

# Control of phosphate balance by the kidney and intestine

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**Abstract** The prevention and correction of hyperphosphatemia are major goals of the treatment of chronic kidney disease (CKD)-bone mineral disorders, and thus, Pi balance requires special attention. Pi balance is maintained by intestinal absorption, renal excretion, and bone accretion. The kidney is mainly responsible for the plasma Pi concentration. In CKD, reduced glomerular filtration rate leads to various Pi metabolism abnormalities, and Pi absorption in the small intestine also has an important role in Pi metabolism. Disturbances in Pi metabolism are mediated by a series of complex changes in regulatory hormones originating from the skeleton, intestine, parathyroid gland, and kidney. In this review, we describe the regulation of type II sodium-dependent Pi co-transporters by the kidney and intestine, including the regulation of Pi transport, circadian rhythm, and the vicious circle between salivary Pi secretion and intestinal Pi absorption in animals with and without CKD.

**Keywords** Proximal tubule · Intestine · Salivary glands · PTH · FGF23 · 1,25(OH)<sub>2</sub>D

## Introduction

Inorganic phosphate (Pi) balance refers to the steady-state difference between Pi input and output over a period of time. When Pi intake exceeds Pi loss (positive balance), Pi accumulates and may lead to growth or weight gain, bone

mineralization, or repletion of stores. When Pi loss exceeds Pi intake (negative balance), Pi stores are progressively depleted, eventually resulting in clinical symptoms of deficiency. The Pi balance is maintained mainly by the kidney, bone, and small intestine. It is dependent on the absorption of dietary Pi in the intestine, and re-absorption and excretion of Pi in the kidney and skeletal pools. Absorbed Pi enters the extracellular space and is distributed in three compartments: blood, soft tissue, and bone [1]. The kidney is mainly responsible for the plasma Pi concentration [2]. In chronic kidney disease (CKD), reduced glomerular filtration rate (GFR) leads to various Pi metabolism abnormalities, and Pi absorption in the small intestine also has an important role in Pi metabolism [3]. Disturbances in Pi metabolism are mediated by a series of complex changes in regulatory hormones originating from the skeleton, intestine, parathyroid gland, and kidney [4]. In this review, we focus the regulation of sodium-dependent Pi co-transporters in a number of tissue, including kidney, intestine, and salivary glands.

## Control of renal Pi balance

Approximately 80% of filtered Pi is reabsorbed in the proximal tubule. Under normal conditions, ~15% of filtered Pi is ultimately excreted. The renal proximal tubules have a regulatory system that functions through sodium-dependent Pi co-transporters (NaPi transporter). At least two types of transporters (NaPi-II and NaPi-III) play important roles in renal Pi re-absorption [5]. The characteristics of NaPi transport function have been reported in detail by Murer's group [5]. In the present review, we described the regulation of NaPi-II transporters: NaPi-IIa (SLC34A1, Npt2a) and NaPi-IIc (SLC34A3, Npt2c),

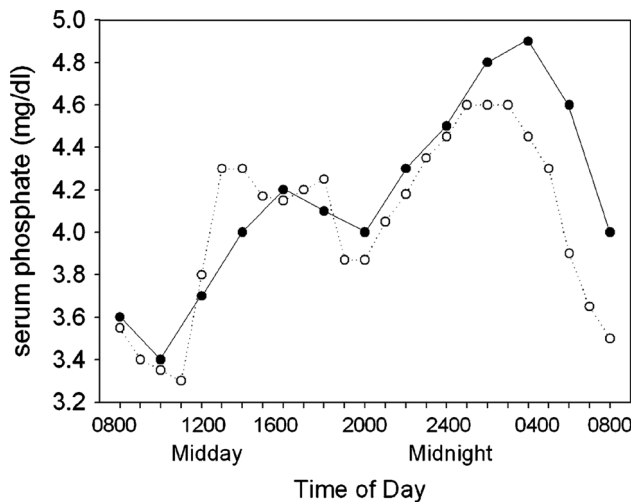
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because earlier gene knockout (KO) studies for renal Na/Pi co-transporters revealed the importance of NaPi-IIa and NaPi-IIc in renal Pi re-absorption [6].

### Many factors affect the Npt2a and Npt2c levels in the proximal tubules

1. Parathyroid hormone (PTH)/fibroblast growth factor 23 (FGF23)—recent excellent reviews have been published regarding the regulation of Pi transport by PTH and FGF23 [7, 8]. PTH is secreted by the parathyroid glands and regulates the endocytosis of NaPi-IIa and NaPi-IIc from the apical membrane of the proximal tubular cells [7, 8]. Scaffolding protein NHERF1 plays an important role in the regulation of NaPi-IIa by PTH and FGF23 [7, 8]. Recent studies demonstrated that the control of phosphorylation of sodium-hydrogen exchanger regulatory factor-1 NHERF1 (by cyclic AMP-dependent protein kinase, protein kinase C, extracellular signal-regulated kinase-1/2 and serine/threonine-protein kinase 1) is a trigger for endocytosis of NaPi-IIa from the apical membrane [7–9]. Furthermore, the bone secretes FGF23, which suppresses Pi re-absorption and vitamin D synthesis in the kidney. Klotho is an essential co-receptor for FGF23 signaling and downregulates the levels of NaPi-IIa protein in the apical membrane [10]. According to recent studies of site-specific klotho deletion in kidney, NaPi-IIa is regulated by FGF23/klotho signal in proximal and distal tubules [11, 12]. The physiologic roles of secreted klotho on phosphatidic activity, however, remain unknown. NaPi-IIc is also regulated by PTH and FGF23 [4, 13]. Our recent studies suggest that recycling endosomes are involved in NaPi-IIc endocytosis. The regulation of NaPi-IIc, however, is largely unclear.
2. Vitamin D—1,25-dihydroxyvitamin D ( $1,25(\text{OH})_2\text{D}$ ), the biologically active form of vitamin D, regulates NaPi-IIa and NaPi-IIc gene expression in the kidney [14, 15]. Administration of  $1,25(\text{OH})_2\text{D}$  to vitamin D-deficient rats upregulates the initial rate of renal Pi uptake [14]. Our previous studies demonstrated that the renal Pi uptake as well as the amounts of renal NaPi-IIa and NaPi-IIc protein are downregulated in the global vitamin D receptor (VDR)-KO mice at growth period [16, 17]. Despite the plasma PTH concentration is abnormally high in VDR-KO mice, renal Pi transport activity is not reduced in adult [16, 17]. The mechanisms of the PTH-resistance for Pi transport in VDR-KO mice remain unknown.
3. Vitamin A/thyroid hormone—the vitamin A metabolite ATRA (all-*trans*-retinoic acid) increases the rate of Pi transport in renal proximal tubular cells [15]. Masuda et al. identified retinoic acid-response elements in both gene promoters [15]. In addition, thyroid hormone increases Pi re-absorption by the elevation of renal NaPi-IIa transcription [18]. Ishiguro et al. identified a thyroid response element in NaPi-IIa gene by 3,5,3-tri-iodothyronine [18].
4. Acute inflammation—Ikeda et al. investigated lipopolysaccharide (LPS)-induced inflammation in mice [19]. LPS administration significantly decreased NaPi-IIa protein expression in the brush border membrane (BBM) after injection. Moreover, tumor necrosis factor- $\alpha$  injection also increased plasma PTH and decreased renal BBM NaPi-IIa expression [19]. The downregulation of NaPi-IIa expression in renal BBM through induction of plasma PTH levels alter Pi homeostasis during LPS-induced acute inflammation [19].
5. NAD/nicotinamide (NAM)—we previously reported that NAM inhibits intestinal and renal Na/Pi transport activity in rodents [20]. Treatment with NAM decreases serum Pi concentration in hemodialysis patients with hyperphosphatemia. Administration of NAM may suppress intestinal Pi absorption [20]. We reported that hepatectomy-induced hypophosphatemia is due to abnormal NAM metabolism, including nicotinamide phosphoribosyltransferase activation in the renal proximal tubular cells [21]. Thus, the liver–kidney axis may be involved in NAD and Pi metabolism [21].
6. Circadian rhythm—the previous studies indicated that the circadian rhythm of serum Pi is remarkably consistent in individuals [22–25]. These studies demonstrated that serum Pi is highest after midnight in normal subjects on normal diets (Fig. 1). In addition, circadian variations in urinary Pi excretion and serum Pi have long been recognized in humans and rodents [24]. The mechanisms involving these diurnal changes remain unclear. The serum Pi peak is observed after midnight between 02:00 and 04:00, and the Pi trough occurs between 08:00 and 10:00 [22, 23, 25]. Because prolonged fasting abolishes the nocturnal peak of Pi, intestinal Pi absorption clearly contributes to this circadian change of serum Pi [22]. Changes in PTH, growth hormone,  $1,25(\text{OH})_2\text{D}$ , or FGF23, however, cannot fully explain the circadian rhythm of serum Pi and the mechanism of this diurnal variation remains to be clarified [24]. Murer and co-workers reported that the diurnal changes in urinary Pi excretion are mainly related to changes in serum Pi and the tubular threshold, but not to NaPi-IIa expression [26]. The diurnal increase in renal Pi excretion is associated with a mild reduction in the sodium-dependent Pi transport rate in proximal tubular brush



**Fig. 1** Circadian variation of serum Pi levels in normal subjects. This figure is shown as Fig. 1 in Becker et al. [24]. The data from Jubiz et al. describe mean serum phosphate concentration in ten normal individuals (8 males, 2 females) aged 22–32 years (closed circles) [22]. Portale et al. reported similar findings in six healthy men aged 26–44 years, on a normal phosphate diet (open circle) [23]. Copyright permission is obtained from Oxford University Press

border membranes [26]. All of the mineral metabolism-associated hormones have a diurnal rhythm. Recently, Kawai et al. investigated the mechanism of the circadian clock and Pi metabolism in a rodent model [27]. They showed that skeletal FGF23 expression is regulated by the time of food intake, suggesting that the timing of food intake, and not the amount of ingested Pi, is a predominant determinant for the circadian profiles of skeletal FGF23 expression. In addition, a recent study in rodents demonstrated a nearly circadian rhythm for bone mineralization following fluctuations in the activity of the clock gene *Per1* [28, 29]. It is possible that variations in the bone mineralization process produce variations in individual serum Pi levels [28, 29]. There is no direct evidence that the bone–kidney axis regulates the circadian rhythm of Pi metabolism. In contrast, considering the importance of the timing of the meal, the intestine–kidney axis, rather than the bone–kidney axis, may provide important factors for the circadian rhythm of Pi metabolism. Further studies are needed to clarify the mechanism of the circadian rhythm of Pi metabolism.

### CKD and Pi excretion

Bricker and co-workers proposed the intact nephron hypothesis that sets the path for kidney research for the next 50 years [30, 31]. The hypothesis states that although the diseased kidney contains fewer nephrons,

the remaining nephrons are functionally normal. To maintain the homeostasis of any given solute, renal function of the diseased kidney must undergo adaptive changes, wherein the excretion rate of each functioning nephron must progressively increase to compensate for the damaged nephrons. There are a number of excellent reviews of Pi metabolism abnormalities in CKD [2, 32].

In the early stages of CKD, the plasma Pi concentration is maintained within the normal range by increase in the serum concentrations of PTH and FGF23, which increases urinary excretion of Pi, and decrease in serum active vitamin D concentration, which decreases intestinal Pi absorption [32]. At the late stage of CKD, the decline in GFR induces relative Pi retention with increased plasma Pi. Both FGF23 and PTH plasma levels are elevated in CKD and lead to a compensatory increase in fractional Pi excretion via inhibition of Pi transport to compensate for the decreased Pi filtration [32]. Experimentally, this is confirmed by a reduction in the NaPi-IIa expression in a classic model of advanced CKD [32]. Recent reports also indicate that FGF23 increases in parallel to albuminuria in immunoglobulin A nephropathy [33]. De Seigneux et al. showed that proteinuria indirectly modulates kidney Pi handling via a toxicity that decreases Klotho expression, which thereby alters FGF23 signaling, or via a direct competitive effect for endocytosis in the proximal tubules [33]. They demonstrated that albuminuria correlates with an increased plasma Pi concentration independently of GFR and that induction of glomerular proteinuria modifies kidney tubular Pi handling by altering FGF23 signaling. Thus, it is possible that many factors are involved in resistance to FGF23 [32].

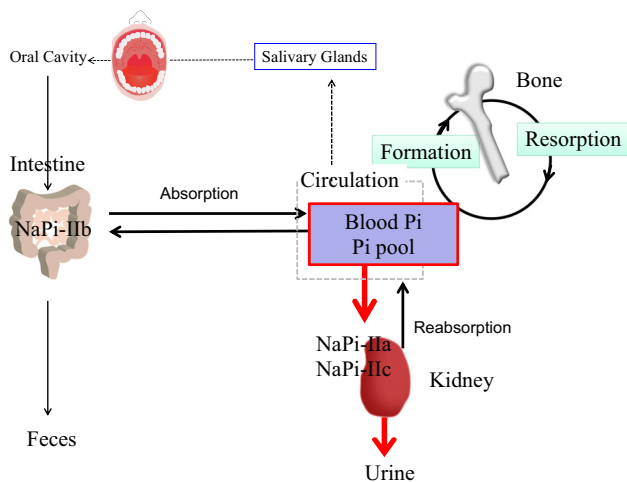
### Intestinal Pi handling

In adults, net Pi absorption typically ranges from 55 to 80% of the customary intake. Dietary Pi and 1,25(OH)<sub>2</sub>D are the most important physiologic regulators of intestinal Pi absorption [34]. There are two pathways for the intestinal Pi absorption, paracellular, and cellular, and at least two mechanisms, passive diffusion and sodium-dependent active transport [3]. The paracellular pathway occurs at tight junctions and utilizes electrochemical gradients. The cellular transport pathway requires sodium-dependent Pi transporters, including NaPi-IIb (SLC34A2, Npt2b), that are expressed in the small intestine [3]. The relative proportion of absorption via each mechanism varies depending on the luminal Pi concentration, with active transport contributing to between 30 and 80%. Intestinal Npt2b levels are regulated by FGF23, 1,25(OH)<sub>2</sub>D, and dietary Pi levels [35–37].

The effect of FGF23 on Npt2b levels in the small intestine is associated with FGF23-mediated decrease in  $1,25(\text{OH})_2\text{D}$  levels [38–40]. Indeed, injecting FGF23 into VDR-KO mice does not further decrease the levels of Npt2b in the small intestine [35]. In addition, the FGF23 injection had no effect on Npt2b levels in Npt2a-KO mice, Npt2c-KO mice, or Npt2a/Npt2c double KO mice, although FGF23 administration attenuates the increase of  $1,25(\text{OH})_2\text{D}$  levels in all these mutant mice [41]. Thus, FGF23-dependent  $1,25(\text{OH})_2\text{D}$  repression appears to be an important factor for Npt2b expression in the small intestine. Furthermore, additional factors may be involved in FGF23-dependent Npt2b downregulation.

### CKD and intestinal Pi absorption

The vicious circle between salivary Pi secretion and intestinal Pi absorption (Fig. 2)—saliva is an important organ for maintaining Pi homeostasis in CKD and dialysis patients [42]. The daily volume of saliva produced by the salivary glands is nearly one-fifth the total plasma volume. Based on the previous data and in accordance with the daily rate of saliva secretion, salivary glands seem to be able to secrete a relevant amount of Pi, possibly ranging from 300 to 600 mg [42]. Savica et al. reported that a high-Pi diet increases saliva Pi levels in CKD and hemodialysis patients [42]. In addition, in dialysis patients, salivary Pi levels correlate with serum Pi and multivariate analysis which confirms that serum Pi is the only independent factor predicting increased salivary Pi secretion [42]. The



**Fig. 2** Pi balance in the kidney, small intestine, and salivary glands. Pi balance is dependent on the absorption of dietary Pi in the intestine, and re-absorption and excretion of Pi in the kidney and bone pools. The ingestion of the Pi secreted in saliva and its subsequent absorption in the small intestine starts a vicious circle between salivary Pi secretion and fasting Pi absorption, thereby worsening hyperphosphatemia

ingestion of the Pi secreted in saliva and its subsequent absorption in the small intestine starts a vicious circle between salivary Pi secretion and fasting Pi absorption, thereby worsening hyperphosphatemia (Fig. 2). Therefore, salivary Pi binding could be a useful approach for reducing serum Pi levels in dialysis patients. Chitosan is a polymer of glucosamine, similar to sevelamer, which would bind Pi present in high concentration in the saliva of CKD patients [43]. Recent studies, however, have been unable to duplicate these results [44, 45]. A systemic analysis data concluded that the amount of chitosan contained in the chewing gum (20 mg) is too little to account for the originally observed reduction in serum Pi [43]. The development of effective Pi binders and the understanding of mechanisms of Pi handling in the salivary glands is essential. The saliva is primarily produced in the acinar compartment, where 85% of the proteins are ultrafiltered from the capillary bed adjacent to the glands. In the ductal system, saliva is converted from an isotonic to a hypotonic solution with lower sodium and chloride compared with the plasma [46]. Npt2b is also located in ductal cells and may be involved in the regulation of the salivary Pi concentration [47]. We speculate that cross-talk of Npt2b in the salivary glands and the small intestine is very important for Pi metabolism in the gastrointestinal tract. Understanding the mechanisms of the cross-talk between saliva and intestinal Npt2b with respect to the dietary Pi load may help us to resolve the Pi metabolism disturbances in CKD.

Intestinal Pi absorption—the physiology of the intestinal regulation of Pi absorption in humans is poorly understood and the regulators of intestinal Npt2b expression in CKD are uncertain. Experiments conducted using the 5/6 nephrectomy rat model revealed that intestinal Pi transport in the duodenum and jejunum remains unaltered [48]. Studies in mice with adenine-induced CKD also indicate that Pi uptake and Npt2b expression in the intestinal brush border membranes are not significantly altered [49]. Inhibitors of intestinal Npt2b and its molecular targets are expected to be effective for treating CKD-associated hyperphosphatemia [50]. Based on studies using Npt2b-KO mice, intestinal Npt2b is an important target for the treatment of CKD-associated hyperphosphatemia [50]. In a renal failure model of adenine administration in intestinal Npt2b-KO mice, the serum Pi concentration was significantly suppressed [50]. The effectiveness of Npt2b inhibitors in humans, however, is not known. Recent reports indicate that treatment with an NHE3 inhibitor is effective for altering intestinal Pi absorption and corrects hyperphosphatemia [51]. The mechanisms of Pi transport inhibition are not well understood [51].

On the other hand, NAM administration suppresses the expression of intestinal Npt2b and corrects

hyperphosphatemia in renal failure models [20]. As described earlier, marked hypophosphatemia is common after major hepatic resection, but the pathophysiologic mechanism remains unknown. We investigated a partial hepatectomy-related hypophosphatemia (PH) model and demonstrated that PH is due to abnormal NAM metabolism, including nicotinamide phosphoribosyltransferase activation in the renal proximal tubular cells [21]. In addition, PH results in marked reduction of intestinal Npt2b expression and possibly suppressed intestinal Pi absorption. We speculate that PH increases intestinal nicotinamide phosphoribosyltransferase activity and suppresses intestinal Npt2b levels. Studies of the NAM metabolism pathway in the liver–intestine axis may contribute to a better understanding of intestinal Pi handling.

### Future perspectives

Pi disturbances begin at an early stage in CKD and are associated with cardiovascular disease. In the course of CKD, FGF23, and PTH levels progressively increase, while Klotho and 1,25(OH)<sub>2</sub>D levels decrease. The following issues remain to be resolved: (1) The molecular mechanisms of the paracellular pathway for intestinal Pi absorption; (2) The factors that determine the circadian rhythm of the serum Pi concentration; (3) The vicious circle between salivary phosphate secretion and intestinal Pi absorption; (4) NAD and Pi metabolism in the liver–kidney axis; and (5) The mechanisms of cross-talk between renal NaPi-IIa (Npt2a) and NaPi-IIc (Npt2c) and intestinal NaPi-IIb (Npt2b).

Finally, the prevention and correction of hyperphosphatemia are major goals of the treatment of CKD-bone mineral disorders, and thus, Pi balance requires special attention. Further studies are necessary to clarify the mechanisms of disturbance of intestinal and renal Pi balance in CKD-bone mineral disorders.

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### Compliance with ethical standards

**Conflict of interest** The authors have declared that no conflict of interest exists.

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