Potassium: From Physiology to Clinical Implications

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\section*{Potassium Homeostasis}

Potassium (K\textsuperscript{+}) is the most abundant intracellular cation, with 98\% of the total pool being located in the cells at a concentration of 140–150 mmol/l, and only 2\% in the extracellular fluid, where it ranges between 3.5 and 5 mmol/l. A fine regulation of the intracellular-extracellular gradient is crucial for life, as it is the main determinant of membrane voltage; in fact, acute changes of K\textsuperscript{+} plasma levels may have fatal consequences.

\section*{Summary}

An integrated system including an 'internal' and 'external' control prevents significant fluctuations of plasma levels in conditions of K\textsuperscript{+} loading and depletion. The internal control regulates the intra-extracellular shift, a temporary mechanism able to maintain a constant K\textsuperscript{+} plasma concentration without changing the total amount of body K\textsuperscript{+}. The external control is responsible for the excretion of the ingested K\textsuperscript{+}, and it has the kidney as the major player. The kidney excretes nearly 90\% of the daily intake. Along the proximal tubule and the thick ascending limb on Henle’s loop, the amount of K\textsuperscript{+} reabsorption is quite fixed (about 80–90\%); conversely, the distal nephron has the ability to adjust K\textsuperscript{+} excretion in accordance with homeostatic needs. The present review analyzes: (1) the main molecular mechanisms mediating K\textsuperscript{+} reabsorption and secretion along the nephron; (2) the pathophysiology of the principal K\textsuperscript{+} derangements due to renal dysfunction, and (3) the effect of ingested K\textsuperscript{+} on blood pressure and renal electrolyte handling. Key Messages: Maintaining plasma K\textsuperscript{+} levels in a tight range is crucial for life; thus, multiple factors are implicated in K\textsuperscript{+} homeostasis, including kidney function. Recent studies have suggested that K\textsuperscript{+} plasma levels, in turn, affect renal salt absorption in animal models; this effect may underlie the reduction of blood pressure observed in hypertensive subjects under K\textsuperscript{+} supplementation.

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(nearly 70 mEq) is located in the extracellular fluids (ECF), where it ranges from 3.5 to 5 mEq/l. The K⁺ gradient between the ICF and the ECF is based on the activity of the Na-K ATPase, located on the plasma membrane of all animal cells. As the intracellular K⁺ pool is much higher than the extracellular pool, clearly changes of total K⁺ content and/or distribution cause more dramatic fluctuations of extracellular rather than intracellular K⁺ concentration [K⁺]. However, several factors modulate K⁺ redistribution between the ICF and the ECF and its excretion in order to maintain the extracellular concentration in a tight range.

It is well known that an ingested K⁺ load does not result in a significant increase in plasma [K⁺]. Theoretically, an intake of 35 mEq of K⁺ would raise its plasma level by 2.5 mEq/l in humans if the distribution was totally extracellular. In contrast, only about a quarter of the ingested K⁺ remains in the ECF, as a large cellular storage reservoir (including muscle, liver and red blood cells) buffers plasma K⁺ upon its intake.

Both internal and external homeostasis contributes to maintaining a normal K⁺ balance, by controlling its distribution (the first) and its excretion (the second) in either conditions of K⁺ loading and K⁺ depletion (fig. 1).

The internal K⁺ homeostasis is mainly dependent on hormones such as insulin and catecholamines; in addition, acid base balance and plasma osmolality regulate cellular K⁺ uptake. This process is extremely efficient and guarantees constant plasma [K⁺] through the regulation of the Na⁺-K⁺ ATPase activity. The kidney is the main determinant of external K⁺ homeostasis, being responsible for the excretion of 90% of the K⁺ ingested, versus 10% eliminated via stools.

Renal Homeostasis of K⁺

A small fraction of plasmatic K⁺ ions may bind nonfilterable proteins, with a consequent restriction to glomerular filtration. Unbound K⁺ is freely filtered across the glomerulus, and the concentration of K in the glomerular filtrate is equal to the plasmatic concentration.

Clearance studies during the 1940s and the 1950s demonstrated that the bulk of tubular K⁺ reabsorption occurs along the proximal tubule (PT) and the thick ascending limb on Henle’s loop (TAL) [1] (fig. 2). In vivo micropuncture experiments in the 1960s showed that ~50% of the filtered load reaches the late PCT in the collections from tubular fluids [2], indicating that the PT reabsorbs the majority of the filtered load. The evidence of a strong association between the rate of fluid and K⁺ reabsorption suggests that, at least in part, K⁺ absorption occurs with a solvent drag mechanism. In addition, a paracellular diffusion from the lumen to the interstitium is driven by the difference of concentration (that is higher in the lumen upon fluid absorption) and by the favorable electrochemical gradient, especially along the latest part of the PT. The role of transcellular transport is partially known. The basolateral Na-K ATPase mediates the electrogenic transport of K⁺ from the interstitium to the cell, generating a high intracellular [K⁺] and a cell-positive electric potential. Along the final part of the PT (segment S3), which includes the fraction of the PT that enters the outer medulla, K⁺ is secreted into the luminal fluid. The presence of K⁺ channels has been demonstrated at both the basolateral and luminal sites. The K⁺ efflux to the lumen and to the interstitium generates a cell-negative gradient that provides the driving force for cation reabsorption. Immunofluorescence studies have demonstrated the presence of KCNE1 and KCNQ1 proteins on the brush-border membrane of PT, which colocalize to form an apical K⁺ channel. KCNE1 −/− mice show hypokalemia and volume depletion [3]. Micropuncture studies showed a strong reduction of K⁺ secretion along the PT. These channels are supposed to mediate membrane repolarization, as Na⁺ substrate absorption leads to membrane de-
polarization. Accordingly, a defective function of the KCNE-KCNQ1 channels, as in KCNE1−/− mice, affects solute absorption leading to osmotic polyuria. Luminal [K⁺] concentration at the tip of the Henle’s loop is 10-fold the plasma concentration.

Along the descending limb, K⁺ is delivered to the tubular fluid. It has been postulated that K⁺ ions are trapped in the medulla by the countercurrent mechanism after being actively absorbed along the TAL and passively enter the lumen along the descending limb. K⁺ reabsorption along the TAL is primary mediated by the Na-K-2Cl cotransporter (NKCC2), following the Na⁺-favorable gradient generated by the Na-K ATPase. The high intracellular [K⁺] favors its recycling back to the lumen through the apical K⁺ channels ROMK. Several lines of evidence demonstrate that K⁺ recycling through ROMK is crucial to guarantee Na-K-2Cl reabsorption; the uptake of these ions is dramatically reduced by either reducing luminal [K⁺] and by blocking ROMK activity [4]. At the basolateral level, the K⁺ efflux is mediated by the K-Cl and K-HCO₃⁻ cotransporters besides the K channels.

Only 10% of the filtered load reaches the distal nephron. The evidence that the amount of K⁺ excreted could exceed the amount of K⁺ filtered suggested the presence of K⁺ secretion along the nephron. The final K⁺ excretion (which is generally 10–20% of the filtered load) is mainly a function of distal secretion. The distal convolute tubule (DCT) has a minor role in K⁺ secretion compared with the connecting tubule (CNT) and the cortical collecting duct (CCD). Microperfusion studies in rats showed that the DCTs are able to secrete K⁺ through the K-Cl cotransporter [5].

At the basolateral level, the presence of the K⁺ channel KCNJ 10 kir4.1 has been shown to mediate K⁺ recycling to the interstitium, safeguarding the activity of the Na-K ATPase pump; genetic defects of this channel lead to salt wasting because of a reduced activity of the Na-K ATPase.

The CNT and collecting duct (CD) are the main segments mediating K⁺ secretion. Along these segments, the epithelium is constituted mainly of principal cells, which represent 70–75% of the total population; these cells mainly mediate Na reabsorption and K⁺ secretion. The latter requires several transmembrane proteins: first, the basolateral Na-K ATPase increasing the intracellular [K⁺] provides the gradient for K⁺ efflux; in addition, apical Na⁺ entry through the ENaC channels provides a favorable electrochemical gradient for K⁺ secretion; eventually, K⁺ exit to the lumen is mediated by the ROMK channels, BK (big potassium) channels and K-Cl cotransporter. Several lines of evidence demonstrate that BK channels mediate the bulk of K⁺ secretion along the CNT and CCD in response to a high urinary flux [6]. Those channels are relatively inactive under normal conditions but are activated by a high urinary flow.

**Fig. 2.** K⁺ reabsorption and secretion along the nephron.
A small fraction of epithelial cells along the distal nephron is represented by intercalated cells that are scattered among the principal cells. These cells are able to reabsorb K⁺ under certain conditions, such as K⁺ depletion. The α-intercalated cells show several isoforms of K-H ATPase. The latter is also expressed along the gastric and colonic mucosa, and it is inhibited by omeprazole. These transporters are activated by hypokalemia and may serve as a compensatory machinery mediating increased K⁺ reabsorption when needed.

As the rate of K⁺ reabsorption along the PT and the TAL is mainly fixed, the final excretion depends mainly on K secretion along the distal nephron. This process is under a fine regulation by several factors.

**Feedback Control of K⁺ Homeostasis**

The feedback control in physiology is, by definition, a regulatory mechanism of a biological function that maintains specific parameters (temperature, pH, electrolyte levels, etc.) in the optimal range through an internal control. The negative feedback implicates the sensation of a stimulus (high plasma [K⁺]), which in turn activates some effectors that finally return the parameter back to normal. Several hormones constitute the effectors of the feedback control of K⁺ homeostasis. A K⁺-rich meal induces the pancreatic secretion of insulin, which stimulates the Na-K pump, minimizing the increase in plasma [K⁺] by promoting K⁺ uptake into the muscle cells [7]. When K⁺ plasma levels increase enough, hyperkalemia induces aldosterone secretion. Aldosterone, in turn, promotes renal K⁺ secretion through the upregulation of both the basolateral Na-K ATPase and the luminal K⁺ channels along the principal cells. In addition, aldosterone enhances K⁺ secretion along the colon, an essential function especially in the presence of kidney disease. Conversely, when the extracellular K⁺ content is reduced because of a deficient intake or an increased urine excretion, K⁺ is released from the intracellular space to the plasma. In this condition, skeletal muscle becomes insulin resistant to K⁺ (but not glucose) uptake, even before plasma [K⁺] decreases. In addition, hypokalemia leads to the downregulation of the skeletal muscle Na-K ATPase isofrom 2, with the consequent leak of K⁺ to the plasma, and to the inhibition of aldosterone release, with the reduction of urinary K⁺ secretion.

**Feedforward Control of K⁺ Homeostasis**

A feedforward loop refers to a control system that responds to a specific stimulus in a predefined manner. Besides the feedback control, several lines of evidence suggest that K⁺ homeostasis is under a feedforward control that is induced when minimal changes in plasma [K⁺] occur. It has been established that the regulation of plasma aldosterone by K⁺ is activated when the plasma [K⁺] is higher or lower than normal, with little effect when it is within the physiological range. In animal models, it has been shown that after an oral K⁺ load, significant urinary K⁺ excretion occurs without changes in plasma aldosterone concentration. However, the existence of a kaliuretic reflex has been suggested arising from an unknown sensor in the splanchnic bed.

The K⁺ adaptation refers to the increased K⁺ secretion along the distal nephron after an ingested K⁺ load, without any significant change in aldosterone plasma levels. In animal models, this adaptation is mediated by an increased ROMK channel, Na-K ATPase and ENaC activity.

Glucagon has been proposed as a potential effector of the feedforward control. This hormone is secreted upon a protein-rich meal, and intraportal glucagon infusion produces significant increases in renal blood flow and the glomerular filtration rate. The feedforward regulation of K⁺ homeostasis may include several mechanisms: (1) insulin release that rapidly induces cellular K⁺ uptake into insulin-responsive tissues; (2) glucagon secretion that promotes renal K⁺ excretion after a protein-rich meal; (3) a yet-unidentified gut sensor that enhances renal K⁺ excretion after ingestion. When plasma [K⁺] increases beyond these layers of control, feedback regulation is activated.

**Hypokalemia: Lessons from the Salt-Losing Tubulopathies**

Genetic studies of inborn errors resulting in renal salt wasting have provided novel insights into the pathophysiology of renal electrolyte handling. Additional studies on engineered mice and cell culture have further highlighted...
the function of specific transport proteins and the molecular pathways regulating their activity.

Bartter (BS) and Gitelman (GS) syndromes are genetic disorders leading to hypokalemic metabolic alkalosis due to renal loss, with a compensatory activation of the RAAS axis. Genetic studies have shed light onto the molecular basis underlying these disorders that have long been considered two forms of the same disease (table 1).

**Bartter Syndrome**

BS is a rare tubular dysfunction caused by a genetic mutation leading to a defective NaCl absorption along the TAL. To date, mutations of five genes have been described. The defect determines fluid and electrolyte loss resulting in hypovolemia, with a compensatory activation of the renal epithelia, leading to hypokalemia and metabolic alkalosis.

**Genetics**

BS type I is caused by mutations of the SLC12A1 gene, encoding the kidney-specific Na-K-2Cl cotransporter NKCC2. BS type II depends on KCNJ1 mutations, leading to a defective function of the K⁺ channel ROMK. Immunostaining studies have shown an apical distribution of ROMK channels along the TAL, DCT and CD in rats, consistent with a role in K⁺ secretion along the renal epithelia. Along the TAL, ROMK channels mediate K⁺ efflux back to the lumen, favoring NaCl reabsorption by supplying K⁺ to the NKCC2, as described above. Thus, it is not surprising that the loss of function of ROMK results in a defective fluid and salt reabsorption. As a consequence of NaCl and fluid loss, a stimulation of K⁺ secretion along the distal nephron leads to hypokalemia, a hallmark of all forms of BS. Interestingly, patients carrying KCNJ1 mutations may show hyperkalemia as an initial sign of the disease, soon after birth; then, this defect is even reversed [8]. Micropuncture [9] and clearance [10] studies in young rats have demonstrated limited ability in K⁺ secretion compared with adults. Accordingly, in microperfused CCD from mammals, K⁺ secretion is low in newborns and cannot be stimulated by tubular flow [11]. This defective adaptation has been linked to the low expression of the apical BK channels that are known to mediate the flux-dependent K secretion.

BS type III depends on the mutation of the CLCKNA gene, encoding the basolateral Cl⁻ channel, CLC-Kb. Two proteins, CLC-Kb and CLC-Ka, encoded by the CLCNKb and CLCNKa genes, respectively, mediate the basolateral Cl⁻ efflux along this segment. Both proteins are associated with the Barttin subunit, essential for protein insertion into the plasma membrane. There is no evidence of CLCKNA mutations in BS. CLCk1 (the ortholog of CLC-Ka)-deficient mice show a phenotype of nephrogenic diabetes insipidus.

The fact that CLC-Kb is very abundant along the DT explains why this variant of BS is less commonly associated with a defect of urinary concentration and with hypercalcioria. Often, those patients show an overlapping phenotype between BS and GS.

BS type IV refers to the BS forms correlated with BSND mutations, encoding for the Barttin subunit of the basolateral chloride channel. The CLC-K/Barttin Cl channels also localize to the cochlea. In fact, patients carrying a BSND mutation show a phenotype of BS associated with hearing defects.

BS type V is caused by gain-of-function mutations of the calcium-sensing receptor (CaSR). The protein is expressed in the parathyroid and in the kidney, and it is mainly involved in calcium homeostasis. Along the TAL, the CaSR is expressed on the basolateral membrane. In the presence of hypercalcemia, the protein inhibits salt absorption, with consequent calcium and magnesium loss. Some case reports show that activating mutations of the CaSR show a BS phenotype.

**Clinical Presentation**

BS may have different clinical presentations:

- Antenatal BS, or hyper-prostaglandin E2 syndrome, refers to the most severe form, characterized by polyhydramnios due to excessive urinary output and premature birth. BS types I and II and sometimes type III are associated with this clinical subtype. After birth, patients commonly show fever, vomiting and lethargy. Biochemical analysis shows metabolic alkalosis, hyperkalemia, iso-hypostenuria and hypercalciuria. Nephrocalcinosis is frequent. A high urinary excretion of prostaglandin E2 or its metabolites is typical. High renin and aldosterone plasma levels are due to volume depletion. The cause of the high urinary and plasmatic prostaglandin levels is still unknown; it is possible that this is due to the defective NaCl reabsorption along the TAL.

- Classic BS is sustained by BS type III. The onset of the disease occurs generally during childhood. The clinical course is milder than in the antenatal subtype; patients show polyuria and polydipsia; nephrocalcinosis is uncommon.

- BS with sensorineural deafness. BS type IV may have a high variable of the severity of hydroelectrolyte imbalance. The clinical form may range from the antenatal form to mild hydroelectrolyte defects. Hearing loss is typical for BS type IV.


Gitelman Syndrome

Genetics

GS is a genetic tubular disorder sharing several analogies with BS. It is caused by mutations of the SLC12A3 gene, encoding the thiazide-sensitive sodium chloride cotransporter (NCC). Patients with heterozygous mutations do not show the disease; some studies have shown that they may have a tendency to exhibit hypotension, while complete GS phenotype occurs solely when the mutation is in homozygosis. A minority of patients show mutations of the CLCNKB gene, which is also responsible for BS type III. The presence of the basolateral Cl– channel along the TAL and DT explains why patients carrying CLCNKB mutations often show a phenotype overlapping between BS and GS.

Pathophysiology and Clinical Presentation

The defective NaCl absorption along the DCT leads to an increased Na+ load to the distal nephron. The latter, together with the activation of the RAAS induced by volume depletion, stimulates Na+ reabsorption along the CD, which in turn favors H+ and K+ secretion.

Besides hypokalemia and metabolic alkalosis, hypomagnesemia and hypocalciuria are other hallmarks of the disease. The renal handling of Ca2+ and Mg2+ is only partially understood in GS.

Normally, Mg2+ is freely filtered by the glomerulus; it is then reabsorbed along the PT by 10% and along the TAL by 50–70% via a paracellular pathway [12]. Along the DT, Mg2+ reabsorption occurs through a transcellular pathway, via the TRPM6 channels. Mice lacking the NCC cotransporter show a downregulation of TRPM6 in either the DCT or in the intestine, leading to stool and urinary Mg2+ loss.

Similar to Mg2+, Ca2+ ions are filtered by the glomerulus and reabsorbed along the PT and TAL in a paracellular manner, following the favorable electrochemical gradient. Transcellular Ca2+ absorption occurs along the DCT, mediated by the voltage-sensitive Ca2+ channel TRPV5 expressed on the apical membrane.

Low urine calcium excretion in GS is attributed to two mechanisms: (1) the volume contraction induced by the defective salt absorption along the CDT leads to a compensatory increased NaCl absorption along the PT, which also drives paracellular Ca2+ absorption; (2) NCC inactivation determines cell hyperpolarization, which activates TRPV5 channels leading to the increased Ca2+ absorption [13].

Hyperkalemia in the Setting of Renal Failure

Hyperkalemia is a potential life-threatening electrolyte imbalance. This condition may result from the disruption of the equilibrium between the ICF and ECF concentrations or from an increased total body content, caused by excessive food intake or defective excretion. The resulting increased plasma [K+] leads to a partial membrane depolarization that may eventually cause cardiogenic and skeletal muscle paralysis.

In the clinical setting, chronic renal failure is the most common risk factor for chronic hyperkalemia [14]. As the GFR declines, plasma [K+] is maintained in the normal range until the latest stages of renal failure. A previous study demonstrated that stage IV CKD patients showed the same increase in K+ secretion after K+ ingestion compared with people with normal renal function, indicating a conserved ability to excrete K+. This is possible because the remaining nephrons enhance their ability to secrete K+ through the upregulation of the entire machinery mediating its secretion along the distal nephron, including the Na–K ATPase, the ENaC and ROMK channels [15]. However, those patients showed a higher increase in plasma [K+] after K+ ingestion. This evidence suggested that CKD patients may have an impaired internal K+ homeostasis. However, this issue has been extensively investigated in animal models and humans, with conflicting results. Thus, whether the tissue K+ distribution is impaired in CKD is still debated [15].

Several pathophysiological factors favor the development of hyperkalemia in CKD patients: (1) the reduced urine output in the latest stage of CKD impairs the flux-dependent K+ secretion; (2) the presence of metabolic acidosis, another common complication of the late CKD, affects K+ distribution between the ICF and ECF by favoring K+ release to the extracellular space; (3) the presence of comorbidities, such as diabetes, with the reduction in insulin plasma levels and plasma hyperosmolality, increases the risk of hyperkalemia; and (4) the use of medications such as ACE inhibitors or ARBs.

Deregulation of the RAS axis has a proven critical role in determining hyperkalemia in CKD patients. Several studies have demonstrated that the use of ACE inhibitors and/or ARBs reduces the intraglomerular pressure, decreases tubulointerstitial damage and slows the progression of kidney disease in animal models of CKD. Accordingly, several trials have demonstrated the beneficial ef-
Effects of ACE and/or ARBs in CKD patients. The RAS inhibitor therapy has been shown to reduce the risk of renal failure and cardiovascular events in CKD patients, rendering RAS inhibitors the first-choice antihypertensive drugs [16]. However, the intensive treatment with RAS inhibitors in CKD patients increases the risk of adverse effects. The risk to develop hyperkalemia in stage 3 or higher CKD patients is low in those taking monotherapy with RAS blockers, unless volume depletion occurs [17]. Conversely, dual RAS blockade, even though it has been proven to reduce proteinuria, does not show significant benefits in slowing the progression of kidney disease or in reducing cardiovascular events, but increases the risk of hyperkalemia. The most recent KDIGO recommendations declare that there are insufficient data to prove a real benefit of dual RAS blockade in slowing CKD progression, so it should not be routinely prescribed.

Effect of Plasma [K+] on Blood Pressure

Epidemiological studies have demonstrated an inverse relationship between hypertension and dietary K+ intake. In the 1980s, Lever et al. [18] demonstrated that exchangeable total Na+ and K+ showed a positive and negative correlation, respectively, with hypertension. Further independent survey studies have shown that low K+ intake correlates with high blood pressure [19]. A recent meta-analysis has demonstrated that K+ supplementation resulted in a reduction of systolic and diastolic blood pressure by 4.7 and 3.5 mm Hg, respectively, with a greater effect in hypertensive patients [20]. Altogether, a number of epidemiological studies strongly suggest that increasing dietary K+ intake has a protective effect on the development of hypertension [21]. Whether K+ may have a regulatory effect on the control of blood pressure is an object of intensive investigations.

A number of potential mechanisms have been implicated, including sodium retention, hormonal activation and direct effect on the vascular smooth muscle function. KCl supplementation has been shown to inhibit, while low KCl diet enhances, NaCl reabsorption along the DCT, via the upregulation of NCC [22].

Dietary K+ loading, as well as increased plasma [K+] by KCl infusion, resulted in increased urinary Na+ excretion in association with reduced NCC phosphorylation in rats [23] (fig. 3). These findings suggest that K+ dietary intake, through changes in plasma [K+], modulate NCC activity, and therefore NaCl balance. Genetic diseases leading to the activation (Gordon syndrome) and inactivation (GS) of NCC cause opposite effects on plasma [K+], namely hyperkalemia and hypokalemia, respectively. However, it is not surprising that extracellular [K+] may modulate NCC function.

Terker et al. [24] have recently demonstrated that extracellular [K+] regulates NCC function directly, without hormonal mediation. They showed that plasma [K+] affects urinary K+ excretion by altering Na+ delivery to the CNT. Low K+ intake, leading to low plasma [K+], hyperpolarizes plasma membrane and causes a reduction of intracellular [Cl−] in the DCT. The latter has been shown to activate NCC via the WNK pathway, inducing a temporary increase in blood pressure in mice. This study indicates that plasma [K+] modulates NCC function in a hormone-independent manner, through changes in intracellular [Cl−] that in turn modulate the function of WNK kinases, known regulatory proteins of several transporters, including NCC.

The opposite effect is supposed to be determined by high K+ intake, which may lead to NCC inhibition with consequent increased Na delivery to the distal nephron, resulting in increased Na+ absorption and K+ secretion.

Conflict of Interest Statement

All authors declared no competing interests.
References


