ClC-5 chloride channel and kidney stones: what is the link?

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Abstract

Nephrolithiasis is one of the most common diseases in the Western world. The disease manifests itself with intensive pain, sporadic infections, and, sometimes, renal failure. The symptoms are due to the appearance of urinary stones (calculi) which are formed mainly by calcium salts. These calcium salts precipitate in the renal papillae and/or within the collecting ducts. Inherited forms of nephrolithiasis related to chromosome X (X-linked hypercalciuric nephrolithiasis or XLN) have been recently described. Hypercalciuria, nephrocalcinosis, and male predominance are the major characteristics of these diseases. The gene responsible for the XLN forms of kidney stones was cloned and characterized as a chloride channel called ClC-5. The ClC-5 chloride channel belongs to a superfamily of voltage-gated chloride channels, whose physiological roles are not completely understood. The objective of the present review is to identify recent advances in the molecular pathology of nephrolithiasis, with emphasis on XLN. We also try to establish a link between a chloride channel like ClC-5, hypercalciuria, failure in urine acidification and protein endocytosis, which could explain the symptoms exhibited by XLN patients.

Is nephrolithiasis a matter of imbalances in mineral metabolism?

Nephrolithiasis (or kidney stones) is one of the most common diseases in countries like the USA, United Kingdom, Japan, and Brazil (1-4). Most of the patients present recurrent kidney stones and metabolic disorders (5-7). Male patients predominate, but about 30% of patients are female (1-3).

Most calculi found in patients with kidney stones are formed by calcium salts, mainly calcium phosphate and calcium oxalate (8-10). Calculi formed by cystine (a dimer of cysteine), urate, and struvite (Mg(NH₄)PO₄) are found less frequently (1). The fact that calculi are mainly formed by calcium salts has led many investigators to speculate about a possible problem in calcium homeostasis (1). Although several investigators have explored this hypothesis, there are no conclusive data about this mechanism of stone formation.

Important advances have been recently made with the identification of inherited forms of nephrolithiasis. Several genes have been identified by genetic mapping of families with a history of kidney stones. Among these genes are the cystine transporter, which is defective in autosomal recessive cysti-
nuria (11), the calcium-sensing receptor (12), and the voltage-gated chloride channel ClC-5 (for a review, see Ref. 13), on which we focus this review.

Data involving ClC-5 in the pathogenesis of nephrolithiasis have been recently reviewed by Scheinman (14). In the present review we point out the postulated roles that ClC-5 could have in the pathogenesis of nephrolithiasis. Also, we try to establish a link between this chloride channel and stone formation.

The ClC-5 chloride channel and X-linked recessive nephrolithiasis: are they just variants or different diseases?

Dent’s disease was first reported by Dent and Friedman in 1964 (15). At that time, they identified this disease as a form of Fanconi’s syndrome due to impairment of renal acidification similar to that observed in such patients (15). Besides the renal acidification impairment, Dent’s disease patients also exhibited rickets, hypercalciuria, nephrocalcinosis, low molecular weight (LMW) proteinuria, and nephrolithiasis (14-17). Progressive renal failure may occur in the late stages of Dent’s disease (14,16,17).

Several reports have identified other syndromes with phenotypes similar to that of Dent’s disease. These diseases have been called X-linked recessive nephrolithiasis in the USA, X-linked recessive hypophosphatemic rickets in Italy, and LMW proteinuria with hypercalciuria and nephrocalcinosis in Japan (14). Although these syndromes have hypercalciuria, LMW proteinuria, and nephrocalcinosis in common, rickets and renal failure are not always present in all of them (16). Table 1 summarizes the common symptoms and their occurrence among the variants. Also, Table 1 shows mutations in the ClC-5 characterizing idiopathic nephrolithiasis in the USA and the characteristics of nephrolithiasis in Brazil.

Pedigree analysis of families with Dent’s disease revealed that this syndrome may be linked to the X chromosome. After 30 years, Thakker and colleagues (18-21) finally identified the locus responsible for the gene defect in the 11.22 region of the X chromosome. Further isolation of the gene encoded for the Brazilian patients. ND, Not determined. (Modified from Ref. 17, with permission).

*Found in just one patient (Ref. 60).

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<th>X-Linked recessive syndrome</th>
<th>XLN</th>
<th>Dent’s disease</th>
<th>X-Linked recessive rickets</th>
<th>LMW proteinuria/ nephrocalcinosis</th>
<th>Idiopathic nephrocalcinosis</th>
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<tbody>
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<td>UK</td>
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tated in Bartter’s syndrome which is characterized by electrolyte loss (24). The localization of ClC-K in the basolateral membrane of the thin ascending limb of Henle’s loop as well as its regulation by dehydration suggest a role in the mechanism of urinary concentration (25).

Lloyd and coworkers (26) showed that the mutant ClC-5 found in patients with Dent’s disease has lower conductance than the wild-type chloride channel. They also characterized several mutations of the ClC-5 gene in patients with X-linked recessive nephrolithiasis, X-linked hypophosphatemic rickets, and LMW proteinuria with hypercalciuria and nephrocalcinosis, showing that these diseases are just variations of a single pathology (26,27). All mutants exhibited loss of function when compared to wild-type ClC-5, suggesting that ClC-5 is needed for normal renal function. These syndromes were then grouped under the same name of X-linked hypercalciuric nephrolithiasis (XLN) by Scheinman (14).

The information that the ClC-5 channel could be the molecular entity responsible for stone formation led several researchers to investigate the molecular basis of the disease. Although mutations were found within the ClC-5 gene, no apparent relation between stone formation and a chloride channel was detected. The literature on ClC-5 and XLN will be discussed in the next sections.

**ClC-5 chloride channel and protein endocytosis**

The majority of patients with XLN also have LMW proteinuria (14,26-28). It is well known that LMW proteins are reabsorbed in the S2 and S3 segments of the proximal tubules, where several proteins filtered by the glomerulus are endocytosed and degraded within the lysosomes, providing the uptake of the amino acids (29).

ClC-5 is found in the S2 and S3 segments of proximal tubules, mainly in immediately early endosomes (30-35). Also, Devuyst et al. (35) demonstrated that proximal tubule cells transfected with ClC-5 are able to endocytose albumin and that the channel colocalizes with the albumin present in the endosomes. These data suggest a possible role of ClC-5 in non-receptor-mediated endocytosis, explaining the LMW proteinuria found in such patients.

Some hypotheses about the role of ClC-5 in the endosomes have been raised. Within the lysosomes several proteases work in an
acidic environment provided by the vacuolar H\(^+\)-ATPase (31-33). The transport of a positively charged ion such as H\(^+\) should be followed by an anion, such as Cl\(^-\), to maintain electroneutrality within the organelle. As long as it is present in the endosomes, CIC-5 may allow the transport of chloride through the endosomal membrane (31,32,34,35). Therefore, loss of CIC-5 function may lead to impairment of H\(^+\)-ATPase activity which would decrease the pH inside the endosome. Moreover, the failure in organelle acidification may lead to a decrease in protein endocytosis, and therefore, to the LMW proteins excreted in urine by XLN patients.

CIC-5 is found in endosomes but not in lysosomes (31-33). Acidic secretion also occurs in lysosomes and therefore they should also have a system for Cl\(^-\) transport. Hence, chloride channels other than CIC-5 may be present in lysosomes (34,35). Thus, it is reasonable to assume that CIC-5 contributes to Cl\(^-\) transport in the late endosome, but it is not clear how the channel works when the endosome becomes a lysosome.

Deficiency in endocytosis is supposed to be lethal due to its role in cellular nutrition. In rats, CIC-5 is expressed in other tissues besides the kidney, such as brain, lungs, liver, endothelial cells, and colon (31-33,36). In addition, the human CIC-5 channel is also expressed in endothelial cells, suggesting that the distribution of CIC-5 may not be restricted to the kidney (37). It is well known that protein endocytosis takes place in liver, endothelial cells, and colon. However, in XLN patients no disorders in endocytosis are observed in any other organs in addition to the kidney. Moreover, neither symptoms due to chloride channel defects nor dysfunction in endocytosis were detected in the endothelium. Thus, it is not understood how this important function could be impaired only in the kidney.

Several investigators have postulated that in the renal tubule the calculi are cleared by endocytosis (8,9,38,39). This mechanism would lead to fewer stones, lesser calculus formation, and thus, amelioration of the symptoms. Thus, besides having a role in protein endocytosis, the CIC-5 channel may also be responsible for the clearance of the calculi formed within the kidney tubules since both protein and calculi are endocytosed by similar mechanisms (29,38,39). Taken together, the data addressing the role of CIC-5 in protein endocytosis indicate that CIC-5 may also be important for calculus clearance. However, further studies are necessary to clarify its role in calculus endocytosis.

### CIC-5 chloride channel and urine acidification

Alkaline urine is a common sign found in Fanconi’s syndrome patients (15). Although Dent’s disease is currently characterized as an independent pathology, a urine acidification disorder was also found in some patients (13-17). It is well known that alkaline urine enhances the precipitation of calcium salts resulting in kidney stones (1,8,9). However, urine alkalization is not as widespread as hypercalciuria and LMW proteinuria in XLN patients (14-17).

Since the first report of CIC-5 mRNA localization several speculations have been made about its role in urinary acidification (32,40-43). CIC-5 mRNA and protein are expressed in the intercalated cells of collecting ducts (32,33,38,44). These cells are responsible for proton (H\(^+\)) secretion and, therefore, urine acidification. Working independently, Gunther et al. (33) and Sakamoto and coworkers (44) showed that the CIC-5 protein colocalizes with H\(^+\)-ATPase on the plasma membranes of type A intercalated cells from collecting ducts.

It is reasonable to assume that there should be Cl\(^-\) conductance to maintain electroneutrality of H\(^+\) secretion as well as within the endosome. To propose that Cl\(^-\) secretion may occur through the CIC-5 channel it is
necessary to show that CIC-5 is also expressed on the luminal plasma membrane, as demonstrated by Gunther and coworkers (33). In agreement with Gunther and coworkers, Friedrich et al. (45) have shown that CIC-5 is indeed expressed as plasma membrane conductance in *Xenopus* oocytes. In addition, Mo et al. (46) demonstrated that CIC-5 is inhibited by a decrease in extracellular pH. Therefore, the CIC-5 channel is sensitive to mild changes in pH, with down-regulation of the channel when the pH decreases. This seems to be an excellent regulatory mechanism in both the intra-endosomal space and the tubular lumen. However, some contradictory data have ruled out this hypothesis.

First of all, Luyckx et al. (34) did not show the expression of CIC-5 in intercalated cells. They showed that this channel is mainly present in subcellular compartments but not on the plasma membrane. Also, animals with lower expression of CIC-5 do not have any problem in urine acidification when compared with normal littersmates (47). In fact, despite the pattern of expression and its regulation by pH, there are no other data consistently relating CIC-5 dysfunction to urine alkalization. Once again, further experiments are necessary to confirm the role of CIC-5 in urine acidification.

**The role of CIC-5 in mineral metabolism**

Hypercalciuria is another major symptom of XLN (14-17). The hypothesis that higher levels of urinary calcium lead to precipitation of calcium salts and then to stone formation is well accepted today (1,2,8,9). Although CIC-5 seems to be expressed mainly in the human kidney (the major site of the disease), it is also expressed in rat colon, brain, and lungs (33,34,36-38,48,49) and in *Xenopus* oocytes (50).

The distribution of CIC-5 mRNA along the rat nephron reveals that this chloride channel is more abundant in the S3 segments of proximal tubules, in the cortical and medullary thick ascending limb of Henle’s loop, and in cortical collecting ducts (CCD) (33,34,36,37). An independent group using *in situ* hybridization further confirmed the expression of the CIC-5 channel on these segments (32). The localization of CIC-5 matches the major sites of calcium reabsorption in the kidney (51). Thus, localization of CIC-5 in the nephron segments responsible for calcium reabsorption and for the hypercalciuric phenotype exhibited by XLN patients suggested a role of CIC-5 in calcium transport. Therefore, a failure in CIC-5 function would lead to hypercalciuria. However, the relationship between CIC-5 and calcium metabolism has only recently been investigated.

Silva et al. (52) have shown that parathyromone (PTH) regulates CIC-5 expression *in vivo* in an animal model of disturbed calcium homeostasis. The authors showed that vitamin D-deficient and thyroparathyroidectomized animals have a lower (about 50%) expression of CIC-5 mRNA in the kidney cortex. The expression of CIC-5 returned to normal values when the animals were treated with PTH. These data are reproducible when the CIC-5 protein is analyzed by Western blots. It is well known that PTH enhances calcium reabsorption by the kidney and that the major site of this action is the kidney cortex (51). Animals with low or almost no expression of CIC-5 have high urinary calcium excretion (52). These data suggest that CIC-5 is necessary for calcium reabsorption in the kidney.

Although it may be not clear how a chloride channel could enhance calcium reabsorption, Gesek and Friedman (49) showed a functional chloride channel in cultured CCD cells which reabsorb calcium under PTH and calcitonin stimulation. PTH, the major hormone increasing calcium reabsorption in the kidney, activates this chloride channel in CCD cells (49). The mechanism of activation of this chloride channel would require...
gene expression. Taken together, these data suggest that ClC-5 could be the chloride channel involved in this mechanism since PTH also enhances its expression in vivo. However, other chloride channels with similar properties, such as the closely related ClC-3, are also expressed in CCD cells (32,53). Therefore, it still is necessary to determine the molecular identity of the channel reported by Gesek and Friedman (49).

ClC-5 protein was detected in the plasma membrane as well as in immediately early endosomes of the S3 segments and on the apical membrane of the CCD cells of rat kidney (33,44). It is well known that these segments participate in the transcellular mechanisms of calcium reabsorption (51).

As mentioned above, the loss of ClC-5's function leads to hypercalciuria, a widespread symptom in XLN (14-17). Although failure of calcium reabsorption is a good hypothesis to explain the hypercalciuria, some clinical findings do not agree with the animal model proposed by Silva and coworkers (52).

Patients with XLN show elevated serum vitamin D (1α,25(OH)2 vitamin D3 or calcitriol) and normal plasma PTH levels, a fact that is not expected for patients with excessive calcium excretion. These clinical observations lead to a speculation that hypercalciuria could be a secondary mechanism due to excessive intestinal reabsorption enhanced by calcitriol (54,55). This hypothesis was explored by Luyckx and coworkers (47) who produced a mouse model with lower expression of the ClC-5 channel. On a calcium-restricted diet these mice have a lower urinary calcium/creatinine ratio. The authors suggest that enhanced calcium reabsorption in the intestine (but not the decrease in calcium reabsorption in the kidney) leads to an increase in calcium filtered by the kidney glomeruli and then hypercalciuria.

In agreement with this hypothesis is the fact that ClC-5 is expressed in the intestine, mainly in rat colon and a human colon-derived cell line HT-29 (37,48,52). It may be possible that ClC-5 acts on intestinal calcium absorption as well as on kidney calcium reabsorption. However, an enhancement in the intestinal absorption of calcium would be followed by an increase in serum calcium compared to control mice. Nevertheless, this enhancement in serum calcium does not take place in mice or in XLN patients. Also, an excess of calcium is excreted in feces by Dent's disease patients (14-17,52,54). Although the intestinal component relating ClC-5 dysfunction to kidney stones may be an important factor for the pathogenesis of nephrolithiasis, further studies are necessary to test this hypothesis.

**Future perspectives and concluding remarks**

Up to now, approximately 40 papers have been published about the ClC-5 chloride channel. Many of them have explored and characterized the mutations responsible for the several variants of XLN (20,21,26,28,30,56-59). Very few papers have focused on the physiological role and regulation of this channel (35,47,49). On the other hand, the structure and function of ClC-5 have not been studied as extensively as for the other members of the ClC family (22,23).

We are aware that many experiments should be done to address all the questions raised since the cloning of ClC-5 by Thakker and coworkers (18,19). The possible roles attributed to ClC-5 need further investigation and represent a growing area of research. Important questions need to be answered: 1) Why does XLN, being an X-linked recessive disease, show symptoms in female carriers? 2) Why do different mutants exhibit different phenotypes? 3) Does hypercalciuria lead to body calcium and bone mass loss, and thus to a rickety phenotype in children?

Knockout mice could be a good model for answering the first question. Except for
the mice with lower CIC-5 expression obtained by Luyckx and coworkers (47), no other animal model has been developed. However, it would not be a surprise if CIC-5 knockout mice did not present any of the symptoms of the human disease. Moreover, the CIC-5 chromosomal localization is still unknown. As long as mice are not syngeneic to humans, it is possible that the murine CIC-5 locus is not located on chromosome X. This would impair a genetic analysis of carrier females and their offspring. This pedigree analysis is essential to investigate the inheritance of genetic diseases and it has been done for XLN as well (24). But if mouse CIC-5 is located on chromosome X it would be a good model for the pattern of inheritance and the variations of symptoms.

Different members of the CIC chloride channel family have been intensively studied in terms of structure and function (22,23). Although CIC-5 may have a structure and function pattern similar to that of other members of the family, such studies have not been done on CIC-5. Similar studies are necessary because they could help clarify why different mutations of CICN5 may result in such different phenotypes (20,21,26,28,30,56-60).

Rickets are not so common as hypercalciiuria and nephrolithiasis. However, the rickets phenotype is common in Europe (20,21,26,27,30,57-59). Recent data suggest a role of CIC-5 in calcium metabolism (47,52). If CIC-5 is actually involved in calcium metabolism it could be expected to be expressed in bone, the major site of calcium storage. So far its expression has not been analyzed in bone or in bone cells. As we mentioned above, CIC-5 has a different pattern of expression varying from a widespread channel expressed in *Xenopus* (51) to an almost tissue-specific channel in humans (19,48,50). It is possible that CIC-5 may also have a role in human bone metabolism if it is expressed in bone. This could represent more evidence of the growing dysfunction observed in some XLN patients. However, if CIC-5 has a role in bone metabolism a growth deficiency would be expected in mice with lower expression of CIC-5, but this is not the case (47). Thus, once again, a fully characterized animal model would be very important for the study of XLN phenotypes, or might perhaps raise more questions about the role of this chloride channel in this complex disease.

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