by which kidney injury–induced modulation of intestinal lymphatics contributes to systemic organ disease progression.

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Nonstandard Abbreviations and Acronyms

LYMPHATIC SYSTEM

Lymphatic Form and Function

The lymphatic system comprises an extensive vascular network punctuated by lymphoid organs, which plays an essential role in interstitial fluid balance, immune surveillance, and lipid transport. Recently, the lymphatic system has also been recognized as a facilitator of interorgan cross talk under physiological and pathological conditions.3,5,6 Divided into 3 classes, lymphoid organs are central components of the body's immune response. Primary lymphoid organs (bone marrow and thymus) contain lymphoid progenitor cells required for lymphocyte differentiation. Secondary lymphoid organs, also called mucosa-associated lymphoid tissues, include lymph nodes, the spleen, tonsils, Peyer patches, and the appendix. These organs facilitate adaptive immune responses by providing the environment in which naive lymphocytes are activated and maintained, resulting in the antigendriven expansion of memory T cells, effector B cells, and plasma cells.7 Tertiary lymphoid organs include collections of immune cells that resemble secondary organs but form in peripheral tissues in response to chronic inflammation or tissue injury.⁸ While lymphoid organs provide cellular components of the body's immune response, lymphatic vessels facilitate their trafficking.

The lymphatic vessel network courses through every vascularized organ of the body, with the exception of bone marrow.⁹ In the interstitium, immune cells, macromolecules, and excess fluid (collectively referred to as lymph) enter highly permeable, blind-ended lymphatic capillaries that measure 30 to 80 μm in diameter. Lymph then drains into larger precollector and collecting vessels.

These vessels differ from lymphatic capillaries in that they are considerably less permeable, become increasingly covered by smooth muscle cells, and contain intraluminal valves that help facilitate unidirectional pumping of lymph toward the ≈600 to 800 lymph nodes dispersed throughout the body. Within the node, lymph is screened for antigens that mediate adaptive immune responses and self-antigen tolerance¹⁰ and then funneled toward the thoracic duct, where it reenters the systemic circulation at the junction of the left subclavian and internal jugular veins.

Lymphatic Endothelial Cells

Lymphatic endothelial cells (LECs) line lymphatic vessels and lymph nodes and form the intraluminal valves of collecting vessels. During development, most LECs are derived from a subpopulation of LYVE-1 (lymphatic vessel endothelial hyaluronan receptor 1)/PROX1 (prospero homeobox 1)-positive venous endothelial cells located in the cardinal vein.11 Discrete nonvenous and hematogenic sources of LEC progenitor cells have also been described and are thought to give rise to portions of the dermal, mesenteric, and cardiac lymphatic vasculature.¹²⁻¹⁴ Once specified, LECs migrate and proliferate, forming rudimentary lymph sacs and a primitive plexus of vessels that undergo extensive remodeling to establish the complex network of lymphatic vessels and organs. The initial stimulus for lymphatic progenitor cell exit from veins is dependent on VEGFC (vascular endothelial growth factor C) signaling.15 VEGFC activates both VEGFR2 (vascular endothelial growth factor receptor 2) and VEGFR3 (vascular endothelial growth factor receptor 3); however, only VEGFR3 is required for developmental lymphangiogenesis.16 In LECs, VEGFR3 mediated signaling can be finely tuned by the formation of VEGFR2 heterodimers and the binding of coreceptors (neuropilin-1 and 2) and other proteins including syndecan 4 and $β1$ integrin.¹⁷ Expressed in both blood and lymphatic endothelium, VEGFR2 plays a pivotal role in angiogenesis but is less important for lymphangiogenesis, primarily acting to indirectly restrict lymphatic vessel development by sequestering VEGFC in blood endothelial cells.¹⁸

Mature LECs are a heterogenous population of cells, genetically distinct from blood endothelial cells, that can be divided into subpopulations based on their location and function within the lymphatic system. Structurally, all LECs express the following endothelial-specific junctional proteins: VE-cadherin (vascular endothelialcadherin), claudin-5, and PECAM-1 (platelet endothelial cell adhesion molecule-1), but the organization of these proteins within cell-cell junctions is variable based on the location of the LEC within the lymphatic network.19 Most capillary-forming LECs have button-like point connections between cells, while LECs that line

collecting vessels have a zipper-like continuous connections between neighboring cells. LEC junctional complexes have a high degree of plasticity and can switch between button and zipper confirmations during development and in pathophysiologic settings as discussed in more detail below.

Single-cell RNA sequencing studies have identified unique transcriptomic signatures for various LEC subpopulations, providing new insights into the specialized functions of each group.²⁰ LECs that line lymph nodes are distinct from LEC populations of peripheral (extranodal) lymphatics in that they express genes involved in attracting and transporting lymphocytes (*ITGA2B*, *MIP-3A*, *MADCAM-1*, and *GLYCAM-1*), screening for microorganisms (*PTX3*), attracting neutrophils to fight off lymph-borne pathogens (*CXCL5*, *CXCL1*, and C-type lectins), and immunological tolerance (*TNFRSF9*, *PDL1*, and *IFNGR*).^{20,21} Similarly, a subset of collecting vessel LECs gives rise to the intraluminal valves that prevent the backflow of lymph. These LECs are unique in that they are highly mechanosensitive and initiate valve formation gene programs in response to the mechanical forces generated by the flow of lymph.22 The molecular and biomechanical pathways involved in LEC development and function are described in Figure 1A and 1B.

Lymphatic Smooth Muscle

As capillaries coalesce into precollector and collecting vessels, they acquire 1 to 3 layers of lymphatic smooth muscle cells (LSMCs) intertwined with collagen and elastin fibers. Additionally, they become studded with the unidirectional, bileaflet valves that delineate the borders of lymphangions. Although aided by extrinsic forces such as skeletal and visceral smooth muscle contractions and respiratory movements, the independent and spontaneous contraction of LSMCs within individual lymphangions serves as the main driver of lymph transport against unfavorable pressure gradients.23,24

Spontaneous lymphangion contractions occur within the greater context of the myogenic tone of the vessel. Similar to the cascade of events needed to generate arterial tone, intracellular calcium accumulation occurs via 2 mechanisms (membrane depolarization–based activation of L-type voltage-dependent calcium channels and inositol 1,4,5-triphosphate–mediated release of sarcoplasmic calcium stores) and drives muscle contraction. Elevated levels of intracellular calcium bind to and activate calmodulin, which in turn activate MLCK (myosin light chain kinase) and subsequent phosphorylation of myosin light chains. This then facilitates myosin and actin interaction and crossbridge cycling, ultimately leading to vasoconstriction (Figure 1C).25,26 Lymphatic tone is regulated by many of the same biochemical vasodilators, vasoconstrictors, and biomechanical factors that modify arteriole tone.²⁷ This has fueled the notion that

lymphatic muscle is a form of vascular smooth muscle. However, lymphatic muscle is a hybrid muscle type that integrates smooth, skeletal, and cardiac muscle contractile proteins.^{28,29} Furthermore, there is regional variability in the expression of specific contractile proteins among lymphatic vessels that can be correlated to differences in pumping capacity. Muthuchamy et al showed that mesenteric vessels expressed higher levels of specific cardiac, visceral, and skeletal actin isoforms compared with thoracic ducts. In addition, mesenteric lymphatic vessels were enriched in SMB (smooth muscle myosin heavy chain B) and the fetal cardiac/skeletal slow twitch β-myosin heavy chain isoform—a non–smooth muscle isoform that confers relatively fast contractile properties. In functional studies, thoracic duct contractions were 30% to 50% less frequent and 60% to 75% less powerful than mesenteric vessel contractions.²⁹ These biochemical and functional variations are likely due to differences in the local environment of each vessel type: the thoracic duct is situated between the aorta and esophagus, each capable of exerting external forces onto the duct that would aid in the movement of lymph, while mesenteric vessels are not subject to the same degree of external compression by external forces and, therefore, must rely on powerful intrinsic contractions to pump lymph.

Lymphatic muscle also has unique electrophysical properties. LSMCs fire spontaneous action potentials that initiate waves of contraction that must be coordinated over the length of a lymphangion. Analogous to the cardiac cycle, LSMC contractions produce a systolic phase where lymph is actively pushed through an open valve into the adjacent lymphangion. This is followed by a diastolic phase where LSMC relaxation and closure of the previously open valve allow the lymphangion to refill. Modulation of membrane potential is integral to generating action potential spontaneous LSMC contractions. Several classes of ion channels including large-conductance calcium-activated potassium channels, inwardrectified potassium channels, delayed rectifier potassium channels, ATP-gated potassium channels, and calciumactivated chloride channels set the resting membrane potential of lymphatic muscle.³⁰⁻³² Similarly, symporters play a role in regulating tone and contractility as inhibition of the NKCC (Na+-K+-2CI− cotransporter) cotransporter with furosemide or bumetanide caused LSMC hyperpolarization, vasorelaxation, and a decrease in contraction frequency. 33,34

The rhythmic nature of lymphatic contractions suggests a pacemaking property. Small transient depolarizations (STDs) are thought to set the timing of lymphatic contractions, although the exact nature of the lymphatic pacemaker is still unresolved. STDs can be generated without neuronal innervation or the presence of an intact endothelium, which suggests that they originate within LSMCs. Another line of evidence that points to the

A

High interstitial pressure

Transmural

flow

Increased

shear stress

LEC

Immune cell

migration

Valve

formation

Increased

permability

Production of

vasoactive

factors

(NO

KrG

LSMC

cAMP/PKA

MI C_K

MLCP

Contraction

 \rightarrow o^oCa²⁺

Calmodulin

IP3

Figure 1. Molecular and biomechanical pathways involved in lymphatic vessel function.

A, High interstitial fluid pressure stretches lymphatic endothelial cells (LECs) opening up gaps between overlapping cells, which promotes influx of fluid, molecules, and cells into the capillary lumen. Transmural flow also alters the organization of LEC junctional proteins and upregulates the leukocyte homing chemokine, CCL21 (C-C motif chemokine ligand 21), and the adhesion molecules, ICAM (intercellular adhesion molecule-1), and E-selectin, thereby increasing leukocyte transmigration into lymphatic capillaries. Lymph accumulation and flow generates shear stress necessary for valve morphogenesis and maintenance, as well as production of LEC-derived vasoactive factors including NO and PGs (prostaglandins), which regulate lymphatic smooth muscle cell (LSMC) contractility. **B**, LECs integrate biomechanical and molecular signals to coordinate transcription factor (PROX1 [prospero homeobox 1], GATA2 [Gata binding protein 2], and FOXC2 [forkhead box C2]) and receptor tyrosine kinase (VEGFR3 [vascular endothelial growth factor receptor 3]) activity necessary for the regulation of lymphatic cell specification, survival, and maintenance. PROX1 is the master regulator of LEC identity and other LEC-related genes including VEGFR3. VEGFR3 can be activated by biomechanical forces or by binding of VEGFC (vascular endothelial growth factor C). In PROX1-positive LECs, VEGFR3 activation is necessary for embryonic and injury-induced lymphangiogenesis. Similarly, high levels of PROX1 and FOXC2 are required for endothelial cell alignment and elongation necessary for intraluminal valve formation. Calcium (Ca^{2+}) signaling has a major role in modulating LEC signaling cascades. Intracellular Ca²⁺ accumulation occurs via cation channel activation (PIEZO1 [Piezo1 type mechanosensitive ion channel component 1] and ORAI1 [ORAI calcium release-activated calcium modulator 1]) or by inositol 1,4,5-triphosphate (IP3)–mediated release of Ca²⁺ from the endoplasmic reticulum. Increased intracellular Ca²⁺ promotes sprouting via inhibition of Notch signaling, valve formation via calcineurin/NFAT (Nuclear factor of activated T cells) signaling, production of eNOS (endothelial-specific NO)-derived NO, and prostaglandin synthesis via PLA₂ (phospholipase A₂)-mediated release of arachidonic acid, which is converted to PG metabolites via COX (cyclooxygenase). LECs can also exert vasoactive effects on neighboring LSMCs via transferred membrane depolarization at myoendothelial junctions. C, Intracellular Ca²⁺ signaling also regulates the magnitude and frequency of LSMC contractions. Ca²⁺ can enter cells via activation of ion channels/transporters including NKCC1 (Na+-K+-2CI− cotransporter 1) that depolarize the membrane, activating voltage-dependent calcium channels (VDCCs). ET-1 (endothelin-1) binding to its receptor promotes IP3-mediated release of Ca²⁺ from the sarcoplasmic reticulum. In contrast, factors including sodium, NO, PGE₂, and histamine inhibit Ca²⁺ accumulation by activating cGMP/PKG (protein kinase G) and cAMP/PKA (protein kinase A) signaling, which activates hyperpolarizing potassium (K⁺) channels, inhibiting VDCCs. Accumulation of intracellular Ca²⁺ binds to calmodulin, which facilitates activation of MLCK (myosin light chain kinase) and phosphorylation of MLC (myosin light chain), allowing for myosin and actin interaction and subsequent muscle contraction. Illustration credit: Sceyence Studios.

pacemaking function of STDs comes via the observation that factors that increase the frequency of lymphatic contractions (eg, histamine, ET-1 [endothelin-1], and the thromboxane mimetic U46619) also increase STD activity. However, STDs can be abolished using the calciumactivated chloride channel blocker niflumic acid, without significantly altering the contraction frequency, making

it unlikely that STDs are the only regulator of lymphatic pacemaking.35 Contraction frequency can adapt to subtle changes in the microenvironment due to alterations in transmural hydraulic pressure, fluid sheer stress, lymph osmolarity, and surrounding tissue temperature.²³ LECs sense and respond to the shear stress generated by lymph flow by releasing vasoactive second messengers

that modulate the basal tone and pacemaker function of adjacent LSMCs.^{23,36,37} For instance, increased shear stress causes release of LEC-derived NO, which affects the basal tone and pacemaker function of LSMCs.²³

Of note, the local interstitial environment exerts different effects on lymphatic vessels based on the quantity and composition of fluid, cells, and solutes leaked from surrounding blood vessels. Numerous diseases are associated with deranged blood vessel endothelial barrier function, increased permeability, and extravasation into the microenvironment surrounding lymphatic vessels. This is especially common in diseases with an inflammatory component where the capillary leakage of bloodderived fluid, cells, and proteins then becomes a critical driver of lymphatic flow and lymphatic dysfunction.^{23,28} Such extravasation may be especially prominent in kidney diseases accompanied by hypoalbuminemia that lessens intravascular oncotic pressure promoting further transudation. In addition to these general principles of lymphatic vessel structure and function, the lymphatics of the gut have unique properties discussed below.

GUT LYMPHATIC HOMEOSTASIS

Lymphatic vasculature is heterogenous and plastic, acquiring specializations based on regional functional requirements and specific properties of the local microenvironment. The lymphatic vessels of the gut have distinctive structural features directly related to the specialized functions of this system, which include absorption of dietary lipids and fat-soluble vitamins, cholesterol transport, and gut immunosurveillance.

Intestinal Lymphatic Anatomy

The intestinal lymphatic network is uniquely patterned and partitions the organ into 2 distinct vascular plexuses: one that runs throughout the outer muscular layers of the gut and the other that is restricted to the inner mucosal and submucosal regions.³⁸⁻⁴⁰ Lymphatic capillaries in the gut, known as lacteals, are exclusively located in the mucosal region, extending into the center of each villus in the small intestine and connecting to underlying vessels in the submucosal space. These, along with the vessels of the muscular layer and vessels stemming from Peyer patches, coalesce and drain into collecting vessels located in the mesentery, which propel lymph toward the thoracic duct and back into the systemic circulation.^{38,39} Of note, this transportation route bypasses the liver, providing a path by which molecules carried in the lymph, including lipid-soluble drugs, can avoid first-pass metabolism and circulate systemically before degradation.40

In contrast to other lymphatic capillary beds, lacteals are surrounded by smooth muscle fibers that extend longitudinally from the submucosal region to the tip of each villus.41 Contraction of these fibers allows lacteals to

actively propel lymph—a process that can be modulated by neurohumoral factors released by the autonomic nervous system.⁴² Furthermore, contraction of visceral smooth muscle in the intestinal wall also aids in this process and may play a larger role in facilitating lymph movement as lipid concentration within the lymph increases. This was demonstrated using a rat model, in which ingesting a high-fat meal resulted in increased lymph viscosity and flow rate, while conversely decreasing the amplitude and frequency of spontaneous mesenteric vessel contractions. One explanation for these seemingly contradictory outputs is that the resulting increase in shear stress generated by increased lymph viscosity and flow triggered a signaling cascade (perhaps NO production) that contributed to the observed decrease in pumping dynamics. This scenario may provide broader mechanistic insight for the lymphatic dysfunction that is often associated with lipid-related pathologies including obesity, diabetes, and other metabolic disorders.43

Lipoprotein Transport and Immunosurveillance

The absorption of dietary fat takes place primarily in the proximal duodenum where lipids pass from the intestinal lumen into enterocytes where they are packaged into triglyceride-rich chylomicrons. Chylomicrons can also contain fat-soluble vitamins and drugs, microbiota components such as bacterial lipopolysaccharide, and food antigens, gut hormones, and chylomicron-specific lipoproteins. Dietary cholesterol is also absorbed by enterocytes and transported into the lamina propria either by incorporation into chylomicrons or by direct secretion in the form of HDL (high-density lipoprotein) bound with the lipoprotein apoAI (apolipoprotein AI). 38,44 Once formed, chylomicrons are released from the enterocyte's basolateral membrane into the lamina propria, where they can enter into lacteals by diffusion through LEC junctions or by intracellular transport across LECs via vesicles.45,46 Chylomicron transport from the lamina propria into lacteals is affected by the rate of lymph formation and flow. As fat absorption occurs, fluid from the intestinal lumen is also absorbed, causing glycosaminoglycans in the lamina propria to become increasingly hydrated. This enhances lymph formation and flow, which helps to facilitate chylomicron transport by either decreasing extracellular matrix resistance to particle movement or increasing the convective fluid movement of chylomicrons through LEC junctions.^{47,48}

Additionally, gut lymphatics play an important role in immunosurveillance and maintaining barrier function. The lumen of the gut serves as an interface between the outside world and the inside of the body. As such, it is continually exposed to vast amounts of antigens originating from pathogens, as well as innocuous substances including food and the commensal bacteria of the microbiome. This presents a complex challenge in which

antigen-presenting cells (namely macrophages and dendritic cells) must strike a balance between inducing an appropriate immune response and developing immune tolerance. Failure to do so can result in chronic inflammatory conditions such as Crohn disease and ulcerative colitis or food allergies and celiac disease. The importance of gut lymphatics to immunosurveillance is further highlighted by the fact that lacteals are required for maintaining villi structure and intestinal barrier function. When lymphatic vessels in the small intestine of transgenic mice were ablated by diphtheria toxin, there was complete collapse of villus architecture, which allowed for pathogen invasion and entry into the systemic circulation, resulting in subsequent septic shock and death.⁴⁹

KIDNEY DISEASE MODIFIES GUT LYMPHATICS

Although the gut is an increasingly recognized player in many diseases, recent studies have identified the intestinal lymphatic network as a key participant in the pathophysiology of diverse conditions ranging from cardiovascular and central nervous system disease to eczema.28,50–54 Disruption and impairment of intestinal lymphatics occurs not only in diseases primarily affecting the gut but also in systemic conditions and diseases in other organ systems. Thus, primary infectious or inflammatory bowel disease stimulate lymphangiogenesis, lymphangiectasia, and intralymphatic stasis that contribute to interstitial intestinal edema, infiltration of fat into the intestinal wall, and a sustained inflammatory response.55–57 In addition, intestinal lymphatic changes have also been documented in obesity, metabolic syndrome, diabetes, peritonitis, and cirrhosis.⁵⁸⁻⁶⁶ Table 1 summarizes the intestinal lymphatic perturbations in extraintestinal diseases. Importantly, kidney injury can now be added to the list of conditions that disrupt intestinal lymphatics.³

Proteinuric kidney injury causes significant expansion of the intestinal lymphatics. Kidney injury also increased the mesenteric lymph flow and modulated pumping dynamics in the mesenteric bed. Mesenteric lymphatic vessels isolated from animals with kidney injury had a significant increase in ejection fraction and a marked decrease in contraction amplitude and end-diastolic diameter compared with vessels from uninjured kidneys. Mesenteric lymph of proteinuric rats had reduced albumin output but increased output of cholesterol, triglycerides, and apoAI—the major structural protein in HDL. These changes in lymph composition could have affected shear stress triggering NO production similar to lymphatic disruption associated with lipid-related pathologies noted above.⁴³ Importantly, the mesenteric lymph of kidney-injured animals was enriched in lipid peroxides, specifically, the highly reactive isolevuglandin **LYMPHATIC SYSTEM IN ORGAN**

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(IsoLG), which cross-links apoAI (see below). The number of Th17 cells and cytokines including IL (interleukin)-6, IL-10, and IL-17 was increased in mesenteric lymph from the kidney-injured animals. S1P (sphingosine-1-phosphate), the immune cell trafficking factor and VEGFC, the prolymphangiogenesis factor also increased. Proteinuric kidney injury altered the transcriptome of intestinal LECs, affecting the expression of genes involved in lymphangiogenesis, vasodilation, and immune cell chemoattraction, for example, CCL21 (C-C motif chemokine ligand 21), eNOS (endothelialspecific NO), SPHK2 (sphingosine kinase 2), and SPNS2 (sphingolipid transporter 2). These are interesting observations in view of the recent findings that the composition of lymph, including lymphatic cells and lipoproteins, is modified during trafficking through the lymphatic network and that the modifications are organ specific.⁷⁸ Thus, kidney injury–induced changes in intestinal lymphatic growth, dynamics, and lymph composition may significantly impact the systemic and organ disorders accompanying kidney disease. Although the mechanisms driving lymphatic remodeling, dysfunction, or composition are unclear, kidney disease presents a fertile circumstance for understanding these pathways.

MECHANISMS BY WHICH KIDNEY DISEASE CAN MODIFY THE LYMPHATIC NETWORK

This section considers several kidney disease–linked complications that can disorder the lymphatic network including disruption of the intestinal epithelial barrier, dysbiosis and production of toxins, altered absorption and modification of lipids, low-grade inflammation, and sodium accumulation (Figure 2).

Disruption of the Intestinal Epithelial Barrier

Kidney disease causes significant proteomic changes to the intestinal epithelium, resulting in reduced expression of proteins required for tight junction formation (claudin-1 and 2) and maintenance of barrier integrity (HSP70 [heat shock protein 70]), which compromise barrier function.⁷⁹ Increasing urea levels occurring with progressive kidney disease directly deplete tight junction proteins including occludin, claudin-1, and zona occludens, further degrading the barrier structure and increasing local levels of oxidative stress.⁸⁰ Progressive disruption of the epithelium and underlying intestinal wall stimulates influx of immune cells, causing local inflammation and increased cytokine production, which results in further retraction and endocytosis of claudins and occludin.81–83 An obvious consequence of increased epithelial permeability is the increased risk of translocation of bacterial products and toxins into

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Table 1. Disease Conditions That Modify Intestinal Lymphatics

Akt indicates serine/threonine-protein kinase; apoAI, apolipoprotein AI; CCL21, C-C motif chemokine ligand 21; eNOS, endothelial-specific NO; ICAM-1, intercellular adhesion molecule-1; IL, interleukin; IsoLG, isolevuglandin; LEC, lymphatic endothelial cell; LTB4, leukotriene B4; oxLDL, oxidised LDL; PGE2, prostaglandin E2; PLA2, phospholipase A2; S1P, sphingosine-1-phosphate; S1PR1, sphingosine 1-phosphate receptor 1; SPHK2, sphingosine kinase 2; TNF-α, tumor necrosis factor alpha; VCAM1, vascular cell adhesion molecule 1; VE-cadherin, vascular endothelial-cadherin; and VEGF, vascular endothelial growth factor.

the systemic circulation. Perhaps equally wide-reaching consequences of barrier degradation are the deleterious effects that local enrichment of invading bacteria and toxins impose on the structure and function of intestinal lymphatic vessels.

Dysbiosis/Toxins

The gut microbiome has emerged as a modulable factor affecting health and disease. Many diseases disrupt the normal gut microbiome, causing dysbiosis that can dictate the disease phenotype. Patients with kidney disease have dysbiosis that includes lower colonization of *Bifidobacteriaceae* families, mainly *Bifidobacterium*, *Lactobacillaceae, Bacteroidaceae*, and *Prevotellaceae* genera and higher intestinal levels of *Enterobacteriaceae*, *Enterobacter*, *Klebsiella*, *Enterococci*, and *Clostridium perfringes*. 84–86 The imbalance of these bacterial species underlies chronic inflammation that prevails across the entire spectrum of kidney disease.^{87,88} Factors linked to adverse transformation of the gut microbiome in chronic kidney disease include (1) reduced

consumption of dietary fibers, (2) slow intestinal transit time that contributes to the imbalance between saccharolytic and proteolytic microbiota, (3) increased urea that supports bacterial families generating specific uremic toxins, (4) gut ischemia, and (5) edema.⁸⁹⁻⁹¹ Critically, the chronic kidney disease patient population is also routinely exposed to medications such as phosphate binders, iron-containing compounds, and especially, antibiotics that further disrupt normal intestinal structure and function, as well as disordering the normal microbiome.

The profound dysbiosis and toxins prevailing in kidney disease patients is relevant because intestinal microbiome is a critical regulator of development and maintenance of gut lymphatics.^{23,56,92,93} Lacteals appear within the intestinal villi at postnatal day 7 and continue to grow and remodel after weaning.⁹⁴ Germ-free mice lacking endogenous microbiota have decreased lacteal length, lower number of Prox1-positive LECs in their villi, and reduced VEGFR3.⁵⁶ The microbiome also regulates junctional plasticity. The proportion of button-like junctions decreased while zipper-like junctions increased between

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Figure 2. Kidney disease disrupts intestinal lymphatics.

Kidney disease damages intestinal architecture and function by causing disruption of epithelial integrity, dysbiosis, dyslipidemic metabolism, inflammatory cell infiltration, and sodium accumulation. The subsequent intestinal damage then impacts lymphatics by (1) increasing lymphatic flow and altering lymph composition; (2) activating lymphatic endothelial cells (LECs) that modulate tight junction organization (button-to-zipper ratio), augmenting production of cytokines/chemokines with vasoactive properties, and altering lymphatic smooth muscle cell (LSMC) contractility and pacemaker functions; (3) stimulating lymphatic growth but negatively affecting interstitial clearance and valve homeostasis. Illustration credit: Sceyence Studios.

LECs of antibiotic-treated mice versus vehicle-treated animals.94 The magnitude of the effect varied by intestinal segment and paralleled the presence of microbiota, being most pronounced in the jejunum and ileum and least in the duodenum, which is the less colonized segment of the intestine.⁵⁶

The microbiome also maintains lacteal length, integrity, and functionality in adulthood. As in immature animals, depletion of microbiota by antibiotic treatment in adults reduced the lacteal length in jejunum and ileum, with little change in the duodenum.⁵⁶ The remodeling was driven by continuous activation of Notch ligand delta-like 4 and adrenomedullin-calcitonin receptor-like receptor signaling.38,95 Lacteal regression was not due to decreased survival of LECs nor a direct toxic effect of the antibiotics on LECs and instead reflected reduced expression of VEGFC, thereby reiterating its critical role in maintaining lacteal integrity.56,96 There are 2 primary sources of VEGFC around lacteals, the longitudinal smooth muscle cells surrounding lacteals and villus macrophages. Antibiotic treatment did not alter abundance or distribution of the smooth muscle cells.⁵⁶ However, a marked reduction in macrophage-expressed VEGFC was observed in antibiotic-treated mice, supporting a critical role of villus macrophages in VEGFC production and maintenance of lacteal structure. It follows then that alterations to the microbiota would also have functional consequences on lacteals. Indeed, depletion of gut microbiota resulted in reduced triglyceride, chylomicron, and free fatty acid

transport attributed to structural defects assessed by intravital imaging.56 The detrimental effects of depleting the intestinal microbiome on lacteal phenotype were reversed by conventionalizing the germ-free animals into a normal environment.⁵⁶ It is notable that only the intestinal lymphatics were affected by the antibiotic depletion of intestinal microbiota as lymphatic vessels in ear skin, trachea, diaphragm, and inguinal lymph nodes showed no discernible effects.56 These findings highlight the unique characteristics of intestinal lymphatics. Unlike LECs in lymphatics of other organs, they have filopodia and positive staining for Ki67 proliferation markers, demonstrating their measurable proliferative capacity and ability to remodel under steady-state conditions.23,97,98

While strong evidence supports microbiome regulation of intestinal lymphatics, there is also evidence for the reverse relationship, whereby lymphatics modulate the gut microbiome.56 In a murine colitis model, supplementation of VEGFC promoted lymphangiogenesis, increased intestinal drainage, and significantly increased the abundance of *Bacterioidate* and decreased the abundance of Firmicutes at phylum level in fecal samples.⁹⁹ In a primate model of Crohn disease, obstruction of lymphatic outflow dramatically altered intestinal microbial subgroups, including greater abundance of *Prevotellaceae* and *Bacteroidetes-Prevotella-Porphyromonas*. 100 These results illustrate a cycle whereby metabolic changes associated with disease, including kidney disease, disrupt the intestinal microbiome, which impairs the lymphatic vascular network, which, in

turn, intensifies dysbiosis and intestinal damage, thus furthering disease.

Dyslipidemia

Lipids have a variety of functions beyond their traditional roles in building cell membranes and energy storage. They also participate in physiologic and pathologic signaling cascades that regulate cellular metabolism, cell fate determination, proliferation, and immune response, and their activity is tied to specific modifications of their basic structure. Kidney disease causes abnormalities in the lipid profile. The characteristics of these abnormalities vary depending on the degree of kidney impairment, the underlying etiology, and whether proteinuria, especially nephrotic syndrome, is present.¹⁰¹⁻¹⁰⁷ Aside from altering lipid levels, a low-grade chronic proinflammatory and high oxidative state prevails across the entire spectrum of kidney disease. This creates a milieu that promotes formation of oxidatively and enzymatically modified lipoproteins and lipid metabolites (eg prostaglandins, lysophosphatidic acid) that directly modulate LEC phenotype and lymphatic growth, barrier function, and contractility.43,108–110

Although interaction between lipids and lymphatics is established, recent studies have uncovered previously unappreciated lipid metabolism in LECs. As in other cells, LECs use mitochondrial fatty acid oxidation for their energy needs. Fatty acid oxidation is also key in promoting LEC proliferation by providing acetyl-CoA that sustains the Krebs cycle and deoxyribonucleotide synthesis needed for cellular proliferation and division.^{111,112} Inhibition of fatty acid oxidation in LECs reduces expression of the rate-limiting enzyme, CTP1A (carnitine palmitoyltransferase 1A), which significantly decreases endothelial cell proliferation and lymphatic differentiation via PROX1-dependent regulation of VEGFR3 expression.^{111,112} These findings suggest a therapeutic potential of lowering fatty acid oxidation to inhibit pathophysiologic lymphangiogenesis (ie, cancer). Conversely, supplementation of fatty acids (eg, acetate) rescues the acetyl-CoA, VEGFR3 expression, and lymphatic sprouting and restores lymphangiogenesis after corneal injury. These findings suggest fatty acids or their metabolites can be used to modulate LECs and lymphatic growth in pathophysiologic settings.

The pathophysiological implication of LEC lipid metabolism was further illuminated by the demonstration that LECs contain lipid droplets. Lipid droplets have traditionally been considered static energy storage deposits that increase with impaired cellular homeostasis.^{113,114} Increasingly, however, lipid droplets are recognized as dynamic and inducible platforms that not only sequester fatty acids and prevent lipotoxicity but also regulate energy metabolism, membrane trafficking, immune

responses, and cytokine production.^{115,116} In view of the specialized function of lymphatic vessels in lipid trafficking, finding lipid droplets in LECs uncovers the cells' potential to regulate fatty acid storage and degradation and, therefore, lymphatic functions. Strong support of this concept comes from a recent study showing that genetic deficiency in lipid droplet degradation by autophagy causes significant LEC damage.117 Autophagy blockade caused accumulation of lipid droplets linked to defective trafficking of lipid droplets to mitochondria. This in turn reduced mitochondrial ATP production, fatty acid oxidation that interrupted the mitochondrial-PROX1–driven transcriptional circuit blunting the normal response to VEGFC. The net result was impaired lymphangiogenesis documented in cultured cells and in an in vivo corneal injury model. It is interesting that ketogenic diets that provide short-chain fatty acids increase lymphangiogenesis and improve lymphatic function after myocardial infarction and in a lymphedema mouse model. $117-119$ Dietary fatty acids also affected accumulation of lipid droplets and cellular injury in proximal tubules of mice with diabetic nephropathy.¹²⁰ Whether the diabetic kidney injury affected the formation of lipid droplets in LECs and whether dietary fatty acids can modulate LEC lipid droplets is currently unknown. Nonetheless, there is strong support for the concept that lipid droplet degradation is an important source of fatty acid oxidation driving lymphangiogenesis and that dietary lipids can maintain LEC responsiveness to VEGFC in physiological conditions and following injury-stimulated lymphangiogenesis.¹¹⁷

Modified lipids have specific roles in regulating lymphatics. This is relevant because kidney disease–associated inflammation promotes lipid oxidation, including oxidation of LDL (low-density lipoprotein) cholesterol, which profoundly affects lymphatic growth and function. Comparison of normal unmodified LDL with oxLDL (oxidized LDL) illustrates the differences: normal LDL stimulated lymphangiogenesis while oxLDL inhibited lymphangiogenesis.109 Oxidized, but not normal, LDL blocked cell cycle progression in cultured LECs, reduced their expression of Akt (serine/threonine-protein kinase) and eNOS, increased p27 (an inhibitor of the cell cycle) expression, and induced intracellular oxidative stress.^{54,109} Knockdown of the lipid transporter, CD36 (cluster of differentiation 36), in LECs prevented oxLDL-induced suppression of lymphangiogenesis via AKT/eNOS pathway and cell cycle. These findings suggest blockade of LEC CD36 signaling may promote lymphangiogenesis and, therefore, clearance of cells and potentially harmful molecules.109

A critical role for oxLDL was recently shown in humans and mice with fatty liver disease. The study revealed hepatic fatty infiltration affects lymphangiogenesis and impairs the ability of LECs to maintain a lymphatic-specific transcriptional profile.110 These changes culminated in diminished lymphatic draining of the liver. The findings complement observations that fatty acid oxidation in

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LECs is critical in promoting lymphatic growth and lymphatic-specific chromatin modifications.112 OxLDL also altered the permeability of the lymphatic vasculature. Single-cell analysis of liver LECs from mouse model of fatty liver disease and cultured LECs exposed to oxLDL skewed the ratio of *Vegfr2/Vegfr3* and the cells became significantly less permeable. Importantly, treatment with recombinant VEGFC (cys156ser) targeting VEGFR3 resulted in increased hepatic lymphatic vessel density but also improved liver inflammation and progressive hepatic fibrosis. Together, these observations show that oxLDL directly impairs LEC phenotype, growth, permeability, and interstitial clearance by the lymphatic vessels, which may present a therapeutic target to lessen disease.

Lipoprotein oxidation generates a variety of lipid peroxides including the most reactive dicarbonyl IsoLGs, which covalently modify proteins, nucleic acids, and other lipids, causing dysfunction in cells and tissues. Among lipoproteins, HDL is especially susceptible to IsoLG adduction. Elevation in IsoLG adducts has been linked to the oxidative damage in atherosclerosis, cardiac arrhythmia, hypertension, kidney damage associated with systemic lypus erythematosus, Alzheimer disease, macular degeneration, and cancer.¹²¹⁻¹²⁵ Different cells, including antigen-presenting immune cells, platelets, cardiac myocytes, and epithelial cells in lung and liver, can generate IsoLG.¹²¹⁻¹²⁸ IsoLG adducts have recently been shown to be increased in gastrointestinal epithelial cells of humans and animal models with gastritis and colitis associated with gastrointestinal carcinogenesis.¹²⁵ We show that kidney injury also stimulates intestinal epithelial production of IsoLG and that the IsoLG colocalizes with apoAI in lymphatic lacteals. Since about one-third of apoAI is synthesized by the ileum, $129,130$ it is possible that enterically produced IsoLG adducts local apoAI and modulates the mesenteric lymphatic bed. Indeed, IsoLGapoAI directly affected lymphatic growth and contractility. Moreover, IsoLG-apoAI increased NOS3 (nitric oxide synthase 3) in cultured LECs, and IsoLG-apoAI directly blunted vasoactivity and increased contraction frequency versus unmodified apoAI. Critically, treatment to scavenge IsoLGs significantly reduced intestinal lymphangiogenesis, albuminuria, and interstitial fibrosis supporting the concept that intestinal lymphatic growth associated with kidney injury is linked to IsoLG.^{3,131} Thus, the mesenteric lymphatic network appears to serve as both target and perpetrator of IsoLG-HDL's effects by augmenting lymphangiogenesis, lymphatic vessel contractions, LEC activation, and increased lymph flow.

Injury releases arachidonic acid from cell membrane phospholipids, providing precursors for bioactive mediators including prostaglandins, thromboxanes, and leukotrienes that have important roles in inflammation, cellular proliferation, fibrosis, and regulation of vascular tone, including lymphatics (Table 2). PGE _s stimulates lymphangiogenesis and increases lymph flow by inducing

VEGFC/D transcription.¹³³ Similarly, thromboxane 2, acting via the TP (thromboxane prostenoid receptor) on macrophages and T cells and not LECs, promotes lymphangiogenesis by inducing VEGFC/D.134 Acting through the TP receptor, thromboxane 2 also increases lymphatic flow rate that reflects expansion of lymphatic density, greater reabsorptive capacity, and stronger pumping by collecting vessels.¹³⁴ Leukotriene modulation of lymphatics depends on their levels whereby lower leukotriene levels prompt lymphangiogenesis while higher concentrations inhibit VEGFR3, lymphatic growth, and Notch signaling, all critical to lymphatic development and maintenance.137,138,146,147 Studies with the nonsteroidal anti-inflammatory drugs ketoprofen and ibuprofen have added to our understanding of the relative effects of prostaglandins and leukotrienes on lymphatics.¹⁴⁸ Ketoprofen inhibits cyclooxygenase enzymes, which generates prostaglandins, and 5-LO (5-lipoxygenase), which generates leukotrienes. Ketoprofen enhanced postsurgical lymphedema, indicating that cyclooxygenase-derived prostaglandins or 5-LO–derived leukotrienes have a role in lymphedema formation. 137 Ibuprofen—a cyclooxygenase inhibitor—exacerbated edema, indicating beneficial effects of prostaglandins in lymphedema.¹³⁷ Indeed, pharmacological inhibitors and genetic targeting of the 5-LO pathway lessened edema and improved lymphatic vascular function.^{137,149,150}

Lysophosphatidic acid, which encompass bioactive phosphoglycerides synthesized from membrane phospholipids, is involved in embryogenic development of lymphatic vessels.^{139,140,151} Lysophosphatidic acid also activates expression of VEGFC that potentiates lymphangiogenesis following injury in adults though NF-κB (nuclear factor kappa B)–mediated induction of IL-8.¹³⁹ Lysophosphatidic acid in human and animal atheroma acts as an LEC chemoattractant to induce lymphangiogenesis.¹⁵²⁻¹⁵⁴

S1P is synthesized from ceramide, phosphorylated by SPHK1 and SPHK2, and carried primarily by HDL. S1P regulates endothelial cell proliferation and migration.141,142,155,156 LYVE-1-driven depletion of SPHK1/ SPHK2 reduces S1P in LECs, leading to abnormal lymphatic morphology and reduced lymphocyte egress from peripheral tissues. This suggests LEC-derived S1P has a key role in lymphatic patterning and lymphocyte trafficking.^{47,143} Indeed, our studies showed that mesenteric lymph of kidney-injured animals contained increased levels of S1P and that intestinal LECs had more SPHK2 mRNA, indicating S1P signaling may be an attractive therapeutic target to improve lymphocyte trafficking.3

Inflammation

Persistent low-grade inflammation prevails across all stages of kidney damage and is evident not only in

Table 2. Lipid-Related Modulators of Lymphatic Vessels

CD36 indicates cluster of differentiation 36; eNOS, endothelial-specific NO; HDL, high-density lipoprotein; IL, interleukin; IsoLG, isolevuglandin; LDL, lowdensity lipoprotein; LDLR, low-density lipoprotein receptor; LEC, lymphatic endothelial cell; MCP-1, monocyte chemoattractant protein-1; PCSK9, proprotein convertase subtilisin kexin type 9; PGE2, prostaglandin E2; PROX1, prospero homeobox 1; ROS, reactive oxygen species; S1P, sphingosine-1-phosphate; SPNS2, sphingolipid transporter 2; Th17, T helper 17 cells; VE-cadherin, vascular endothelial-cadherin; and VEGF, vascular endothelial growth factor.

plasma and kidney parenchyma but also in extrarenal tissues. Immune cells and their cytokines are powerful mediators of lymphatic vessel growth, permeability, and pumping. Proteinuric kidney injury led to a 3-fold increase in the number of CD68-positive cells in the ileum compared with normal controls.³ Macrophages in particular are an important source of VEGFC,¹⁵⁷ and macrophage depletion reduced lymphangiogenesis in acute colitis,¹⁵⁸ hypertension,¹⁵⁹ and proteinuric kidney injury.¹⁶⁰ Dendritic cells and neutrophils have also emerged as sources of lymphangiogenic signals.161 In addition to VEGFC, IL-17 and IL-8 have also been shown to induce lymphangiogenesis.162,163 IFN-γ (interferon gamma) produced by activated T cells decreased lymphatic vessel formation.164,165 Inflammatory mediators impact other lymphatic functions. IL-6, TNFα (tumor necrosis factor alpha), and IFN-γ increase lymphatic permeability in association

with reduced expression of VE-cadherin.¹⁶⁶ Inflammatory cytokines also modulate lymphatic pumping activity. Prostaglandins, IL-1β, IL-6, and TNF-α reduced lymphatic pumping frequency.¹⁶⁷ Blocking TNF- α increased lymphatic contractions and reduced joint inflammation.¹⁶⁸ Substance P—a neuropeptide also secreted by inflammatory cells—increases the pumping frequency of rat mesenteric vessels.169

Sodium

Sodium accumulation accompanies kidney disease and other edema-forming diseases such as congestive heart failure, cirrhosis, and hypertension.¹⁷⁰ Over the last 2 decades, studies have suggested that sodium accumulation within the interstitium is linked to lymphangiogenesis. The sodium-lymphatic relationship has been most

extensively studied in the skin of hypertensive animals and humans and involves transcription factor TonEBP (tonicity-responsive enhancer protein)-induced macrophage secretion of VEGFC secreted by infiltrating macrophages.159,171,172 Depletion of mononuclear phagocytes or blockade of VEGFC during high salt intake increased density and hyperplasia of lymphatic capillaries in the skin in association with hypertension, although recent studies from the same groups have questioned the singular effect of lymphangiogenesis on blood pressure.^{159,171,173} It is, therefore, notable that aside from lymphatic growth, sodium can also modulate lymphatic dynamics, which may have a role in hypertension and other diseases.^{5,174,175} Mizuno et al¹⁷⁵ showed high-sodium diet suppressed the amplitude, ejection fraction, and stroke volume that markedly reduced pumping activity in afferent lymphatics while efferent lymphatics were resistant to the effects high-salt diet and preserved pumping reflected increased contraction frequency. High-salt diet or DOCA treatment increased skin and muscle lymphatic vessel contraction frequency while reducing contraction amplitude.¹⁷⁴ Similarly, we showed that direct exposure of lymphatic vessels to a high-sodium environment increases the frequency of contraction of collecting lymphatic vessels and reduced the contraction amplitude and, to a lesser extent, the ejection fraction.⁵ Currently, there are no studies specifically assessing the effects of sodium on mesenteric lymphatics; however, it is known that this lymphatic bed adapts to physiologic and pathophysiologic stimuli. For example, oxidative stress, fructose feeding, aging, and intestinal inflammation impair intrinsic contractility of the mesenteric collecting lymphatic vessels.^{74,176,177} These are important observations because, unlike blood vessels, which depend on the heart to pump blood, lymph flow is propelled by forces in the surrounding tissues and intrinsic contractions of the lymphatic vessels, which are exquisitely sensitive to the microenvironment.^{37,178} Thus, local sodium concentration may be a significant regulator of lymphatic dynamics.

Sodium can also function as a signaling molecule. We show the Na-K-2Cl cotransporter NKCC1, but not NKCC2, is expressed in lymphatic vessels.³⁴ Thus, while NKCC2 is best known for its actions on tubular epithelial cells responsible for maintenance of sodium homeostasis, we find NKCC1 modulates the tone in collecting lymphatic vessels.34 Of potential clinical relevance are data showing that the NKCC1 inhibitor, furosemide, dilates lymphatic vessels and decreases contraction frequency, amplitude, and ejection fraction. High-sodium environment also decreases lymphatic vessel contraction amplitude and ejection fraction, reduces phosphorated NKCC1, phosphorylated SPAK (SPS1-related proline/ alanine-rich kinase), an upstream kinase, and phosphorylated eNOS, a downstream vasoactive factor, and blunts lymphatic response in injured kidneys.⁵ These results suggest sodium accumulation accompanying kidney disease can contribute to heretofore unrecognized impairment in lymphatic function that may accompany sodium-avid states and impaired lymphatic function. These findings also suggest that resistance to the diuretic effects of furosemide sometimes observed in sodium-avid conditions may reflect dampening of the contractile and pacemaker functions of lymphatic vessels.

CLINICAL IMPLICATIONS AND THERAPEUTIC INTERVENTIONS

Kidney disease is associated with many complications including hypertension, CVD, bone disease, neurologic and cognitive dysfunctions, and progression of kidney scarring, even when the original inciting disease has abated. Each of these clinical conditions has been linked to changes in the gut and led to new nomenclature to encompass the concept of interorgan cross talk via terms such as gut-hypertension axis, gut-CVD axis, gut-bone axis, gut-cognition axis, and gut-kidney axis. Mediators in these interorgan communication networks include intestinally originating metabolites, immune cells activated within the intestinal wall, and lipid particles absorbed from the diet or generated within the intestines and then modified within the local environment. Until now, blood vessels and nerves were thought to be the primary pathways by which metabolites and toxins affect distant organs. It now appears that intestinal lymphatics constitutes an additional pathway in the gut-organ axis (Figure 3).

Currently, therapeutic focus for kidney disease is the pathophysiologic processes affecting the kidneys with additional therapies aimed at treating specific systemic complications such as anemia and bone disease. Here, we focus on the lymphatic network and the possibility that future therapies may target the absorptive capacity, contractile dynamics, and clearance competence of the intestinal lymphatics, which could lessen the adverse consequences associated with kidney disease. Heretofore, no drug has been developed for clinical use that specifically targets lymphatics. However, the increasing number of Food and Drug Administration–approved drugs are being recognized to have effects on lymphatic vessel number or function (Table 3). Pharmacotherapeutic agents that modify the lymphatic network can be broadly categorized by their proposed mechanism or functional target, for example, VEGFC activators/ inhibitors to modulate lymphangiogenesis, ROCK (Rho kinase) inhibitors to modulate lymphatic integrity, ATPgated potassium antagonists to modulate lymphatic contractility, anti-inflammatory/antioxidant agents to modify the lymphatic environment.

VEGFC/VEGFR3 is a critical axis in lymphangiogenesis. Much of the pharmacologic efforts have focused on stimulation or inhibition of lymphangiogenesis in disease models such as inflammatory bowel disease,

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Figure 3. Intestinal lymphatics mediates the adverse consequences of kidney disease in other organ systems. Kidney disease causes intestinal damage and lymphatic dysfunction due to epithelial disruption, alterations of the microbiome and lipid profile, inflammation, and sodium accumulation. This results in the production of disease-associated molecules (bioactive metabolites, modified lipoproteins, activated immune cells, and cytokines) that can then be dispersed throughout the body via the lymphatic vasculature where they have secondary pathogenic effects. This model represents a novel mechanism by which lymphatics mediates the production and distribution of kidney disease–derived factors that subsequently contribute to disease progression in other organ systems. Illustration credit: Sceyence Studios.

atherosclerotic/congestive cardiac failure, joint and skin inflammation, and cancer. Lymphactin—an adeno-associated virus vector–based overexpression of VEGFC gene therapy—is currently in a phase II clinical trial in patients with breast cancer-associated secondary lymphedema.¹⁹² Lymphangiogenesis can also be induced by retinoic acid via the FGFR (fibroblast growth factor receptor) pathway. The 9-cis retinoic acid—a Food and Drug Administration– approved drug for treating Kaposi sarcoma and chronic eczema—stimulates LEC migration and differentiation in vitro and reduces lymphedema via increased lymphangiogenesis in preclinical animal models.181,193 Since VEGFC also binds VEGFR2, VEGFC modulators can also affect blood vessels.194 Future therapeutics will need to consider ways that limit the potentially undesirable vascular side effects such as angiogenesis or capillary leakage.

Lymphatic integrity is another contributor and target of lymphatic dysfunction. For example, the anti-inflammatory agent dexamethasone, acting through glucocorticoid receptors, inhibits VEGFC–induced lymphangiogenesis and also promotes zipper to button junction formation, shown to resolve inflammation.^{182,183,195} FOXC1 (Forkhead box C1) or FOXC2 (Forkhead box C2) induces

hyperactivation of contractile stress fibers in LECs, which is rescued by the ROCK inhibitor Y27632.196 ROCK inhibition also enhances lacteal zipper junction formation and improves LEC barrier integrity in mesenteric collecting vessels, which has been proposed as a therapeutic approach to treat obesity/metabolic dysfunction.⁶⁰ In type 2 diabetes, reduced NO bioavailability and consequent PDE3 (phosphodiesterase 3) activation cause a lymphatic permeability defect.⁶³ Oral administration of cilostazol-a selective PDE3 inhibitor approved by the Food and Drug Administration for treating intermittent claudication reversed genetically and surgically induced lymphedema in mice, which was attributed to both the lymphangiogenic and lymphatic vascular integrity effects.¹⁸⁴ These examples illustrate that therapeutic targets can affect multiple lymphatic functions and different lymphatic segments, for example, capillaries and collecting vessels that may have complementary or opposing effects on the lymphatic network.

Ion channels and transporters are significant mediators of the membrane potential in lymphatic vessels. This is important because many currently used medications belong to this class of drugs. A recent review of >200

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COX-2 indicates cyclooxygenase 2; NKCC1, Na+-K+-2CI− cotransporter 1; PDE3, phosphodiesterase 3; ROCK, Rho kinase; VEGF, vascular endothelial growth factor; and VEGFR, vascular endothelial growth factor receptor.

drugs with potential lymphatic vessel effects found that the most frequently used drug with inhibitory effects on lymphatic pumping was the calcium channel blocker.¹⁹⁷ Notably, most of the identified drugs dampen lymphatic contractile functions.197 These observations highlight the possibility that adverse consequences such as refractory edema with diuretic use may be linked to effects on lymphatics. We found that NKCC1 inhibition by the diuretic furosemide decreased both lymphatic vessel tone and contractility.³⁴ Furosemide has been reported to impair human thoracic duct contractility in vitro.¹⁸⁶ ATP-gated potassium openers, such as pinacidil and diazoxide used to treat hypertension, induce severe peripheral edema, which is contributed by the hyperpolarization in the lymphatic muscle cell and reduced lymphatic contractility and lymph flow.30,198 Conversely, glibenclamide (Food and Drug Administration–approved ATP-gated potassium antagonist) rescued lymphatic contractile dysfunction in a mouse model with gain-of-function mutations in the Kir6.1 gene, which mimics the lymphoedema observed in patients with Cantú syndrome.187 Glibenclamide also partially restored the mesenteric lymphatic contractility in the metabolic syndrome model.⁶¹ Cumulatively, these studies illustrate the previously underrecognized effects of current drugs on lymphatics. The observations also highlight the need for therapies that specifically target the lymphatic function.

Since local inflammation is a key modulator of lymphatics, anti-inflammatory interventions have been used to affect lymphatic functions. The antibiotic doxycycline not only impeded monocyte recruitment and polarization of alternatively activated macrophage but also inhibited lymphangiogenesis via VEGFC/VEGFR3 signaling.^{188,189} In a clinical study of 162 patients with filarial lymphedema, doxycycline treatment also improved lymphedema compared with amoxicillin or placebo.¹⁹⁰ The benefit of doxycycline treatment was independent of ongoing infection and linked to activation of the VEGFC/ VEGFR3 pathway. Inhibition of proinflammatory metabolites of arachidonic acid such as leukotrienes affects lymphatics in animals and humans. Bestatin—a selective leukotriene B4 antagonist—increased lymphatic function and restored lymphatic architecture in the murine tail model of lymphedema.^{137,191} A phase II randomized and placebo-controlled study in patients with lower extremity lymphedema is underway, and results are pending.^{137,191} Because inflammatory modulation of lymphatic function is local, there is increasing focus on delivery of antiinflammatory therapeutics into the relevant tissue.

Recognition of the importance of the lymphatic system on disease progression has led to new strategies to access lymphatics and design biomaterials that specifically enter lymphatics. Currently, this approach includes the route of administration and drug formulation. Gastrointestinal, subcutaneous, or pulmonary administration appear to provide better access to the lymphatic network than intravenous administration of a therapeutic. Here, we focus on gastrointestinal lymphatics. Targeting the gastrointestinal lymphatics increases target specificity by bypassing first-pass metabolism in the liver, which reduces systemic concentrations of a drug. For example, compared with intravenous administration, drug-containing nanoparticles delivered to the intestines were shown to have a $>$ 20-fold increase in drug bioavailability, a 30-fold increase in the elimination half-life, and less accumulation in the heart, lungs, spleen, and kidneys.^{199,200} This approach also reduces off-target effects, which is a serious problem in patients with kidney disease. Specific drug formulations can also target lymphatic entry. For example, while small molecules (<5 nm in diameter) are easily taken up by blood capillaries, middle-size molecular species (5–100 nm) are shuttled to lymphatics.201,202 Lymphatic uptake can be improved by making

lipophilic nanomaterial formulations of small-molecule drugs.203 A nucleoside modified VEGFC mRNA delivered in lipid nanoparticles significantly increased lymphangiogenesis without blood vessel proliferation in an experimentally induced lymphedema.204 Another approach is to use drug-triglyceride formulations that undergo hydrolysis into a drug-monoglyceride, which is then assembled into a lipoprotein that enters the mesenteric lymph. The approach was used for a celecoxib prodrug that linked celecoxib to a glyceride backbone and enhanced drug transport into mesenteric lymph and concentrated celecoxib at a site of lymph leakage.⁵⁸ Compared with nontargeted celecoxib, lymph-targeted celecoxib prodrug was more effective in reversing mesenteric lymphatic vessel branching and lymph leakage, reduced visceral obesity and inflammation, and restored glycemic control in obese mice. Nanomaterial-based strategy can be more narrowly directed to target the lymphatic endothelium when the carrier is conjugated with agents targeting expressed receptors, such as LYVE-1.205

PERSPECTIVES

New evidence indicates that kidney injury causes intestinal lymphangiogenesis, disrupted lymphatic contractility, altered composition of lymph, and activated LECs lining these vessels. These observations underscore the distinct qualities of the lymphatic pathway including the capacity to take up and convey large bioactive metabolites, modified lipoproteins, and activated immune cells generated within the intestinal lumen and wall and disperse them throughout the body, contributing to the adverse consequences associated with chronic kidney disease. Therefore, we propose a novel paradigm of disease progression in which a cycle of detrimental organ cross talk, fueled by kidney injury–induced modulation of intestinal lymphatics, contributes to systemic organ disease progression, which then perpetuates intestinal anomalies (Figure 3). An additional implication of the intestinal-lymphatic axis is its emergence as a novel avenue for drug delivery in the treatment of the serious clinical complications associated with kidney disease. Specifically, the intestinal lymphatic network may provide new drug targets and added advantage by bypassing liver-mediated degradation before delivery. Thus, in addition to specific treatments for kidney disease, future therapies may also target the absorptive capacity, contractile dynamics, and clearance competence of the intestinal lymphatics that could lessen the associated adverse consequences.

ARTICLE INFORMATION

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Disclosures

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