

# Molecular pathogenesis of Bartter's and Gitelman's syndromes

Principal discussant: IRA KURTZ

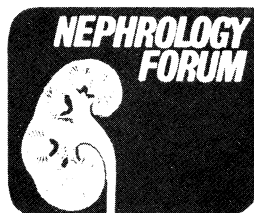
UCLA School of Medicine, Los Angeles, California, USA

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## DISCUSSION

DR. IRA KURTZ (*Chief, Division of Nephrology, Max Factor Professor of Medicine, Center for Health Sciences, UCLA School of Medicine, Los Angeles, California*): When Dr. Jay Stein presented a Nephrology Forum on Bartter's syndrome (BS) in this journal 13 years ago [1], he noted that "several questions remain to be clarified: Is Bartter's syndrome the consequence of a single defect, or rather the clinical phenotype of a variety of tubular defects? Is the syndrome caused by a defect in sodium chloride transport per se or by a primary potassium-wasting state? Are syndromes associated with magnesium and calcium loss, as well as potassium wasting, pathophysiologically linked?" These prescient questions can now be answered with more certainty because of the recent advances in our understanding of the molecular basis for Bartter's/Gitelman's syndromes.

The patient being discussed today presented with the following electrolyte abnormalities: hypokalemia with renal  $K^+$  wasting (high fractional excretion); a chloride-resistant metabolic alkalosis, elevated plasma renin, elevated plasma aldosterone (relative to the serum  $K^+$  concentration), hypomagnesemia with renal magnesium wasting (high fractional excretion), decreased urine calcium excretion, and normal blood pressure. These findings are compatible with either diuretic abuse due to thiazide-like diuretics or Gitelman's syndrome (GS). Because of the family history and the negative diuretic screen, a diagnosis of Gitelman's syndrome was made. In the absence of marked symptoms over many years, we elected not to treat the patient's electrolyte disturbances.

The patient under discussion has the classic features of GS [2, 3]. Patients with this disorder are normotensive and have a chloride-resistant metabolic alkalosis, increased plasma renin and aldosterone/ $K^+$  ratio, hypokalemia, hypomagnesemia, and hypocalciuria. The syndrome was first described in 1966 by Gitelman et al in three patients, two of whom were sisters [2]. The metabolic abnormalities in Gitelman's syndrome shared many of the features of BS, a disorder first reported by Bartter et al four years earlier and with which it has subsequently been frequently confused in the literature [4]. Interestingly, the first case of BS might have been reported in 1957, when Rosenbaum and Hughes

## CASE PRESENTATION

A 42-year-old man presented to UCLA with a history of chronic hypokalemia and hypomagnesemia. His medical history was unremarkable except for occasional cramps in his calves. He had no history of laxative or diuretic abuse or of vomiting, and he was taking no medications. His 47-year-old brother also had a history of chronic hypokalemia and hypomagnesemia and severe intermittent muscle cramps. He was being treated with amiloride and magnesium supplementation. Both parents and a third brother were asymptomatic and had no electrolyte abnormalities.

Physical examination revealed a well-developed white male in no distress. His blood pressure was 110/70 mm Hg; heart rate, 72 beats/min; jugular venous pressure was estimated to be 7 cm; his chest was clear to auscultation. Cardiovascular examination disclosed a regular rate and rhythm, and no murmurs, rubs, or gallops; his abdomen was nontender without hepatosplenomegaly. Neurologic examination revealed cranial nerves 2-12 intact, no sensory or motor deficit, and no Chvostek or Trousseau sign. He did not have peripheral edema.

Laboratory values were: serum sodium, 136 mEq/liter; potassium, 2.4 mEq/liter; chloride, 94 mEq/liter; total  $CO_2$ , 28 mEq/liter; venous pH, 7.47 and  $PCO_2$ , 38 mm Hg; BUN, 10 mg/dl; serum creatinine, 0.7 mg/dl; ionized calcium, 1.25 mmol/liter; total calcium, 9.2 mg/dl; phosphorus, 3.2 mg/dl; magnesium, 1.1 mg/dl; supine plasma renin, 16.6 ng/ml/hr; supine aldosterone, 168 pg/ml; PTH, 13 pg/ml;  $1,25(OH)_2$  vitamin D, 42 pg/ml. Urine chemistries were: sodium, 64 mEq/liter; potassium, 42 mEq/liter; chloride, 58 mEq/liter; calcium, 1 mg/dl; magnesium, 4.3 mg/dl; phosphorus, 67.4 mg/dl; and creatinine, 81 mg/dl. A diuretic screen was negative. The patient was discharged on no medications and has remained asymptomatic.

**Key words:** metabolic alkalosis, hyperprostaglandin E syndrome, hypokalemia, hypomagnesemia, hypocalciuria, nephrocalcinosis.

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**Table 1.** Clinical differences between Bartter's and Gitelman's syndromes

	Neonatal Bartter's syndrome	Adult Bartter's syndrome	Gitelman's syndrome
Polyhydramnios	+	–	–
Failure to thrive	+	–	–
Growth retardation	+	–	–
Polyuria	+	+	–
Polydipsia	+	+	–
Muscle cramps/spasm	–	–	±
Chondrocalcinosis	–	–	±

described an infant with hypokalemia, metabolic alkalosis, polyuria, growth failure, and recurrent episodes of volume depletion [5]. The patients reported by Bartter et al shared many of the features of GS in that they had a hypokalemic, chloride-resistant metabolic alkalosis, with elevated plasma renin levels and normal blood pressure. Unlike GS, the patients with BS had a urinary concentrating defect and growth failure and subsequently were shown to be hypercalciuric [6]. Despite these known differences, most clinicians fail to distinguish between these syndromes. Indeed, it appears that most patients reported in the literature with BS in fact have GS [7].

### Clinical features

Bartter's and Gitelman's syndromes can be distinguished clinically (Table 1) as well as on the basis of differences in divalent cation metabolism [8–14]. In BS, symptoms are dictated by the patient's age at presentation. Polyuria and polydipsia are always present in children and adults with BS and can be the only presenting symptoms. Most cases of BS present in neonates [15–21]. Maternal polyhydramnios with premature labor is a common finding, presumably because of excessive urine production in utero [6, 21, 22]. Some of the infants have an appearance typical of BS, including a prominent forehead, triangular facies with drooping mouth, and large eyes and pinnae (Fig. 1) [23–25]. Subsequent difficulties in the neonate include failure to thrive and growth retardation [21]. Vomiting, diarrhea, and fever have been reported, and a variant of neonatal BS has been described with sensorineural deafness [25]. Neonatal BS can be diagnosed prenatally by finding elevated amniotic fluid chloride and aldosterone concentrations [26, 27].

Seyberth and colleagues have suggested that the severe neonatal form of BS be called the hyperprostaglandin E<sub>2</sub> syndrome [17, 28, 29]. Indeed, an increased rate of PGE<sub>2</sub> excretion is a common feature in BS [30–33]. Indomethacin has been used to decrease the incidence of polyhydramnios in pregnant patients with and without BS, although complications in the neonate, including acute renal failure, tricuspid regurgitation, and patent ductus arteriosus have been reported [19, 34–36]. In addition, indomethacin appears to improve the rate of growth in these children [34].

Given that the neonatal form seems to differ only in severity from the childhood and adult disorder, the term neonatal BS rather than hyperprostaglandin E<sub>2</sub> syndrome appears justified.

In contrast to BS, GS is a milder disease and doesn't present symptomatically in the neonatal period [7, 37]. Patients with GS are not volume depleted clinically. They can have a plasma aldosterone concentration that is not markedly elevated, as in the patient presented today, because their hypokalemia suppresses adrenal aldosterone production. Furthermore, patients with GS are often asymptomatic, or they only have neuromuscular complaints consisting of intermittent muscle spasms, cramps, or tetany [2, 3, 37–39]. Polyuria and polydipsia are not a feature of GS. In a subset of patients with electrolyte abnormalities indistinguishable from those of GS, arthritis in several joints due to chondrocalcinosis has been reported both in sporadic cases [40–43], and recently in a familial form [44].

### Divalent cation disturbances in BS and GS

Several differences in divalent cation metabolism allow the two syndromes to be easily distinguished based on simple laboratory measurements (Fig. 2). Patients with BS have fasting hypercalciuria, whereas those with GS have a low urinary calcium excretion [6, 9–12, 14, 45]. In addition to hypercalciuria, medullary nephrocalcinosis is usually present in BS but is always absent in GS. The hypercalciuria in BS can predate the appearance of overt nephrocalcinosis. In addition to hypocalciuria, all patients with GS characteristically have hypomagnesemia and a high fractional excretion of magnesium [2, 10]. The patient presented today had a normal  $U_{mg}/U_{cr}$  of 0.05. In GS, urinary magnesium excretion reflects the dietary intake and only exceeds the normal excretion rate when patients are ingesting magnesium supplements [2, 10]. In BS, both the absolute urinary calcium excretion and the fractional excretion rate are increased. Despite the hypocalciuria in GS, the serum total and ionized calcium concentrations are normal, as in the patient presented today [10]. Colussi et al also reported that serum PTH levels are in the low normal range in GS, possibly because of the inhibitory effect of hypomagnesemia on PTH secretion [10]. In contrast, hypomagnesemia is not a typical finding in BS. A modest or transient decrease in serum magnesium has been reported in some neonates with BS. However, these infants generally are extremely ill, and whether the hypomagnesemia is due to extrarenal causes (vomiting, diarrhea) has not been excluded. Hypomagnesemia in adults diagnosed with BS in most cases is due to the failure to diagnose GS [9, 46]. Patients with BS and GS also differ in that the former have elevated 1,25(OH)<sub>2</sub>D levels [8, 10, 12, 47] and PGE<sub>2</sub> excretion rates [30, 31, 33, 48]. The increased serum 1,25(OH)<sub>2</sub>D in patients with BS occurs in the presence of normal 25-OHD, 24,25(OH)<sub>2</sub>D, PTH, and phosphorus levels [47]. The mechanism for elevated serum 1,25(OH)<sub>2</sub>D levels in BS is



Fig. 1. Two patients with neonatal Bartter's syndrome described by Landau et al [25]. Both patients have the typical facial appearance with prominent forehead, triangular face, drooping mouth, and large eyes and pinnae.

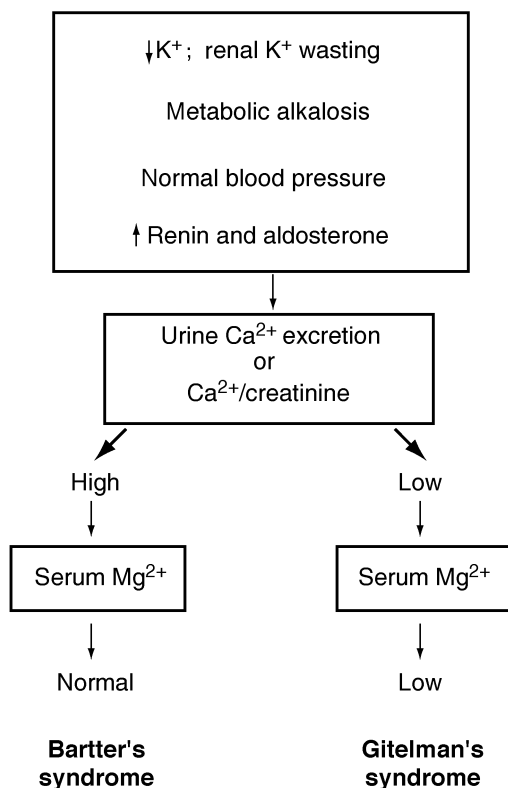


Fig. 2. Approach to the clinical differentiation between Bartter's syndrome and Gitelman's syndrome based on divalent cation metabolism. Note that some patients with type-3 BS have a normal urine  $\text{Ca}^{2+}$ /creatinine ratio.

unknown. Previous studies have shown that  $\text{PGE}_2$  stimulates the hydroxylation of 25-(OH)D in thyroparathyroidectomized rats and in chick and rat kidney cells [49–51].

Restrepo de Rovetto et al have argued that  $\text{PGE}_2$  stimulates the activity of  $1\alpha$ -hydroxylase, which results in elevated  $1,25(\text{OH})_2\text{D}$  levels in BS [47]. Indomethacin therapy reduces the elevated  $1,25(\text{OH})_2\text{D}$  levels,  $\text{Ca}^{2+}$  excretion, and  $\text{PGE}_2$  excretion in BS [8, 47]. A second unproven possibility is that the renal calcium leak per se increases plasma  $1,25(\text{OH})_2\text{D}$  levels.

#### Role of PGE in BS/GS

The BS/GS literature is replete with additional metabolic abnormalities that have contributed to the confusion regarding the pathogenesis of these syndromes. Most disputed has been the role of excess prostaglandin excretion in the pathogenesis of the electrolyte abnormalities [29–32, 52–56]. Indeed 17 years ago, a Nephrology Forum presented by Dr. Michael Dunn was devoted to the role of prostaglandins in BS [33]. Patients with GS usually have normal  $\text{PGE}_2$  excretion rates [7, 57], although elevated excretion rates also have been reported in GS (that is, in patients misdiagnosed as having BS) [9, 58, 59]. Although cyclooxygenase inhibitors ameliorate some of the electrolyte disturbances and growth retardation in BS [34, 60], these drugs are less effective in GS [57, 61]. Excess  $\text{PGE}_2$  excretion in BS is thought to be a secondary phenomenon [37]. This conclusion was reached prior to the description of the specific transport defects in BS/GS on the basis of the following considerations: (1) mepacrine, a potent phospholipase inhibitor, inhibits  $\text{PGE}_2$  excretion without increasing plasma potassium and decreasing plasma renin [46]; (2) loop diuretics, which mimic several of the features of BS, increase  $\text{PGE}_2$  production and/or excretion [62, 63]; and (3) hypokalemia is not a feature of prostaglandin-producing tumors [64].

### Clinical assessment of the tubular transport defects in BS/GS

Early clearance studies suggested that the transport defect in BS was located in the diluting segment [65, 66]. Initially this finding was attributed to abnormal thick ascending limb function, although later studies suggested that abnormal NaCl transport in the distal convoluted tubule also resulted in abnormal urine diluting ability in some patients misdiagnosed with BS and in GS [9, 13, 61, 67–69]. The finding that patients with neonatal BS have a urine concentrating defect in addition to a diluting defect, fail to respond normally to furosemide, yet have a normal response to thiazides, suggested that a transport abnormality in the thick ascending limb of Henle was involved [70]. Similarly, the finding that patients with GS concentrate their urine normally, have hypocalciuria, fail to respond appropriately to thiazide diuretics, and have a greater-than-normal response to loop diuretics suggested that these patients have a transport abnormality in the distal convoluted tubule [9, 13, 61, 67–69, 71]. We know now that the mere presence of a diluting defect does not distinguish BS from GS, since both defective thick ascending limb and distal convoluted tubule NaCl transport lead to impaired urine diluting ability [61].

### NaCl and calcium transport in the distal convoluted tubule

The characterization of the NaCl transport processes in the thick ascending limb of Henle (TAL) and the distal convoluted tubule (DCT) has proven invaluable in our understanding of the transport abnormalities in BS and GS. I will begin with a brief outline of normal DCT transport. The distal tubule comprises three distinct segments: the distal convoluted tubule, the connecting tubule, and the initial cortical collecting duct [72]. Most in-vivo data have been obtained from rats in which delineation among these segments is difficult. In-vivo transport studies, and more recent studies of immortalized mouse distal tubule cells, have shown that the DCT plays an important role in electroneutral NaCl absorption and PTH-responsive  $\text{Ca}^{2+}$  absorption [72, 73]. The molecular basis for electroneutral NaCl transport in the DCT was defined in 1993, when Gamba et al isolated a ~3.7 kb cDNA, *fTSC* (thiazide-sensitive cotransporter, also known as NCCT (NaCl cotransporter), that encoded the NaCl cotransporter in flounder (*Pseudopleuronectes americanus*) bladder [74]. A fragment of *fTSC* then was used to screen a rat renal cortical cDNA library; the rat NaCl cotransporter (*rTSC1*), a ~4.4 kb cDNA encoding a ~110 kD protein with 12 putative membrane-spanning domains, was isolated and identified [75]. Injection of oocytes with *rTSC1* cRNA significantly increased  $\text{Cl}^-$ -dependent, thiazide-inhibitable  $^{22}\text{Na}^+$  uptake. In-situ hybridization and immunocytochemistry studies combined with functional data indicate that the

NCCT gene (human gene mapping workshop approved symbol SLC12A3) encodes the apical membrane thiazide-sensitive NaCl cotransporter in the distal convoluted tubule [76–78]. In the rat kidney, polyclonal antibodies against *rTSC1* intensely label the apical membranes in the distal nephron beginning in the early distal convoluted tubule and terminating within the connecting tubule; the intensity of labeling diminished from early to late sites along the distal tubule [78]. In addition to the DCT, evidence indicates NCCT expression in osteoblast-like cells [79] and peripheral blood mononuclear cells [80]. The physiologic role NCCT plays in these extrarenal sites remains to be determined.

We have known for three decades that thiazide diuretics decrease urinary calcium excretion [81]. Micropuncture and microperfusion experiments identified the distal convoluted tubule as the site where thiazides stimulate calcium absorption [82]. Stimulation of  $\text{Ca}^{2+}$  absorption has been attributed to (1) decreased apical  $\text{Na}^+$  uptake driving basolateral  $\text{Na}^+/\text{Ca}^{2+}$  exchange with subsequent increased apical  $\text{Ca}^{2+}$  uptake; and (2) decreased intracellular chloride concentration, which causes hyperpolarization of the apical cell membrane and stimulates apical  $\text{Ca}^{2+}$  uptake via dihydropyridine-sensitive  $\text{Ca}^{2+}$  channels [82, 83]. Three alternatively spliced isoforms of the  $\text{Na}^+/\text{Ca}^{2+}$  exchange gene, NCX1–NACA2, NACA3, and NACA6–have been identified in these cells derived from the mouse DCT, which might mediate  $\text{Na}^+$ -dependent  $\text{Ca}^{2+}$  efflux from the DCT [84]. The relative role of the basolateral  $\text{Na}^+/\text{Ca}^{2+}$  exchange isoforms and  $\text{Ca}^{2+}$ -ATPase in mediating cellular calcium extrusion is currently unknown; both transporters are co-expressed with *rTSC1* in the late DCT and connecting tubule [78].

### Molecular basis for Gitelman's syndrome

It was an “annum mirabilis” for this field in 1996, when Lifton's group uncovered the molecular basis for GS [85] and BS [86, 87] and allowed many of Dr. Stein's questions to be answered. Simon et al first demonstrated complete linkage of GS to the NCCT gene locus (SLC12A3) on chromosome 16q13, and specified GS as an autosomal recessive disorder with 99% penetrance [85]. The linkage to chromosome 16q13 has been confirmed by others [88–91]. Simon et al identified a wide variety of mutations distributed throughout the NCCT gene consistent with loss-of-function alleles. Additional potential loss-of-function mutations also were reported in 1996 [90, 92]. Missense mutations are most frequent and are usually localized to the intracellular domains of the protein rather than to the extracellular or transmembrane domains. Many of the patients described thus far are compound heterozygotes. This finding, together with the documentation of independent mutant alleles within the same kindred, suggests that mutant alleles are not rare. In the Swedish and Italian populations studied, the prevalence of heterozygotes based

Normal	GVMIRCMLNI	WGVILYLRLP	WITAQAGIVL	TWIIILSVT	VTSITGLSIS	190
F	GVMIRCMLNI	WGVILYLRLP	WITAQAGIDE	KTEAQRREM	VGRMVNE	187
M	GVMIRCMLNI	WGVILYLRLP	WITAQAGIVL	TWIIILSVT	VTSITGLSIS	190
Normal	MDQERKAIIS	LLSKFELGFH	EVHILPDINQ	NPRAEHTKRF	EDMIAPFRLN	930
F	-----	-----	-----	-----	-----	187
M	MDQERKAIIS	LLSKFELGFH	EVHILPDINQ	NPRAEHPFGR	-----	920

**Fig. 3. Predicted amino acid sequence of paternal (F) and maternal (M) NCCT polypeptides in the presented patient with GS.** Messenger RNA was isolated from peripheral blood mononuclear cells obtained from normals and from the patient under discussion with GS. Following reverse transcription with AMV reverse transcriptase, NCCT cDNA was amplified by PCR and sequenced. Genomic DNA from the patient and his parents also was sequenced to determine the inheritance of each allele. Two different mutations were detected in the patient's NCCT cDNA (compound heterozygote). The cDNA from the patient's paternal allele had an additional 119 bp insertion between exon 3 and 4, generating a predicted premature stop codon after amino acid 187. In cDNA from the maternal allele, exon 24 was deleted, resulting in a predicted premature stop codon after amino acid 920.

on phenotypic expression is approximately 1% [14, 85, 93]. Simon et al have estimated a sporadic prevalence of 0.001 and a mutant allele frequency of 1/200 [85].

The patient presented today is a compound heterozygote (Fig. 3). Two different mutations were detected in the patient's NCCT cDNA obtained from his peripheral blood mononuclear cells. In cDNA derived from the patient's maternal allele, exon 24 was deleted and resulted in a premature stop codon (after amino acid 920). The cDNA derived from the paternal allele had an additional 119 bp insertion between exon 3 and exon 4, generating a premature stop codon (after amino acid 187). The patient's genomic DNA had a previously reported 5' splice mutation in intron 24, GGT → GTT (maternal allele) [85], and a previously unreported 3' splice site mutation in intron 3, CAG → CAA (paternal allele), which resulted in the activation of a nearby cryptic splice site in intron 3 [80].

The severity of symptoms has varied widely in families with Gitelman's syndrome. The phenotypic consequences of the mutations described thus far are unknown. Whether patients with homozygous frameshift mutations have a more severe phenotype than do those with homozygous or compound heterozygous missense mutations remains to be determined [92]. Interestingly, the patient under discussion was essentially asymptomatic, whereas his brother, who has identical mutations in the NCCT gene, had frequent muscle cramps and spasms. The electrolyte abnormalities in the brother were essentially the same, that is, serum sodium, 140 mEq/liter; potassium, 2.3 mEq/liter; chloride, 94 mEq/liter; total CO<sub>2</sub>, 27 mEq/liter; BUN, 13 mg/dl; serum creatinine, 1.0 mg/dl; ionized calcium, 1.21 mmol/liter; total calcium, 9.1 mg/dl; phosphorus, 3.7 mg/dl; and magnesium,

1.1 mg/dl. These findings suggest that other biochemical abnormalities or psychogenic or environmental factors are modifying the severity of the symptoms in the two brothers. Alternatively, a background gene effect could modify the electrolyte abnormalities and/or symptoms in patients with identical mutations in the NCCT gene locus. The creation of knockout mice with various NCCT mutations will be required to correlate specific mutations with the severity of specific GS phenotypes (biochemical abnormalities, symptoms). In addition, comparing identical mutations in several strains of mice will help determine the role of background genes in altering the phenotypic severity in GS.

In 1979, Bauer et al reported a syndrome with electrolyte abnormalities identical to GS but which differs from the classic disorder in that patients have arthritis due to chondrocalcinosis [40]. As I mentioned earlier, several sporadic cases have since been described [41–43, 94]. Smilde and colleagues reported a familial form of the disorder in which 7 affected family members had radiologic abnormalities involving their wrists, elbows, shoulders, knees, and symphysis pubis [44]. Magnesium therapy appeared to improve the radiologic appearance of the joint abnormalities in one patient over a 10-year followup. Magnesium, a known cofactor in the conversion of pyrophosphate to inorganic phosphate, increases the solubility of calcium pyrophosphate *in vitro* [95]. However, if hypomagnesemia is causally related to the arthritis in these patients, it is unclear why most patients with GS and patients with other hypomagnesemic disorders don't have chondrocalcinosis. Whether a separate gene locus accounts for this syndrome or whether background genes alter the typical GS phenotype is unknown. The gene for a separate

familial form of chondrocalcinosis associated with intermittent benign seizures and lacking the metabolic abnormalities in GS has been localized to chromosome 5p [96].

### Na-K-2Cl cotransport

Clearance studies suggested that the thick ascending limb (loop of Henle) was the site of the transport abnormalities in BS [65]. The characteristics of  $\text{Na}^+$ ,  $\text{K}^+$ , and  $\text{Cl}^-$  transport in the cortical thick ascending limb (cTAL) and medullary thick ascending limb (mTAL) were first demonstrated by Rocha and Kokko [97] and by Burg and Green [98]. Transport studies in intact nephrons and vesicle preparations have established that the predominant mode of apical  $\text{Cl}^-$  absorption is via a furosemide-sensitive Na-K-2Cl cotransporter [99]. Sodium chloride transport in the TAL is stimulated by hormones that activate cAMP, and is inhibited by extracellular  $\text{Ca}^{2+}$ ,  $\text{PGE}_2$ , ANF, and cytochrome P-450 arachidonate metabolites [100]. In 1994, using a homology-based screening strategy and functional expression in oocytes, Gamba et al reported the isolation of a ~4.7 kb cDNA encoding the rat apical Na-K-2Cl (NKCC2) cotransporter [75]. Payne and Forbush reported several splice variants in rabbit and mouse kidney, which they have called NKCC2A, NKCC2B, and NKCC2F [101]. Using PCR, Yang et al have identified a fourth isoform, NKCC2AF, in rat kidney [102]. These isoforms, which arise from alternative splicing of cassette exons, result in a variable 96-bp region located in the second transmembrane domain. Amplification by PCR of DNA derived from specific nephron segments revealed segment-specific expression: the A isoform is expressed in the cTAL, mTAL, and macula densa; B isoform is expressed in the cTAL and macula densa; F isoform is expressed in the mTAL and outer medullary collecting duct (OMCD); and AF is expressed in the mTAL. Whether these splice variants differ in their functional characteristics is unknown. Immunoblot studies show that the major NKCC2 protein is approximately 150 kD [103]. The topology of this transporter is similar to NCCT in that both proteins have 12 putative membrane spanning domains and a central hydrophobic core region. The overall sequence identity between the transporters is approximately 50%. Immunocytochemistry studies have localized the NKCC2 protein to the apical membrane of the cTAL and mTAL [103]. Other members of the Na-K-2Cl cotransport family have been cloned, including the secretory form of the Na-K-2Cl cotransporter, NKCC1, first cloned from dogfish shark rectal gland [104], and subsequently cloned from human colonic cells [105] and murine inner medullary collecting duct [106]. More recently, the KCl cotransporter KCC1 encoding a 1085 amino acid polypeptide has been cloned [107]. The KCC1 cotransporter shares approximately 25% homology with other members of the cation-chloride cotransporter family, is expressed in several tissues, and might represent the housekeeping form responsible for cellular volume

regulation. In the renal tubule, this transporter might play a role in basolateral KCl efflux in the rabbit proximal tubule [108] and cTAL [109].

### Mechanism for $\text{K}^+$ secretion in the TAL and CCD

Low conductance (30 pS) inward rectifying  $\text{K}^+$  channels play a significant role in mediating  $\text{K}^+$  secretion in the thick ascending limb (TAL), the cortical collecting duct (CCD), and various tissues [110]. Luminal  $\text{K}^+$  efflux in the TAL ensures an adequate supply of  $\text{K}^+$  for the apical Na-K-2Cl cotransporter. In the CCD,  $\text{K}^+$  secretion by principal cells ensures adequate renal excretion of dietary  $\text{K}^+$  intake. Efforts at identifying channel molecules responsible for renal  $\text{K}^+$  secretion resulted in the expression cloning of ROMK1 (rat outer medulla  $\text{K}^+$  channel) from rat renal outer medulla [111], and subsequently in the cloning of several splice variants in rats (ROMK1-3,6 [112]) and humans (ROMK1-5 [113]). This divergence of ROMK mRNA results in the formation of channels with variable lengths of amino termini. The ROMK channels are members of a superfamily of structurally and functionally related  $\text{K}^+$  channel proteins [110]. The common functional properties shared by these channels is their inwardly rectifying current voltage relationship. Rectification can be weak or strong and is due to a voltage-dependent block of the channel pore by intracellular magnesium and polyamines. The ROMK channels expressed in *Xenopus* oocytes form a weakly inwardly rectifying potassium-channel current with a single channel conductance of 35–45 pS [114]. The channel requires magnesium and ATP to open, exhibits a high open probability, and is inhibited by millimolar concentrations of ATP, low intracellular pH, and arachidonic acid [110, 114]. These properties are comparable to those of the low-conductance potassium channel electrophysiologically identified in the apical membrane of thick ascending limb and cortical collecting duct cells [110]. In rats, ROMK1 transcripts are expressed in the CCD, OMCD, and inner medullary collecting ducts; ROMK2 is located in the mTAL, cTAL, DCT, connecting tubule (CNT), and CCD; and ROMK3 is found in the mTAL, cTAL, DCT, and CNT [115, 116]. No information exists regarding isoform distribution in the human nephron. No functional differences have been found thus far in human ROMK1-5 when expressed in *Xenopus* oocytes [117]. The ROMK2 potassium channel and the cystic fibrosis transmembrane regulator (CFTR) can form a complex, and co-expression of ROMK2 and CFTR increases the sensitivity of ROMK2 to the  $\text{K}^+$  channel inhibitor glibenclamide [118]. However, it has been suggested that the ROMK6 isoform is more likely to associate with CFTR and form the epithelial ATP-sensitive potassium channel, at least in the rat [112].

**Table 2.** Genetic differences between Bartter's and Gitelman's syndromes

Clinical syndrome	Gene locus	Chromosome
Bartter's syndrome		
Type-1	SLC12A1	15q15-q21
Type-2	KCNJ1	11q24
Type-3	CLCNKB	1p36
Gitelman's syndrome	SLC12A3	16q13

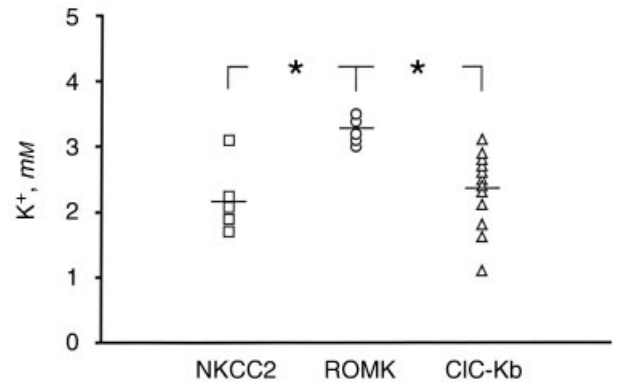
### Mechanism for basolateral Cl<sup>-</sup> transport in the TAL

Chloride is absorbed across the basolateral membrane of the TAL in part via chloride channels [119]. Three structural classes of chloride channels have been described: (1) CFTR (cystic fibrosis transmembrane regulator); (2) glycine and GABA<sub>A</sub> receptors; and (3) the CLC family of Cl<sup>-</sup> channels [120]. It is the last group of chloride channels that plays an important role in mediating basolateral Cl<sup>-</sup> transport in the TAL. Jentsch et al identified the first of the voltage-gated CLC genes (ClC-0) by expression cloning from the marine ray *Torpedo marmorata* [121]. Nine CLC family members (ClC-1 through ClC-7, ClC-Ka and ClC-Kb) are known to play a role in transport, volume regulation, and muscle and neuronal excitability in many tissues. ClC-Ka and ClC-Kb are expressed exclusively in the human kidney [122, 123]. Expression of ClC-Ka, ClC-Kb, and the rat homologues ClC-K1 and ClC-K2 has been localized by RT-PCR in various nephron segments. The precise localization throughout the nephron of these ClC chloride channels at the protein level is currently unknown [122, 124, 125].

### Molecular basis for Bartter's syndrome: mutations in NKCC2, ROMK, and ClC-Kb

Simon et al demonstrated linkage to the NKCC2 gene locus (SLC12A1) in 9 patients from five families with neonatal BS [86]. The authors mapped the gene to chromosome 15q15-q21 using a (GT)<sub>n</sub> dinucleotide repeat sequence. The NKCC2 protein is encoded by 26 exons, with the coding region spanning 80 kb of genomic DNA. Six mutations in NKCC2 were identified, including non-conservative missense and frameshift mutations. In the five families studied, all affected offspring were homozygous for alleles of markers used for genotyping. In four of the five families, affected individuals were offspring of a consanguineous union. The latter finding differs in patients with GS, who are most often compound heterozygotes and offspring of a non-consanguineous union, as was the patient discussed today. A possible explanation for the difference is that heterozygotes in the population with NCCT mutations are more common. We now refer to patients with BS due to NKCC2 mutations as having type-1 BS.

The genetic heterogeneity of neonatal BS was demonstrated recently (Table 2) [87, 126]. Some patients with neonatal BS have mutations of the ROMK potassium-



**Fig. 4.** The severity of hypokalemia in type-1 (NKCC2, □) versus type-2 (ROMK, ○) and type-3 (ClC-Kb, △) Bartter's syndrome mutations. Mean serum potassium in type-1 BS was  $2.2 \pm 0.24$  mEq/liter,  $3.3 \pm 0.08$  mEq/liter in type-2 BS, and  $2.4 \pm 0.12$  mEq/liter in type-3 BS. The data in patients with ROMK mutations were obtained from Derst et al [117]. The data in patients with NKCC2 mutations came from Simon et al [86]. The data in patients with ClC-Kb mutations were from Ref. 128. \*  $P < 0.001$ .

channel gene locus (KCNJ1) on chromosome 11q24. Non-conservative missense mutations, frameshift mutations, and mutations predicted to cause premature truncation of the protein were identified. We refer to patients with ROMK mutations as having type-2 BS. As I said earlier, 5 ROMK splice variants have been identified in human kidney, although their relative distribution in the distal portion of the nephron is unknown. All 5 isoforms have a common 372 amino acid core encoded by exon 5. The mutations described thus far affect exon 5 and involve all ROMK isoforms. Derst and colleagues analyzed the channel activity of 5 ROMK channel mutations by mutating wild-type rat ROMK1 transfected into COS-7 cells [117]. In cells expressing mutated ROMK1, channel currents were low to absent. Mutations in the N- or C-terminus had low but significant channel currents.

Given the importance of ROMK channel activity in mediating both K<sup>+</sup> recycling in the TAL and K<sup>+</sup> secretion in the CCD, one might expect that hypokalemia would be less severe in patients with BS who have ROMK mutations. Impaired K<sup>+</sup> secretion by the CCD would be predicted to ameliorate the K<sup>+</sup> wasting that occurs because of impaired TAL function. Furthermore, complete loss of ROMK function might not affect TAL transport to the same extent as mutations causing complete loss of function of NKCC2, because of the presence of 70 ps K<sup>+</sup> channels in the TAL, which mediate approximately 80% of the apical K<sup>+</sup> conductance under control conditions [110]. This channel might partially restore K<sup>+</sup> recycling in patients with ROMK defects. A comparison of patients with BS with NKCC2 mutations (type-1 BS) versus ROMK defects (type-2 BS) indicates that the difference in the degree of hypokalemia is an important distinguishing clinical feature in some patients (Fig. 4). In addition, it would be expected that patients with type-2 BS would have a lower transtubular

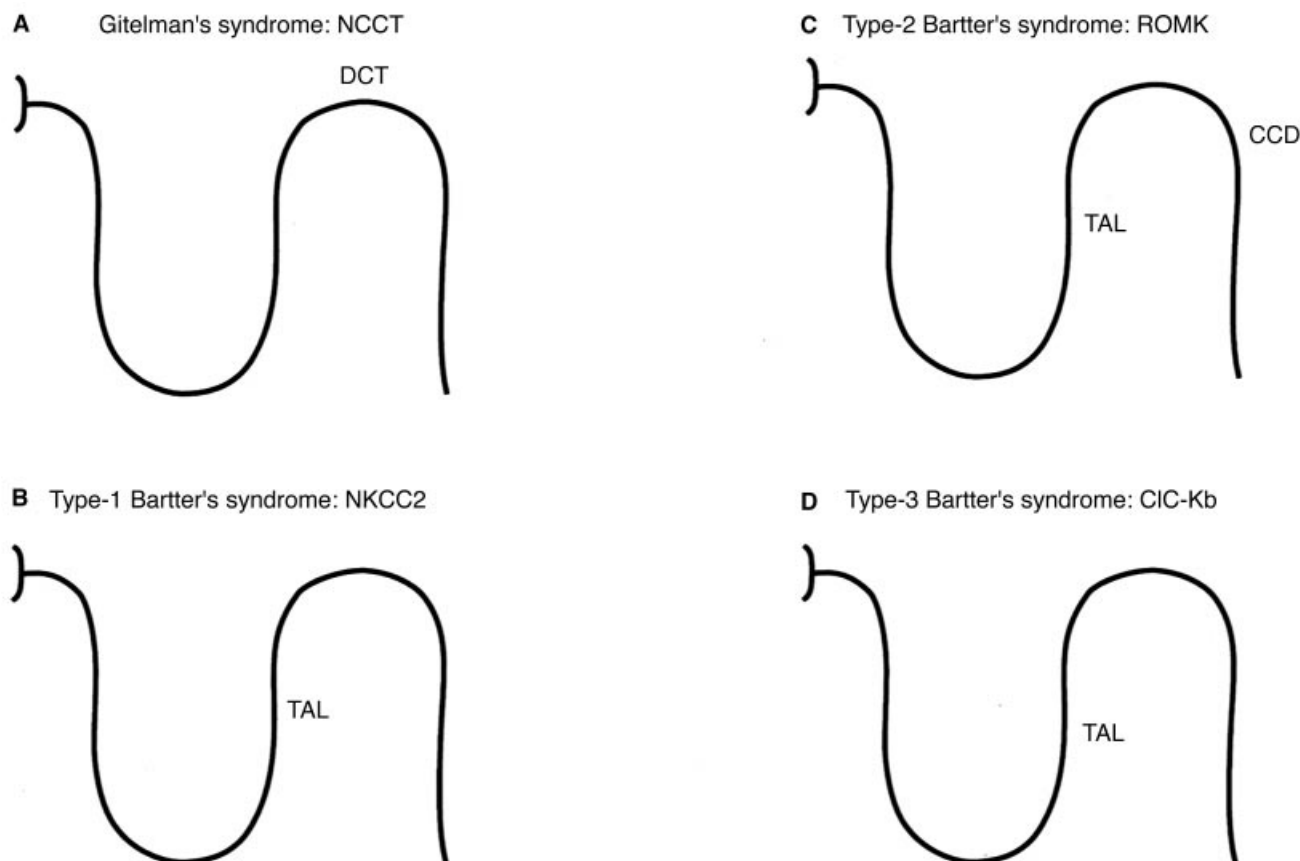


Fig. 5. Localization of transport abnormalities in Gitelman's and Bartter's syndromes.

potassium gradient (TTKG) and less of a decrease in urinary  $K^+$  excretion in response to amiloride administration. However, not all patients with ROMK mutations have mild hypokalemia [87]. In a study in rats by Wang et al, the  $K^+$  channel blocker U37883A inhibited  $Na^+$  and  $K^+$  absorption in the TAL and  $K^+$  secretion in the CCD; a natriuresis ensued without a change in urinary  $K^+$  excretion [127]. This could be an interesting model for studying the effect of certain lack-of-function ROMK mutations on blood and urine chemistries prior to the availability of ROMK knockout mice.

Simon et al recently reported that mutations in the *CLCNKB* gene locus on chromosome 1p36 encoding the CIC-Kb protein also lead to a BS phenotype (type-3 BS) [128]. Eleven different mutations including large deletions and missense and nonsense mutations were characterized in 17 kindreds. As shown in Figure 3, patients with type-3 BS have more severe hypokalemia than do patients with type-2 BS, and they have hypokalemia comparable to that in patients with type-1 BS. Interestingly, none of the patients with type-3 BS had nephrocalcinosis, and 6 of the 17 index cases had urine  $Ca^{2+}$ /creatinine ratios in the normal range. The absence of nephrocalcinosis distinguishes these patients from those with *NKCC2* or ROMK

defects. Whether patients with *CLCNKB* mutations have microscopic nephrocalcinosis detectable by renal biopsy remains to be determined. Of interest is the discovery that mutations in another member of the *CLC* family, *CLCN5*, causes X-linked hypercalciuric nephrolithiasis (XLHN) [129]. Unlike type-3 BS, these patients have nephrocalcinosis. The localization of the transport abnormalities in GS and BS is depicted in Figure 5.

#### Related uncharacterized syndromes

The literature contains reports of several patients with normotensive hypokalemic metabolic alkalosis but additional features that prevent them from being classified as having GS or BS. These disorders likely are due to different renal tubular transport defects. Runeberg and coworkers described a 6-year-old boy with tetany and arthritis in the knees and ankles [130]. He had hypokalemia, metabolic alkalosis, normal blood pressure, hypocalcemia, hypercalciuria, a high fractional excretion of magnesium, chondrocalcinosis, and nephrocalcinosis. Renal biopsy showed juxtaglomerular cell hyperplasia and interstitial fibrosis. Oral magnesium therapy slowed the progression of the nephrocalcinosis and decreased the intra-articular calcifications. Güllner et al reported 3 siblings, ages 10 to 14 years, who



presented with mild growth retardation, hypokalemia, metabolic alkalosis, normal blood pressure, elevated plasma renin activity, and resistance to angiotensin II [131, 132]. The serum magnesium was normal ( $1.51 \pm 0.06$  mEq/liter). One child had polyuria, polydipsia, and neuromuscular complaints, whereas the others were asymptomatic. The PGE<sub>2</sub> excretion was elevated in all the patients. The urinary calcium excretion was not measured. Clearance studies revealed normal distal fractional Cl<sup>-</sup> absorption, and the renal biopsy showed interstitial fibrosis and abnormalities restricted to proximal tubules without juxtaglomerular cell hyperplasia. Mehrotra et al recently described a 44-year-old man with hypokalemia, metabolic alkalosis, normal blood pressure, mild renal insufficiency, hypermagnesemia, normocalcemia, a low fractional excretion of magnesium, hypocalciuria, glucosuria, and elevated plasma renin activity and plasma aldosterone levels [133]. Clearance studies showed a high fractional delivery of Cl<sup>-</sup> to the diluting segment and a high distal fractional reabsorption of Cl<sup>-</sup>. The patient's appropriate response to furosemide and chlorothiazide administration suggested intact TAL and DCT function. A renal biopsy was not performed; however, a clinical diagnosis of chronic interstitial nephritis was made. The authors postulated a defect in proximal tubule NaCl absorption. Williams et al reported a neonate whose prenatal course was complicated by polyhydramnios [18]. The infant, who was volume depleted, required mechanical ventilation and developed generalized seizures at 6 days of age, had hypokalemia, hypomagnesemia, metabolic alkalosis, hypophosphatemia with renal phosphate wasting, polyuria, hypercalciuria, hypermagnesuria, glucosuria, severe hyponatremia, and adrenal insufficiency. Nephrocalcinosis was not present. The infant died at 4 months and autopsy revealed juxtaglomerular cell hyperplasia and extensive tubular dilation (primarily of the proximal tubules) with poorly developed brush borders. The authors suggested that a proximal tubule defect could account for the electrolyte abnormalities. It is of interest that despite a proposed defect in proximal tubule transport, bicarbonate absorption in this segment appears to have been spared, given that the patients are alkalemic rather than having proximal renal tubular acidosis. Whether mutations in KCC1 are responsible for any of these uncharacterized syndromes with abnormal proximal tubule transport is unknown.

### Summary of diagnostic considerations

As I said, patients with Gitelman's and Bartter's syndromes differ in their clinical presentation and laboratory findings (Table 1, Fig. 2). Patients with BS usually present as neonates; patients with GS present in their teens or as adults. Second, the predominant symptoms in BS are polyuria and polydipsia; patients with GS usually complain of muscle spasm and cramps. The degree of hypokalemia is

equally as severe in patients with GS and those with type-1 BS (NKCC2 defect) and type-3 BS (ClC-Kb defect). Patients with type-2 BS (ROMK defect) can have less severe hypokalemia. Differences in divalent cation metabolism clearly distinguish GS from BS. Patients with GS are hypomagnesemic and hypocalciuric. In contrast, patients with BS usually have a normal serum magnesium, are hypercalciuric, and develop nephrocalcinosis. Interestingly, patients with type-3 BS fail to develop nephrocalcinosis and are not uniformly hypercalciuric.

In summary, loss of function mutations in NCCT causes GS. In this disorder, there is defective NaCl transport in the distal convoluted tubule. Patients with BS (type-1, type-2, and type-3) have loss of function mutations in the transporters that are required for the normal absorption of Na<sup>+</sup>, K<sup>+</sup>, and Cl<sup>-</sup> in the TAL (NKCC2, ROMK, and ClC-Kb).

### QUESTIONS AND ANSWERS

DR. NICOLAOS E. MADIAS (*Chief, Division of Nephrology, New England Medical Center, Boston, Massachusetts*): Could you tell us whether any of the mutated proteins of the Na-K-2Cl cotransporter or the NaCl cotransporter have been expressed in cell systems, and whether the in-vitro findings correlate with disturbed function?

DR. KURTZ: At present, no studies have addressed the effect of specific mutations in type-1 BS on NKCC2 function in vitro. Derst et al have analyzed five recently described ROMK channel mutations causing type-2 BS [117]. The authors expressed mutated rat ROMK1 in COS-7 cells and were able to demonstrate the effect of specific type-2 BS mutations on channel currents. Whether these in-vitro findings will correlate with a particular clinical phenotype is unknown. The number of patients studied to date is too small to determine whether the severity of polyuria, hypokalemia, and hypercalciuria/nephrocalcinosis correlates with specific mutations. It is becoming increasingly evident that background genes also can alter a given phenotype, so it might be overly simplistic of us to expect a close correlation between a specific mutation and the clinical severity of BS in all patients [134–136]. With regard to GS, Kunchaparty et al have recently expressed in *Xenopus* oocytes the mouse NaCl cotransporter incorporating a GS mutation (premature stop codon at amino acid 968), which abolished function [137]. In addition recently we showed that peripheral blood mononuclear cells (obtained from the patient with GS discussed today) lack chlorothiazide-inhibitable <sup>22</sup>Na uptake [80]. Our further understanding of the physiologic processes in the kidney that result in the clinical differences between these disorders likely will depend on the creation of knockout mice with GS and BS phenotypes. This approach also will allow the role of background genes and several other candidate genes to be tested.

DR. MADIAS: Could you summarize for us the current

understanding of genotypic-phenotypic relationships in these disorders? To what do you attribute the phenotypic variability between patients with these syndromes? Also, could you address the abnormalities of heterozygote individuals?

DR. KURTZ: Mutations in the SLC12A3 gene locus are the only known cause of Gitelman's syndrome. Whether patients with chondrocalcinosis and arthritis as well as with the electrolyte abnormalities found in GS have SLC12A3 mutations will require further studies. In Bartter's syndrome, because three genes are known to be involved, one might expect a wider spectrum of clinical and laboratory abnormalities. In addition, no patients with GS have been found to have NKCC2, ROMK2, or ClC-Kb defects. Conversely, no patients with the clinical features of BS have NCCT mutations.

You raise the interesting issue of heterozygotes. One might speculate that these individuals have subtle but detectable salt wasting and potassium wasting, which generally is not apparent clinically. It is unknown, for example, whether in either syndrome heterozygotes have more negative salt balance when placed on a zero Na<sup>+</sup> intake, and more negative potassium balance when placed on a zero K<sup>+</sup> intake. Studies addressing these issues would be of interest.

DR. MADIAS: It would be interesting to see in expression studies whether a mutated protein might cause increase of function.

DR. KURTZ: It is certainly possible, although as yet unproven, that certain patients with hypertension have gain-of-function mutations. This issue is currently an active area of investigation. Virtually all transport proteins known to mediate Na<sup>+</sup> transport directly or indirectly in the kidney are being studied as candidate genes for gain-of-function mutations in patients with idiopathic hypertension.

DR. DAVID LEE (*Chief, Division of Nephrology, Sepulveda Veterans Administration Hospital, and Professor of Medicine, UCLA, Los Angeles, California*): Could you expand on the role that prostaglandins play in Bartter's syndrome? The neonatal form of Bartter's syndrome is sometimes called hyperprostaglandin E syndrome. In addition, do all patients respond to indomethacin, or only a subgroup?

DR. KURTZ: The role of prostaglandins in BS is controversial. Lüthy and colleagues have suggested that, in most patients with Gitelman's syndrome, prostaglandin E<sub>2</sub> excretion is normal and indomethacin is less effective in correcting the hypokalemia [57]. However, some patients originally diagnosed with BS in retrospect actually had GS; they had abnormal prostaglandin excretion and electrolyte abnormalities that were ameliorated by indomethacin [46, 67]. In BS, PGE<sub>2</sub> excretion usually is increased. As I mentioned, Seyberth et al suggested that the neonatal form of the disorder be called the hyperprostaglandin E syn-

drome [29]. Cyclo-oxygenase inhibitors do ameliorate many of the features of BS, although the electrolyte abnormalities are not completely normalized. Excess PGE<sub>2</sub> excretion in BS can directly inhibit Na-K-2Cl cotransport [100, 138], and arachidonic acid inhibits ROMK1 channels [139]. These effects would be predicted to exacerbate the underlying transport abnormalities in BS. Furthermore, as I discussed, PGE<sub>2</sub> might play a role in increasing 1,25(OH)<sub>2</sub>-vitamin D in BS, thereby possibly contributing to the hypercalciuria and nephrocalcinosis [47]. On the other hand, Delaney et al have shown that mepacrine, a phospholipase inhibitor, normalized PGE<sub>2</sub> excretion in their patients without altering body weight or plasma potassium [46]. This finding argues against a role for prostaglandins. Furthermore, Zusman et al reported a patient with a prostaglandin-producing renal cell carcinoma who did not have the electrolyte abnormalities of BS [64].

DR. MADIAS: You referred to other syndromes with high plasma prostaglandin levels. Normal pregnancy is such a situation. High prostaglandin levels are associated with elevated plasma renin and aldosterone, hyporesponsiveness to vasoconstrictors, and low urinary calcium excretion, that is, a physiologic model of GS.

DR. KURTZ: The high prostaglandin levels you describe would be characteristic of BS, whereas hypocalciuria is indeed reminiscent of GS. It is of interest that when hypocalciuria occurs in pregnancy, it is usually in the context of pre-eclampsia [140]. The mechanism does not appear to be decreased intestinal uptake, and likely it is mediated by the kidney [141]. Furthermore, in pre-eclampsia, the hypocalciuria is associated with low 1,25(OH)<sub>2</sub>-vitamin D levels [142]. Despite elevated urinary aldosterone excretion in pregnancy [143], hypokalemia, hypomagnesemia, and a chloride-sensitive metabolic alkalosis rarely are present.

DR. CHARLES KLEEMAN (*Attending Physician, West Los Angeles Veterans Administration Hospital, and Professor of Medicine, UCLA*): What drives the hyperreninemic hyperaldosteronemic state? What is the fundamental cause of this state in both GS and BS?

DR. KURTZ: It is currently believed that the elevated plasma renin activity is due to various degrees of volume depletion resulting from defective renal NaCl absorption. This scenario is analogous to that seen in patients taking diuretics, that is, mild volume depletion in GS and severe volume depletion in neonatal BS. In many patients, however, the aldosterone level is not elevated because of the suppressive effect of concomitant hypokalemia on adrenal aldosterone production.

DR. MADIAS: Is the ROMK channel expressed outside of kidney, and are ROMK mutations associated with pathology elsewhere?

DR. KURTZ: By Northern analysis, ROMK message has been detected only in the kidney [113]. However, with more

sensitive PCR techniques, one can detect ROMK isoforms in various extrarenal tissues [112, 113]. We do not know whether functional abnormalities in various ROMK channels outside the kidney play a role in the manifestations of type-2 BS.

DR. LEE: You mentioned that the thiazide-sensitive NaCl cotransporter is located in the distal tubule. Were the results you reported regarding expression of this transporter in lymphocytes performed at the protein level?

DR. KURTZ: Thus far we have shown by Northern analysis that a 4.4 kb transcript of the thiazide-sensitive NaCl cotransporter is present in human peripheral blood mononuclear cells [80]. We also recently demonstrated an absence of thiazide-inhibitable  $\text{Na}^+$  transport in cells obtained from 2 patients with GS [80]. In regards to Dr. Madias' previous question regarding extrarenal ROMK expression, one could speculate that the dermatitis reported in some patients with GS [2] is somehow related to abnormal lymphocyte NaCl cotransport.

DR. ARNOLD FELSENFELD (*Attending Physician, West Los Angeles Veterans Administration Hospital, Professor of Medicine, UCLA*): Dr. Kurtz, you made the analogy between the loop diuretics and BS, and you referred to the elevated calcitriol levels in BS. Does the literature contain data regarding elevation of calcitriol levels in patients receiving loop diuretics?

DR. KURTZ: Furosemide increases  $1,25(\text{OH})_2\text{D}$  in a chick renal tubule suspension, possibly via prostaglandin  $\text{E}_2$ -mediated stimulation of  $1\alpha$  hydroxylase activity [62]. In addition, chronic furosemide administration increases serum PTH, in contrast to thiazides, which decrease PTH levels [144–146]. I am unaware of any in-vivo data demonstrating an effect of furosemide on  $1,25(\text{OH})_2$  vitamin D. Your question addresses the more general issue as to whether the effects of loop diuretics mirror all the electrolyte and metabolic abnormalities of type-1 BS (NKCC2 mutations). One important and as-yet-unresolved finding is the absence of hypomagnesemia in patients with NKCC2 mutations despite the known inhibitory effect of loop diuretics on magnesium transport. It is perhaps significant that magnesium absorption in the thick ascending limb of Henle's loop is confined to the cortical portion [147]. Net magnesium transport in the medullary portion is essentially zero. Given the known distribution of NKCC2 isoforms that I discussed, if only those isoforms expressed in the medullary thick limb (NKCC2F and NKCC2AF) were mutated in type-1 BS, one might not expect to find abnormal loop magnesium transport, because cortical thick ascending limb function would remain unaffected. However, the NKCC2 mutations described thus far would be expected to affect the cortical and medullary thick ascending limb, as do loop diuretics. Therefore, isoform differences cannot explain the lack of hypomagnesemia in type-1 BS. In type-2 and type-3 BS, hypomagnesemia would also be expected.

In GS, the converse question arises. Thiazide diuretics do not characteristically cause more hypomagnesemia than do loop diuretics [148]. Therefore why is hypomagnesemia universally present in GS despite the lack of significant magnesium absorption in the distal convoluted tubule and collecting duct? One unproven explanation for the severe hypomagnesemia in GS is that defective sodium transport in the early distal convoluted tubule enhances sodium absorption more distally, resulting in an inhibition of putative basolateral  $\text{Na}^+/\text{Mg}^+$  exchange. It is unclear why thiazide diuretics don't cause more severe hypomagnesemia; the same inhibition of apical  $\text{Na}^+/\text{Mg}^{2+}$  exchange should occur. Future studies in knockout mice will address the one or more mechanisms for the lack of hypomagnesemia in BS and the presence of hypomagnesemia in GS. This is one of several unresolved issues that will be the focus of future investigations: (1) What are the mechanisms for normal serum magnesium in BS and hypomagnesemia in GS? (2) Why is  $1,25(\text{OH})_2$  vitamin D elevated in BS? (3) Can patients with NKCC2, ROMK, and ClC-Kb mutations be clinically distinguished? (4) Why do patients with type-3 BS not have nephrocalcinosis? (5) Are other transporters with loss-of-function mutations involved in BS (KCC1)? (6) Is GS with chondrocalcinosis also due to mutations in the SLC12A3 gene locus? (7) What are the mechanisms for differences in clinical manifestations among patients who are members of the same family?

DR. KLEEMAN: Has active transport of magnesium in the thick ascending limb of Henle been reported?

DR. KURTZ: Studies of mouse and rabbit cortical thick ascending limb have shown that magnesium transport is a passive process driven by the lumen-positive potential difference [147]. Hormonal stimulation of passive magnesium absorption results from increases in transepithelial voltage and paracellular permeability. Active magnesium transport has not been reported in the TAL.

DR. MADIAS: I have seen a few patients who, after gentamicin administration, have persistent hypokalemia, hypomagnesemia and magnesuria, chloride-resistant metabolic alkalosis, and high renin and aldosterone levels with normal renal function. Has anyone else here encountered similar patients?

DR. KURTZ: Many years ago, Keating et al reported magnesium wasting in approximately 5% of patients treated with aminoglycosides [149]. The transport abnormality abated weeks to months after the drug was discontinued. Histologically, the proximal tubule is affected, as in cis-platinum toxicity. However, an isolated proximal tubule defect would not be expected to result in severe magnesium wasting. It is possible that aminoglycosides are inhibiting TAL transport by their ability to bind to the basolateral  $\text{Ca}^{2+}$ -sensing receptor (CaR) [150].

*Reprint requests to Dr. I. Kurtz, UCLA Division of Nephrology, 10833 Le Conte Avenue, Room 7-155 Factor Building, Los Angeles, California 90095-1689, USA*

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