

Tubular Sodium Transport

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Na⁺, the primary extracellular cation, is of critical importance to the maintenance of extracellular fluid volume. The kidneys play the dominant role in regulating Na⁺ excretion. Each day, the glomeruli filter roughly 25,000 mEq of Na⁺. From this quantity, almost 10 times the total exchangeable Na⁺ in the body, the kidneys typically absorb over 99%. A remarkable feature of the Na⁺ absorptive process is the precision with which it is regulated. An individual consuming a typical diet containing 6 g of Na⁺ will excrete 260 mEq of Na⁺ per day. The same individual placed on a 2-g Na⁺-restricted diet will promptly reduce Na⁺ excretion to 87 mEq per day. Although the fraction of filtered Na⁺ absorbed by the kidney changes from 99.0% on a standard diet to 99.6% on a Na⁺-restricted diet, this small change represents the addition or removal of over 1 L per day to extracellular fluid volume. Thus, the kidneys absorb large amounts of filtered Na⁺ with remarkably precise control.

The exquisitely sensitive regulation of Na⁺ absorption by the kidneys relies on sequential actions of the various nephron segments, each with highly specialized transport capabilities. Figure 5.1 provides an overview of Na⁺ transport along the nephron. In general, the absolute rates of Na⁺ reabsorption are greatest in the proximal tubule and fall as the tubular fluid proceeds from proximal to distal segments. Conversely, the ability to transport Na⁺ against steep tubular fluid to blood gradients and its physiologic control increase along the nephron. For example, the proximal tubule reabsorbs the bulk (60% to 70%) of the filtered Na⁺ load, but as will be detailed later, does so against at most small electrochemical gradients. Moreover, the ability to alter Na⁺ transport in the proximal tubule, in relative terms, is rather limited, usually varying by less than 25%. The collecting duct, in contrast, reabsorbs only a minor fraction (~2% to 4%) of the filtered Na⁺ load. However, the collecting duct can transport Na⁺ against a large electrochemical gradient to produce urine, which is almost Na⁺ free (<10 mEq/L). In addition, the rate of Na⁺ transport in the collecting duct can vary over a wide range (tenfold) in response to physiologic stimuli. The different nephron segments thus permit both

high rates of Na⁺ transport (proximal segments) and highly regulated Na⁺ transport (distal segments).

Substantial progress has been made recently in identifying the proteins that mediate Na⁺ transport in each nephron segment and in defining their interactions and regulation within each segment. Many Na⁺ transport proteins have been linked to specific genetic disorders (Table 5.1). Given the primacy of renal Na⁺ transport to the control of extracellular fluid volume, it is not surprising that the majority of these genetic disorders are characterized by either hypotension or hypertension. An updated and curated database of these genetic disorders is maintained at the Online Mendelian Inheritance in Man website (www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=OMIM).

This chapter considers the transepithelial transport of Na⁺ by the various nephron segments. The discussion of each nephron segment begins with a description of the general features of Na⁺ transport in that segment along with pertinent structure–function relations. The mechanism of Na⁺ transport is then considered on a cellular or subcellular level, with emphasis on recent electrophysiologic, biochemical, and molecular findings. Finally, each section includes a consideration of the factors that regulate Na⁺ transport in the individual segments.

PRINCIPLES OF MEMBRANE TRANSPORT

This section describes physical principles that underlie the movement of ions across individual membranes and epithelia. However, it is not intended to be an extended treatment of the thermodynamic aspects of membrane transport processes.

Diffusion Processes

Solute transport across membranes may occur by diffusion or convection, or by a mediated process. Diffusion is the random Brownian motion of a molecule with respect to adjacent molecules and occurs as the consequence of thermal energy.¹ Because the diffusional movement of an

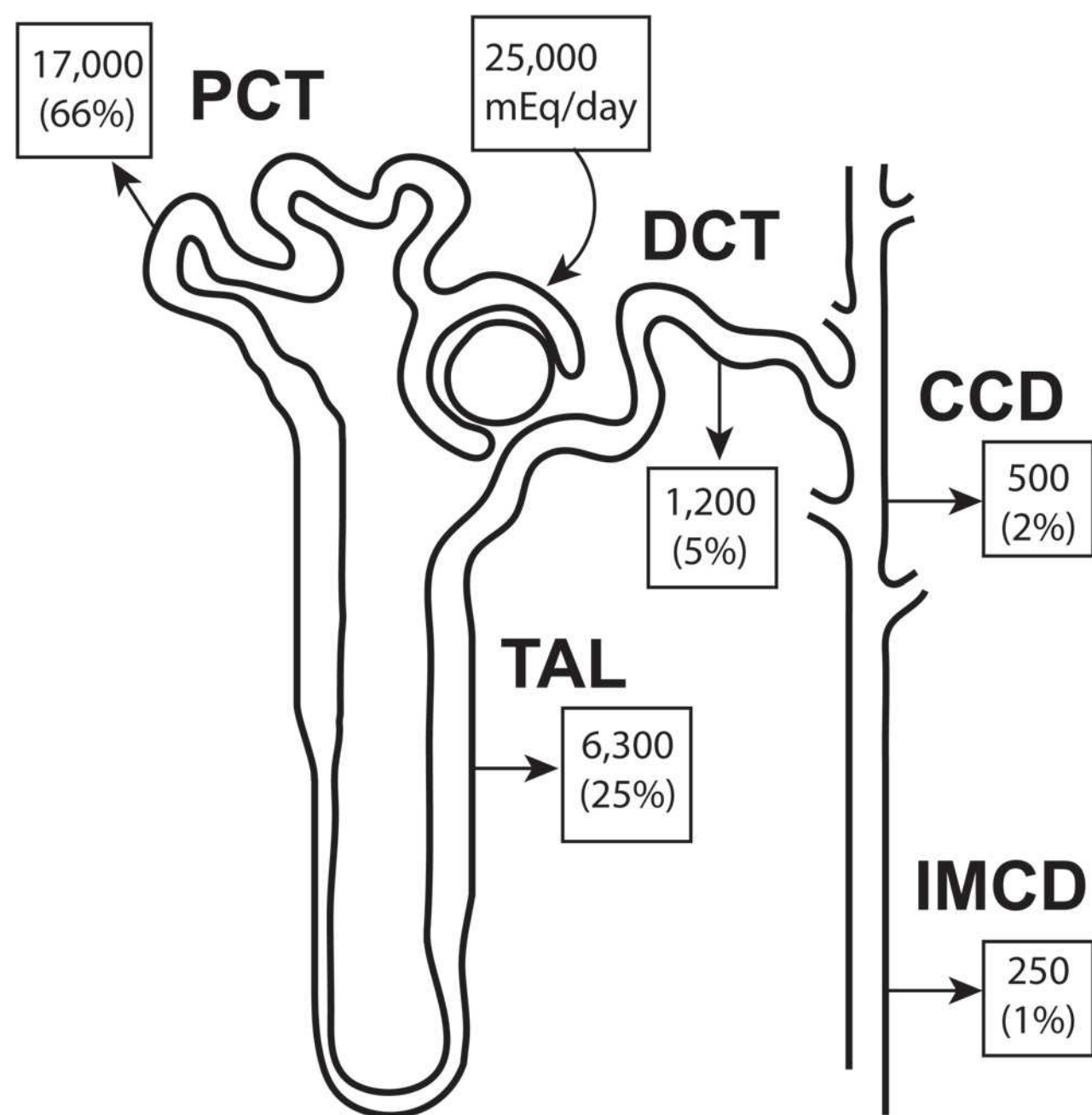


FIGURE 5.1 The contribution of various nephron segments to Na⁺ transport. *PCT*, proximal convoluted tubule; *DCT*, distal convoluted tubule; *CCD*, cortical collecting duct; *TAL*, thick ascending limb; *IMCD*, inner medullary collecting duct.

individual molecule is random, a concentration gradient is required for any net transfer of molecules to occur across a membrane. Thus, the concentration gradient represents the driving force for net transport.

For charged solutes, the driving force for transport is the sum of the chemical and electrical potential gradients. The Nernst equation describes the equilibrium condition for a membrane permeable only to a single ionic species:

$$V_m = V_2 - V_1 = -(RT/ZF) \ln (C_2/C_1) \quad (1)$$

where R is the gas constant, T is the absolute temperature, Z is the valence of the solute, F is the Faraday constant, and C and V are concentration and electrical potential terms, respectively. At equilibrium, then, the voltage (V_m) across an ideally selective membrane is defined by the concentrations of the permeant ion on both sides of the membrane, C_2 and C_1 , respectively. For systems containing more than one permeant ion, the equilibrium voltage can be described by the Goldman–Hodgkin–Katz (GHK) equation^{2,3}:

$$V_m = -RT/F \ln [(P_{Na}C_{2Na} + P_KC_{2K} + P_{Cl}C_{1Cl}) / (P_{Na}C_{1Na} + P_KC_{1K} + P_{Cl}C_{2Cl})] \quad (2)$$

where P_x is the permeability of the respective solutes, in this case, Na⁺, K⁺, and Cl⁻. Thus, in a system containing multiple charged solutes, the transmembrane voltage is a function of the relative concentrations and permeabilities of each solute on the two sides of the membrane.

Convective Processes

Convection is the vectorial movement of an ensemble of molecules and is driven by an externally imposed force (e.g., hydrostatic pressure). Examples of convective transport include glomerular filtration and solvent drag, a process in which solute movement is coupled to water movement.

Bulk water flow may be driven by hydrostatic pressure and/or osmotic pressure. The familiar Starling equation:

$$J_v = K(\Delta P - \Delta \pi) \quad (3)$$

describes net volume flow (J_v) in response to hydrostatic (ΔP) and osmotic ($\Delta \pi$) pressure differences. The equivalence of osmotic and hydrostatic pressure is explicit in the Starling equation. The degree to which a solute exerts an osmotic pressure depends on the degree to which it permeates membranes. The ratio of the observed osmotic pressure to that predicted if a solute were excluded absolutely from a membrane is termed the reflection coefficient, σ :

$$\sigma = \Delta \pi_{\text{obs}} / \Delta \pi_{\text{theoretical}} \quad (4)$$

For impermeant solutes, $\sigma = 1$; for highly permeable solutes, σ approaches 0.

For solutes with $\sigma < 1$, transmembrane solute flux will be accelerated in the direction of volume flow. This acceleration is known as solvent drag.⁴ Thus, the net passive flux of a permeable solute across a membrane may be driven by both diffusion and entrainment with solvent flow (i.e., solvent drag).

Facilitated Diffusion

Biologic membranes are composed primarily of lipid bilayers. Because the permeability of many hydrophilic solutes through lipid membranes is low, membranes contain proteins that facilitate the transport of certain solutes. Transport proteins, often termed carriers or transporters, have a high degree of specificity for the transported solute. Flux through the limited number of transporters saturates as the solute concentration is increased. An example of carrier-mediated facilitated diffusion is the entry of glucose into renal tubular cells mediated by the hexose transporter, GLUT-1.⁵ The movement of ions through ion channels represents another form of facilitated diffusion. In this case, integral membrane proteins containing several membrane-spanning domains form pores in cell membranes through which ions permeate. Ion channels generally have a high degree of specificity for the ions being transported and very high transport rates. Facilitated diffusion mechanisms, like enzymes, serve only to accelerate the rate of transport, but do not affect the equilibrium distribution of solutes. In other words, facilitated diffusion, like simple diffusion and convection, is a passive process that tends to dissipate transmembrane gradients.



Active Transport Processes

Active transport is a special case of facilitated transport in which chemical bond energy is supplied to the transport process so that the final distribution of the solute is remote from equilibrium. The coupling of solute transport to the energy source can take two forms. In primary active transport, solute transport is coupled directly to an energy-yielding reaction.

The most widely recognized example of primary active transport is the transport of Na^+ and K^+ by the Na^+, K^+ -ATPase. This enzyme, often referred to as the sodium pump, couples the extrusion of cellular Na^+ to cellular K^+ uptake.⁶ In renal tubules, this enzyme is localized to the basolateral membrane. In general, segments with high rates of active Na^+ transport have high Na^+, K^+ -ATPase activity.⁷ The hydrolysis of each ATP molecule ordinarily pumps three sodium ions out of the cell coupled to two potassium ions moving inward.⁸ Therefore, the pump is electrogenic. The Na^+, K^+ -ATPase is responsible for maintaining the cell Na^+ activity at a low level, which provides the energy for the Na^+ -coupled transport of many other solutes. Thus, the inhibition of Na^+, K^+ -ATPase (e.g., by peritubular ouabain addition) causes a significant rise in the cell Na^+ activity.⁹ The affinity constant (K_m) of the pump for intracellular sodium, about 15 to 30 mM,¹⁰ is similar to the intracellular sodium activity measured in proximal tubule cells.⁹ Therefore, the pump is unsaturated with respect to sodium, and pump activity is very sensitive to changes in the intracellular sodium concentration activity.

In secondary active transport, solute movement against its electrochemical gradient is energized by the movement of another solute down its own gradient.¹¹ Na^+ , because of its steep inward electrochemical gradient maintained by the sodium pump, often participates in the transport of other solutes, either in the same (cotransport or symport) or opposite (exchange or antiport) direction. Thus, by coupling solute transport with sodium movement into cells, cellular metabolic energy generated by the Na^+, K^+ -ATPase is stored in the form of a Na^+ concentration gradient, analogous to a battery, and then dissipated in the transport of a variety of different solutes. Some examples of symport processes include Na^+ -glucose and Na^+ -amino acid cotransport; Na^+ -proton and Na^+ -calcium exchange are two examples of antiport processes.

PROXIMAL TUBULE

General Features

The proximal tubule is the major site for Na^+ absorption within the kidney and serves two major purposes. First, the proximal tubule protects the extracellular fluid volume by reclaiming the bulk, approximately 60% to 80%, of the glomerular filtrate.^{12,13} The proximal tubule with its well-developed brush border membrane is optimally designed to perform the reabsorption of such a large fraction of the

filtrate.¹⁴ Second, the absorption of sodium in the proximal tubule provides, by way of coupled processes, the driving force for the absorption of other solutes, such as bicarbonate, glucose, phosphate, and amino acids.

Under most circumstances, fluid at any given point along the proximal tubule has virtually the same Na^+ concentration and osmolality as plasma.¹³ The isosmotic nature of proximal tubule fluid absorption derives from the high water permeability of this segment,¹⁵ which effectively clamps the osmolality of the tubular fluid at that of plasma. Although Na^+ transport in the proximal tubule occurs in the absence of large electrical or chemical gradients, the bulk of Na^+ absorption in the proximal tubule involves active transport. For example, Na^+ can be reabsorbed against both concentration^{13,16} and electrical¹⁷ gradients. In addition, fluid absorption and sodium transport cease when Na^+, K^+ -ATPase activity is inhibited or when cell metabolism is slowed.^{18,19}

A significant amount of proximal Na^+ transport also occurs passively.²⁰ For example, in the late convoluted and straight tubules (S2 and S3 segments), Na^+ diffuses passively out of the tubule driven by the lumen-positive electrical potential difference in those segments. This potential difference derives from a Cl^- concentration gradient across the tubule wall. Even in this case, however, it is the active transport of Na^+ in upstream portions of the proximal nephron that ultimately accounts for these gradients and potentials.

Nephron Heterogeneity

Analyses of proximal tubular Na^+ transport are complicated by two factors: a nonhomogeneous nephron population and axial changes in fluid composition. There is considerable heterogeneity of both morphologic and functional characteristics along the proximal tubule. The S1 segment cells have extensive basal interdigitations, numerous mitochondria, and a well-developed luminal brush border.²¹ S3 segment cells, in contrast, are flatter and have fewer mitochondria, lower brush borders, and much less extensive basolateral membranes than S1 cells.²¹ The Na^+, K^+ -ATPase activity of the S3 segment is only 25% of that for the S1 segments.²² As might be expected on the basis of these observations, the net rates of the Na^+ and fluid transport in the S3 straight segment are, in general, lower than in the S1 convoluted tubule.²³

Juxtamedullary proximal convoluted tubules have higher rates of volume and bicarbonate absorption than their superficial counterparts,^{23,24} although this disparity has not been noted between juxtamedullary and superficial straight segments.²⁵

Glomerular ultrafiltrate in the early proximal tubule undergoes axial composition changes. Figure 5.2 illustrates that the chloride concentration rises as a consequence of the preferential absorption of NaHCO_3^- over NaCl in this segment.²⁶ Glucose, amino acids, and other organic compounds are also absorbed avidly in association with Na^+ so that their luminal concentrations in this

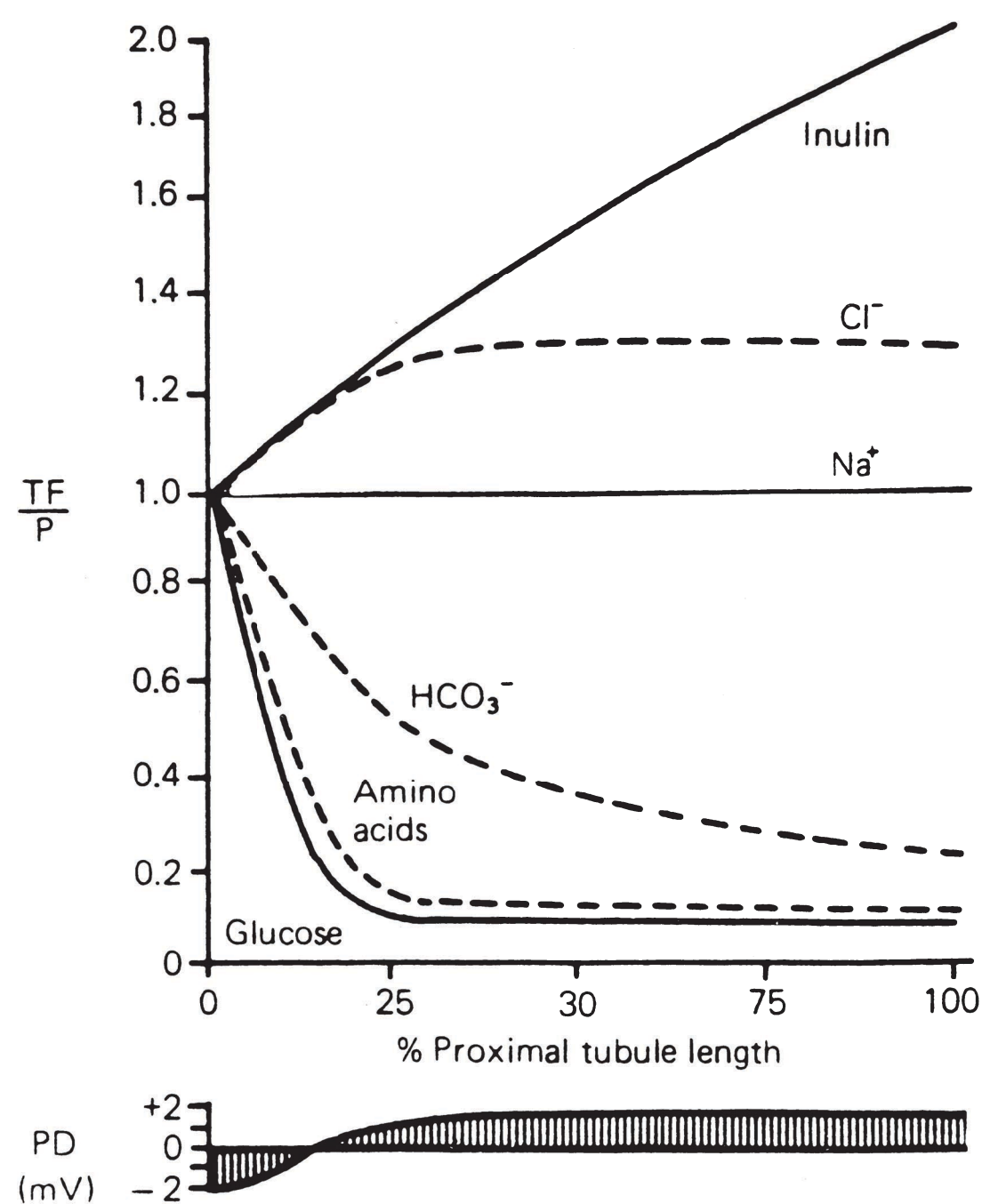


FIGURE 5.2 The profile of transepithelial voltage and solute concentrations along the mammalian proximal tubule. TF/P , tubular fluid/plasma concentration ratio; PD , potential difference. (From: Rector FC Jr. Sodium, bicarbonate, and chloride absorption by the proximal tubule. *Am J Physiol.* 1983;244:F461, with permission.)

segment approach zero.^{26,27} The omission of glucose and amino acids from luminal fluids reduces both the potential difference and the volume absorptive rate.^{24,28} In the S3 segment, on the other hand, the omission of glucose and alanine has no effect either on the potential difference or on the fluid absorptive rate,^{25,29} although the deletion of all organic solutes does reduce volume absorption by 50%.³⁰ The rates of transport of glucose, amino acids, phosphate, and Na^+ in the early proximal convoluted tubule exceed those in the proximal straight tubule.^{31–33} These rates correlate well with the relative basolateral membrane areas of the respective segments.³⁴

Electrophysiology of the Proximal Tubule

Transepithelial Potential Difference

The electrogenic nature of the transport of Na^+ coupled to glucose and amino acids creates a lumen-negative transepithelial electrical potential difference in the early proximal convoluted tubule. The deletion of glucose and alanine from the luminal fluid reduces the potential difference from about -5.0 mV, lumen-negative, nearly to zero.^{24,28} This transepithelial potential difference becomes lumen-positive ($+2$ to 4 mV) when the tubular fluid to plasma chloride concentration ratio is approximately 1.3.³⁵ This lumen-positive voltage is probably a diffusion potential arising from the Cl^- concentration gradient.

Electrical Resistance

The electrical resistance of the mammalian proximal tubule is remarkably low, making this tubule a classic example of a leaky epithelium, with resistances of 5 to $10 \Omega\text{-cm}^2$.^{36,37} In the proximal tubule, the total cellular resistance (i.e., the sum of the apical and basolateral resistance) is 20- to 70-fold greater than the transepithelial resistance. This indicates that the paracellular resistance is low and that the predominant route for passive ion flows in the proximal tubule involves the paracellular pathway.

Ionic Selectivity

The initial convolution of the superficial proximal tubule is Na^+ selective; thereafter, the superficial convoluted and straight tubules are Cl^- selective.^{29,38} In contrast, juxtamedullary proximal tubules are Na^+ selective throughout their course.³⁸ Because the Cl^- concentration rises as fluid flows along the convoluted tubule (Fig. 5.2), oppositely directed gradients for Cl^- and HCO_3^- in late convoluted and straight segments give rise to a lumen-positive transepithelial potential difference.²⁶ The latter indicates a higher permeability for Cl^- than HCO_3^- , both in superficial and juxtamedullary proximal tubules.^{23,39}

MECHANISMS OF SODIUM REABSORPTION

Apical Membrane Sodium Entry

In the proximal tubule, Na^+ entry into the cell may be coupled to the movement of other solutes, such as glucose, chloride, or protons; or Na^+ may enter independently. In either case, the driving force for Na^+ entry is the steep electrochemical gradient favoring Na^+ influx.

The intracellular Na^+ activity ranges from 15 to 35 mM.^{10,40} The entry of Na^+ into cells appears to be rate limiting for transepithelial Na^+ transport. Amphotericin B, a polyene antifungal, increases the permeability of the luminal membrane to Na^+ and causes a large rise in net sodium absorption.⁴¹

Na^+/H^+ Exchange

Directly coupled Na^+/H^+ exchange in the proximal tubular brush border is responsible for most proton secretion and for a large fraction of Na^+ reabsorption in the proximal tubule.⁴² The mechanism whereby Na^+/H^+ exchange effects Na^+ reabsorption is presented in Figure 5.3. Briefly, entry of Na^+ is coupled to extrusion of a proton into the lumen. The proton titrates a filtered HCO_3^- molecule to form carbonic acid. Carbonic acid subsequently is dehydrated to CO_2 in a reaction catalyzed by carbonic anhydrase IV in the brush border membrane.⁴³ Within the cell, the reverse process occurs: carbonic acid formed by the hydration of CO_2 dissociates into H^+ and HCO_3^- . The H^+ is extruded into the lumen by Na^+/H^+ exchange or by the vacuolar $H^+-ATPase$ to repeat another cycle while the HCO_3^- is transported into the blood via a $1Na^+:3HCO_3^-$ cotransport process (vide infra).

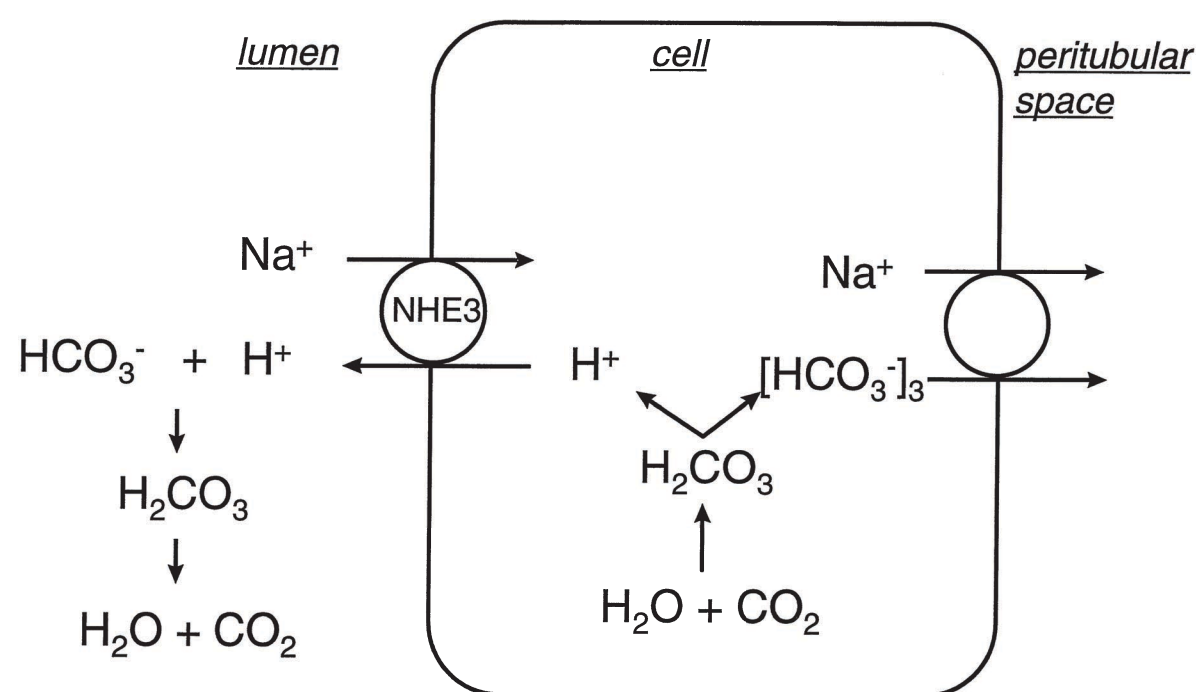


FIGURE 5.3 The scheme of NaHCO_3 transport mediated by Na^+/H^+ exchange. See text for explanation.

The mammalian Na^+/H^+ exchanger (NHE) is electro-neutral with a stoichiometry of one proton for one sodium.⁴⁴ The exchanger is reversibly inhibited by high concentrations of amiloride.⁴⁵ Intracellular protons, via an internal activator site,⁴⁶ increase Na^+/H^+ exchange in response to intracellular acidosis.⁴⁷

The apical and basolateral membranes of kidney cells contain different forms of NHEs with different affinities for amiloride.⁴⁸ The apical Na^+/H^+ exchanger is involved in urinary acidification and has a low amiloride affinity, whereas the basolateral exchanger has a high affinity for amiloride. The sensitivity of NHE1 to amiloride⁴⁹ and its basolateral localization⁵⁰ suggest that it represents the “housekeeping” NHE. In contrast, NHE3 is in the brush border membrane of proximal tubule cells.⁵¹ NHE3 knockout mice have significantly reduced rates of Na^+ and HCO_3^- transport in proximal tubules.⁵² In addition, pharmacologic inhibitors of NHE3 reduce proximal tubule Na^+ reabsorption by about one third.⁵³ However, a significant rate of amiloride-sensitive HCO_3^- transport still persists in the proximal tubules of NHE3 knockout mice,⁵⁴ indicating that the NHE3 is responsible for much, but not all, proximal tubular Na^+ -coupled luminal acidification (vide infra). NHE8 is also expressed in the apical membranes of cortical tubules and may contribute to these processes.⁵⁵

Sodium–Glucose Cotransport

Electrophysiologic studies in kidney proximal tubules show that apical membranes depolarize with addition of glucose to luminal fluids.⁵⁶ The depolarization occurs because the Na^+ -glucose transporter is electrogenic. The Na^+ -glucose transporter is specific for the D-stereoisomers of glucose, galactose, and α -methyl-D-glucoside.⁵⁷ The Na^+ -glucose cotransporter has little affinity for cations other than Na^+ .⁵⁸ Phlorizin inhibits Na^+ -glucose cotransport by competing with glucose for its binding site.⁵⁹

The rate of glucose transport by the early proximal tubule is greater than in late proximal segments.⁶⁰ In the proximal straight tubule, the K_m for D-glucose is 5 to 20 times lower than in the proximal convoluted tubule. More-

over, the transporter in the early cortical proximal tubule has a 1:1 Na^+ to glucose stoichiometry,⁶¹ whereas the transporter in the straight medullary segment has a 2:1 stoichiometry.⁶² By coupling the energy from two Na^+ ions moving down their electrochemical gradient to the transport of each glucose molecule, the medullary transporter is able to establish a much greater cellular to extracellular glucose concentration ratio than a 1:1 Na^+ :glucose transporter.²⁷ The 2 Na^+ :1 glucose transporter is, therefore, well suited to the straight segment, where tubular fluid glucose concentrations have already been reduced by glucose absorption in the more proximal segments.

A Na^+ -glucose cotransporter (SGLT-1), which belongs to the SLC5 gene family,⁶³ mediates high affinity Na^+ -glucose cotransport with a sodium-to-glucose coupling ratio of 2:1, whereas SGLT-2⁶⁴ shares 59% homology to SGLT-1 and mediates low-affinity Na^+ -glucose cotransport with a sodium-to-glucose coupling ratio of 1:1.⁶⁵ In situ hybridization revealed high levels of a SGLT-2 message in the S1 segment of the proximal tubule.⁶⁶ Recently, better antibodies have confirmed that, in rat kidney, SGLT-1 immunolocalizes to the brush border membrane of all three segments of the proximal tubule.⁵ By immunohistochemistry, SGLT-2 was detected at the brush border of the early proximal tubule in mice, which was absent in SGLT-2 knockout animals.⁶⁷ Thus, it appears that SGLT-2 may represent the low-affinity, high-capacity sodium–glucose cotransporter in the early proximal tubule, whereas SGLT-1 may represent the high-affinity, low-capacity transporter of the proximal straight tubule.⁶⁵

Mutations in SGLT-2 form the basis for renal glycosuria (Table 5.1), an inherited condition characterized by a lowered threshold for tubular reabsorption of glucose.⁶⁸ In contrast, the dominant clinical manifestations of inactivating mutations of SGLT-1⁶⁹ relate to the failure to absorb sugars in the intestinal tract (glucose–galactose malabsorption). These findings suggest that SGLT-2 plays a much more significant role, quantitatively, than SGLT-1 in proximal tubule glucose reabsorption. SGLT-2 inhibitors are currently under investigation as potential therapeutic agents for the treatment of diabetes.⁷⁰

Sodium–Amino Acid Cotransport

The proximal tubule reabsorbs amino acids from the tubular fluid via an active transport step at the luminal membrane.³¹ Samarzija and Frömter,⁷¹ using double-perfusion micro-puncture techniques, observed a depolarization of the luminal membrane during amino acid transport, and they were able to identify five classes of amino acid transporters in the luminal membrane. Over the last decade, many of the transport proteins that mediate the different amino acid transport systems have been identified in kidney and intestine. This topic has been recently reviewed.⁷²

Both Na^+ -dependent and Na^+ -independent amino acid uptake pathways have been characterized in the kidney.⁷² Neutral amino acid transport appears to involve at least

three separate transport systems,⁷³ one that transports all neutral amino acids, one specific for imino acids, and one for the β -amino acids. Glycine may also have a specific transporter.⁷⁴ In the kidney, neutral amino acid transport is driven by a Na^+ gradient, as supported by experiments in slices, perfused tubules, and brush border membranes. The neutral amino acid transporter B⁰AT1 (SLC6A19) cotransports one Na^+ per amino acid.⁷⁵ The K_m of the substrate decreases with an increasing cosubstrate concentration and vice versa. The initial step for transport involves the binding of the amino acid to B⁰AT1, and this binding affinity increases under hyperpolarizing conditions.⁷⁶

The acidic and basic amino acid groups each have their own transport systems.^{77,78} At least one amino acid transporter, a Na^+ -independent transporter for neutral and dibasic amino acids, has been cloned from the kidney.^{79,80}

NaCl Transport

Two basic mechanisms account for NaCl reabsorption in the proximal tubule. In simple electrogenic Na^+ entry, sodium is transported actively through the cell, thereby creating a lumen-negative potential difference. Cl^- reabsorption then proceeds through the paracellular pathway driven by the lumen-negative potential difference. In electrically neutral NaCl transport, both Na^+ and Cl^- move through the cell at equal rates, such that no transepithelial potential and, hence, no driving force for paracellular Cl^- movement is generated.

Neutral NaCl Transport

Several lines of evidence indicate that a sizable fraction of proximal NaCl transport is transcellular and electroneutral.⁸¹ First, by virtue of the coupling of Cl^- entry to apical Na^+ entry, the intracellular Cl^- activity of proximal tubule cells is greater than predicted from an equilibrium distribution.⁸² Second, Cl^- absorption persists even when the driving force for passive, paracellular movement is abolished.⁸³ Conversely, Cl^- reabsorption is inhibited by cyanide in the absence of any change in the passive driving forces for Cl^- movement.⁸³ Finally, the luminal application of SITS,⁸⁴ an anion-exchange inhibitor, or removal of chloride from the tubule perfusate,⁸⁵ reduces net Na^+ reabsorption.

In principle, electroneutral NaCl transport across the apical membrane of proximal tubule cells could occur as directly coupled NaCl cotransport or as parallel Na^+/H^+ and Cl^- /base exchangers. There is no good evidence for the former process in the mammalian proximal tubule.⁸¹ However, considerable evidence supports the view that electroneutral NaCl transport in the proximal tubule involves parallel exchangers. The coupling of Na^+ absorption to Cl^- absorption in this case occurs because of the relation between cell pH and concentration of base within the cell. With reference to Figure 5.4, the extrusion of H^+ in exchange for Na^+ results in the liberation of the base for participation in Cl^- /base exchange. The uphill entry of Cl^- , then, is indirectly coupled to the downhill entry of Na^+ , because both are coupled to the transport of an acid–base pair. The model illustrated

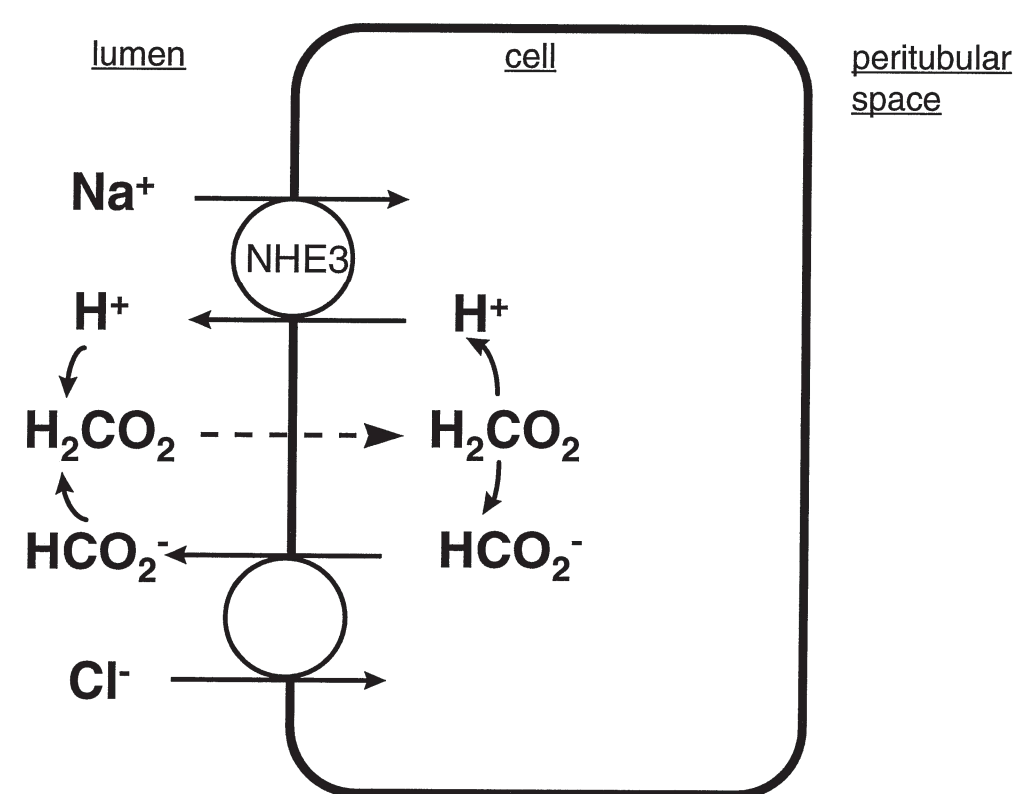


FIGURE 5.4 The scheme of neutral NaCl transport mediated by the parallel action of $\text{Na}^+ - \text{H}^+$ exchange and formate/ Cl^- exchange. Formate (HCO_2^-) combines with H^+ in the tubular lumen to form formic acid (H_2CO_2), which reenters the cell by non-ionic diffusion. A similar scheme applies for oxalate– Cl^- exchange.

in Figure 5.4 uses formate/ Cl^- exchange as the anionic component of electroneutral NaCl transport.

As indicated previously, there is abundant evidence for a Na^+/H^+ exchanger in proximal tubule brush border membranes. With respect to NaCl transport, the inhibition of Na^+/H^+ exchange by high concentrations of amiloride⁸⁶ or more specific inhibitors of NHE3⁵³ results in a dramatic fall in transcellular NaCl transport. Likewise, knockout of NHE3 also reduces NaCl and fluid reabsorption in the proximal tubule.⁵² Several Cl^- /base exchangers have been implicated in NaCl transport. Recent interest has focused on the role of Cl^- /formate (HCO_2^-) and Cl^- /oxalate ($\text{C}_2\text{O}_4^{2-}$) exchange in NaCl transport.

A Cl^- /formate exchanger is present in brush border membrane vesicles.⁸⁷ A role for Cl^- /formate exchange in neutral NaCl transport is suggested by the finding that the addition of formate to the luminal perfusate increases the rate of NaCl reabsorption in rabbit proximal tubules.⁸⁸ As depicted in Figure 5.4, formate is presumed to leave the cell in exchange for Cl^- . The secreted formate then combines with a proton, which was transported by the Na^+/H^+ exchanger to form formic acid. The formic acid then reenters the cell by nonionic diffusion and dissociates to supply substrate for the continuation of both exchange processes. CFEX (SLC26A6), a homolog of pendrin, is a protein capable of mediating Cl^- /formate exchange and is present in apical membranes of the proximal tubule.⁸⁹

Cl^- /oxalate ($\text{C}_2\text{O}_4^{2-}$) exchange has also been demonstrated in brush border membrane vesicles.⁹⁰ It has been suggested that Cl^- /oxalate exchange may mediate neutral NaCl transport in a manner analogous to that described for Cl^- /formate exchange. It has also been suggested that NaCl absorption proceeds via the operation of three parallel transporters: the Na^+ -sulfate cotransport, the sulfate/oxalate exchange, and the Cl^- /oxalate exchange.⁹¹ Indeed, the CFEX protein, in addition to Cl^- /formate exchange, is also able to mediate Cl^- /oxalate, oxalate/formate, oxalate/oxalate, and oxalate/sulfate exchange.⁹²

Other Na^+ -dependent transport processes have been described in the apical membrane of the proximal tubule. However, they do not contribute significantly to Na^+ reabsorption because of the low concentrations of substrate present.

Simple Electrogenic Na^+ Entry

The classic Ussing model for salt reabsorption involves passive entry of Na^+ across apical membranes and extrusion across the basolateral membrane by Na^+, K^+ -ATPase. A problem in assessing the contributions of electrogenic processes to Na^+ transport in the proximal tubule is the presence of other mechanisms of Na^+ entry. However, when the contributions of Na^+/H^+ exchange and Na^+ cotransport to net Na^+ absorption are minimized by deleting glucose, amino acids, and bicarbonate from the perfusate, a fraction of fluid absorption in isolated perfused straight segments persists and the transepithelial potential is -1.0 mV .³⁰ These results indicate that in the proximal straight tubule simple electrogenic Na^+ transport constitutes a mechanism for Na^+ absorption. A conductive Na^+ pathway has been demonstrated in brush border membrane vesicles.⁹³ Unlike the Na^+ channel found in the distal nephron segments, the Na^+ channel in the proximal tubule is not blocked by amiloride.⁹³

In the proximal convoluted tubule, however, the deletion of glucose, bicarbonate, and amino acids completely abolishes fluid absorption.²³ Consequently, simple electrogenic proximal Na^+ transport may be limited to straight segments.

Passive NaCl Absorption

The rise in tubular fluid Cl^- concentration, and the attendant lumen-positive voltage (Fig. 5.2), provides a mechanism for passive NaCl absorption in late regions of the proximal nephron. In the Cl-selective superficial pars recta, approximately one-third of net NaCl absorption can be accounted for by this mechanism.²⁰

Basolateral Membrane

The proximal tubule, particularly the S1 segment, possesses high Na^+, K^+ -ATPase activity in the basolateral membrane. The Na^+, K^+ -ATPase pumps Na^+ , which entered cells apically, across basolateral membranes. In other words, the pump keeps the cell Na^+ activity low and maintains the electrochemical gradient for Na^+ entry across the apical membrane. Consequently, the inhibition of Na^+, K^+ -ATPase activity with ouabain decreases transepithelial Na^+ reabsorption and increases the intracellular Na^+ activity in the proximal tubule.⁹⁴

Na^+ also exits across the basolateral membrane in concert with HCO_3^- . Studies in intact tubules and in membrane vesicles have demonstrated an electrogenic, stilbene-sensitive $\text{Na}^+/\text{HCO}_3^-$ cotransporter in the basolateral membrane of rat and rabbit proximal tubules.^{95,96} The cotransporter transfers two net negative charges across the basolateral membrane. The stoichiometry of this process is $1 \text{ Na}^+ : 1 \text{ HCO}_3^- : 1 \text{ CO}_3^{2-}$ (or SO_3^{2-}).⁹⁷ Thus, this transport moiety for Na^+ extrusion, $[\text{Na}^+(\text{HCO}_3^-)_3]^{2-}$ is electronegative, with the lumen-negative cell interior providing a major driving

for Na^+ extrusion. A number of $\text{Na}^+/\text{HCO}_3^-$ cotransporters have been cloned, along with different splice variants.⁹⁸ One of these, NBC1 (encoded by SLC4A4 and subsequently renamed NBCe1-A) is localized to the basolateral membrane of the S1 and S2 segments of the proximal tubule.⁹⁹ As illustrated in Figure 5.3, the net result of these steps is the reabsorption of Na^+ and HCO_3^- , accounting for the bulk of HCO_3^- reabsorption and about 20% of Na^+ reabsorption in the proximal tubule. It is believed that this cotransport process accounts for the basolateral transport of most of the bicarbonate reclaimed from the luminal fluid.⁹⁷ Mutations in NBCe1-A cause proximal (type II) renal tubular acidosis and other defects in the eye, teeth, and mental development (Table 5.1). The regulation and role of NBC in acid-base transport in the proximal tubule is an area of active research that has been reviewed recently.¹⁰⁰

The pathways for Cl^- exit across the basolateral membrane are less well defined. Several pathways for Cl^- exit across the basolateral membrane have been proposed: conductive Cl^- channels, KCl cotransport, and $\text{Na}^+(\text{HCO}_3^-)_2/\text{Cl}^-$ exchange (Fig. 5.5). Studies of rat proximal convoluted tubules¹⁰¹ and rabbit convoluted and straight proximal tubule segments^{102,103} using intracellular microelectrodes have indicated that the proximal tubule cell has a very low Cl^- conductance. Thus, in normal proximal convoluted tubule and proximal straight tubule segments, conductive Cl^- efflux across basolateral membranes appears to play a minor role in NaCl absorption. Under hypotonic conditions, however, cell swelling dramatically increases the basolateral membrane Cl^- conductance.¹⁰⁴

Because the chemical gradient for K^+ to leave cells exceeds that for Cl^- entry, KCl cotransport can mediate basolateral Cl^- exit from proximal tubule cells. Ion-selective microelectrode studies have demonstrated KCl cotransport in basolateral membranes of rabbit proximal tubule cells.^{103,105}

Stilbene-sensitive, Na^+ -dependent $\text{Cl}^-/\text{HCO}_3^-$ exchange has been demonstrated in rat¹⁰⁶ and rabbit^{107,108} proximal tubules. In this case, the entry of 1 Na^+ and 2 HCO_3^- across the basolateral membrane is coupled to the efflux of Cl^- . The Na^+

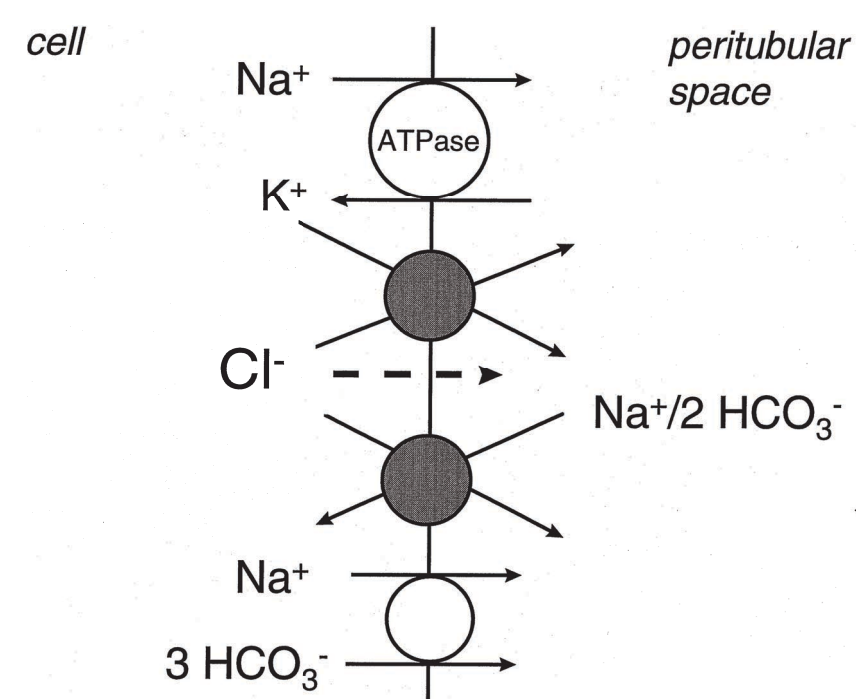


FIGURE 5.5 The transport pathways for Na^+ and Cl^- absorption across the basolateral membrane of proximal tubular cells. Cl^- can leave the cell via KCl cotransport, $\text{Na}^+ - 2\text{HCO}_3^-/\text{Cl}^-$ exchange, and Cl^- channels (minor). Na^+ exits via the Na^+, K^+ -ATPase and $\text{Na}^+ - 3(\text{HCO}_3^-)$ cotransport.

and HCO_3^- that enter the cell are thought to be recycled through the $[\text{Na}^+(\text{HCO}_3^-)_3]^{2-}$ cotransporter (see previous). Indeed, $\text{Na}^+(\text{HCO}_3^-)_2/\text{Cl}^-$ exchange may account for much more Cl^- movement than KCl transport.¹⁰⁸ Na^+ -independent $\text{Cl}^-/\text{HCO}_3^-$ exchange is also present in the basement membrane of proximal tubules.^{106–108} However, under physiologic conditions, this process mediates net Cl^- influx and does not contribute to net NaCl absorption.

CONTROL OF PROXIMAL TUBULAR SODIUM REABSORPTION

Glomerulotubular Balance (GTB)

The proximal tubule responds to an increase in glomerular filtration with an increase in the absolute rate of fluid absorption (APR) to minimize variations in the fractional proximal fluid absorption. This phenomenon is termed glomerulotubular balance (GTB). The efficiency of GTB—that is, the extent to which APR/GFR remains constant—is subject to physiologic and pathologic control. The prime factor modulating GTB in vivo is the effective circulating volume. Thus, at a constant or near constant GFR, volume expansion and volume contraction decrease and increase, respectively, the absolute rate of proximal Na^+ absorption. In other words, volume expansion and volume contraction reset GTB upward and downward, respectively.^{109,110} This section considers some of the factors that modulate proximal Na^+ absorption.

Peritubular capillary oncotic pressure is one of the factors regulating the rate of salt and water absorption from the proximal tubule.¹¹¹ The oncotic pressure of the peritubular proteins favors the movement of fluid across the basement membrane, whereas capillary hydrostatic pressure retards this movement. Thus, at a given renal blood flow, an increase in the glomerular filtration rate and, hence, the filtration fraction, will cause an increase in the oncotic pressure in the postglomerular peritubular capillaries. At constant single nephron glomerular filtration rates (SNGFR), the perfusion of efferent capillaries with hypo-oncotic fluids decreases the absolute rate of proximal fluid absorption, whereas perfusion of the capillaries with hyperoncotic fluids increases proximal absorption.¹¹² The effects of the peritubular protein concentration can also be demonstrated in isolated perfused tubules.¹¹³

It is not precisely clear how the peritubular protein concentration modulates proximal fluid absorption. The effect is not simply because of the oncotic pressure exerted by the proteins, because changes in absorption do not occur when active transport is inhibited or from comparable changes in the transtubular hydrostatic pressure.¹¹⁴ The prevailing views are that the peritubular protein may directly affect transcellular Na^+ transport¹¹⁵ or the back leak of Na^+ through the paracellular pathway.^{113,116}

Luminal factors also contribute to glomerulotubular balance. The flow dependence of proximal absorption has been investigated in the convoluted¹⁵ and straight segments³⁰ of

isolated, perfused rabbit nephrons. The key observations are (1) the volume absorptive rate is clearly dependent on the perfusion rate, (2) the flow dependence persists in the absence of active transport when anion gradients are present, and (3) the flow dependence is abolished in the absence of active transport and anion gradients.

An explanation for these results lies in a consideration of axial versus radial changes in fluid composition along the tubule.¹¹⁷ The rate of passive Na^+ absorption in the late proximal tubule is dependent on the magnitude of the chloride gradient between the lumen and the bath. At low axial perfusion rates, the radial Cl^- gradient tends to dissipate as a function of distance along the tubule, but this dissipation is minimized at higher axial perfusion rates. Hence, the integrated driving force for passive NaCl absorption increases with the rate of tubule perfusion. In addition, the availability of solutes, such as glucose, amino acids, and bicarbonate, is also partly responsible for the flow dependence of proximal fluid absorption.¹¹⁸

The flow dependence of reabsorption in the proximal tubule has also been tested in the mouse proximal tubule. In this experimental model, which included the use of the NHE3 knockout mouse, the data supported the hypothesis that the “brush border” microvilli act as mechanosensors that transmit fluid dynamic torque to the actin cytoskeleton and thus modulate Na^+ absorption.¹¹⁹

Catecholamines

Renal denervation reduces proximal tubule Na^+ and fluid absorption.¹²⁰ Both α - and β -adrenergic receptors exist in the proximal tubule.^{121,122} The rate of salt and water reabsorption in the proximal tubule is stimulated by α - and β -adrenergic agonists.¹²³ α -Adrenergic agonists increase apical Na^+ entry via the stimulation of Na^+/H^+ exchange¹²² and also increase basolateral Na^+ efflux via Na^+,K^+ -ATPase activity in rat proximal tubules by a pathway that involves the activation of calcineurin.¹²⁴ The effects of β -adrenergic agonists on Na^+,K^+ -ATPase activity are less clear, with one study showing that they increase Na^+,K^+ -ATPase activity via protein kinase C (PKC).¹²⁵ However, others have found that β -adrenergic agonists inhibit Na^+,K^+ -ATPase activity via PKA.¹²⁶

Dopamine, which is produced by proximal tubular cells,¹²⁷ inhibits Na^+ reabsorption. Dopamine inhibits Na^+,K^+ -ATPase activity via its receptors DA-1 and DA-2.¹²⁸

Parathyroid Hormone

Parathyroid hormone (PTH) causes a 30% to 50% reduction in proximal tubular Na^+ and phosphate absorption.¹²⁹ PTH stimulates adenylyl cyclase, cAMP production, and PKA, which in turn inhibits Na^+/H^+ exchange in several proximal tubule systems.¹³⁰ Weinman et al.¹³¹ demonstrated that phosphorylation by PKA inhibits Na^+/H^+ exchange activity via a PDZ domain-dependent interaction with the NHE regulatory factor (NHERF). In the absence of NHERF, cAMP

does not inhibit the exchange activity of NHE3.¹³² The current model for this inhibition is that NHE3 associates with PKA indirectly via NHERF and the cytoskeletal protein ezrin. PKA, when active, phosphorylates NHE3 at serines 552 and 605, which mediates the inhibition of the exchanger.¹³³ NHE3 is directly phosphorylated by other protein kinases, including calmodulin-dependent protein kinase II, which inhibits Na^+/H^+ exchange activity and PKC, which stimulates the exchanger.¹³⁴

Angiotensin II

The systemic administration of low doses of angiotensin II (Ang II) inhibits the excretion of Na^+ ,¹³⁵ whereas inhibitors of Ang II increase Na^+ excretion.¹³⁶ Systemic Ang II causes changes in renal blood flow, aldosterone secretion, filtration fraction, and catecholamine release from renal sympathetic nerve endings.¹³⁷ Low concentrations of Ang II ($<10^{-9}$ M) cause an increase in proximal tubule fluid and bicarbonate reabsorption, effects that are mediated by the AT1 subtype of Ang II receptors present in both the brush border and basolateral membranes of the proximal tubule.¹³⁸ Higher concentrations ($>10^{-8}$ M) depress fluid and bicarbonate absorption, presumably via counterbalancing effects mediated by the lower affinity AT2 receptor.¹³⁸ Studies have demonstrated that the stimulatory effect of Ang II on fluid and bicarbonate reabsorption occurs via enhanced apical Na^+/H^+ exchange via NHE3 and basolateral $\text{Na}^+-\text{HCO}_3^-$ cotransport via NBC-1 in the proximal tubule.¹³⁹ The physiologic effects of Ang II may involve the coupling of these receptors to both phospholipase A_2 and inhibitory G proteins.^{140,141}

Thyroid Hormone

On a clinical level, hypothyroidism is associated with a decreased cardiac output, renal blood flow, and glomerular filtration rate (GFR). Clearance studies in hypothyroid rats have documented decreases in GFR, renal Na^+ reabsorption, and renal Na^+,K^+ -ATPase activity.¹⁴² These changes are reversible after thyroid hormone replacement.¹⁴³ The thyroid hormone may exert direct effects to stimulate proximal tubular salt and fluid reabsorption via increased basolateral K^+ permeability¹⁴⁴ and/or direct stimulation of Na^+/H^+ exchange through an increase in NHE3 transcription.¹⁴⁵

Corticosteroids

Although mineralocorticoids do not have an effect on proximal tubular sodium reabsorption,¹⁴⁶ there is evidence for glucocorticoid receptors in the proximal tubule.¹⁴⁷ Dexamethasone inhibits apical membrane Na^+ -phosphate cotransport in cultured proximal tubular cells via PKC activation.¹⁴⁸ Dexamethasone also enhances the activity of apical NHE3 and the mRNA expression and functional activity of basolateral NBC-1.¹⁴⁹ The resulting increase in proximal tubule HCO_3^- reabsorption could contribute to the maintenance of the metabolic alkalosis that is associated with increased glucocorticoid production in vivo.

Nitric Oxide

Various forms of nitric oxide synthase (NOS) are expressed in the proximal tubule.¹⁵⁰ Low basal production of nitric oxide (NO) by the proximal tubule is boosted dramatically by lipopolysaccharide (LPS) and cytokines. Even under basal conditions, the proximal tubule may be affected by NO produced by adjacent cells, such as endothelium or other nephron segments. The overall effect of NO on proximal tubule Na^+ transport is controversial and may be biphasic, with acute inhibition and chronic stimulation of Na^+ reabsorption as assessed by pharmacologic agents and genetic knockouts of NOS, respectively.¹⁵¹ In vitro, NO decreased Na^+/H^+ exchange and Na^+,K^+ -ATPase activity in cultured proximal tubule cells.¹⁵²

MECHANISM OF ISOTONIC FLUID ABSORPTION

Proximal tubule water absorption is coupled tightly to solute absorption, because the measured osmolality of the tubular fluid is generally identical to plasma. Three general mechanisms of solute-solvent coupling have been suggested to account for this isotonic absorption: lateral interspace hypertonicity, effective osmotic gradients because of different reflection coefficients for solutes in the tubular and peritubular fluids, and luminal fluid hypotonicity.

The standing gradient theory argues that active transport of salt into the lateral intercellular space raises the osmolality of the space, thus providing an osmotic gradient for fluid transport from the lumen to the interspace.¹⁵³ The tight junctions in this model are presumed to be impermeable to water, so that the osmotic flow of water from the cell into the hypertonic interspace raises the hydrostatic pressure in this compartment and forces fluid across the basement membrane.

An alternative explanation¹¹⁷ proposes that an effective osmotic driving force for fluid absorption can exist between solutions of identical osmolalities if the reflection coefficients of the membrane for the solutes in the solutions differ. Specifically, the elevated tubular fluid-to-plasma Cl^- concentration found in the late proximal convolution and in the pars recta provides an effective osmotic driving force for fluid absorption because σHCO_3^- exceeds σCl^- . That is, the bicarbonate in the peritubular fluid is a more “effective” osmotic agent than is the chloride in the tubular fluid, and, thus, net water flows out of the tubule. Although this mechanism may be applicable to the proximal straight tubule, the reflection coefficients for NaCl and NaHCO_3 measured across the rabbit proximal convoluted tubule are virtually identical,¹⁵⁴ so oppositely directed Cl^- and HCO_3^- gradients may make only a negligible contribution to fluid absorption in convoluted segments.

Finally, because of the high osmotic water permeability of the proximal tubule, only small degrees of absolute luminal hypotonicity are needed to provide a sufficient driving

force to account for the observed rates of fluid reabsorption.²⁰ Experimental evidence supports the view that absolute luminal hypotonicity is a significant driving force for fluid reabsorption in the proximal tubule. Thus, when proximal tubules are perfused and bathed by symmetric NaCl solutions, the luminal fluid becomes slightly hypotonic.¹⁵⁵ The development of luminal hypotonicity can be amplified by maneuvers that decrease the water permeability of the proximal tubule. The aquaporin 1 (AQP1) water channel is abundantly expressed in the proximal tubule. In AQP1 knockout mice, the osmolality of tubular fluid at the end of the proximal tubule is significantly lower than in normal mice.¹⁵⁶ As the luminal fluid becomes more hypotonic, the resorbate becomes more hypertonic, and the degree of resorbate hypertonicity correlates with the rate of volume reabsorption by the tubules.¹⁵⁷

THE LOOP OF HENLE

The dissociation of salt and water absorption by the loop of Henle is ultimately responsible for the capacity of the kidney, either to concentrate or to dilute the urine. The active absorption of NaCl in the water-impermeable thick ascending limb of Henle (TALH) serves both to dilute the urine and to supply the energy for the “single effect” of countercurrent multiplication. A functionally similar segment, known as the diluting segment, is found in amphibians and teleosts.¹⁵⁸

The mammalian loop of Henle contains the descending thin limb (DTL), the ascending thin limb (ATL), and the thick ascending limb (TAL). The loop of Henle absorbs about 25% to 40% of the filtered Na^+ load.¹⁵⁹ Furthermore, the fluid leaving the loop is dilute, indicating that more NaCl is absorbed in the loop than water.

SALT TRANSPORT BY THE THIN DESCENDING AND THIN ASCENDING SEGMENTS

As the tubular fluid enters the descending thin limb and flows toward the tip of the renal papilla, it becomes more concentrated.¹⁶⁰ Passive models for urinary concentration indicate that this increase in osmotic pressure is attributable to water extraction rather than solute entry.¹⁶¹ Current observations indicate that the aquaporin water channel AQP1 mediates water movement across the luminal surface of the DTL.¹⁶² According to some reports, however, AQP1 expression is significantly lower in short loop nephrons than long loop nephrons,¹⁶³ suggesting that (1) not all DTLs in the kidney extract water to the same degree, and/or (2) water movement in short loop DTLs is facilitated by alternative water channels or via the paracellular route. In contrast to the DTL, *in vitro* microperfusion studies demonstrate that the ATL is relatively impermeable to water.¹⁶⁴

The study of NaCl transport by the DTL and ATL has been complicated by the fact that the transport characteristics of these two nephron segments exhibit significant

interspecies heterogeneity.¹⁶⁵ Although the Na^+ and Cl^- permeabilities appear to vary widely among rabbits, hamsters, and rats, one consistent finding was that the DTL is relatively less permeable to NaCl than the ATL. Coupled with the DTL's high permeability to water, the relative lack of solute permeability ensures that the osmotic pressure of fluid entering the renal papilla is greater than that of the fluid leaving it.¹⁶⁶ Thus, the formation of dilute urine by the loop of Henle begins in the ATL.

The decrease in osmolality in the ATL is due primarily to a fall in the NaCl content of the luminal fluid. The electroneutral transport of NaCl appears to occur through two key mechanisms. The transepithelial movement of sodium from the ATL lumen occurs via the paracellular pathway,¹⁶⁷ whereas Cl^- diffusion occurs through a transcellular route. Yoshitomi et al.¹⁶⁸ detected conductive pathways for Cl^- in both the apical and basolateral membranes of ATL cells. The transcellular movement of Cl^- in the ATL is regulated, because the basolateral Cl^- conductance is inhibited at low pH¹⁶⁸ and by low intracellular Ca^{2+} concentrations.¹⁶⁹ Uchida et al.¹⁷⁰ cloned a Cl^- channel in 1993 from the rat renal medulla, ClC-K1 (termed ClC-Ka in humans), which represents the major mediator of transcellular Cl^- movement in the thin ascending limb. This channel, which belongs to the ClC family of Cl^- channels, is expressed exclusively within the kidney and has been localized by immunohistochemistry to both the apical and basolateral membranes of the thin ascending limb of Henle.¹⁷¹ Its activity is dependent on the coexpression of barttin, an accessory protein that forms a complex with ClC-K1, increases its abundance at plasma membranes, and modifies channel gating.^{172–175} The expression of ClC-K1 is increased by dehydration.¹⁷⁰ Genetic knockout of the ClC-K1 gene in mice produced a urinary-concentrating defect, confirming the role of passive NaCl transport in the thin ascending limb in the urinary concentrating mechanism.¹⁷⁶

NaCl ABSORPTION IN THE THICK ASCENDING LIMB

General Features

The studies of Rocha and Kokko¹⁷⁷ and Burg and Green¹⁷⁸ were the first to investigate salt absorption in the thick ascending limb, and their work defined several key features of this unique epithelium. First, salt absorption in the medullary and cortical TAL generates a lumen-positive transepithelial voltage, which is sensitive to furosemide. Second, the transport of Cl^- occurs against both electrical and chemical gradients and involves an active transport process that is dependent on intact basolateral Na^+, K^+ -ATPase activity.¹⁷⁹ A final important feature of the TAL is that this segment consists of a tight epithelium, which despite its high ionic conductance, possesses a very low permeability to water. The apical membrane of the TAL constitutes the major barrier to transcellular and paracellular water flow.¹⁸⁰ The high

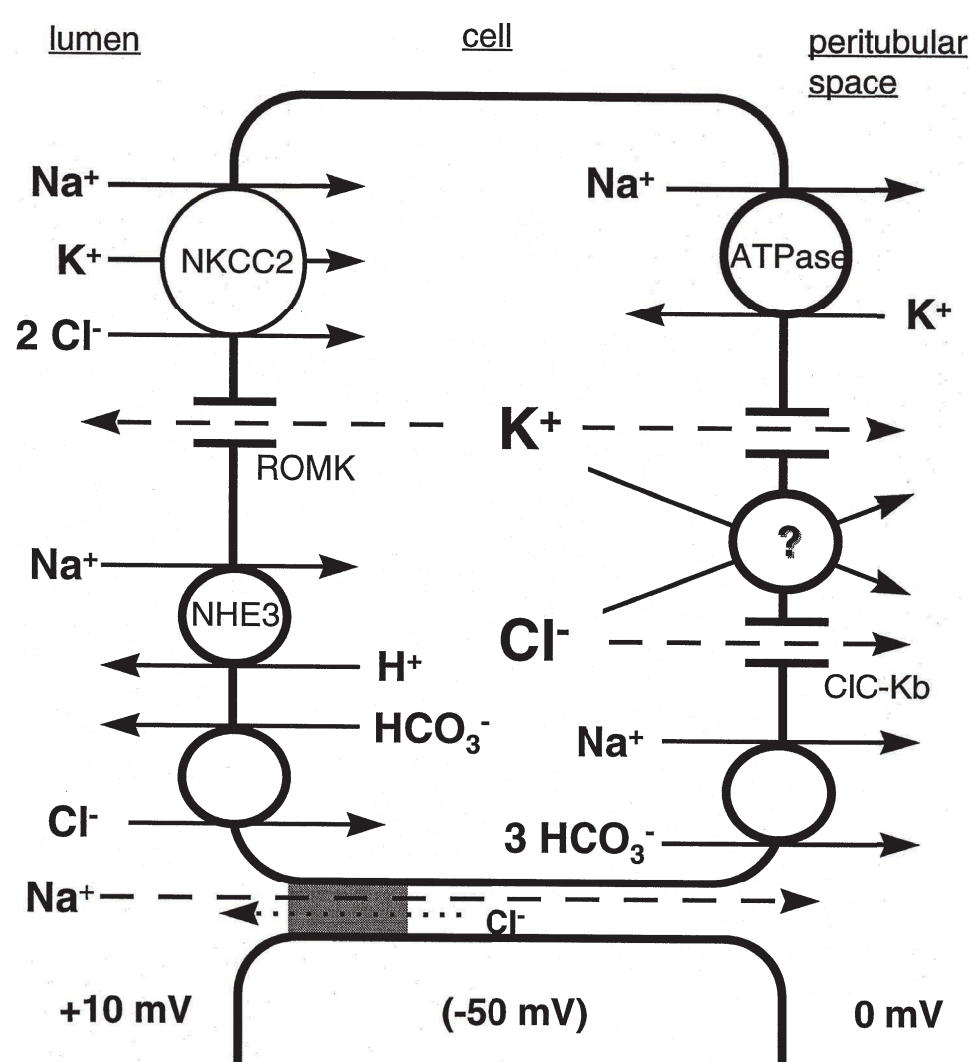


FIGURE 5.6 A model depicting the major elements of the mechanism of NaCl absorption by the thick ascending limb. *Dashed lines* indicate passive ion movements down electrochemical gradients. *ROMK*, renal outer medullary K⁺ channel; *ClC-Kb*, chloride channel Kb.

ionic conductance and low water permeability effectively further dilutes fluid entering the TAL from the ascending thin limb.

Figure 5.6 integrates the results of several electrophysiologic and biochemical studies to provide a model of salt reabsorption in the thick ascending limb. According to this model, net Cl⁻ absorption by the TAL is a secondary active transport process. Luminal Cl⁻ entry into the cell is mediated by an electroneutral Na⁺-K⁺-2Cl⁻ cotransport process driven predominantly by the favorable electrochemical gradient for Na⁺ entry.¹⁸¹ Because the Na⁺ gradient is maintained by the continuous operation of the basolateral membrane Na⁺,K⁺-ATPase pump, the apical entry of Cl⁻ via the cotransporter ultimately depends on the operation of the basolateral Na⁺,K⁺-ATPase.

In contrast to the electroneutral entry of Cl⁻ across the apical membrane, the majority of Cl⁻ efflux across the basolateral membrane proceeds through conductive pathways.^{182,183} A favorable electrochemical gradient for Cl⁻ efflux through dissipative pathways has been demonstrated by Greger et al.¹⁸⁴ in the rabbit cTAL. Intracellular Cl⁻ is maintained at concentrations above electrochemical equilibrium by the continued entry of Cl⁻ via the apical Na⁺-K⁺-2Cl⁻ cotransporter.¹⁸⁵

According to the model in Figure 5.6, K⁺ that enters TAL cells via the Na⁺-K⁺-2Cl⁻ cotransporter recycles, to a large extent, across the apical membrane via a K⁺-conductive pathway. This apical K⁺ recycling serves several purposes. First, it ensures a continued supply of luminal K⁺ to sustain Na⁺-K⁺-2Cl⁻ cotransport. Without recycling, the luminal K⁺ concentration would fall rapidly as a consequence of K⁺ entry via Na⁺-K⁺-2Cl⁻ cotransport and would limit net NaCl absorption. Second, the apical membrane K⁺ current

provides a pathway for net K⁺ secretion by the TAL. In mouse TAL, for example, the rate of K⁺ secretion amounts to about 10% of the rate of net Cl⁻ absorption.¹⁸² K⁺ secretion in this segment is an active process, ultimately driven by the Na⁺,K⁺-ATPase, proceeding in the face of a lumen-positive transepithelial potential. Third, under open circuit conditions, the transcellular and paracellular pathways form a current loop in which the currents traversing the two pathways are of equal size, but which traverse in the opposite direction. The potassium current from cell to lumen polarizes the lumen and causes an equivalent current to flow from the lumen to the bath through the paracellular pathway.¹⁸⁶ Because the paracellular pathway is cation selective ($P_{Na}/P_{Cl} = 2$ to 6), the majority of the current through the paracellular pathway is carried by Na⁺ moving from the lumen to the interstitium. This paracellular absorption of Na⁺ increases the efficiency of Na⁺ transport by the TAL.¹⁸⁷ With reference to Figure 5.6, for each Na⁺ transported through the cell and requiring the use of ATP, one Na⁺ is transported through the paracellular pathway without any additional energy expenditure. Finally, the apical K⁺ current satisfies the continuity requirement imposed by a high degree of conductive Cl⁻ efflux across basolateral membranes.¹⁸²

A small component of Na⁺ transport by the TAL is accounted for by NaHCO₃ absorption.¹⁸⁸ In the rat TAL, the rate of NaHCO₃ absorption is roughly 5% to 10% of that for NaCl absorption. NaHCO₃ absorption appears to be mediated by an apical membrane amiloride-sensitive Na⁺/H⁺ exchanger and a basolateral membrane electrogenic Na⁺-3(HCO₃⁻) cotransporter.¹⁸⁸

The following sections will describe the individual components of the mechanism for TAL salt transport (Fig. 5.6) in greater detail.

Apical Na⁺-K⁺-2Cl⁻ Cotransport

Studies of Cl⁻ transport across apical membranes of intact TAL segments¹⁸⁹ and in isolated membrane vesicle preparations¹⁹⁰ established that the predominant mode for Cl⁻ entry into the TAL cell is via a Na⁺-K⁺-2Cl⁻ cotransporter. A characteristic feature of this transporter is its sensitivity to inhibition by furosemide, bumetanide, and other 5-sulfamoylbenzoic acid derivatives.¹⁹¹ The measurement of isotope flux into TAL cells or membrane vesicles prepared from the inner stripe of the outer medulla yielded a stoichiometry of 1 Na⁺:1 K⁺:2 Cl⁻ cotransport.¹⁹⁰ K⁺-independent NaCl cotransport has also been described under certain conditions.¹⁹²

The proteins that mediate the Na⁺-K⁺-2Cl⁻ cotransport have been cloned. An absorptive form of the Na⁺-K⁺-2Cl⁻ cotransporter, referred to as NKCC2 or BSC1, was initially cloned by Gamba et al.¹⁹³ based on sequence homology to the thiazide-sensitive Na⁺-Cl⁻ cotransporter (see the following). A second Na⁺-K⁺-2Cl⁻ cotransporter, NKCC1, was cloned by Payne et al.¹⁹⁴ NKCC2 (BSC1) is the primary mediator of apical salt entry in the thick ascending limb. In situ hybridization and single-nephron reverse transcriptase polymerase

chain reaction (RT-PCR) studies demonstrated the expression of NKCC2 in the MTAL and CTAL,¹⁹⁵ and immunohistochemical studies indicate that NKCC2 is localized to the apical membrane of these nephron segments.¹⁹⁶ The importance of NKCC2 in mediating salt reabsorption in the TAL is illustrated by the fact that loss-of-function mutations of NKCC2 cause Bartter syndrome (Table 5.1),¹⁹⁷ a Mendelian salt-wasting disorder characterized by hypokalemia, metabolic alkalosis, hyperaldosteronism, and normal-to-low blood pressure, results from a defect in salt absorption by the thick ascending limb.

The NKCC2 cDNA encodes a glycoprotein containing ~1100 amino acids and having a predicted molecular weight of 115 to 120 kDa.¹⁹³ The full-length protein contains 12 transmembrane domains containing a sizable extracellular loop with N-glycosylation sites positioned between transmembrane segments 7 and 8, and large intracellular amino and carboxy termini flank the transmembrane regions. NKCC2 belongs to the SLC12A family of cation chloride cotransporters, which is part of the amino acid polyamine organocation cotransporter (APC) superfamily.¹⁹⁸ Based on the homology to other crystallized APC family members, the cotransporter structure probably consists of two clustered groups of five transmembrane helices that are positioned in a symmetric, inverted orientation.¹⁹⁹ The details regarding how this fold facilitates the three-ion cotransport remain obscure but surely will provide an initial framework for more detailed structure-function studies in the coming years.

The two cytoplasmic domains of NKCC2 mediate specific regulatory functions. The amino terminus is believed to be unstructured and contains several cytosol-exposed serines and threonines, which are phosphoacceptor sites. These residues are phosphorylated by at least three protein kinases that stimulate NKCC2 activity and/or plasma membrane expression (discussed in detail below).²⁰⁰ The carboxy terminus is large and comprises ~40% of the total NKCC2 sequence. It may also contain phosphorylation sites, although to date this has not been explored in detail. It is clear, however, that the NKCC2 C-terminus serves as a hub for interactions with proteins that regulate its trafficking, including the glycolytic enzyme aldolase²⁰¹ and secretory membrane carrier protein 2 (SCAMP2).²⁰² It also serves as the interface for the formation of NKCC2 homodimers,²⁰³ which was confirmed recently when the crystal structure of the C-terminus of the related prokaryotic cation chloride cotransporter MaCCC was solved.¹⁹⁹

At least six isoforms of NKCC2 have been identified.²⁰⁰ These isoforms are the result of alternative splicing of two regions of the NKCC2 gene: the first region is a 96 base pair region that encodes part of the second transmembrane domain, whereas the second region encompasses the extreme C-terminus. Three variants of the 96 base pair region are encoded by different versions of exon 4 of the NKCC2 gene. These exons are differentially spliced into NKCC2 pre-mRNAs to generate three distinct isoforms (A, B, and F), which alter the amino acid composition of the second transmembrane

domain. Each of the A, B, and F isoforms can have either a long C-terminus, or a truncated C-terminus; although to date, the short isoforms have only been described in the murine TAL.²⁰⁴ In addition, several “tandem” transcripts have been described in the human kidney; these contain combinations of exons 4A, 4B, and 4F spliced alongside one another into the NKCC2 pre-mRNA.²⁰⁵ Transcripts containing exons 4A/4F, 4B/4A, and 4B/4A/4F have been reported. Because these tandem transcripts contain redundant sequences encoding for the second transmembrane domain, they probably cause the misfolding of NKCC2, resulting in the formation of nonfunctional isoforms. Because these isoforms may still form oligomers with NKCC2, they likely exert a dominant-negative effect on NKCC2 function and inhibit its activity.²⁰⁶

The A, B, and F isoforms show differential expression within the thick ascending limb. In the rat nephron, the A isoform was found in both the cortical and medullary TAL, the B isoform is restricted to the cortical TAL, whereas the F isoform is present in the medullary, but not the cortical, the TAL, and to a lesser extent, in the outer medullary collecting duct. Although some interspecies discrepancies have been noted, similar findings have generally been observed in the embryonic mouse and human kidney.^{205,207,208}

When the 4A, 4B, and 4F exons are spliced into transcripts containing a long C-terminus, all three products are capable of mediating $\text{Na}^+\text{-K}^+\text{-2Cl}^-$ cotransport. However, the A, B, and F isoforms have different transport properties that may have physiologic relevance. Isoforms A and B have higher affinities for Na^+ , K^+ , and Cl^- than the F isoform. The A isoform possesses the highest transport capacity of all three isoforms. Based on the known distribution of the A, B, and F isoforms in the TAL, it is currently thought that the A isoform accounts for the high transport capacity of the medullary TAL, whereas the presence of the more active A and B isoforms in the cortical TAL allows for the continued reabsorption of salt to take place, even though the tubular fluid in this segment is more dilute than plasma. Supporting this is experimental evidence demonstrating that the NKCC2 A and B isoforms can both be strongly activated by Na^+ , K^+ , and Cl^- at concentrations that are much more dilute than the composition of tubular fluid in the cortical TAL.²⁰⁸

Apical K^+ Conductance

An important feature of the luminal membrane of the TAL is a barium-inhibitable potassium conductance.¹⁸⁵ This apical membrane K^+ conductance allows K^+ to be recycled from the cell back into the luminal fluid to support further NaCl absorption via the NaCl cotransporter. Using measured values of intracellular K^+ activity (rabbit cTAL),²⁰⁹ apical membrane conductance, and intracellular voltage, it can be shown that the measured apical membrane conductance is sufficient to provide for the recycling of all of the potassium uptake via the $\text{Na}^+\text{-K}^+\text{-2Cl}^-$ cotransporter.

Three types of K^+ conductances have been characterized in the apical membrane of the thick ascending limb: a high-conductance (150 pS) calcium-activated K^+ channel, which

does not contribute to net K^+ transport,²¹⁰ an intermediate-conductance (70 pS) K^+ channel,²¹¹ and a low-conductance (30 to 35 pS) K^+ channel.^{211,212} The latter two conductances account for the majority of the apical K^+ secretion in the TAL.²¹³ Both the intermediate- and the low-conductance channels are inhibited by intracellular ATP and are increased by high dietary potassium.²¹⁴

A major breakthrough in elucidating the molecular mechanism of K^+ secretion in the TAL was made when Ho et al.²¹⁵ cloned the ATP-dependent K^+ channel ROMK from rat kidney. This channel is the prototype for a large family of inward-rectifying K channels (Kir channels), and hence, is also called Kir 1.1. Based on its biophysical properties and regulation, investigators had long suspected that ROMK was the sole mediator of the 30-pS K^+ conductance in the TAL. This hypothesis was confirmed when studies of the thick ascending limb in the ROMK knockout mouse revealed the absence of a 30-pS channel.²¹⁶ Subsequent studies of the ROMK knockout mouse demonstrated that these mice also lack a 70pS conductance.²¹⁴ Thus, it is likely that ROMK also comprises at least a portion of the intermediate K^+ secretory conductance in this nephron segment. It has been proposed that the intermediate-conductance channel may be a heteromeric protein containing ROMK and other, as yet, unidentified subunits.

The notion that ROMK is the predominant channel responsible for apical K^+ recycling in the thick ascending limb is also supported by the finding of mutations in the ROMK gene in families with type II Bartter syndrome.^{217,218} These mutations have generally been confirmed to result in defective K^+ -channel function.²¹⁹ As noted, Bartter syndrome results from a defect in thick ascending limb salt transport. Thus, the presence of ROMK mutations as a cause of Bartter syndrome indicates the important role of ROMK in net salt absorption by the thick ascending limb (Table 5.1).

Basolateral Membrane Cl^- Transport

Cl^- exit across the basolateral membrane of TAL cells is largely conductive, proceeding down its electrochemical gradient through Cl^- -selective channels in the basolateral membrane.²²⁰ The primary mediator of basolateral chloride transport in the TAL in humans is thought to be the chloride channel ClC-Kb, the sequence of which is closely related to the aforementioned human ClC-Ka channel expressed in the ATL (see previously). The name for the corresponding rodent ortholog of ClC-Kb is ClC-K2. In contrast to the relatively narrow expression of ClC-K1, ClC-K2 is expressed broadly throughout the basolateral membranes of multiple segments in the rodent nephron, including the TAL, the distal convoluted tubule, the connecting tubule, and the collecting duct.^{171,221,222} Inherited mutations of the ClC-Kb cause type III Bartter syndrome, which can manifest as a mixed Bartter/Gitelman phenotype, possibly owing to the expression of ClC-K2/ClC-Kb in the distal convoluted tubule (DCT) and TAL.²²³

Similar to ClC-K1/ClC-Ka channels in the ATL, ClC-K2/ClC-Kb channels require barttin accessory subunits to be fully functional. Barttin was originally identified by positional cloning of the BSND gene, which is responsible for type IV Bartter syndrome (Table 5.1), a severe form of hereditary salt wasting that is accompanied by sensorineural deafness.²²⁴ Barttin is located in the basolateral membranes of the thin and thick ascending limb, distal nephron, and also in the stria vascularis of the inner ear, where it is believed to play a role in K^+ secretion into the endolymph. The regulation of ClC-K channels by barttin appears to be multifaceted, because the accessory subunit influences channel protein stability, subcellular localization, and gating.^{172,173,175} The mutations of barttin identified in Bartter syndrome patients generally impair the ability of barttin to produce a Cl^- conductance when expressed with ClC-Kb.^{172,173}

Although less completely studied, there do appear to be additional transport pathways that mediate basolateral Cl^- flux in the TAL. For example, a barium-sensitive transcellular K^+ - Cl^- cotransport mechanism has been proposed (Fig. 5.6).²²⁵ The expression of two KCl cotransporters that belong to the same family of SLC12 electro-neutral cotransporters as NKCC2, KCC1,²²⁶ and KCC4²²⁷ have been observed. Thus, both of these cotransporters may participate in the extrusion of K^+ and Cl^- from cells in the cortical and medullary TAL. Studies in knockout animals, however, indicate that KCC4 is not the primary mediator of basolateral Cl^- flux in the TAL, because these mice do not develop a salt-wasting phenotype. Rather, the KCC4 knockout mouse develops metabolic acidosis and sensorineural deafness, suggesting that KCC4 plays important physiologic functions in the acid-secreting intercalated cells of the collecting duct and in the cells of the inner ear.²²⁸

Synchronous $Na^+/H^+:Cl^-/HCO_3^-$ Exchange

Friedman and Andreoli²²⁹ found that net Cl^- absorption and the transepithelial voltage were doubled when CO_2 and HCO_3^- were added to the external solutions bathing cortical TAL segments. Because the $(CO_2 + HCO_3^-)$ -stimulated rate of NaCl absorption did not result in net CO_2 transport and could be abolished by the lipophilic carbonic anhydrase inhibitor ethoxzolamide or by the luminal addition of the anion-exchange inhibitor SITS or DIDS, it was proposed that the apical membrane of the mouse cortical TAL contains parallel, near synchronous $Na^+/H^+:Cl^-/HCO_3^-$ exchangers in addition to a $Na^+-K^+-2Cl^-$ cotransporter. The addition of CO_2 and HCO_3^- to the bathing solutions had no effect on net NaCl transport in either the rabbit cTAL or the mouse mTAL. Both the rat and mouse medullary TAL do contain Na^+/H^+ exchangers in their apical membranes. However, in these segments, Na^+/H^+ exchange plays a role in net HCO_3^- transport and cell pH regulation rather than transcellular NaCl absorption.²³⁰ The NHE3 isoform of the Na^+/H^+ exchanger is the major isoform expressed in the apical membrane of the thick ascending limb.^{231,232}

REGULATION OF SALT ABSORPTION IN THE TAL

Vasopressin

The peptide hormone arginine vasopressin (AVP, also known as antidiuretic hormone, ADH) remains the most extensively characterized stimulatory hormone for NaCl reabsorption in the TAL. The reabsorption of NaCl in the TAL is crucial for efficient urinary concentration, because it is this process that plays a key role in maintaining a hypertonic medullary interstitial solute gradient for water reabsorption in more distal portions of the nephron.²³³ Thus, from a teleologic perspective, vasopressin ought to be an important regulator of this process. The cognate receptor for vasopressin, the V2 receptor (V2R), is expressed in both the cortical and medullary TAL, where it participates in signaling cascades that stimulate NKCC2 activity.²³⁴

Binding of AVP to V2R results in increased intracellular levels of cAMP in the TAL. The increase in cAMP levels ultimately drives the translocation of NKCC2 from intracellular subapical vesicles to the luminal plasma membrane. In addition, cAMP serves as a signal that increases the phosphorylation of residues in the NKCC2 amino terminus that stimulate cotransporter activity *in vitro*.^{235,236} Although this process may be mediated upstream by the cAMP-dependent protein kinase A (PKA), it is unclear whether PKA directly phosphorylates the N-terminal residues that stimulate its activity. Rather, it appears that two other kinases in particular are important downstream mediators of this process. These kinases, the Ste20 SPS1-related proline alanine-rich kinase (SPAK) and oxidative stress responsive kinase 1 (OSR1), are structurally homologous, activated by vasopressin, bind to a defined docking site harbored within the NKCC2 N-terminus, and directly phosphorylate these previously identified stimulatory residues (Fig. 5.7).^{237–239}

The importance of SPAK and OSR1 in NKCC2 function has recently been established in transgenic and knockout models.^{237,240,241} A complete knockout of SPAK mRNA and

protein expression yields a mild salt-wasting phenotype that approaches Gitelman syndrome, likely owing to the elimination of its activity in the DCT (see the following). However, the absence of a Bartter-like phenotype in these animals does not refute the importance of SPAK and OSR1 in the TAL. Rather, the Gitelman-like salt-wasting phenotype in these animals appears to be due to two factors. First, knockout of the SPAK gene not only ablates full-length kinase-active SPAK, but also eliminates the expression of a truncated kidney-specific kinase-defective SPAK isoform (KS-SPAK) that suppresses baseline SPAK and OSR1 activity in the TAL.²³⁷ Second, in the SPAK knockout mouse, TAL-expressed OSR1 appears to compensate for the absence of SPAK activity. Both of these effects ultimately result in increased rather than decreased NKCC2 abundance and phosphorylation, which probably compensates for the lack of DCT salt reabsorption and gives rise to a relatively mild salt-wasting phenotype. The dominance of OSR1 over SPAK in the TAL was recently verified in studies of the OSR1 knockout mouse, which, in contrast to the SPAK knockout animal, exhibits a Bartter-like phenotype.²⁴¹ On the other hand, knock-in mice bearing a mutation that ablates a key catalytic activation site in SPAK do exhibit decreased NKCC2 phosphorylation; presumably, this inactivating mutation exerts a dominant-negative effect on NKCC2 phosphorylation by preventing the binding of stimulatory kinases such as OSR1 to the NKCC2 N-terminal docking site.²⁴² Although it is clear that SPAK and OSR1 are important kinases that are linked to tubular salt reabsorption in the TAL, it is unclear how they directly connect to the well-established upstream cAMP-dependent signaling cascades, which are mediated by vasopressin. Nevertheless, given the fact that vasopressin activates SPAK and OSR1 and stimulates NKCC2 phosphorylation at key SPAK/OSR1 phosphorylation sites, it would appear that these kinases are important downstream intermediaries of the vasopressin signaling pathway.

In addition to its effects on apical $\text{Na}^+\text{-K}^+\text{-2Cl}^-$ cotransport, vasopressin increases the transcellular electrical conductance of the mouse medullary TAL, and this conductance increase is a major element in the mechanism for the hormone-dependent increase in the rate of net salt absorption.^{243,244} The available evidence suggests that both the apical and basolateral membrane conductances are increased by vasopressin. In apical membranes, AVP increases conductance by increasing the functional number of K^+ channels¹⁸²; this increase occurs even when net salt absorption is abolished by furosemide. Ecelbarger et al.²⁴⁵ showed that this increase in apical K^+ conductance was at least in part due to a dramatic upregulation in ROMK abundance and apical localization. Thus, the machinery for apical recycling of K^+ is increased in the TAL, which would aid in augmenting NKCC2-mediated NaCl reabsorption.

The predominant portion of the ADH-induced increase in cellular conductance is accounted for by an increase in the basolateral membrane Cl^- conductance.¹⁸² Two mechanisms have been suggested for the hormone-

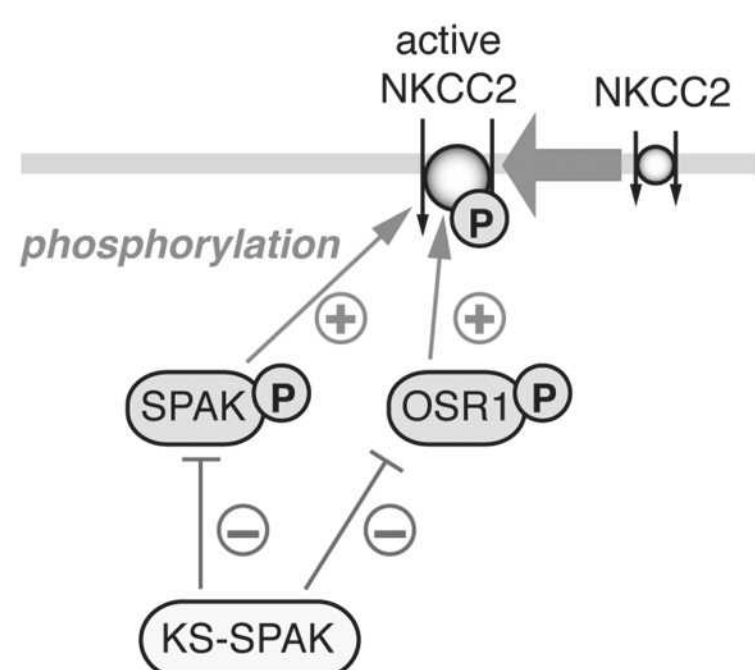


FIGURE 5.7 The current model for SPS1-related proline alanine-rich kinase /oxidative stress responsive kinase 1 (SPAK/OSR1) and kidney-specific (KS)-SPAK regulation of NKCC2 via phosphorylation in the thick ascending limb of the loop of Henle.

dependent increase in basolateral Cl^- conductance. Schlatter and Greger¹⁸³ have proposed that the ADH-induced increase in intracellular cAMP results in a direct increase in Cl^- channel activity. Such a mechanism has been amply demonstrated in Cl^- -secreting epithelia, such as the trachea and intestine. Alternatively, ADH might enhance Cl^- conductance indirectly by increasing apical membrane Cl^- entry.¹⁸² According to this proposal, an ADH-dependent activation of apical membrane NKCC2 and ROMK leads to an increase in intracellular Cl^- concentration. Because the activity of basolateral Cl^- channels is exquisitely sensitive to changes in intracellular Cl^- concentration,²⁴⁶ this increase will translate into an increase in the basolateral membrane Cl^- conductance.

Prostaglandins

Prostaglandin 1 (PGE_2), the major product of prostaglandin synthesis in the renal medulla, participates in a local negative-feedback system that modulates the rate of NaCl absorption by the TAL. PGE_2 resulted in a 50% reduction in ADH-stimulated net Cl^- absorption in isolated perfused mammalian mTAL segments, but had no effect in the absence of ADH.²⁴⁷ It is likely that PGE_2 inhibits ADH-stimulated generation of cAMP in the mTAL by activating an inhibitory G protein, G_i .²⁴⁷ Although interstitial cells are a major source of PGE_2 production in the renal medulla, thick ascending limb cells can metabolize arachidonic acid through at least two pathways. Escalante et al.²⁴⁸ demonstrated that purified medullary thick ascending limb cells produce 20-hydroxy-eicosatetraenoic acid (20-HETE) via the cytochrome P450 enzyme, ω -hydroxylase. 20-HETE was subsequently shown to inhibit NaCl transport in the thick ascending limb at steps, which include the apical K^+ channel²⁴⁹ and the $\text{Na}^+/\text{K}^+-2\text{Cl}^-$ cotransporter.²⁴⁸ Thick ascending limb cells, particularly in the macula densa (MD), also express COX-2.²⁵⁰ COX-2 expression in these cells may be coupled to renin secretion. Thus, COX-2 expression is increased by salt restriction, diuretics, and, in Bartter syndrome, all conditions characterized by hyperreninemia.²⁵¹ Moreover, COX-2 inhibitors reduce renin secretion in these settings.²⁵²

Osmolality

Peritubular osmolality is a third factor regulating mTAL salt absorption. In isolated mouse and rabbit TAL segments, increases in peritubular osmolality rapidly and reversibly inhibit the ADH-stimulated rate of net Cl^- absorption.²⁴³ Molony and Andreoli²⁵³ determined that hypertonicity inhibits the basolateral membrane Cl^- conductance. This inhibition of transcellular salt absorption occurs at a locus beyond the generation of cAMP, because supramaximal concentrations of either ADH or cAMP are unable to reverse the hypertonicity-mediated effect.²⁵⁴ Thus, increasing the absolute magnitude of interstitial osmolality provides a negative feedback signal, which can reduce ADH-dependent salt absorption by the medullary TAL.

Hypercalcemia

Hypercalcemia often results in an ADH-resistant urinary concentrating defect, that is, nephrogenic diabetes insipidus.²⁵⁵ At least part of this concentrating defect results from the inhibition of ADH-stimulated cAMP production in the TAL by calcium.²⁵⁶ Preincubation of tubule segments with pertussis toxin abolishes the effect of hypercalcemia on cAMP generation, indicating that the inhibition of cAMP generation is mediated through the activation of G_i .²⁵⁷ The effects of hypercalcemia are mediated by a G protein-coupled calcium-sensing receptor (CaSR) present on the basolateral membrane of TAL cells.²⁵⁸ Activation of this receptor also inhibits the activity of the 70-pS apical K channel (ROMK, see previous) via the production of 20-HETE, a P450 metabolite of arachidonic acid.^{259,260} Even changes in serum calcium within the physiologic range can alter NaCl absorption via the CaSR.²⁶¹ This may help to explain the hypotensive effect of high calcium intake in salt-sensitive hypertensive individuals.²⁶²

Modulation of NaCl Absorption by Other Peptide Hormones

In addition to ADH, a number of other peptide hormones stimulate adenylate cyclase activity in the TAL. In the mouse and rat, glucagon stimulates NaCl reabsorption and increases the transepithelial potential in isolated microperfused mTAL segments. Calcitonin and PTH stimulate sodium transport in the cortical, but not the medullary, portions of the TAL.²⁶³

Ang II receptors are present in the thick ascending limb.²⁶⁴ Ang II has been reported to both stimulate and inhibit sodium transport in the thick ascending limb.^{265,266} Chronic infusion of Ang II increased the abundance of NKCC2 in the rat outer medulla by 87%.²⁶⁷ In a model of heart failure, in which Ang II levels are elevated, NKCC2 expression was also increased and this increase could be prevented by treatment with an angiotensin receptor antagonist.²⁶⁸

Adrenergic Agents

β -Agonist-sensitive adenylate cyclase activity is present in the rat, but not the rabbit TAL.²⁶⁹ Likewise, β -adrenoceptors have been detected along the rat TAL by autoradiographic localization.²⁷⁰ The physiologic effects of adrenergic agents have been tested in micropuncture and in *in vitro* microperfusion studies. DiBona and Sawin²⁷¹ demonstrated an enhancement of loop NaCl absorption during low-frequency renal nerve stimulation. Acute renal denervation, on the other hand, depressed NaCl absorption by the loop of Henle.²⁷²

Micromolar concentrations of isoproterenol stimulate net Cl^- absorption by *in vitro* perfused mouse medullary and cortical TAL.²⁷³ The effects of isoproterenol on NaCl absorption in these segments can be blocked by propranolol.

Nitric Oxide

Acute administration of nitric oxide donor or L-arginine, the substrate for NOS, decreases NaCl absorption in isolated perfused thick ascending limb segments.²⁷⁴ The effect of

L-arginine on NaCl transport can be blocked by L-NAME, an inhibitor of NOS, indicating that endogenous production of NO mediates the effect of L-arginine. The inhibitory effect of NO on net NaCl absorption appears to involve, at least in part, the inhibition of NKCC2 activity.²⁷⁵ The thick ascending limb expresses all three isoforms of NOS.²⁷⁶ Plato et al.²⁷⁷ used mice deficient in the various NOS isoforms to determine that the effect of L-arginine is mediated by eNOS rather than iNOS or nNOS. In contrast to its effect on NaCl absorption, NO stimulates NaHCO_3^- absorption in the thick ascending limb.²⁶⁶ Finally, although short-term exposure to NO inhibits NaCl absorption, chronic exposure increases NKCC2 expression,²⁷⁸ which could translate into increased NaCl absorption.

Sodium Balance

Dietary Na^+ restriction in rats results in a transient decrease in NKCC2 expression,²⁷⁹ whereas high Na^+ intake has no major effect on NKCC2 expression.²⁸⁰ Chronic treatment with furosemide in conjunction with a high sodium diet increases NKCC2 expression.²⁸¹ The latter phenomenon may account for the development of diuretic resistance and the interdose rebound in sodium absorption in patients chronically treated with loop diuretics.

THE DISTAL NEPHRON

Anatomic Considerations

The distal nephron may be divided into three segments: the DCT, the connecting tubule (CNT), and the collecting duct. Perhaps owing to the nature of the original micropuncture studies characterizing the distal nephron, the DCT was initially thought to be a segment consisting of a homogeneous population of epithelial cells. However, more recent work clearly indicates that the DCT can be further divided into two functionally distinct subsegments, referred to as the “early” and “late” DCT (DCT1 and DCT2, respectively).²⁸² One of the primary features that distinguishes between the early and late portions of the DCT is the differential sensitivity of these segments to the mineralocorticoid hormone aldosterone. More specifically, the late DCT expresses the enzyme 11 β -hydroxysteroid dehydrogenase 2 (11 β -HSD2), which metabolizes cortisol, thereby rendering mineralocorticoid receptors sensitive to aldosterone.²⁸³ For this reason, the late DCT, CNT, and the cortical collecting duct (CCD) are collectively termed the aldosterone-sensitive distal nephron (ASDN).

Na^+ Transport in the Distal Convoluted Tubule and the Connecting Segment

General Characteristics

The DCT absorbs roughly 5% to 10% of the filtered Na^+ load.²⁸⁴ Fluid enters the DCT with a Na^+ concentration of 25 to 30 mM, but salt is added along the initial 20% of the

DCT, so that the Na^+ concentration averages 50 mM at a point 200 to 300 μm from the macula densa.²⁸⁵ From there, tubular Na^+ concentration decreases along the DCT to a value of approximately 30 mM at the end.²⁸⁶ Tubular fluid to plasma Na^+ ratios as low as 0.10 have been observed during stationary microperfusion.²⁸⁷ This finding, together with the presence of the lumen-negative potential difference (see the following), clearly establishes the active nature of Na^+ absorption in this segment.

Na^+ absorption by the DCT is load dependent. That is, over a wide range of delivery rates, the proportion of Na^+ absorbed by the DCT remains constant at 80%.²⁸⁶ At high tubular fluid flow rates, the fall in luminal Na^+ concentration along the tubule is attenuated; thus, more Na^+ is available to distal Na^+ absorptive sites at high rather than at low flow rates.

Electrophysiologic Considerations

Depending on the type of electrodes that were used, measurements of the transepithelial voltage in the earliest loops of the DCT vary from slightly lumen negative to slightly lumen positive.^{288–291} Consistent among all measurements, however, is the observation that the transepithelial voltage becomes progressively more lumen negative as tubular fluid passes to the end of the DCT and into the CNT and CCD.

The progressively lumen-negative transepithelial electrical potential is primarily due to a change in Na^+ transport pathways from the early DCT to more downstream nephron segments. As shown previously by Ellison et al.,²⁹² Na^+ reabsorption in the early DCT is largely mediated by an electroneutral, thiazide-sensitive NaCl cotransporter, whereas Na^+ absorption in the late DCT also involves an amiloride-sensitive electrogenic pathway.²⁸² These and other mechanisms of Na^+ absorption are discussed in detail in the following sections.

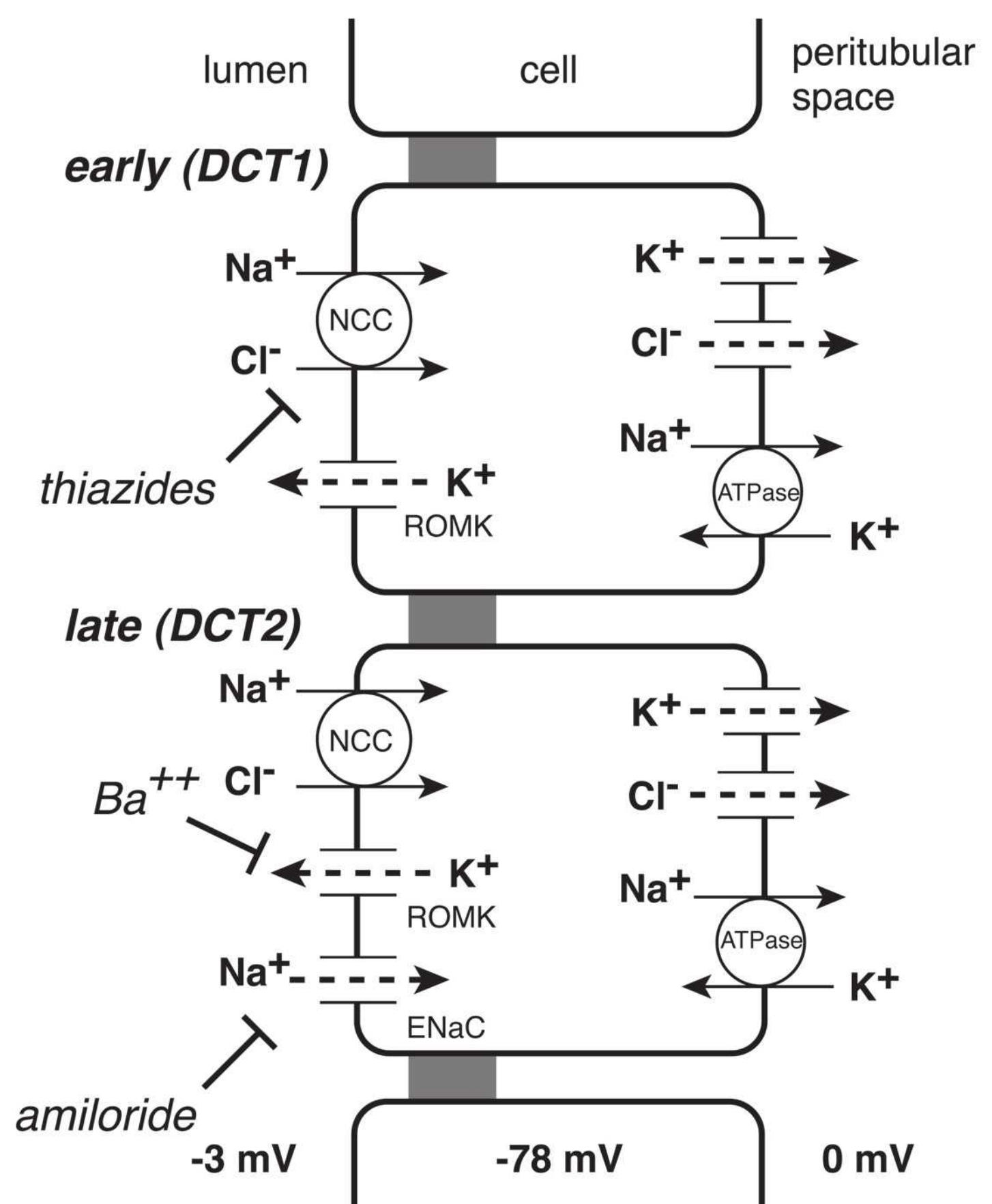
The Mechanism of Na^+ Absorption

A general model for these mechanism by which Na^+ is reabsorbed in the early and late DCT is presented in Figure 5.8.

The Apical NaCl Cotransport. The absorption of Na^+ and Cl^- in the early DCT are coupled.²⁹³ The coupling of Cl^- to Na^+ entry provides a mechanism to maintain the intracellular Cl^- activity above its electrochemical equilibrium.²⁹⁴ The early distal tubule is the site of action of thiazide diuretics.²⁹⁵ Autoradiographic studies²⁹⁶ and immunocytochemical studies²⁹⁷ have demonstrated thiazide-binding sites in the apical membranes of DCT and connecting tubule cells.

The thiazide-sensitive NaCl cotransporter (NCC, TSC, SLC12A3) mediates electroneutral NaCl reabsorption in the early and late DCT (Fig. 5.8).²⁹⁸ The cotransporter shares considerable sequence homology to the bumetanide-sensitive $\text{Na}^+ - \text{K}^+ - 2\text{Cl}^-$ cotransporter (NKCC2) present in the TAL,¹⁹³ yet exhibits markedly different inhibitor sensitivity and ionic requirements. NCC transports NaCl with a

FIGURE 5.8 A model of NaCl absorption by cells of the early and late distal convoluted tubule (DCT). The early and late DCT express NCC, a thiazide-sensitive, electro-neutral NaCl cotransporter. The late DCT also expresses the amiloride-sensitive Na^+ channel ENaC. Basolateral K^+ channels include Kir4.1 , mutated in EAST/SeSAME (epilepsy, ataxia, sensorineural deafness, and tubulopathy/seizures, sensorineural deafness, ataxia, mental retardation, and electrolyte imbalance) syndrome, a hereditary salt-wasting disorder. ROMK, renal outer medullary K channel; NCC, thiazide-sensitive NaCl cotransporter.



1:1 stoichiometry, is K^+ -independent, and is inhibited by thiazide diuretics.²⁹⁹ NCC is expressed in the apical membrane of DCT cells and extends, in most species, into the connecting segment.³⁰⁰

Gitelman Syndrome. Loss-of-function mutations in the NCC gene have been linked to the pathogenesis of Gitelman syndrome.²¹⁸ This syndrome resembles Bartter syndrome (hypokalemia, alkalosis, sodium wasting), except that urinary calcium excretion is reduced in Gitelman syndrome and is elevated in most cases of Bartter syndrome. Thus, physiologically, Gitelman syndrome mimics the effects of thiazide diuretics. Several studies have evaluated the consequences of mutations causing Gitelman syndrome on NCC function, and in most cases, these mutations alter the NCC coding sequence in such a way that they reduce NCC expression and trafficking to the apical plasma membrane of the DCT.^{301,302} In this regard, most of these disease-causing mutations result in the conformational misfolding of NCC, resulting in the recognition of mutant NCC by chaperone-dependent endoplasmic reticulum (ER) quality control machinery that targets the cotransporter for proteasomal degradation.^{301,303}

Familial Hyperkalemic Hypertension. Studies of another inherited disorder of distal Na^+ transport, familial hyperkalemic hypertension ([FHHt] type 2 pseudohypoaldosteronism,

Gordon syndrome), have yielded additional insights into the regulation of NCC function. FHHt is the phenotypic opposite of Gitelman syndrome and is characterized by hypertension, hyperkalemia, and metabolic acidosis (Table 5.1). The disorder is largely corrected by the infusion of sodium with poorly reabsorbable anions, or by treatment with thiazide diuretics.³⁰⁴ These features suggested that the hypertension in FHHt is chloride dependent, and that an increase in NCC activity may be involved in the pathogenesis of PHA II. Positional cloning demonstrated that FHHt is caused by mutations in either of two with-no-lysine (WNK) serine-threonine kinases, WNK1 or WNK4.³⁰⁵ These kinases have been intensely studied since their linkage to FHHt in 2001, and are well-established regulators of NCC, but the mechanisms by which they affect NCC function are complex. Both WNK1 and WNK4 have been shown to regulate NCC plasma membrane trafficking and activity (Fig. 5.9). In the current model it is believed that WNK4 acts as an inhibitor of NCC trafficking at the baseline state because it prevents the cotransporter from trafficking from the biosynthetic pathway to the cell surface.^{306–308} In contrast to this inhibitory effect on NCC traffic, it has been hypothesized that under certain physiologic states that favor NaCl reabsorption, WNK4 stimulates NCC phosphorylation. Because NCC is structurally similar to NKCC2, it is activated by the serine-threonine kinases SPAK and OSR1 (see previous section). WNK4 and WNK1 activate SPAK and OSR1, which then can

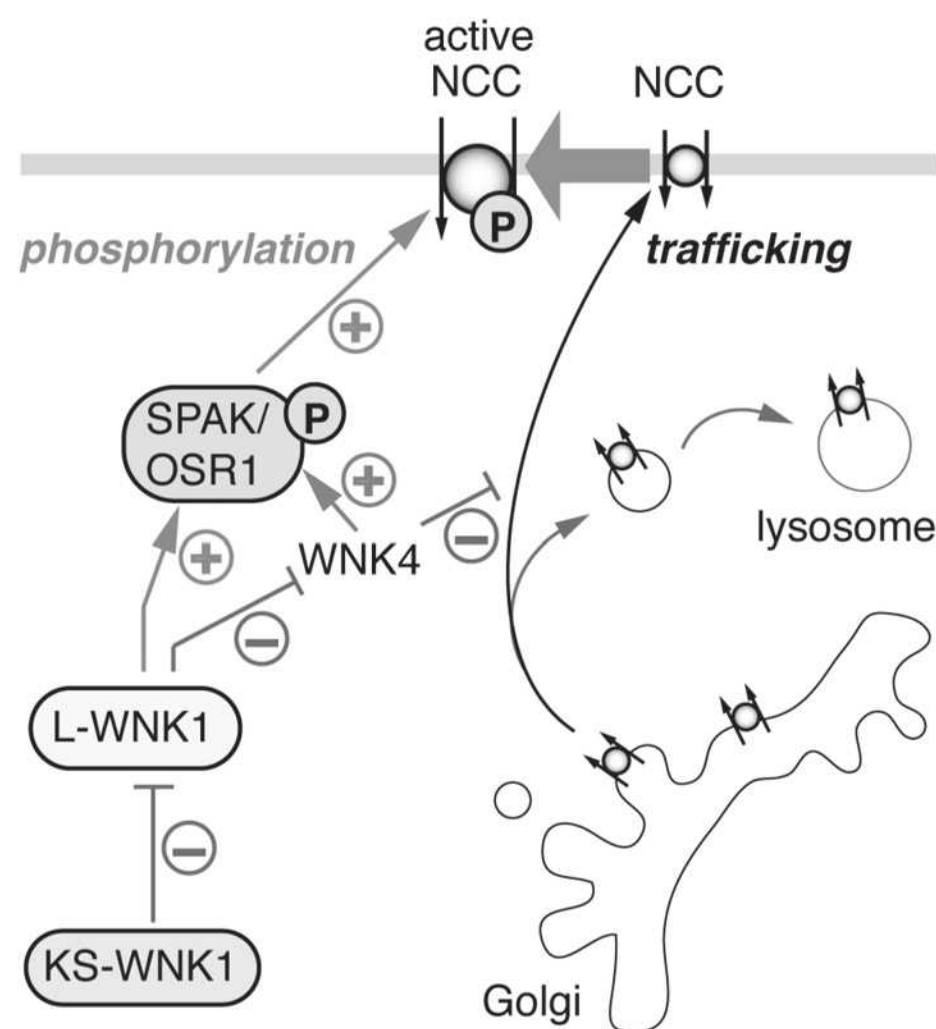


FIGURE 5.9 A model of thiazide-sensitive cotransporter (NCC) regulation by the with-no-lysine SPS1-related proline alanine-rich kinase/oxidative stress responsive kinase 1 (WNK-SPAK/OSR1) signaling pathway. With-No-Lysine (WNK) kinases regulate NCC trafficking and phosphorylation. *L-WNK1*, full-length kinase-active (“long”) WNK1; *KS-WNK1*, short kinase-defective “kidney-specific” WNK1.

bind to NCC and stimulate its activity by phosphorylating a cluster of serines and threonines in its NH_2 -terminus.³⁰⁷ In contrast to the TAL, only the full-length kinase active SPAK is expressed in the DCT (i.e., no inhibitory KS-SPAK isoforms are expressed).²³⁷ Due to the observations that WNK4 inhibits NCC under some experimental conditions and stimulates it under others, several have proposed that the WNK signaling pathway acts as a “switch” that converts the DCT from a salt-wasting to a salt-reabsorptive segment, although more work is needed to confirm this hypothesis.^{309–311} In FHHt, missense mutations of WNK4 stimulate NCC activity through two mechanisms. First, mutant WNK4 is incapable of reversing the inhibitory effect on NCC at the baseline; this results in increased NCC surface expression.^{308,312} Second, the FHHt-associated mutant increases NCC phosphorylation, presumably through enhanced SPAK/OSR1 activity. Thus, through incompletely defined mechanisms, mutant WNK4 appears to “lock” the DCT into a high Na^+ reabsorptive state by increasing the total number of active NCC molecules at the cell surface, resulting in thiazide-sensitive hypertension.³¹³

In contrast, WNK1 appears to influence NCC traffic indirectly by suppressing the inhibitory effect of WNK4 (Fig. 5.9). This effect requires intact WNK1 kinase activity and releases NCC from intracellular retention, thereby facilitating NCC delivery to the cell surface.^{307,308} Unlike WNK4, WNK1 mutations that cause FHHt do not alter the kinase’s coding sequence. Rather, WNK1 mutations are large intronic deletions that increase the mRNA abundance of WNK1 isoforms.³⁰⁵ Kinase-active WNK1 is capable of phosphorylating and activating SPAK and OSR1.³¹⁴

Thus, the WNK1 gene mutations that cause FHHt presumably do so by reversing the inhibitory effect of WNK4 on NCC traffic and by stimulating NCC phosphorylation.³¹⁵ As with SPAK gene expression in the TAL (see previous), long and short WNK1 isoforms are expressed in the DCT. The long form of WNK1 (*L-WNK1*) is kinase active and stimulates NCC activity and trafficking through the mechanisms described previously. A short kidney-specific kinase-defective product (*KS-WNK1*), in contrast, suppresses *L-WNK1* activity (Fig. 5.9).³¹⁶ The balance of these two isoforms has been postulated to act as a “switch” that controls overall *L-WNK1* kinase activity in the DCT, which should in turn regulate NCC activity.³¹⁰ Consistent with this idea, selective knockout of *KS-WNK1* expression increases NCC activity, whereas overexpression of the kidney-specific isoform in *KS-WNK1* transgenic mice causes salt-wasting through NCC inhibition.³¹⁷

The Basolateral Electrogenic Na^+ Pump. In both subsegments, the basolateral extrusion of Na^+ from the cytosol into the peritubular space (and eventually, the plasma) occurs via the electrogenic Na^+, K^+ -ATPase. The activity of this pump results in the generation of a constant transepithelial voltage across the basolateral membrane of -60 to -90 mV.^{318,319} A reduction in the luminal sodium concentration causes V_{bl} to depolarize, whereas increases in the sodium concentration hyperpolarizes V_{bl} . In addition, V_{bl} depolarizes after ouabain treatment.³²⁰ These observations are consistent with the notion that apical Na^+ entry stimulates the electrogenic Na^+, K^+ -ATPase system in the basolateral membrane.

Basolateral K^+ Efflux. Recent insights from rare Mendelian diseases highlight the importance of basolateral K^+ transport in tubular sodium transport in the distal nephron. In 2008, two groups identified that patients with mutations in the K^+ channel gene *KCNJ10* (*Kir4.1*) develop hereditary salt wasting.^{321,322} Patients with these mutations develop a complex constellation of neurologic defects in addition to the renal salt wasting. The disease has been named EAST syndrome (epilepsy, ataxia, sensorineural deafness, and tubulopathy) by some investigators and SeSAME syndrome (seizures, sensorineural deafness, ataxia, mental retardation, and electrolyte imbalance) by others (Table 5.1). The salt-wasting phenotype in these patients is reminiscent of Gitelman syndrome, suggesting that the disorder results in an impairment of salt transport in the DCT. Indeed, *Kir4.1* is expressed on the basolateral membrane of DCT cells, and patients with EAST/SeSAME syndrome mutations develop markedly reduced infoldings of the DCT basolateral membrane. This deficiency in total basolateral membrane content results in a decrease in the number of surface Na^+, K^+ -ATPase molecules, resulting in decreased sodium pump activity and impaired salt reabsorptive capacity.³²³ Additionally, it is thought that loss-of-function mutations of *Kir4.1* impair K^+ recycling across the basolateral membrane, which reduces the efficacy of Na^+, K^+ -ATPase (Fig. 5.8).³²¹

Basolateral Cl^- Transport. As mentioned previously, ClC-Kb channels and barttin are both expressed in the DCT, where they mediate basolateral Cl^- reabsorption.¹⁷² Because the apical transport mechanism in the DCT via NCC occurs through the coupled reabsorption of Na^+ and Cl^- , basolateral Cl^- transport is essential to the development of a gradient for Na^+ entry and to reduce the intracellular Cl^- concentration, which stimulates SPAK/OSR1-mediated NCC phosphorylation.³²⁴ Mutations in either ClC-Kb or the accessory subunit result in impaired distal salt reabsorption. Because barttin is required for adequate basolateral Cl^- reabsorption in the loop of Henle and DCT (see previous), patients with barttin mutations develop a more severe form of Bartter syndrome.¹⁷²

Apical Conductive Na^+ Channels. The entry of sodium into the *Amphiuma* distal tubular cell³²⁵ and late rat DCT cell³²⁶ is inhibited by amiloride, a Na^+ channel blocker. A Na^+ channel in the apical membrane would serve to depolarize the membrane and create the observed lumen-negative transepithelial potential. This transepithelial voltage, in turn, is a driving force for passive Cl^- reabsorption. Na^+ channel subunits have been found by immunolocalization in the late DCT in mouse and rat kidney,^{327,328} but not the human kidney.³²⁹

The Regulation of NaCl transport in the Distal Convoluted Tubule

Na^+ Delivery. NaCl reabsorption in the DCT is dependent on the delivered load of NaCl.²⁸⁶ The DCT responds to chronic increases in the delivery of NaCl with an increase in the capacity for NaCl transport,³³⁰ as well as marked ultrastructural changes in the DCT cell. These morphologic changes include an increase in the size of the DCT cell, an increase in the basolateral membrane surface area, and an increase in the size of mitochondria.³³⁰ Accompanying the functional and morphologic changes are an increase in Na^+, K^+ -ATPase activity and an increase in thiazide-binding sites.³³¹ These effects appear to result from an increase in Na^+ entry into the DCT cell rather than the increase in distal NaCl delivery or changes in plasma aldosterone or ADH levels that occur with chronic furosemide treatment. Inhibition of NaCl entry into DCT cells with chronic thiazide treatment resulted in a loss of cell height, loss of normal polarity, and apoptosis of the DCT cells.³³² The cellular mechanisms whereby NaCl entry affects transport function and morphology are not known.

Dietary Na^+ . Studies in rats and rabbits have yielded conflicting results regarding the effects of increased dietary Na^+ on DCT morphology and Na^+ transport. In rats, no consistent effect of a high Na^+ diet on either cell morphology, transport rates, or thiazide-receptor density could be demonstrated.^{330,331,333} Rats fed a low Na^+ diet²⁷⁹ or treated with thiazide diuretics,²⁸¹ however, demonstrated

a significant increase in NCC expression, which may have been because of the stimulation of mineralocorticoid secretion.³³⁴ In contrast, rabbits fed a high Na^+ diet developed an increased rate of DCT Na^+ reabsorption and an increase in Na^+, K^+ -ATPase activity.³³⁵

Steroid Hormones. Currently available evidence suggests that adrenal steroid hormones regulate Na^+ transport in the DCT. The presence of both mineralocorticoid and glucocorticoid receptors in the DCT has been demonstrated by immunohistochemistry and by hormone binding.^{283,336,337} In addition, an adrenalectomy resulted in a decrease in Na^+, K^+ -ATPase activity in the DCT.³³⁸ The Na^+, K^+ -ATPase activity could be restored by replacement doses of glucocorticoids, but not by mineralocorticoids.^{338,339} Microperfusion studies of superficial distal tubules (containing both DCT and CNT), however, demonstrated an increase in Na^+ transport in animals receiving aldosterone infusions.^{340,341} Both the thiazide-sensitive and the thiazide-insensitive components of Na^+ transport were increased by aldosterone.³⁴¹ The former may reflect neutral NaCl cotransport in the DCT, whereas the latter reflects electrogenic Na^+ absorption in the late DCT or CNT. Aldosterone infusion also resulted in an increase in thiazide-binding sites in the renal cortex, as determined by [^3H]metolazone binding, an increase in the natriuretic response to thiazide diuretics, and a large increase in NCC protein.³⁴¹⁻³⁴³ These findings establish NCC as an aldosterone-regulated transporter. By combining immunohistochemical and in situ hybridization techniques, Bostanjoglo et al.²⁸³ determined that DCT cells coexpress NCC, mineralocorticoid receptors, and 11 β -hydroxysteroid dehydrogenase type 2, an enzyme typically found in mineralocorticoid target sites. Thus, DCT cells, particularly those in the late portions of the DCT, express the key elements required for selective mineralocorticoid actions.

In 1998, Kim et al.³⁴³ showed that aldosterone increases the total protein abundance of NCC. This effect did not correlate with changes in NCC mRNA abundance, suggesting that aldosterone regulates NCC abundance by posttranslational mechanisms. Recent work indicates that at least two molecular mechanisms account for this effect. Aldosterone rapidly induces and activates the serine-threonine kinase serum and glucocorticoid-regulated kinase 1 (SGK1), which directly phosphorylates and inactivates two inhibitors of NCC trafficking, WNK4 (described previously), and neural precursor cell-derived, developmentally downregulated 4-2 (Nedd4-2), an E3 ubiquitin ligase.^{344,345} SGK1-mediated phosphorylation of two residues located at the C-terminus of WNK4 releases NCC from inhibition. This results in WNK4 inactivation and diverts the cotransporter away from intracellular degradation pathways, allowing NCC to traffic directly to the plasma membrane from the biosynthetic pathway.³⁴⁴ By phosphorylating Nedd4-2, SGK1 suppresses the ability of the E3 ligase to attach ubiquitin molecules to NCC; this

reduces NCC degradation and increases its plasma membrane expression.³⁴⁵ Both of these effects provide an explanation for the aldosterone-induced increase in NCC total protein abundance that was initially seen by Kim et al.³⁴³ Aldosterone also appears to stimulate NCC transport activity by promoting NCC phosphorylation, an effect that occurs independently of the effects of mineralocorticoids on NCC trafficking.^{346,347} The mechanism by which this effect occurs remains unclear but appears to correlate directly with increased SPAK/OSR1 activity, suggesting that additional connections between aldosterone and the WNK/SPAK/OSR1 signaling pathway may exist.

As mentioned previously, glucocorticoids increase Na^+, K^+ -ATPase activity following an adrenalectomy in the DCT.^{338,339} This effect was not blocked by spironolactone, a mineralocorticoid receptor antagonist, suggesting that glucocorticoids were acting via glucocorticoid receptors rather than mineralocorticoid receptors.³³⁹ In addition, dexamethasone infusions increased thiazide-sensitive NaCl transport and [^3H]metolazone binding sites in adrenalectomized rats.^{341,342} Nevertheless, the role of glucocorticoids in the physiologic regulation of Na^+ transport in the DCT remains unclear.

Gonadal steroid hormones may also influence NaCl transport in the DCT. Chen et al.³⁴⁸ reported gender differences in the density of thiazide receptors and in the natriuretic response to thiazides in rats. Female rats had higher levels of thiazide-binding sites in the renal cortex than males. The levels in females fell following ovariectomy, whereas levels rose in males following orchiectomy. Moreover, the increase in urinary Na^+ excretion caused by thiazides was greater in females than in males, suggesting that the differences in thiazide-binding sites were reflective of differences in thiazide-sensitive salt transport in vivo. Likewise, using antibodies against NCC, Verlander et al.³⁴⁹ found that estrogen treatment increased NCC expression in the DCT. These results are consistent with the view that male sex hormones (e.g., testosterone) may downregulate NCC expression and salt transport, whereas estrogens increase NCC expression and salt transport in the DCT. The authors are not aware of gender differences in the response of humans to thiazide diuretics.

Vasopressin. Recent work indicates that vasopressin is a potent activator of the thiazide-sensitive cotransporter.^{350,351} Two groups have shown that the vasopressin analog deamino-Cys-1, d-Arg-8 vasopressin (dDAVP) stimulates increases in NCC abundance, trafficking to the plasma membrane, and activation through SPAK/OSR1-mediated phosphorylation of its amino terminus. This effect is probably dependent on cyclic AMP and protein kinase A (PKA), well-established intermediaries of vasopressin-dependent signaling.³⁵² The effect on NCC abundance may be a Nedd4-2-dependent process, because PKA can phosphorylate and inactivate Nedd4-2 through mechanisms similar to SGK1³⁵³; however, this hypothesis is yet to be tested.

Na^+ TRANSPORT IN THE CORTICAL COLLECTING DUCT

General Considerations

The transport processes in the collecting duct mediate final adjustments in urinary composition. The collecting duct is a major locus of action of mineralocorticoid hormones and plays a major role in K^+ homeostasis and acid–base balance. Quantitatively, it is a minor site of Na^+ absorption, reclaiming only about 2% to 4% of the filtered Na^+ load.³⁵⁴

Electrophysiologic Aspects

The transepithelial voltage in the CCD varies widely from +10 to −100 mV,^{295,355,356} which largely results from differences in mineralocorticoid levels at the time of measurement in the animals. The reported values for transepithelial resistance also vary widely depending on apical Na^+ and K^+ channel activities, as shown by the effects of luminal amiloride and barium, respectively, on transepithelial resistance.^{356,357} The basolateral membrane is conductive to K^+ and, at least in rabbits, Cl^- .³⁵⁷

Mechanisms of Salt Absorption in Collecting Ducts

A proposed model for Na^+ absorption and K^+ secretion in the CCD is presented in Figure 5.10. As indicated previously, apical membranes of CCD principal cells possess conductive pathways for Na^+ and K^+ .^{358,359} Na^+ enters principal cells through Na^+ channels in the apical membrane down its electrochemical gradient. Na^+ is then pumped across the basolateral membrane by the Na^+, K^+ -ATPase in exchange for K^+ . The Na^+ current across the apical membrane depolarizes

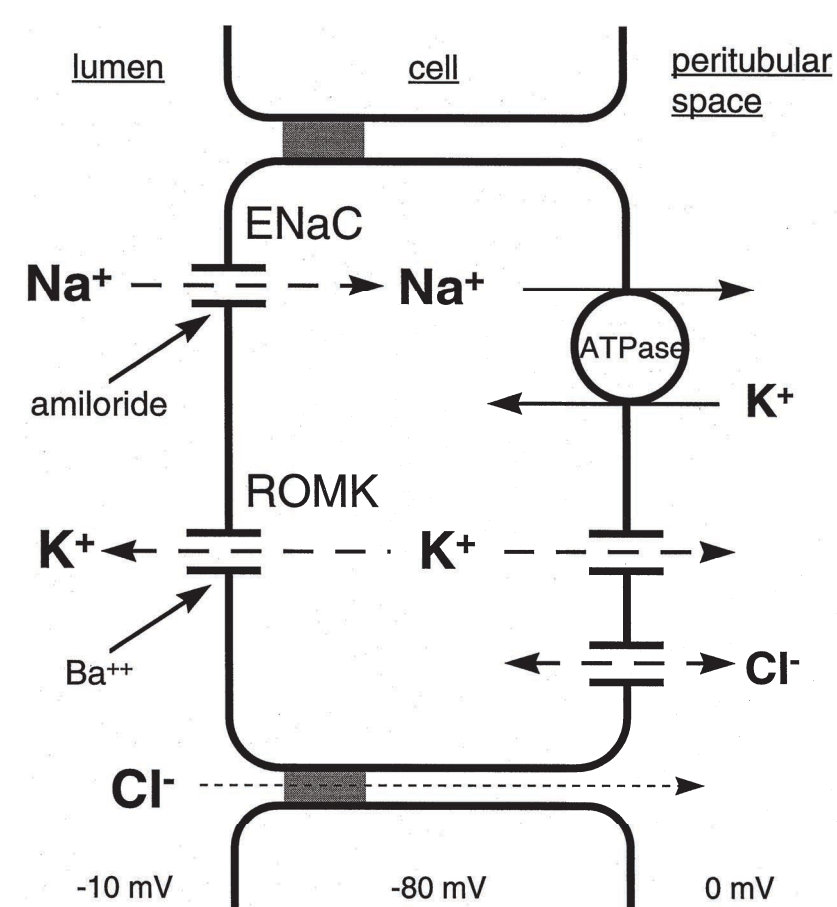


FIGURE 5.10 A model of salt transport by the principal cell of the cortical collecting duct. Apical Na^+ entry proceeds via amiloride-sensitive Na^+ channels (ENaC). Apical K^+ channels (ROMK) mediate K^+ secretion by this segment. Cl^- absorption is driven by the lumen-negative voltage through the paracellular pathway.

the cell, so that the cellular K^+ is above its equilibrium concentration, and thus leaves the cell through conductive pathways in the apical or basolateral membrane.³⁶⁰

Apical Na^+ Channels

Apical Na^+ entry depolarizes the apical membrane relative to the basolateral membrane, causing a lumen-negative transepithelial voltage, which in turn provides the driving force for Cl^- reabsorption through the paracellular pathway. Patch-clamp studies of collecting duct cells and urinary bladder cells provided some details regarding the electrophysiologic properties and regulation of apical membrane Na^+ channels.³⁶¹ The amiloride-sensitive epithelial Na^+ channel (ENaC) has been cloned,^{362,363} has a single-channel conductance to Na^+ of 4 to 5 pS, and is highly selective for Na^+ over K^+ ($P_{Na}/P_K > 20$). ENaC exhibits slow gating, with openings and closings lasting several seconds. Amiloride blocks ENaC at submicromolar concentrations.³⁶⁴

ENaC consists of three homologous subunits (α , β , and γ).³⁶³ The subunits share a common structure consisting of two transmembrane domains, intracellular N- and C-termini, and a large extracellular loop.^{365,366} Although the α subunit of ENaC can form Na^+ channels on its own,³⁶² coexpression of all three subunits dramatically increases the membrane Na^+ conductance.³⁶³ Based on homology to the related acid-sensing ion channel (ASIC1), the crystal structure of which was recently solved,³⁶⁷ native ENaC likely exists as an α , β , and γ heterotrimer. The single channel properties of the expressed channel closely resemble those of the 4 to 5 pS highly selective channel studied in native tissues.³⁶³ The large extracellular domains of the α and γ subunits can get proteolytically cleaved by furin in the biosynthetic pathway.^{368,369} Other proteases like prostatic and plasmin may cleave the γ subunit.³⁷⁰ These cleavage events release small inhibitory fragments of the α and γ subunits, causing an increase in the open probability of ENaC.³⁷⁰ ENaC activity is also enhanced by increasing luminal flow rates, as the extracellular domain of ENaC subunits respond to laminar shear stress.^{371,372} Thus, collecting duct Na^+ absorption is enhanced with increased distal fluid delivery via ENaC, a mechanosensitive channel.³⁷³

The key importance of ENaC for Na^+ reabsorption in the collecting duct is highlighted by naturally occurring mutations in ENaC that are responsible for Liddle syndrome, an autosomal dominant form of hypertension, and pseudohypoaldosteronism type 1, an autosomal recessive form of salt wasting (see Table 5.1).³⁷⁴ In Liddle syndrome, mutations in the ENaC subunits result in an increase in amiloride-sensitive Na^+ channel activity with a consequent increase in sodium reabsorption and volume-mediated hypertension.^{375–377} Most Liddle mutations occur in the cytoplasmic C-termini of the γ (SCNN1G) and β (SCNN1B) subunits.³⁷⁵ These mutations affect a conserved PY motif in the C-terminus, which is necessary for interaction with the ubiquitin ligase Nedd4-2 and subsequent internalization and degradation of ENaC.³⁷⁸ Liddle syndrome mutations

cause excessive ENaC accumulation at the apical plasma membrane, increasing Na^+ absorption. Pseudohypoaldosteronism type 1 (PHA-1), the clinical opposite of Liddle syndrome, is caused by homozygous inactivating mutations in the ENaC channel, resulting in a syndrome of Na^+ -wasting, hypotension, and hyperkalemia.³⁷⁹ The majority of mutations causing PHA-1 are frameshift or nonsense mutations that result in truncated, nonfunctional ENaC proteins.³⁷⁴

Apical Electroneutral Na^+ Transport

It was first observed in rats that a portion of Na^+ entry across the apical membrane in perfused CCD segments was sensitive to luminal hydrochlorothiazide, which inhibited Na^+ and Cl^- absorption without changing the transepithelial voltage.³⁸⁰ In addition, amiloride inhibited Na^+ transport in this segment by only 50%, and the effects of amiloride and hydrochlorothiazide were additive. It was thus concluded that the CCD may possess two parallel transport pathways for Na^+ : an electrogenic pathway involving amiloride-sensitive Na^+ channels and a thiazide-sensitive neutral NaCl cotransport pathway.³⁸⁰ More recently, a Na^+ -dependent Cl^-/HCO_3^- exchanger (NDCBE/SLC4A8) was identified in intercalated cells and found to mediate the amiloride-resistant, thiazide-sensitive electroneutral Na^+ reabsorption in the CCDs of mice.³⁸¹ That NCC knockout mice also exhibited significant natriuresis following treatment with thiazide diuretics suggests the importance of this pathway. The mechanism for net NaCl reabsorption in this setting involves the parallel activity of the NDCBE and Na^+ -independent Cl^-/HCO_3^- exchange via pendrin/SLC26A4 at the apical membrane of type B intercalated cells. The basolateral reabsorption pathways for the Na^+ and Cl^- that enter at the apical membrane in this model are the Na^+, K^+ -ATPase and ClC-K Cl^- channels. Thus, a novel target of thiazide diuretics distinct from NCC has been identified in the collecting duct with the implication that Cl^- transport by intercalated cells plays an important role in blood pressure regulation.³⁸² Because pendrin is a key mediator of bicarbonate secretion, and thus acid-base regulation, these findings also suggest a strong link between acid-base and volume/blood pressure regulation.

Control of Na^+ Absorption in the Cortical Collecting Duct

Aldosterone

Aldosterone is one of the key regulators of Na^+ transport in the collecting duct, where it increases the rates of Na^+ absorption and K^+ secretion.^{383,384} The major target site for mineralocorticoid effects is the principal cell of the CCD, although actions in the DCT have also been documented (see previous). Mineralocorticoid effects are produced by the binding of either mineralocorticoids or glucocorticoids³⁸⁵ to mineralocorticoid receptors found predominantly in the CCD.^{336,337} In addition, the binding of glucocorticoids to glucocorticoid receptors can also produce mineralocorticoid responses.³⁸⁵ This lack of specificity results from two

factors: first, mineralocorticoid receptors do not discriminate between aldosterone and glucocorticoids,³⁸⁶ so that either class of steroids can bind to and activate the receptor. Second, the DNA binding domains of mineralocorticoid and glucocorticoid receptors are highly conserved such that both receptors can activate many of the same genes.³⁸⁷ The specificity for mineralocorticoids *in vivo* is provided by the selective degradation of glucocorticoids, but not mineralocorticoids, by the enzyme 11 β -hydroxysteroid dehydrogenase.³⁸⁶ 11 β -Hydroxysteroid dehydrogenase activity is high in CCD segments.^{388,389} Illustrating the important role of this enzyme in regulating access of hormones to the mineralocorticoid receptor, genetic deficiency of this enzyme produces a syndrome, apparent mineralocorticoid excess, resembling hyperaldosteronism (hypertension, hypokalemia, metabolic alkalosis), except that aldosterone levels are low.³⁹⁰ The clinical manifestations result from the stimulation of mineralocorticoid receptors by circulating glucocorticoids.

In isolated perfused CCDs from mineralocorticoid-treated rabbits, there is an increase in both the Na^+ and K^+ conductance of the apical membrane and an increase in basolateral Na^+, K^+ -ATPase activity.³⁵⁶ The functional changes after aldosterone treatment are accompanied by morphologic changes in principal cells. The basolateral membrane length of principal cells falls by 35% after an adrenalectomy, and the administration of aldosterone, but not dexamethasone, restores the membrane length to control levels.³⁹¹

An early effect of aldosterone, occurring within a few hours of exposure, is an increase in the sodium permeability of the apical membrane of the CCD principal cell. Results from electrophysiologic and immunologic studies support the view that the early aldosterone-induced increase in apical sodium permeability is because of the activation of quiescent Na^+ channels rather than the synthesis and/or the insertion of new Na^+ channels into the membrane.³⁹²⁻³⁹⁴ Several pathways have been implicated in the aldosterone-mediated increase in Na^+ channel activity. An important downstream mediator of aldosterone is the serum and glucocorticoid-regulated kinase SGK1.³⁹⁵ SGK1 is expressed in the thick ascending limb, DCT, connecting segment, and cortical collecting tubules.³⁹⁶ Aldosterone increases the transcription of SGK1 *in vitro*,³⁹⁵ although the effects of aldosterone on SGK1 protein expression *in vivo* appear to be minor.³⁹⁶ Coexpression of SGK1 with ENaC results in markedly greater sodium currents than seen with ENaC alone.^{395,397} SGK1 is required for the stimulation of sodium transport by aldosterone both *in vitro*³⁹⁷ and *in vivo*.³⁹⁸ SGK1 increases sodium currents by increasing cell surface expression of ENaC³⁹⁹ and also increasing the open probability of individual ENaC channels.⁴⁰⁰ The effect of SGK1 on cell surface expression of ENaC is largely mediated by Nedd4-2. Specifically, Nedd4-2 is a substrate for SGK1. Upon phosphorylation by SGK1, the affinity of Nedd4-2 for the PY domains of ENaC is diminished,⁴⁰¹ whereas Nedd4-2 binding to 14-3-3 scaffolding proteins is increased.^{402,403} Thus, 14-3-3 proteins act to sequester Nedd4-2 and prevent it from binding to ENaC.

As discussed earlier, the binding of Nedd4-2 to ENaC induces internalization and degradation of the channel.

Recently, additional aldosterone-induced proteins have been discovered, including the glucocorticoid-induced leucine zipper protein (GILZ1), the transcription of which is rapidly induced by aldosterone in collecting duct principal cells. GILZ1 stimulates ENaC-mediated Na^+ transport and ENaC surface expression by inhibiting Raf-1 in the extracellular signal-regulated kinase (ERK) signaling pathway.^{404,405} Soundararajan and colleagues⁴⁰⁶ have also identified another aldosterone-induced protein, CNK3 (connector enhancer of kinase suppressor of Ras 3), which also stimulates ENaC and appears to be a scaffolding protein. The emerging story is that a multiprotein complex exists in CCD cells that promotes context-specific aldosterone signal transduction to induce ENaC activity and apical Na^+ conductance. Specifically, aldosterone stimulates ENaC activity through synergistic aldosterone-dependent activation by SGK1 via Nedd4-2 and by GILZ1 via Raf-1 and CNK3.

Nongenomic effects of aldosterone (rapid effects that are mineralocorticoid receptor-independent) have also been described in many tissues, including the kidney.⁴⁰⁷ Elucidating the mechanisms of such effects is an area of active investigation and appears to include the mitogen-activated protein kinase (MAPK) pathway and the methylation of proteins and lipids. Aldosterone also increases the activity of a number of cellular methyltransferase enzymes.⁴⁰⁸ Moreover, the β subunit of ENaC itself is a substrate for methylation, and, when methylated, exhibits increased Na^+ transport activity.⁴⁰⁹ The effects of aldosterone on Na^+ channel activity may also involve small GTP-binding proteins, for example, Ras,⁴¹⁰ and phosphatidylinositol 3-kinase.⁴¹¹ SGK1 is a downstream mediator of phosphoinositide-3-kinase.⁴¹²

Sometime after the increase in apical membrane Na^+ conductance occurs, the basolateral membrane Na^+, K^+ -ATPase activity and pump current increase.⁴¹³ This increase is due initially to the effect of increased cell sodium activity on existing pump units.⁴¹⁴ Later, aldosterone induces the synthesis of additional Na^+, K^+ -ATPase pump subunits.⁴¹⁵ Increased apical membrane Na^+ entry may promote the late synthesis of Na^+, K^+ -ATPase, because the inhibition of sodium entry by amiloride markedly reduced the aldosterone-induced increase in Na^+, K^+ -ATPase activity.⁴¹⁶

Aldosterone also exerts an additional, late effect on the amiloride-sensitive Na^+ conductance. Patch-clamp studies in rats exposed to high levels of aldosterone for several days demonstrated a large increase in the amiloride-sensitive whole cell Na^+ conductance,⁴¹⁷ which correlated with an increase in the number of active Na^+ channels in the apical membrane.⁴¹⁸ Increases in ENaC mRNA⁴¹⁹ and protein³³⁴ levels in aldosterone-treated tissues suggest that the synthesis of new ENaC channels may contribute to the late aldosterone-induced increase in Na^+ conductance. SGK1, in addition to increasing the cell surface expression of ENaC, also regulates the transcription of ENaC subunits, principally α .⁴²⁰

Antidiuretic Hormone (ADH)

Exposure of rat CCD segments *in vitro* to ADH results in a sustained stimulation of Na^+ absorption.⁴²¹ ADH increases the transepithelial potential, depolarizes the apical membrane, and increases the conductance of the apical membrane of principal cells.⁴²² These changes are entirely reversed by luminal amiloride, indicating that ADH increases the apical membrane sodium conductance of the principal cell. These effects of ADH are mediated intracellularly by cAMP.⁴²³ Moreover, the effects of ADH on Na^+ transport in the rat CCD are enhanced by prior treatment of the animals with mineralocorticoids.⁴²¹

ADH and aldosterone increase apical membrane ENaC activity through both overlapping and distinct mechanisms of action. Like aldosterone via SGK1, ADH induces enhanced PKA-dependent phosphorylation of Nedd4-2 at phosphorylation sites that overlap those of SGK1.³⁵³ As described previously, these phosphorylation events promote sequestration of Nedd4-2 through enhanced binding to 14-3-3 proteins. In contrast to aldosterone, which can activate quiescent channels, ADH, via cAMP, promotes the insertion of additional Na^+ channels into the apical membrane.⁴²⁴ In addition, PKA-mediated phosphorylation may also directly stimulate the activity of ENaC channels already present in the apical membrane, potentially both directly⁴²⁵ and indirectly via Nedd4-2 inhibition, causing enhanced channel residency time at the membrane and thus proteolytic cleavage.³⁷⁰ Long-term exposure to ADH may also increase the capacity of the collecting duct for Na^+ transport by increasing the expression of both Na^+, K^+ -ATPase and ENaC.⁴²⁶

Other Agents

Bradykinin is produced in the connecting duct⁴²⁷ and binds to specific receptors in the CCD.⁴²⁸ An intrarenal infusion of bradykinin produces a diuresis. Bradykinin has been reported to reduce Na^+ absorption in rat CCD segments.⁴²⁹ More recently, bradykinin was shown to acutely and reversibly decrease the open probability of ENaC in patch-clamp studies performed *ex vivo* on split open CCDs isolated from rats. This effect appears to be mediated specifically through the bradykinin B_2 receptor.⁴³⁰

As will be discussed in the following section, the major site of action of atrial natriuretic peptide (ANP) is the inner medullary collecting duct. ANP stimulates cGMP production in the CCD,⁴³¹ and inhibits the hydro-osmotic actions of ADH in the CCD.⁴³² Biphasic effects of ANP on conscious, sedated rats have been observed with acute water and sodium excretion within minutes, followed by retention after 90 minutes, which was associated with increased apical membrane expression of α - and γ -ENaC. These delayed effects may represent a compensatory response to increase sodium and water reabsorption and prevent volume depletion in response to prolonged ANP infusion.⁴³³

α 2-Adrenergic agonists inhibit sodium reabsorption in the rat CCD.⁴³⁴ This inhibition is associated with an increase

in the apical membrane resistance, reflecting decreased Na^+ entry through Na^+ channels. These changes appear to result from an inhibition of adenylate cyclase and antagonism of ADH by α 2 agonists.⁴³⁵

Prostaglandin E_2 exerts diuretic and natriuretic effects on the kidney. Part of this action is mediated by an inhibition of Na^+ absorption in the CCD.⁴³⁶ The application of PGE_2 to the basolateral surface of perfused rabbit CCD segments reversibly inhibits the negative transepithelial voltage and net sodium absorption.⁴³⁶ The effect of PGE_2 on sodium transport is coupled to a rise in intracellular $[\text{Ca}^{2+}]$ and is dependent on the activation of PKC.⁴³⁷ Four PGE_2 receptor subtypes, designated EP1, EP2, EP3, and EP4, have been characterized. Studies using receptor subtype-specific agonists and antagonists suggest that the EP1 receptor mediates PGE_2 -dependent inhibition of Na^+ transport in the CCD.⁴³⁸ This inhibition of Na^+ transport is associated with depolarization of the apical membrane voltage, consistent with inhibition of the basolateral Na^+, K^+ -ATPase.⁴³⁹ In contrast to the inhibitory effect of basolateral PGE_2 on Na^+ transport, luminal PGE_2 increases transepithelial voltage and presumably Na^+ absorption via the EP4 receptor.⁴⁴⁰

Epidermal growth factor (EGF) reduces Na^+ absorption in rabbit CCD by about 50% via the inhibition of apical electrogenic Na^+ entry via ENaC.^{441,442} More recent studies performed on cultured mouse CCD cells suggest that the effects are mediated by the ErbB2 EGF receptor and that there is actually a biphasic effect of EGF on ENaC activity.⁴⁴³ Acutely (<4 h), EGF treatment increases the ENaC current, an effect that appears to be mediated via the PI-3-kinase pathway. Chronically (>8 h), the ENaC current was inhibited via effects through the MEK/ERK pathway.⁴⁴³

Endothelin-1 (ET-1) is heavily secreted from the basolateral aspect of CD epithelial cells where it binds to basolateral ET_B receptors and may thus act in an autocrine fashion to inhibit both Na^+ and water reabsorption in this segment, as shown *in vitro*.⁴⁴⁴ The physiologic relevance of these findings has been confirmed *in vivo*, because mice with CD-specific knockout of ET-1 or ET receptors are hypertensive on a normal Na^+ diet and have exacerbated Na^+ retention and hypertension when placed on a high- Na^+ diet. ET-1 inhibits ENaC activity through Src- and MAPK-dependent pathways.⁴⁴⁴

Nitric oxide is also a downstream mediator of ET-1,⁴⁴⁴ where it decreases Na^+ transport in rat CCD segments by 40% to 80%.⁴⁴⁵ The addition of nitric oxide to tubules decreased the intracellular $[\text{Na}^+]$, but did not affect the activity of basolateral Na^+, K^+ -ATPase, suggesting that the primary effect of nitric oxide is to inhibit apical Na^+ entry via ENaC. The inhibitory mechanism involving NO remains to be determined.⁴⁴⁴

Dopamine inhibits ADH-dependent Na^+ transport and transepithelial voltage in rat and rabbit CCD segments, although the particular basolateral dopamine receptor subtype may be species specific.^{446,447}

Ang II stimulates sodium channel activity in rabbit and mouse CCDs.⁴⁴⁸ Chronic Ang II infusion also increases the abundance of the α subunit of ENaC, the rate-limiting subunit for ENaC assembly.⁴⁴⁹ Both the acute effects of Ang II on ENaC activity and the chronic effects on ENaC abundance are mediated by the AT1 receptor.^{448,449} A more recent study performed on rats suggests that the Ang II-induced ENaC stimulation downstream of the AT1 receptor involves a Ca^{2+} -independent PKC pathway that induces superoxide generation. The blocking of arachidonic acid-induced inhibition of ENaC may also play a role.⁴⁵⁰

Bicarbonate (HCO_3^-) changes in the CCD lumen, which may reflect changes in total-body acid-base status or local HCO_3^- secretion from neighboring intercalated cells via the apical Na^+ -independent $\text{Cl}^-/\text{HCO}_3^-$ exchanger, pendrin, have been shown recently to modulate Na^+ reabsorption in the CCD.^{451,452} Specifically, the bicarbonate-stimulated soluble adenylyl cyclase (sAC) appears to stimulate both basal and agonist-stimulated Na^+ reabsorption in the kidney collecting duct, acting to enhance Na^+, K^+ -ATPase catalytic activity.⁴⁵¹ Another study suggests that pendrin increases ENaC abundance and activity, at least in part by increasing luminal $[\text{HCO}_3^-]$.⁴⁵²

The elucidation of other cellular signaling pathways and kinases that are important in the regulation of ENaC, and Na^+ transport in the CCD is a very active area of research and involves signaling pathways downstream of diverse stimuli, including inflammation (IkappaB kinase/NF-kappaB/NF- κ B pathway)⁴⁵³ and metabolic stress (AMP-activated protein kinase).⁴⁵⁴ A full treatment of these signaling pathways is beyond the scope of this chapter, but the interested reader is referred to a recent review.⁴⁵⁵

NA⁺ TRANSPORT IN THE OUTER MEDULLARY COLLECTING DUCT

The transport properties of the outer medullary collecting duct (OMCD) have been studied by in vitro perfusion of isolated tubule segments. The functional properties of the OMCD differ depending on the location of the segment within the outer medulla. Segments within the outer stripe of the outer medulla (OMCD_o) exhibit electrophysiologic properties resembling the cortical collecting duct; that is, a lumen negative transepithelial voltage and electrogenic apical Na^+ entry.⁴⁵⁶ Compared to the CCD, the OMCD_o displays a less negative transepithelial voltage, much lower ionic permeabilities, and a lower rate of active reabsorption of Na^+ .⁴⁵⁷ As the collecting duct descends into the medulla, principal cells, which mediate Na^+ and K^+ transport in the CCD (see previous), are replaced by cells the electrical properties of which are similar to intercalated cells of the CCD (i.e., the apical membrane lacks a demonstrable Na^+ or K^+ conductance).⁴⁵⁸ Within the inner stripe of outer medulla (OMCD_i), principal cells are virtually absent and no net Na^+ absorption occurs.⁴⁵⁷

NA⁺ TRANSPORT IN THE INNER MEDULLARY COLLECTING DUCT

Mechanism of Na⁺ Transport

The analysis of salt transport by the inner medullary collecting duct (IMCD) has been confounded by problems of axial tubule heterogeneity, species variability, and differences in experimental approach. Based on morphologic factors, the IMCD has been divided into three subsegments: IMCD₁, IMCD₂, and IMCD₃.⁴⁵⁹ This morphologic heterogeneity is paralleled, to some extent, by functional heterogeneity. For example, the urea permeability and its responsiveness to ADH increases from IMCD₁ to IMCD₃ with increasing medullary depth.⁴⁶⁰ Further complicating the analysis is the observation that similar subsegments from different species exhibit different properties relating to salt transport.^{461,462} Finally, for unclear reasons, studies examining IMCD function in vivo (e.g., by microcatheterization)⁴⁶³ have yielded markedly different results than have in vitro studies of isolated perfused tubules.^{461,464}

Microelectrode impalement studies of IMCD segments from rats demonstrated that the apical membrane constituted the major cellular resistance, and the luminal application of amiloride increased the apical membrane voltage and resistance and decreased the transepithelial voltage.⁴⁶¹ These results, and others,⁴⁶⁵ are consistent with the presence of an amiloride-sensitive sodium conductance in the apical membrane of IMCD cells. Patch-clamp studies of cultured rat IMCD cells indicate that Na^+ entry is mediated by a 20 to 30 pS amiloride-sensitive, nonselective, cGMP-gated cation channel in the apical membrane.^{465,466} The basolateral membrane of IMCD cells contains the Na^+, K^+ -ATPase, a K^+ conductance and a HCO_3^- conductance.⁴⁶¹

The results discussed previously can be combined into a model for sodium transport in the IMCD (Fig. 5.11). Na^+

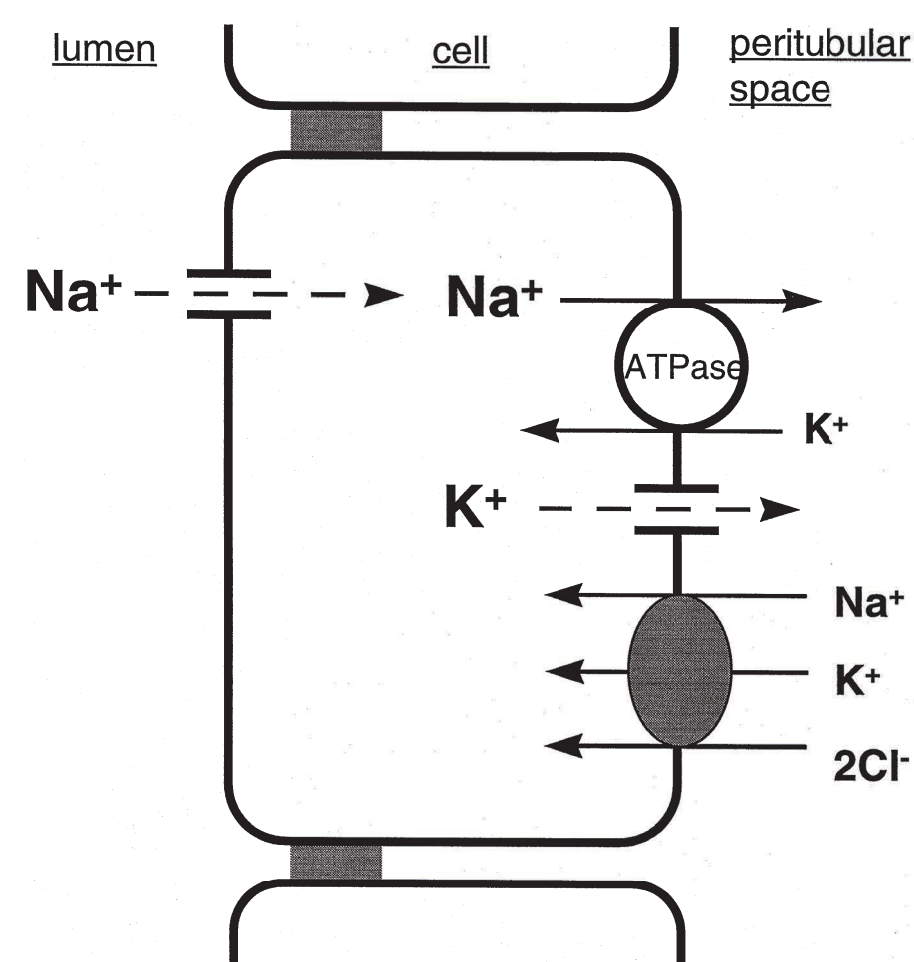


FIGURE 5.11 A model of Na^+ transport in the inner medullary collecting duct. Apical Na^+ entry proceeds via a nonselective amiloride-sensitive cation channel. Basolateral Na^+, K^+ -ATPase and $\text{Na}^+, \text{K}^+, 2\text{Cl}^-$ cotransport may be involved in Na^+ secretion.

entry across the apical membrane occurs down its steep electrochemical gradient through amiloride-sensitive Na^+ channels. Na^+ is extruded across the basolateral membrane by the Na^+, K^+ -ATPase. The basolateral membrane K^+ conductance serves to recycle the K^+ that enters via the Na^+, K^+ -ATPase. The K^+ conductance also hyperpolarizes the cell, thereby favoring Na^+ entry across the apical membrane.

There is some evidence for additional electroneutral Na^+ entry pathways in rat IMCD cells, although their importance in net Na^+ absorption by the IMCD in other species remains unclear. Furosemide and thiazide diuretics both inhibit a portion of Na^+ absorption by rat IMCD segments in vivo.^{467,468} However, Zeidel and colleagues⁴⁶⁹ failed to demonstrate sensitivity of rabbit IMCDs to loop diuretics and found that conductive Na^+ entry at the apical membrane predominated. A $\text{Na}^+ - \text{K}^+ - 2\text{Cl}^-$ cotransporter appears to be present in the basolateral membrane of terminal IMCD segments.⁴⁷⁰ The cloned $\text{Na}^+ - \text{K}^+ - 2\text{Cl}^-$ cotransporter from cultured IMCD cells represents the “secretory” isoform of the $\text{Na}^+ - \text{K}^+ - 2\text{Cl}^-$ cotransporter (NKCC1) rather than the absorptive isoform present in the apical membrane of the thick ascending limb.⁴⁷¹ Finally, the $\text{Na}^+ - \text{HCO}_3^-$ cotransporter NBCn1 has been found in the basolateral membrane of rat IMCDs, which may contribute to cellular defense against both acidification and volume changes in this segment.⁴⁷²

Regulation of Na^+ Transport

The IMCD appears to be the major target site for the potent diuretic hormone ANP.⁴⁷³ This hormone, working through cGMP,⁴⁷⁴ inhibits Na^+ entry via apical Na^+ channels. In cell-attached and excised patches of cultured inner medullary cells, both ANP and dibutyryl cGMP inhibited the activity of the cation channel.⁴⁷⁵

In addition to inhibiting Na^+ reabsorption, ANP may stimulate Na^+ secretion in the IMCD.⁴⁷⁰ In isolated perfused IMCD segments, ANP increased the bath to lumen flux rate of Na^+ and Cl^- , an effect that was inhibited by peritubular furosemide and by omission of either counterion, thus suggesting the role of a basolateral membrane $\text{Na}^+ - \text{K}^+ - 2\text{Cl}^-$ cotransporter.⁴⁷⁶ Sands et al.,⁴⁶⁴ however, found no effect of ANP on the Na^+ permeability of the rat IMCD.

In one study,⁴⁷⁷ ADH stimulated amiloride-sensitive Na^+ absorption by terminal IMCD segments perfused in vitro. Other studies, however, failed to find an effect of ADH or cAMP on the amiloride-sensitive cation channel in apical membranes of cultured IMCD cells.^{464,465}

Micropuncture studies by Ullrich and Papavassiliou⁴⁷⁸ demonstrated that mineralocorticoids increased net Na^+ absorption in the terminal IMCD, which was attributed to a decrease in the passive Na^+ permeability of the tubule leading to a decrease in back leak of NaCl into the lumen. Mineralocorticoids may also increase active Na^+ reabsorption in the IMCD. For example, aldosterone produced a three- to seven-fold stimulation of electrogenic Na^+ transport in cultured rat IMCD cells,⁴⁷⁹ and chronic mineralocorticoid exposure in vivo increased the activity of Na^+, K^+ ATPase in the IMCD.⁴⁸⁰

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