Modulation of intestinal calcium and phosphate transport in young goats fed a nitrogen- and/or calcium-reduced diet

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Abstract

Feeding ruminants a reduced N diet is a common approach to reduce N output based on rumino-hepatic circulation. However, a reduction in N intake caused massive changes in Ca and inorganic phosphate (P_i) homoeostasis in goats. Although a single dietary Ca reduction stimulated intestinal Ca absorption in a calcitriol-dependent manner, a concomitant reduction of Ca and N supply led to a decrease in calcitriol, and therefore a modulation of intestinal Ca and Pi absorption. The aim of this study was to examine the potential effects of dietary N or Ca reduction separately on intestinal Ca and P_i transport in young goats. Animals were allocated to a control, N-reduced, Ca-reduced or combined N- and Ca-reduced diet for about 6–8 weeks, whereby N content was reduced by 25% compared with recommendations. In Ussing chamber experiments, intestinal Ca flux rates significantly decreased in goats fed a reduced N diet, whereas P_i flux rates were unaffected. In contrast, a dietary Ca reduction stimulated Ca flux rates and decreased P_i flux rates. The combined dietary N and Ca reduction withdrew the stimulating effect of dietary Ca reduction on Ca flux rates. The expression of Ca-transporting proteins decreased with a reduced N diet too, whereas P₁-transporting proteins were unaffected. In conclusion, a dietary N reduction decreased intestinal Ca transport by diminishing Ca-transporting proteins, which became clear during simultaneous N and Ca reduction. Therefore, N supply in young ruminant nutrition is of special concern for intestinal Ca transport.

key words: Flux rates of calcium and phosphate: Goats: Na⁺-dependent P_i transporter IIb: Transient receptor potential vanilloid channel type 6: Ussing chambers

Feeding ruminants a N-reduced diet is preferable for economic and environmental reasons. Dietary crude protein (CP) concentrations of 11-12% were recommended to meet the requirements of growing goats⁽¹⁾. A reduced dietary N supply was associated with a significant reduction in urinary N excretion⁽²⁾ due to increased expression of renal urea transporters, and thus greater renal urea re-absorption⁽³⁾ in goats. In addition, the urea transporting capacity of the ruminal epithelium was increased due to dietary N reduction⁽⁴⁾. By possessing such efficient recycling mechanisms, ruminants such as goats are able to maintain rumen microbes' N supply, and therefore a sufficient synthesis of microbial protein as the most important source for host protein, provided that energy supply to the rumen flora is also adequate for microbial protein synthesis.

As monogastric species do not have the similar potential to utilise N efficiently, a low-protein diet changes metabolic pathways seriously. A reduction of dietary protein leads to changes in Ca and inorganic phosphate (Pi) homoeostasis in monogastric animals and humans, including decrease in intestinal Ca absorption, reduced urinary Ca excretion and

diminished plasma calcitriol and insulin-like growth factor 1 (IGF1) concentrations⁽⁵⁻⁸⁾.

Despite the N-recycling mechanisms of ruminants, our own previous studies have shown that feeding an N-reduced diet with an adequate Ca supply to young goats resulted in a decrease in plasma calcitriol concentrations⁽⁹⁾. In addition, the concomitant reduction of dietary N and Ca also resulted in decreased plasma calcitriol concentrations and increased intestinal Ca and Pi absorption associated with decreased expression of the involved transport proteins including the apical transient receptor potential vanilloid channel type 6 (TRPV6), the major intestinal apical Ca channel, and Na⁺-dependent P_i transporter IIb (SLC34A2, NaPiIIb), the major intestinal P_i transporter in young goats^(10,11). In contrast, a solitary dietary Ca reduction stimulated renal calcitriol synthesis and intestinal Ca absorption in adult female goats⁽¹²⁾. Therefore, it was assumed that calcitriol, as it is known for monogastric animals, is a potent modulator of transcellular intestinal Ca and Pi transport in young goats. In monogastric species, several studies have shown that the expressions of the major intestinal

Abbreviations: BBM, brush-border membranes; CaBPDOK, Ca-binding protein D9K; CaR, Ca-sensing receptor; CP, crude protein; IGF1, insulin-like growth factor 1; IGF1-R, insulin-like growth factor 1 receptor; NaPiIIb, Na⁺-dependent P_i transporter IIb; PBST, PBS containing 0.1% Tween 20; P_i, inorganic phosphate; PiT1, Na⁺-dependent P_i transporter 1; PMCA, plasma membrane Ca²⁺ ATPase; TRPV6, transient receptor potential vanilloid channel type 6; VDR, vitamin D receptor.

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Ca transporting structures such as TRPV6, the intracellular Ca-binding protein D9K (CaBP_{D9K}) and the basolateral located plasma membrane Ca2+ ATPase (PMCA), as well as the most abundant intestinal P_i transporter NaPiIIb, were regulated by calcitriol^(13,14). Based on the known decrease in plasma calcitriol concentrations in response to an N-reduced diet in young goats, it was hypothesised that a dietary N reduction with a sufficient Ca supply modulates intestinal transcellular Ca and Pi absorption in young goats. Therefore, the aim of the present study was to determine Ca and Pi flux rates of intestinal epithelia in young goats during a single dietary N reduction compared with goats receiving diets sufficient in N and Ca content (control group), diets with reduced Ca content or diets insufficient in N and Ca contents. Therefore, the effects of a reduced dietary N supply on functional mechanisms in caprine intestinal epithelia were characterised. Furthermore, a more detailed characterisation of the modulation of transcellular and paracellular Ca and P_i transport processes in caprine intestinal epithelia was carried out.

In addition, it was assumed that potential changes in intestinal Ca and P_i absorption during dietary N reduction were based on altered expression levels of the described corresponding Ca and P_i-transporting proteins. For this reason, the expressions of *TRPV6*, *CaBP*_{D9K} and *PMCA*, as well of *NaPiIIb*, in caprine intestinal epithelia were examined. The expression and activity of intestinal basolateral Na^+/K^+ -*ATPase*, which provides the driving force for Na⁺-coupled P_i transport, were analysed. The molecular characterisation of an additional intestinal Na⁺-dependent P_i transporter 1 (*SLC20A1*, *PiT1*)⁽¹⁵⁾ and the determination of expressions of potential modulators of intestinal Ca and P_i transport, nuclear vitamin D receptor (*VDR*)^(16,17), Ca-sensing receptor (*CaR*)⁽¹⁸⁾ and insulin-like growth factor 1 receptor (*IGF1-R*)⁽¹⁹⁾, were carried out in goats' intestinal epithelia, representing further structures that might be affected by the dietary interventions.

Interestingly, in rats, a low-Ca diet induced a stimulation of active transcellular Ca absorption in the ileum^(20,21), which was thought to be incapable of active, vitamin D-dependent Ca transport. Therefore, we hypothesised that a reduction in dietary N and/or Ca supply, as applied in the present study, leads to a shift in intestinal Ca and/or P_i absorption sites, besides the fact that their predominant intestinal absorption sites are present in the proximal and mid-jejunum. Therefore, three different localisations of the goat small intestine were examined.

Methods

The protocols of the animal feeding and handling experiments were approved by the Animal Welfare Commissioner of the University of Veterinary Medicine Hannover (Hannover, Germany) and was in line with the German Animal Welfare Law.

Animals and feeding regimens

A total of twenty-six male, coloured, German goats (about 1-week old) were fed a commercial milk replacer for 6 weeks and were offered wheat straw *ad libitum* during this period. After weaning, all the animals were maintained on a pelleted control diet containing 21 % CP and 1 % Ca for 1 week to adapt

them to the pelleted diet. Subsequently, the goats with an initial weight of $16 \cdot 1$ (SEM $1 \cdot 76$) kg were allocated into four feeding regimens: (1) receiving a control diet (21 % CP, 1 % Ca), (2) an N-reduced diet (8 % CP, 1 % Ca), (3) a Ca-reduced diet (22 % CP, $0 \cdot 4 \%$ Ca) and (4) a combined N- and Ca-reduced diet (8 % CP, $0 \cdot 3 \%$ Ca) for 6–8 weeks. Goats of the same feeding regimen were housed together in groups of six (N–/Ca+ and N+/Ca–) or seven animals (N+/Ca+ and N-/Ca-) with water available *ad libitum*. The pelleted concentrates were fed 3 times/d, and the amount per animal was 70 g/kg⁰⁻⁷⁵. In addition, the animals received 25 % of the concentrate weight as chopped wheat straw. To estimate the mean intake of nutrients and minerals per animal, all offered and refused feeds were monitored daily. Animals were weighed weekly.

Diets

The feed content of DM, crude ash, crude fibre, crude fat and CP was determined by Weende analysis (proximate analysis), the standard procedure of the Verband Deutscher Landwirtschaftlicher Untersuchungs- und Forschungsanstalten⁽²²⁾. The amount of acid-detergent fibre and neutral-detergent fibre was measured by a method described by Van Soest *et al.*⁽²³⁾. The four diets were isoenergetic containing approximately 12.5 MJ metabolisable energy/kg DM. Table 1 shows the components and composition of the diets. To adjust the weight of reduced N diets, Sipernat 22S, a fine particle silica – which cannot be metabolised and which is commonly used as a digestibility marker due to its inert structure⁽²⁴⁾ – was added.

Blood and saliva samples

Blood samples (9 ml each) of the goats were collected shortly before slaughtering by puncturing the vena jugularis with both EDTA-coated and lithium heparinate-coated syringes as well as serum syringes (Sarstedt). Plasma was separated by centrifugation (2000 g at room temperature, 15 min). In addition, saliva samples were collected immediately after blood collection with a sponge from the oral cavity after 5 min of incubation according to the method of Boehnke *et al.*⁽²⁵⁾. Plasma, serum and saliva samples were stored at -20° C for subsequent analysis.

Intestinal tissue, ruminal and abomasal fluid samples

With finishing experimental feeding after 6–8 weeks, goats were slaughtered after captive bolt stunning by exsanguination. To avoid circadian effects, slaughtering was performed always at the same time in the morning. For technical reasons (Ussing chamber experiments), one goat/d was killed. To avoid significant time effects, animals from each feeding group were slaughtered in an alternating manner. Samples of the ruminal and abomasal fluid were collected during slaughtering.

Segments of the proximal and mid-jejunum as well as that of the ileum (50 cm each) were removed within 5-min postmortem, beginning 1-m distal from the pylorus, and were rinsed with ice-cold saline (0.9 % NaCl, w/v). Table 1. Components and composition of wheat straw and pelleted concentrate diets*

	Wheat straw	N+/Ca+	N–/Ca+	N+/Ca-	N–/Ca–
Components (as-fed basis) (g/kg)					
Soyabean meal	-	130	80	122	101
Urea	BDL	30	-	35	-
Wheat starch	-	376	366	434	432
Beet pulp	-	420	400	250	250
Molasses	-	-	10	54	60
Soyabean oil	-	10	31	28	35
Mineral-vitamin mix†	-	10	10	10	10
Sipernat 22S‡	-	-	69	50	95
MgHPO ₄ .3H ₂ O	-	9	10	10	10
NaH ₂ PO ₄ .2H ₂ O	-	7	7	7	7
CaCO ₃	-	16.5	17.5	-	-
Composition					
DM (g/kg)	911	881	886	898	897
Nutrients (g/kg DM)					
Crude ash	82.3	68·1	133-2	96.9	141.6
CP	36.2	209	82.4	216	78
Acid-detergent fibre	500.5	99.9	83.5	73·5	70.2
Neutral-detergent fibre	757.4	151	142.2	103.6	99.2
Crude fat	18.7	25	42.9	41.2	46.8
Urea	BDL	33.4	2.3	45.4	4.5
Ca	3.6	12.4	12.4	4	3.5
Р	0.9	4.9	4.7	4.8	4.7
Vitamin D ₃	ND	ND	ND	ND	ND
ME (MJ/kg DM)	8.0	13.3	11.9	13.1	11.8

BDL, below detection level; CP, crude protein; ND, not detected; ME, metabolisable energy.

* Composition expressed as fed (analysed by the Association of German Agricultural Investigation and Research Center).

† Mineral-vitamin mix per kg: 12 1 g Ca; 1 9 g Na; 2 2 g Mg; 400 mg (1 200 000 IU) vitamin A; 0.3 mg (12 000 IU) vitamin D3; 10 g vitamin E, 6335 mg Zn; 3000 mg Mn; 201 mg Co; 201 mg I; 15 mg Se.

‡ Sipernat type 22S (Evonik Industries AG) is a fine particle silica with high oil absorption capacity. It is widely used as a flow regulator, anti-caking and dusting agent especially in the food and feed industry.

For RNA isolation, preparation of crude membranes, brushborder membranes (BBM) and nuclear extracts, the mucosa of the middle section of each intestinal segment was stripped off and immediately frozen in liquid N_2 and stored at -80° C until further preparation.

For the Ussing chamber experiments, intestinal segments were opened along the mesenteric line, rinsed with ice-cold saline (0.9 %, w/v) and were maintained in a glucose-containing Krebs–Henseleit buffer solution aerated with carbogen (95 % O_2-5 % CO_2) until the epithelia were mounted in the Ussing chambers.

Incubation of epithelial tissues and measurement of intestinal calcium and inorganic phosphate flux rates in Ussing chambers

After serosal and muscle layers had been stripped from the mucosal layer, the intestinal epithelia were mounted between the two halves of the incubation chambers with an exposed serosal area of 1.13 cm². Thus, the chambers were separated into serosal and mucosal compartments. On both the sides, the intestinal tissue was incubated with 10 ml of a 38°C warm buffer solution that was continuously aerated with carbogen and maintained at pH 7.4. Both the buffers contained (mM) 113.6. NaCl, 5.4 KCl, 1.2 MgCl₂.6H₂O, 21.0 NaHCO₃, 1.2 CaCl.2H₂O, 1.2 NaHPO₄.2H₂O, 1.2 mannitol, 0.01 indomethacin and additionally 10.0 mM glucose and 7.0 mM-HEPES in the case of the serosal buffer and 20.0 mM-HEPES in the case of the mucosal buffer. After an equilibration time of 20 min, about

148 kBq of ⁴⁵Ca, ³²P and (³H)-mannitol (PerkinElmer GmbH) as radio isotopic tracers was added to each chamber to the serosal or mucosal side; (³H)-mannitol was used as a marker of the paracellular transport⁽²⁰⁾. At intervals of 15 min, samples were taken and immediately replaced by equal volumes of the respective buffer solution. Radioactivity of the samples was determined using a liquid scintillation counter (Wallac 1410; PerkinElmer GmbH). Unidirectional Ca, P_i and mannitol flux rates from the mucosal to the serosal (J_{ms}) and from the serosal to the mucosal (J_{sm}) sides were calculated from the rate of tracer appearance on the unlabelled side using standard equations⁽²⁶⁾. To determine net flux rates (J_{net}), J_{sm} were subtracted from respective J_{ms} of paired tissues.

In those chambers containing mid-jejunal epithelia, flux measurements were carried out before and after adding 10mm-Na⁺-arsenate (Sigma-Aldrich) to the mucosal side in order to characterise the transcellular part of the intestinal P_i transport by competitive inhibition of Na⁺-dependent P_i transport, with the mid-jejunum representing the major intestinal segment of Na⁺-dependent P_i transport⁽²⁷⁾.

Biochemical determinations

Plasma urea concentrations were measured using a commercial kit (R-Biopharm; inter-assay CV 3.6%; intra-assay CV 5.8%). Ionised Ca concentrations were determined in whole blood samples using an ion-sensitive electrode (Chiron Diagnostics GmbH; inter-assay CV 2%; intra-assay CV 1%). Concentrations of total Ca and inorganic P_i were measured colorimetrically in

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plasma, saliva, ruminal and abomasal fluids by standard spectrometric techniques^(28,29) (inter-assay CV 8.2% (Ca and P_i): intra-assay CV 4.9% (Ca), 1.7% (Pi)). Serum calcidiol concentrations were measured using a competitive ELISA kit (Immundiagnostik AG; inter-assay CV < 13.2%; intra-assay CV < 10.7 %). Calcitriol concentrations were measured using a commercial radioreceptor assay kit (Immundiagnostik AG; inter-assay CV < 20 and <15% for samples with calcitriol concentrations of 10 and 60 pg/ml, respectively; intra-assay CV < 15 and <10% for these two concentrations). The calcitriol assay had a detection limit of 2 pg/ml. Both calcidiol and calcitriol assays had been used before to determine respective hormone concentrations in goats^(10,17,30). Total plasma IGF1 and serum concentrations of thyroid hormones triiodthyronine (T3) and thyroxine (T4) were analysed in the Clinic for Cattle, Endocrinology Laboratory, University of Veterinary Medicine, Hannover, Germany, by ELISA and competitive chemiluminescence immunoassays, respectively (inter-assay CV 8.5% (IGF1); intra-assay CV 3.5% (IGF1), 7.0-13.2% (T3), 4·4-10·8% (T4)).

Total RNA isolation and reverse transcription

Total RNA was isolated using the RNeasy Mini-Kit (Qiagen) according to the manufacturer's protocol. The RNA concentrations

were measured by UV absorbance (BioPhotometer plus; Eppendorf AG). The quality and integrity of the extracted RNA were assessed using an RNA 6000 nanoassay for an Agilent 2100 Bioanalyzer (Agilent Technologies).

Using a random hexamere, oligo-dt primers and TaqMan Reverse-Transcription Reagents (Applied Biosystems), 200 ng of isolated RNA was reverse-transcribed for further analysis according to the manufacturer's protocol.

Intestinal expressions of TRPV6, CaBP_{D9K}, PMCA, NaPillb, PiT1, Na⁺/K⁺-ATPase, VDR, CaR and IGF1-R mRNA

For quantification of the expressions of glyceraldehyde-3phosphate dehydrogenase (*GAPDH*), *CaR*, *PMCA* and the *TRPV6*, caprine-specific TaqMan[®] primers and probes (Table 2) were purchased from TIB MOLBIOL. Reaction mixtures (20 µl) contained TaqMan Universal PCR Master Mix (Applied Biosystems), 300 nM specific primers, 100 nM specific probe and 16 ng reverse-transcribed complementary DNA (cDNA). The PCR product was amplified (50°C, 2 min; 95°C, 10 min; forty cycles of 95°C, 15 s and 60°C, 1 min) and analysed using a real-time PCR cycler (CFX96TM; Bio-Rad). Expressions of *CaBP*_{D9K}, *IGF1-R*, *Na⁺/K⁺-ATPase*, *NaPiIIb*, *PiT1* and *VDR* were determined using SYBR Green[®] PCR assays. For *CaBP*_{D9K}, *Na⁺/K⁺-ATPase* and *VDR*, specific primers (Table 3) were

Table 2. Primers and	probes	used for	TaqMan	assays
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Genes	Primers and probes $(5' \rightarrow 3')$	References
GAPDH	Sense: CAAGGTCATCCATGACCACTTT	Wilkens et al.(31)
	Antisense: CGGAAGGGCCATCCACA	
	FAM-CTGTCCACGCCATCACTGCCACCC-TMR	
CaR	Forward: ATCGGGTCCTCCTCGTGTTT	Elfers et al. ⁽¹⁸⁾
	Reverse: GGACTTGAAGGCGAAGAAGAA	
	FAM-TGTCATCTGTGCCATTTGGCTCAACAC-TMR	
PMCA	Sense: GGTATTGCTGGAACTGATGTAGCTAA	Wilkens et al. (32)
	Reverse: CGTCCCCACATAACTGCTTT	
	FAM-CAATGCTTGTAAAATTGTCATCCGTGAGA-BBQ	
TRPV6	Forward: TGATGGGAGACACTCACTGG	Wilkens <i>et al.</i> ⁽³¹⁾
	Reverse: GCAGCTTCTTCTCCAGCATC	
	FAM-TGGCTACAACCTGCGCCCT-BBQ	

GAPDH, glyceraldehyde-3-phosphate dehydrogenase; CaR, Ca-sensing receptor; TRPV6, transient receptor potential vanilloid channel type 6.

Genes	Primers $(5' \rightarrow 3')$	References
CaBP _{D9K}	Sense: GCCAAAGAAGGTGATCCAAA	This study
	Antisense: CCAACACCTGGAATTCTTCG	
IGF1-R	Forward: GGAGAAIAAICCAGICCIAGCACC	This study
		This study
Na /K -ATPase	Antisense: GCCACTCGGTCCTGATATGT	This study
NaPillb	Sense: CGGTCCAAAACAAAGTATGATCAAG	This study
	Reverse: AGCCAAAGGGGTAAGGGAA	
PiT1	Sense: ATTCATCCTCCGTAAGGCAGATC	This study
	Antisense: CAGCAATGGTGCTCCAGTATACA	
VDR	Sense: GCACTTCCTTACCTGACCCC Antisense: CCGCTTGAGGATCATCTCCC	Herm <i>et al.</i> ⁽⁵⁹⁾

CaBP_{D9K}, Ca-binding protein D9K; IGF1-R, insulin-like growth factor 1 receptor; NaPillb, Na⁺-dependent P_i transporter IIb; PiT1, Na⁺-dependent P_i transporter 1; VDR, vitamin D receptor.

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purchased from Life Technologies, and for IGF1-R, NaPiIIb and PiT1 primers were purchased from TIB MOLBIOL. Reaction mixtures (20 µl) contained KAPA SYBR FAST Universal Master Mix (PEQLAB Biotechnologie GmbH), 200 nm specific primers and 16 ng reverse-transcribed cDNA. PCR products were amplified (95°C, 3 min; forty cycles of 95°C, 10 s and 60°C, 30 s) and detected using a real-time PCR cycler (CFX96TM; Bio-Rad). The thermal profile for melt curve determination began with an incubation of 10 min at 55°C with a gradual increase in temperature (0.5°C/10 s). Absolute copy numbers were determined using calibration curves generated with cloned PCR fragment standards⁽³¹⁾. Specificity of the amplicons was verified by sequencing (GATC) and using NCBI Blast (http://blast.ncbi.nlm.nih.gov/Blast.cgi). Expression of genes of interest were normalised to GAPDH as a constant expressed housekeeping gene. The reactions were carried out twice and included no template control.

Intestinal expressions of TRPV6, CaBP_{D9K}, PMCA, NaPillb, PiT1, Na⁺/K⁺-ATPase, VDR, CaR and IGF1-R protein

The protein expressions of the described transport proteins were exclusively examined in the mid-jejunal epithelia. This was carried out with regard to the results of the Ussing chamber experiments, indicating the mid-jejunum to be the most important intestinal segment for Ca transport. Concerning P_i transport, the mid-jejunum was the segment showing the second highest net flux rates following the ileum in goats, underlying its physiological importance in intestinal P_i transport. Furthermore, our own preliminary experiments have shown protein expressions of *TRPV6* and *NaPiIIb* to be partly below the detection level in the proximal jejunum and ileum.

Mucosa samples of the mid-jejunum were homogenised in an ice-cold homogenisation buffer, and crude membranes as well as cytosolic proteins were isolated as described by Wilkens et al.⁽³²⁾. For isolating BBM fractions of the intestinal epithelia, a protocol that was previously described by Wilkens *et al.*⁽³¹⁾ was used. Nuclear extracts from the caprine mid-jejunum were prepared using a method described by Muscher et al.⁽¹⁷⁾. Protein concentrations of all the preparations were measured by the Bradford method (Bio-Rad). Immunoblot assays detecting the expressions of CaBPDOK, NaPiIIb, VDR and CaR proteins in the intestinal segments were performed as described elsewhere $^{(33,34,18)}$. In brief, 15–30 µg of cytosolic preparations (CaBP_{D9K}), BBM fractions (NaPiIIb), nuclear extracts (VDR) or crude membranes (CaR) were separated by SDS-PAGE (16% tricine-SDS-gel in the case of CaBP_{D9K}, 8.5 % SDS gel in the case of NaPiIIb, 10% SDS gel in the case of VDR and CaR) and transferred to nitrocellulose membranes (GE Healthcare) using a tank-blotting system (Bio-Rad). Specificity of the CaBP_{D9K} antibody (Swant; diluted 1:2500 in PBS containing 0.1% Tween 20 (PBST) and 5 % fat-free milk powder) in goats was validated by cross-reactivity with ovine tissue as reported by the manufacturer. The anti-NaPiIIb (gift from Professor Dr J. Biber, Institute of Physiology, University of Zurich, Switzerland; diluted 1:2000 in PBST) and anti-CaR (Enzo Life Sciences GmbH; diluted 1:250 in PBST containing 1% bovine serum albumin (BSA; Sigma-Aldrich)) antibodies were successfully

pre-incubated with the corresponding antigenic peptide⁽³⁴⁾ (data not shown). In the case of the VDR antibody (Enzo Life Sciences GmbH; diluted 1:500 in PBST), alignment with the corresponding recombinant protein verified the specificity of the detected band in caprine intestinal tissue (data not shown). For detecting TRPV6 protein content, BBM fractions (25 µg) were incubated at room temperature in a loading buffer containing 5 mM dithiothreitol (DTT) for 5 min and were separated by 8.5% SDS-PAGE and transferred to a nitrocellulose membrane. The TRPV6 antibody (Alomone; diluted 1:500 in PBST) was successfully pre-incubated with the corresponding antigenic peptide (data not shown). For abundance of PMCA, 40 µg of crude membrane fractions were separated by 7 % SDS-PAGE and transferred to a nitrocellulose membrane. For Na^+/K^+ -ATPase abundance, 10 µg of crude membranes were heat denatured in a loading buffer containing 5 mm-DTT for 20 min at 70°C and separated by 10% SDS-PAGE and subsequent transfer onto nitrocellulose membranes. For both antibodies (anti-PMCA (diluted 1:2000 in PBST) and anti-Na⁺/ K⁺-ATPase (diluted 1:10 000 in PBST) from Enzo Life Sciences), cross-reactivity with ovine tissue was reported by the manufacturer. For detecting PiT1 (anti-PiT1; Thermo Fisher Scientific: diluted 1:500 in PBST) and IGF1-R (anti-IGF1-R: New England Biolabs; diluted 1:500 in Tris-buffered saline (TBS) containing 0.1% Tween (TBST) and 5% BSA), 40 µg of BBM fractions (*PiT1*) or 50 µg of crude membrane fractions (*IGF1-R*) were incubated in loading dye containing 5 mm-DTT and separated on an 8.5% SDS-PAGE without previous heating in the case of PiT1 and after heat denaturation (70°C, 20 min) in the case of crude membrane fractions for IGF1-R detection.

Membranes were blocked overnight at 4°C in *PBST* and 5% fat-free milk powder. In the case of *IGF1-R*, *PBST* was replaced by *TBST*. Immunodetection of electrotransferred proteins was performed according to standard procedures. After washing with *PBST*, or with *TBST* in the case of *IGF1-R*, and incubating with the corresponding secondary antibody, the bound antibody was visualised using enhanced chemiluminescence (SuperSignal; Thermo Fisher Scientific) according to the manufacturer's protocol and ChemiDoc system (Bio-Rad).

Quantification of proteins was carried out using Quantity One software 4.4 and Image Lab 5.2.1 software (Bio-Rad). Values of the investigated proteins were normalised to the amount of β -actin (anti- β -actin, AC-15; Sigma-Aldrich) in the case of *TRPV6*, *PMCA*, *NaPiIIb*, *PiT1*, *Na*⁺/K⁺-ATPase, *VDR*, *CaR* and *IGF1-R*. In case of *CaBP*_{D9K}, *GAPDH* (anti-*GAPDH*; Merck Millipore) as the internal standard with a stable expression level was used.

Measurement of intestinal Na⁺/K⁺-ATPase activity

For measuring the activity of the basolateral located Na^+/K^+ -*ATPase*, 0.5 g of stripped intestinal epithelial tissue from the proximal and mid-jejunum as well as from the ileum was homogenised in 5 ml ice-cold homogenisation buffer containing 20 mM-Tris base, 250 mM-sucrose, 5 mM-sulphuric acid, 5 mMethylene glycol tetraacetic acid and 0.8 M-phenylmethanesulphonyl fluoride. After centrifugation (10 min, 600 **g**, 4°C) and resuspension of the supernatant, Na^+/K^+ -*ATPase* activity was measured following the method of Mircheff & Wright⁽³⁵⁾.

Statistical analysis

All the data are given as means with their standard errors if not stated otherwise and number of animals (n). Data were analysed using GraphPad Prism version 6.05 (GraphPad Software; www.graphpad.com) by two-way ANOVA with Tukey's multiple comparisons test.

Potential relationships between the measured parameters were analysed by Pearson's correlation and linear regression. For comparison of Ca and P_i net flux rates, as well as of Na^+/K^+ -*ATPase* activity in the different intestinal segments, one-way ANOVA was used; P < 0.05 was set to be significantly different, and P < 0.1 was used to define trends.

Table 4. Mean daily intakes of DM, concentrate, nitrogen, calcium and inorganic phosphate and feed efficiency of growing goats receiving a nitrogen and/or calcium-reduced diet (Mean values; number of animals)

Items	N+/Ca+	N–/Ca+	N+/Ca-	N–/Ca–
n	7	6	6	7
DM intake (g/d)	763	652	741	702
Concentrate intake (g/d)	691	614	661	642
Feed efficiency (kg/kg)	0.19	0.16	0.17	0.12
N intake (g/d)	21.19	7.71	20.85	7.69
Ca intake (g/d)	8.08	7.08	2.91	2.37
P _i intake (g/d)	3.11	2.66	2.97	2.73

Results

Intake, body weight and daily weight gain

The animals were clinically healthy throughout the study. Mean daily DM, concentrate, N, Ca and P_i intakes were estimated from group mean values for each animal. Feed efficiency was calculated as the difference between the final and initial weight divided by the estimated individual feed intake during this time period. The results are summarised in Table 4.

Daily energy supply and P_i supply of all the feeding groups and Ca supply in the (N+/Ca+) and (N-/Ca+) groups covered the recommendations of the Society of Nutrition Physiology (GfE) for young ruminating goat kids⁽¹⁾.

Daily weight gain was not affected by one of the dietary interventions and ranged from 0.15 to 0.18 (sem 0.04) kg/d (Table 5). Goats receiving an N-reduced diet had a significant lower final body weight of about 11% compared with goats in the (N+/Ca+) and (N+/Ca-) group (Table 5).

Blood parameters

The following paragraphs describe the results of the different blood parameters for the different feeding regimens, whereby results are described in detail where differences between the feeding regimens were determined. All the data are summarised in Table 6.

 Table 5. Effects of a reduced nitrogen and/or calcium diet on initial and final body weight and weight gain of young goats (Mean values with their pooled standard errors; number of animals)

							P: two-way ANOVA	
Items	N+/Ca+	N-/Ca+	N+/Ca-	N–/Ca–	SEM	N reduction	Ca reduction	Interaction
n	7	6	6	7				
Initial body weight (kg)*	15.6	15.5	16.6	16.7	1.76	0.97	0.22	0.91
Final body weight (kg)†	23.93	21.17	23.58	21.07	2.48	0.04	0.86	0.92
Body weight gain (kg/d)	0.18	0.15	0.16	0.15	0.04	0.24	0.55	0.62

* Initial body weight was determined as the weight at the beginning of experimental feeding at the age of 8 weeks.

+ Final body weight was determined as the weight at the time of slaughter at the age of 12-17 weeks.

Table 6. Effects of a reduced nitrogen and/or calcium diet on blood parameters of young goats (Mean values with their pooled standard errors; n 5 - 7 animals)

								P: two-way ANOVA		
Items	N+/Ca+	N–/Ca+	N+/Ca-	N-/Ca-	SEM	N reduction	Ca reduction	Interaction		
Total blood										
Ionised Ca (mм)	1.39 ^ª	1.25 ^b	1.37 ^a	1.18 ^b	0.05	<0.001	0.09	0.41		
рН	7.43	7.44	7.43	7.44	0.03	0.33	0.81	0.76		
Plasma										
Urea (mм)	7.10 ^a	0.83 ^b	6.29 ^a	1.60 ^b	0.58	<0.001	0.94	0.01		
Calcidiol (nm)	36.33 ^a	75.48 ^b	32.65 ^a	52.57 ^{a,b}	13.18	<0.001	0.06	0.16		
Calcitriol (pmol/l)	54·86 ^{a,b}	37.49 ^a	105·9 ^c	81.94 ^{b,c}	19.31	0.04	<0.001	0.74		
Са (тм)	2.40 ^{a,c}	2.19 ^{b,d}	2.35 ^{a,c,d}	2.11 ^b	0.09	<0.001	0.15	0.68		
IGF1 (nmol/l)	75.19	55.15	66.65	53.18	15.19	0.04	0.50	0.67		
Р, (mм)	2.30	2.37	1.86	2.72	0.43	0.04	0.84	0.08		
T3 (nmol/l)	2.29	2.43	2.25	1.83	0.42	0.53	0.14	0.19		
T4 (nmol/l)	44·13	56.84	48.69	55.77	8.04	0.02	0.67	0.49		

IGF1, insulin-like growth factor 1; T3, triiodthyronine; T4, thyroxine.

^{Lb.c,d} Mean values within a row with unlike superscript letters were significantly different; Tukey's multiple comparisons test (P<0.05).

able 7.	Effects	of a reduc	ed nitrogen	and/or calcium	diet on calciu	im and inorganic	phosphate	concentrations	in saliva,	ruminal	and at	oomasal	fluid i
oung g	oats												

(Mean values with their pooled standard	errors; <i>n</i> 4–7 animals)
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Items						P: two-way ANOVA			
	N+/Ca+	N-/Ca+	N+/Ca-	N–/Ca–	SEM	N reduction	Ca reduction	Interaction	
Saliva Ca (mm)	0.26	0.25	0.25	0.22	0.05	0.54	0.42	0.78	
Saliva P _i (m _M)	34.03 ^{a,b}	26.09 ^a	38·84 ^b	35.18 ^{a,b}	3.89	0.06	0.03	0.47	
Ruminal fluid Ca (mm)	0.79*	0.71	0.55†	0.86	0.26	0.38	0.75	0.14	
Ruminal fluid P _i (m _M)	37.29*	41.71	40·08†	38.54	11.80	0.81	0.98	0.62	
Abomasal fluid Ca (mm)	13.97 ^a	10⋅12 ^c	5.79 ^{b,} †	4·17 ^{b,} ‡	1.65	<0.01	<0.001	0.15	
Abomasal fluid P _i (m _M)	27.95 ^{a,b}	20.88 ^a	33·21 ^{b,} †	31-46 ^{b,} ‡	3.89	0.04	0.001	0.19	

^{a,b,c} Mean values within a row with unlike superscript letters were significantly different; Tukey's multiple comparisons test (P<0.05).

* For one animal, it was not possible to obtain ruminal fluid at the time point of slaughter (n 6).

+ For two animals, it was not possible to obtain ruminal and abomasal fluids at the time of slaughter (n 4).

‡ For one animal, it was not possible to obtain abomasal fluid at the time of slaughter (n 6).

Plasma urea concentrations decreased significantly due to dietary N reduction. Both total and ionised plasma Ca concentrations were significantly reduced in goats fed a reduced N diet and remained unchanged due to dietary Ca reduction. Plasma Pi concentrations were increased when dietary N supply was reduced and remained unaffected by dietary Ca reduction. Plasma concentrations of calcidiol increased (P < 0.001). whereas plasma calcitriol concentrations decreased (P=0.04) when dietary N was reduced (Table 6). In contrast, dietary Ca reduction tended to decrease plasma calcidiol concentrations (P=0.06; Table 6) and resulted in a significant increase in plasma calcitriol concentrations (P < 0.001; Table 6). Plasma IGF1 concentrations showed a significant decrease in goats fed the N-reduced diet. Serum concentrations of T4 significantly increased in goats fed the N-reduced diet and did not change due to a different dietary Ca supply.

Concentrations of inorganic phosphate and total calcium in saliva, ruminal and abomasal fluids

Salivary Ca concentrations remained unaffected by a reduction of dietary N and Ca, respectively (Table 7), whereas salivary P_i concentrations followed a trend of dropping due to the N-reduced feeding regimen and significantly increased due to dietary Ca reduction (Table 7). Concentrations of P_i and Ca in ruminal fluids remained unaffected by any of the feeding regimens (Table 7). In abomasal fluids, concentrations of P_i and Ca were significantly reduced due to dietary N reduction, whereby the reduction of dietary Ca content and a combined reduction of N and Ca even led to a greater decrease in the Ca concentrations and to an increase in P_i concentrations in abomasal fluids (Table 7).

Flux rates of calcium and inorganic phosphate across intestinal epithelia

Net flux rates (J_{net}) of Ca were the highest in the mid-jejunum (average of 22.61 nmol/cm²×h), followed by the proximal jejunum (average of 19.08 nmol/cm²×h) and the ileum (average of 2.20 nmol/cm²×h), with significant differences between the proximal jejunum and the ileum (P < 0.01; data not shown) as well as between the mid-jejunum and the ileum

(P < 0.001; data not shown). In all the three intestinal segments investigated, an N-reduced diet led to a significant decrease in Ca J_{net} flux rates (Table 8). In the proximal jejunum, the J_{ms} flux rates of Ca remained unaffected by a dietary N reduction, whereas J_{sm} flux rates were elevated due to the N-reduced feeding regimen, resulting in decreased J_{net} flux rates (Table 8). In the mid-jejunum, the decrease in Ca Jnet flux rates was due to significantly lower Ca Jms flux rates and unchanged Ca Jsm flux rates (Table 8). Furthermore, in the mid-jejunum, the reduction in dietary Ca content resulted in an increase in Jms flux rates of Ca without affecting the corresponding Jsm flux rates, leading to an increase in Ca J_{net} flux rates (Table 8). In the ileum, due to a significant interaction of N and Ca, the stimulating effect of a Ca-reduced diet on Ca Jnet flux rates seen in the N+/Cagroup, which was the only group showing significant J_{net} flux rates of Ca, was no longer detectable in the combined reduced feeding group (N-/Ca-; Table 8). Single dietary N reduction led to (numerically) reduced Ca absorption in this intestinal segment, which was not statistically significant different from the control group (N+/Ca+; Table 8).

The Na⁺-dependent P_i transport was measured in all the three intestinal segments too, and was the highest in the ileum (average of $293 \cdot 2 \text{ nmol/cm}^2 \times h$), followed by the mid-jejunum (average of $113.5 \text{ nmol/cm}^2 \times h$) and proximal jejunum (average of $17.43 \text{ nmol/cm}^2 \times h$). Net flux rates of P_i were significantly higher in the ileum compared with the proximal and mid-jejunum (P < 0.001; data not shown) as well as in the mid-jejunum compared with the proximal jejunum (P < 0.01; data not shown). The J_{net} P_i flux rates remained unaffected by dietary N reduction in the three intestinal segments investigated, whereas the reduction of dietary Ca content led to a significant decrease in J_{net} flux rates of P_i in the proximal (P=0.03) and mid-jejunum (P=0.03) (Table 8). The decrease in P_i net flux rates in the proximal and mid-jejunum was due to a significant decrease in unidirectional J_{ms} flux rates of P_i (P=0.03 and P = 0.02), whereas J_{sm} flux rates were not influenced (Table 8). The addition of Na+-arsenate to the mucosal side led to a 60 % decrease in J_{net} P_i in the mid-jejunum, irrespective of the feeding regimen (data not shown).

The applied feeding regimens had no impact on mannitol flux rates in any of the three intestinal segments investigated (Table 8). Significant correlations between J_{sm} mannitol and J_{sm}

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Table 8. Calcium, inorganic phosphate and mannitol (Man) flux rates of the different intestinal epithelia of young goats as affected by different dietary nitrogen and calcium supply†

(Mean values with their pooled standard errors; n 5 - 7 animals)

							P: two-way ANOVA	
Items	N+/Ca+	N-/Ca+	N+/Ca-	N–/Ca–	SEM	N reduction	Ca reduction	Interaction
Proximal jejunum								
J _{ms} Ca	51.19	45.70	54.89	45.67	15.17	0.34	0.81	0.81
J _{sm} Ca	25.96	38.98	25.38	31.83	9.07	0.04	0.40	0.48
J _{net} Ca	25.23*	6.73*	29.51*	13.83*	12.53	0.01	0.37	0.82
J _{ms} P _i	42.91 ^{a,b}	46.33 ^a	20·45 ^b	40.49 ^{a,b}	12.30	0.07	0.03	0.19
J _{sm} P _i	18.15	26.65	18·07	21.57	7.19	0.11	0.48	0.50
J _{net} P _i	24·77* ^a	19.68* ^{a,b}	2.38 ^b	18⋅91* ^{a,b}	10.06	0.26	0.03	0.04
J _{ms} Man	32.81	43.73	24.24	34.03	12.10	0.10	0.15	0.93
J _{sm} Man	23.76	29.77	23.11	26.03	9.09	0.34	0.63	0.74
J _{net} Man	9.04*	13.96*	1.13	7.99	7.72	0.14	0.09	0.80
Mid-jejunum								
J _{ms} Ca	51.04 ^{a,b}	40·41 ^a	70·04 ^b	48.44 ^{a,b}	12.81	0.02	0.05	0.40
J _{sm} Ca	28.39	32.36	29.96	27.77	7.24	0.81	0.68	0.40
J _{net} Ca	22.66* ^{a,b}	8.05 ^a	40·08* ^b	20.67* ^{a,b}	13.00	0.02	0.03	0.72
J _{ms} P _i	163.70	157-2	93.07	111.0	46.53	0.81	0.02	0.61
J _{sm} P _i	18.95	23.03	18·48	18.87	6.42	0.49	0.48	0.57
J _{net} P _i	144.7*	134.2*	74.6	92·16*	47.88	0.89	0.03	0.56
J _{ms} Man	43.95	41.09	40.31	35.25	7.41	0.30	0.21	0.77
J _{sm} Man	22.52	26.45	24.79	23.01	6.27	0.74	0.85	0.37
J _{net} Man	21.42*	14.64*	15.52*	12.24*	7.62	0.20	0.29	0.65
lleum								
J _{ms} Ca	17·33 ^a	22·37 ^a	36·77 ^b	19·76 ^a	4.09	0.008	0.001	<0.001
J _{sm} Ca	16.83	23.76	21.42	22.96	5.58	0.15	0.50	0.35
J _{net} Ca	0.51 ^a	-1·39 ^a	15·35* ^b	-3·20 ^a	4.59	<0.001	0.01	0.002
J _{ms} P _i	255-1	286.0	378.0	294.2	113.50	0.65	0.26	0.32
J _{sm} P _i	6.31	10.42	7.53	7.79	2.86	0.14	0.63	0.19
J _{net} P _i	248.8*	275.6*	370.4*	286.5*	114.22	0.62	0.26	0.34
J _{ms} Man	12.52	18.27	17.89	22.62	8.83	0.25	0.28	0.91
J _{sm} Man	11.01	16.80	12.59	13.19	3.28	0.06	0.54	0.13
J _{net} Man	1.51	1.47	5.30*	9.42	8·15	0.62	0.16	0.62

a.b Mean values within a row with unlike superscript letters were significantly different; Tukey's multiple comparisons test (P<0.05).

* Significantly different from zero (one sample *t* test).

+ J_{ms}: mucosal-to-serosal flux rate in nmol/cm² x h; J_{sm}: serosal-to-mucosal flux rate in nmol/cm² x h; net flux rates in nmol/cm² x h (J_{net}=J_{ms}-J_{sm}).

Ca flux rates (Fig. 1(a)-(c)) as well as between J_{sm} mannitol and $J_{sm} P_i$ flux rates (Fig. 2(a)–(c)) in all the three intestinal segments investigated are shown by linear regression. A positive correlation could also be shown between J_{ms} flux rates of Ca and mannitol in the ileum (P = 0.008, r 0.52; Fig. 1(c)) and between J_{ms} flux rates of P_i and mannitol in the proximal jejunum (P < 0.001; r 0.74, Fig. 2(a)). A weak positive correlation was shown for Ca J_{ms} flux rates with corresponding mannitol flux rates in the mid-jejunum (P=0.03; r 0.42, Fig. 1(b)) and for J_{ms} of P_i and mannitol flux rates in the mid-jejunum (P=0.04; r 0.40, Fig. 2(b)). No correlation could be detected between Ca J_{ms} flux rates and corresponding mannitol flux rates in the proximal jejunum (data not shown) or for the J_{ms} of P_i and mannitol flux rates in the ileum (data not shown). In the proximal and mid-jejunum, Jnet flux rates of Pi correlated negatively with calcitriol plasma concentrations (data not shown; P(prox.) = 0.01, r = 0.50; P(mid) = 0.048, r = 0.40).

Intestinal expressions of TRPV6, CaBP_{D9K}, PMCA, NaPillb, PiT1, Na⁺/K⁺-ATPase, VDR, CaR and IGF1-R mRNA

The integrity of the isolated RNA of all intestinal epithelia, which were used for quantitative PCR, expressed as RNA integrity number was at least 8.5 (data not shown).

In all the three intestinal segments investigated, the combination of dietary N and Ca reduction (N-/Ca-) led to a lower expression of TRPV6 mRNA in comparison with the N+/Ca- group (Tables 9-11), and therefore withdrew the expression-stimulating effect detected in the N+/Ca- group compared with the N+/Ca+ group. This interaction between the two feeding regimens tended to be significant in the proximal jejunum (Table 9) and was significant in the mid-jejunum and ileum (Tables 10 and 11). Due to this interacting effect, the mRNA expression of TRPV6 was not significantly reduced in the N-/Ca+ group compared with the N+/Ca+ group in the intestinal segments investigated, although numerically TRPV6 mRNA expression was lower in the N-/Ca+ group (Tables 9-11). In contrast, in all the three intestinal segments investigated, animals fed the Ca-reduced diet (N+/Ca-) showed significantly higher expression levels of TRPV6 mRNA compared with the control group (Tables 9-11). For the mRNA expression of intracellular CaBP_{D9K}, a reducing effect of dietary N reduction could be detected in the proximal and mid-jejunum (Tables 9 and 10). In the ileum, an interacting effect of N and Ca reduction was detectable, leading to the withdrawal of the expression-stimulating effect of dietary Ca reduction on CaBP_{D9K} mRNA expression in the combined reduction group (N-/Ca-; Table 11). Expression of the basolateral located PMCA (a)

150

8

20

40

Mannitol flux rate (nmol/cm²×h)

60

80

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Fig. 1. Linear regression of unidirectional mucosal-to-serosal (J_{ms} ; \bullet of (N+/Ca+), (N-/Ca+) and (N-/Ca-); J_{ms} ; \bullet of (N+/Ca-), (J_{ms} ; \bullet , ——), (J_{ms} ; \bullet , ——) or serosal-to-mucosal (J_{sm} ; O, ------) flux rates of calcium with the corresponding mannitol flux rates in proximal jejunum (a) (J_{sm} Ca=(0.89 ± 0.14), J_{sm} mannitol=(8.30 ± 3.79); r^2 0.64, P < 0.001), mid-jejunum (b) (J_{sm} Ca=(1.12 ± 0.06), J_{sm} mannitol=(2.51 ± 1.42); r^2 =0.94, P < 0.001; J_{ms} Ca=(0.633 ± 0.37), J_{ms} mannitol=(19.16 ± 15.08); r^2 =0.18, P=0.03) and ileum (c) (J_{sm} Ca=(1.06 ± 0.22), J_{sm} mannitol=(9.40 ± 4.89); r^2 =0.47, P=0.008) of goats fed different nitrogen and calcium supply. Calculations are only given when significance was obtained by linear regression. Regression line is only presented when Pearson's r > 0.50.

was decreased in the mid-jejunum due to the reduction of dietary N (P=0.05; Table 10), and it tended to increase as a result of dietary Ca reduction (P=0.06; Table 10). There was no



Fig. 2. Linear regression of unidirectional mucosal to serosal (J_{ms}; , _____) or serosal to mucosal (J_{sm}; O,) flux rates of inorganic phosphate (P_i) with the corresponding mannitol flux rates in proximal jejunum (a) (J_{sm} P_i=(0.71±0.08), J_{sm} mannitol=(2.68±2.26); r^2 =0.76, P<0.001), J_{ms} P_i=1.04±0.20), J_{ms} mannitol=(4.17±6.42); r^2 =0.55, P<0.001), midjejunum (b) (J_{sm} P_i=(0.98±0.04), J_{sm} mannitol=-(3.49±0.97); r^2 =0.96, P<0.001; J_{ms} P_i=(2.74±1.29), J_{ms} mannitol=(28.09±53.00); r^2 =0.16, P=0.04) and ileum (c) (J_{sm} P_i=(0.76±0.07), J_{sm} mannitol=-(2.21±0.98); r^2 =0.83, P<0.001) of goats fed different nitrogen and calcium supply. Calculations are only given when significance was obtained by linear regression. Regression line is only presented when Pearson's r>0.50.

detectable effect of the different diets on *PMCA* expression in the proximal jejunum and ileum (Tables 9 and 11).

The mRNA expression of *NaPiIIb* was unaffected by a dietary reduction of N and/or Ca in the proximal and mid-jejunum

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Table 9. Relative amounts of CaBP Dek, CaR, IGF1-R, Na⁺/K⁺-ATPase, NaPillb, PiT1, PMCA, TRPV6 and VDR mRNA expression normalised to GAPDH in the proximal jejunum of goats fed a nitrogen- and/or calcium-reduced diet (Mean values with their pooled standard errors; number of animals)

Items	N+/Ca+	N-/Ca+	N+/Ca-	N-/Ca-	SEM	P: two-way ANOVA		
						N reduction	Ca reduction	Interaction
n	7	6	6	7				
TRPV6	0.026 ^a	0.005 ^a	0.085 ^b	0.030 ^a	0.02	<0.001	<0.001	0.08
CaBP _{D9K}	8.71 ^{a,b}	5·87 ^{a,c}	10.98 ^b	7.34 ^{a,b}	2.35	0.01	0.13	0.74
PMCA	0.21	0.18	0.21	0.19	0.05	0.30	0.95	0.64
NaPillb	0.013	0·26e - 03	0.48e-03	0.67e-03	0.01	0.34	0.36	0.33
PiT1	0.18e-02	0·14e-02	0.16e-02	0·15e-02	0·27e – 03	0.15	0.94	0.32
Na ⁺ /K ⁺ -ATPase	0.086	0.090	0.10	0.099	0.02	0.86	0.17	0.74
VDR	0.057	0.055	0.059	0.055	0.007	0.38	0.76	0.80
CaR	0·12e – 02	0.11e-02	0·2e – 02	0·2e-02	0.6e - 03	0.81	0.01	0.83
IGF1-R	0·27e-02	0·26e - 02	0·27e-02	0·23e – 02	0·53e – 03	0.30	0.54	0.58

GAPDH, glyceraldehyde-3-phosphate dehydrogenase; TRPV6, transient receptor potential vanilloid channel type 6; CaBP_DGK, Ca-binding protein D9K; NaPillb, Na*-dependent Pi transporter IIb; *PIT1*, Na⁺-dependent P, transporter 1; *VDR*, vitamin D receptor; *CaR*, Ca-sensing receptor; *IGF1-R*, insulin-like growth factor 1 receptor. ^{a.b.c} Mean values within a row with unlike superscript letters were significantly different; Tukey's multiple comparisons test (*P*<0.05).

Table 10. Relative amounts of CaBP_Dek, CaR, IGF1-R, Na+/K+-ATPase, NaPillb, PiT1, PMCA, TRPV6 and VDR mRNA expression normalised to GAPDH in the mid-jejunum of goats fed a nitrogen- and/or calcium-reduced diet (Mean values with their pooled standard errors; number of animals)

Items	N+/Ca+	N-/Ca+	N+/Ca-	N-/Ca-	SEM	P: two-way ANOVA		
						N reduction	Ca reduction	Interaction
n	7	6	6	7				
TRPV6	0.55e – 02 ^a	0.89e-04 ^a	0.044 ^b	0.28e - 02 ^a	0.02	0.01	0.03	0.05
CaBP _{D9K}	1.19	0.034	1.21	0.039	0.80	0.01	0.98	0.98
PMCA	0.089	0.083	0.155	0.087	0.04	0.05	0.06	0.09
NaPillb	0.059	0.096	0.087	0.092	0.05	0.41	0.64	0.53
PiT1	0.36e - 02	0.38e - 02	0.28e-02	0·37e – 02	0.75e – 03	0.18	0.30	0.34
Na ⁺ /K ⁺ -ATPase	0.007	0.081	0.084	0.087	0.01	0.27	0.10	0.52
VDR	0.026	0.026	0.024	0.027	0.006	0.56	0.84	0.67
CaR	0·57e – 03	0.62e - 03	0·16e-02	0·11e-02	0.43e-03	0.67	0.01	0.51
IGF1-R	0·35e – 02	0-40e-02	0·38e-02	0·37e-02	0.60e-03	0.54	0.96	0.24

GAPDH, glyceraldehyde-3-phosphate dehydrogenase; TRPV6, transient receptor potential vanilloid channel type 6; CaBPDak, Ca-binding protein D9K; NaPillb, Na*-dependent Pi transporter IIb; PiT1, Na*-dependent P, transporter 1; VDR, vitamin D receptor; CaR, Ca-sensing receptor; IGF1-R, insulin-like growth factor 1 receptor. ^{a b} Mean values within a row with unlike superscript letters were significantly different; Tukey's multiple comparisons test (P < 0.05).

Table 11. Relative amounts of CaBP_Dek, CaR, IGF1-R, Na⁺/K⁺-ATPase, NaPillb, PiT1, PMCA, TRPV6 and VDR mRNA expression normalised to GAPDH in the ileum of goats fed a nitrogen- and/or calcium-reduced diet (Mean values with their pooled standard errors; number of animals)

Items	N+/Ca+	N-/Ca+	N+/Ca-	N-/Ca-	SEM	P: two-way ANOVA		
						N reduction	Ca reduction	Interaction
n	7	6	6	7				
TRPV6	0.80e - 03 ^a	0.36e - 03 ^a	0.011 ^b	0.11e-02 ^a	0.002	<0.001	<0.001	<0.001
CaBP _{D9K}	0.24e-04 ^a	0·25e – 04 ^a	0·22e – 03 ^b	0.42e-04 ^a	0.53e-04	0.004	0.001	0.004
PMCA	0.11	0.10	0.12	0.12	0.03	0.60	0.33	0.75
NaPillb	0.057	0.076	0.16	0.14	0.06	0.94	0.01	0.57
PiT1	0.018	0.015	0.017	0.016	0.002	0.24	0.86	0.39
Na+/K+-ATPase	0.20	0.16	0.23	0.21	0.05	0.23	0.10	0.64
VDR	0.027	0.024	0.037	0.032	0.01	0.38	0.06	0.78
CaR	0.082	0.082	0.042	0.073	0.05	0.51	0.29	0.49
IGF1-R	0·22e-02	0·24e-02	0·29e - 02	0·22e-02	0·58e – 03	0.44	0.48	0.16

GAPDH, glyceraldehyde-3-phosphate dehydrogenase; TRPV6, transient receptor potential vanilloid channel type 6; CaBPDak, Ca-binding protein D9K; NaPillb, Na*-dependent Pi transporter IIb; PiT1, Na*-dependent Pi transporter 1; VDR, vitamin D receptor; CaR, Ca-sensing receptor; IGF1-R, insulin-like growth factor 1 receptor.

^{a,b} Mean values within a row with unlike superscript letters were significantly different; Tukey's multiple comparisons test (P<0.05)

(Tables 9 and 10), but increased in the ileum due to dietary Ca reduction (P=0.01; Table 11). A reduction of dietary N and/or Ca had no impact on the mRNA expression of the other potential PiT1 in the intestinal segments investigated (Tables 9-11). Expression patterns of the Na^+/K^+ -ATPase differed between the feeding groups in the ileum, where a Ca-reduced diet tended to increase the mRNA expression (P = 0.10; Table 11). In the case of *VDR* mRNA expression, a stimulating effect could be detected in the ileum due to a dietary Ca reduction (P = 0.06; Table 11), but no effect was identifiable in the proximal and mid-jejunum (Tables 9 and 10). The mRNA expression of *CaR* increased in the proximal and mid-jejunum in the Ca-reduced feeding groups (Tables 9 and 10) but remained unaffected in the ileum (Table 11). No effect of the diets was detectable concerning mRNA expression of the intestinal *IGF1-R* in the intestinal segments investigated (Tables 9–11).

Intestinal expressions of TRPV6, CaBP_{D9K}, PMCA, NaPillb, PiT1, Na⁺/K⁺-ATPase, VDR, CaR and IGF1-R protein in the mid-jejunum

Regarding the protein expression of *TRPV6*, a significant interaction of dietary N and Ca reduction was detectable, whereas N+/Ca– goats showed increased *TRPV6* protein expression, demonstrated as a significant effect of dietary Ca reduction, but in combination with a concomitant N reduction (N–/Ca–) this stimulating effect was withdrawn (Fig. 3(a)).

For protein expressions of both the intracellular $CaBP_{D9K}$ and the basolateral *PMCA*, a decreased expression could be detected in cytosol preparations ($CaBP_{D9K}$) and crude membranes (*PMCA*), respectively, due to the N-reduced feeding (Fig. 3(b) and (c)).

The reduction of dietary N and Ca led to a lower expression of NaPiIIb in the N-/Ca- group compared with the N+/Ca- and the N-/Ca+ groups (not statistically significant), and therefore to a significant interaction of both N- and Ca-reduced feeding regimens (P=0.01). Individually, dietary N or Ca reduction had no impact on the protein expression of NaPiIIb (Fig. 3(d)). Protein expression of PiT1 was not affected by the diets (Fig. 3(e)). Protein expression of basolateral Na^+/K^+ -ATPase was reduced when dietary Ca content was reduced (Fig. 3(f)), whereas a dietary N reduction had no impact on the protein expression of the Na^+/K^+ -ATPase. Expression of VDR protein was not affected by dietary N and/or Ca reduction (Fig. 3(g)). Protein expression of CaR was significantly reduced in goats fed N-reduced diets (Fig. 3(h)), whereas no effect of dietary Ca could be detected. The IGF1-R protein expression showed no differences between the feeding groups (Fig. 3(i)).

Intestinal Na⁺/K⁺-ATPase activity

Activity of the basolateral located Na^+/K^+ -ATPase in the proximal jejunum, mid-jejunum and ileum are presented in Table 12. The Na^+/K^+ -ATPase activity diminished in the mid-jejunum due to dietary N reduction (P=0.02), whereas there was no effect of the diets in the proximal jejunum and the ileum. In principle, the activity of the Na^+/K^+ -ATPase in the proximal and mid-jejunum was significantly higher compared with the activity in the ileum (P < 0.001; data not shown).

Discussion

The aim of the present study was to determine separately the effects of dietary N and Ca reduction on intestinal Ca and $P_{\rm i}$

absorption in young goats and the potential overlapping effects of a concomitant reduction of N and Ca, as well as the characterisation of the underlying molecular mechanisms. It has been shown for the first time that a dietary N reduction under adequate dietary Ca supply modulates intestinal Ca absorption in the caprine small intestine.

The limitations of the present study were the group feeding of animals and a reduction of final body weights of the goats fed the N-reduced diets. Nevertheless, energy supply was sufficient for all animals independent of the feeding group according to the recommendations of the GfE. Similar weight gain as well as unaffected plasma T3 concentrations indicated sufficient energy intake of all goats, with T3 representing an energy-dependent hormone, which dropped due to energy deprivation in adult sheep⁽³⁶⁾. Elevation of plasma T4 concentrations due to dietary N reduction cannot be explained yet, but also indicates sufficient energy supply of the animals⁽³⁶⁾.

In all the three caprine intestinal segments investigated, with the proximal and mid-jejunum representing the major absorption sites for Ca in small ruminants^(11,37), Ca net flux rates diminished due to the N-reduced diet. As net flux rates of mannitol, a marker of paracellular transport⁽²⁰⁾, remained unchanged due to dietary interventions, the N-reduced diet with or without additional Ca reduction most probably led to a decrease of active transcellular Ca transport in the small intestine. This assumption is also corroborated by the fact that J_{ms} flux rates of Ca did not or only weakly correlate with corresponding mannitol flux rates in the proximal and midjejunum (Fig. 1(a) and (b)). This is comparable with the results from studies on rats fed protein-reduced diets, which had a significantly decreased uptake of Ca into intestinal BBM vesicles (BBMV), potentially based on altered expressions of apical Ca channels⁽⁷⁾. In contrast, in the N+/Ca- group, active transcellular Ca absorption was stimulated in the mid-jejunum and ileum similar to goats fed the Ca-restricted diet⁽¹²⁾ and in rats fed a low-Ca diet^(20,21). The fact that in the ileum only J_{net} Ca of the N+/Ca- group was significantly different from zero (Table 8) indicated the increase in epithelial transporting capacity, and therefore an extended absorption site for Ca at least in vitro due to this dietary intervention. Plasma Ca concentrations demonstrated that goats fed an adequate N supply were able to compensate for the low Ca intake by increasing absorption efficiency, in contrast to goats fed the N-reduced diet, showing significantly reduced total and ionised plasma Ca concentrations (Table 6), which was seen in young goats fed N- and Ca-reduced diets too^(9,10).

In former studies by Muscher *et al.*⁽¹¹⁾, a simultaneous dietary N and Ca reduction led to an increase in intestinal Ca absorption in the mid-jejunum of young goats. This may have been due to a lower body weight gain, and therefore lower Ca requirement of goats, in the control group in this former study, which was then stimulated by reducing the amounts of Ca and N.

The underlying molecular mechanisms for the changes in intestinal Ca absorption during dietary change could be due to the altered expressions of the involved transporting molecules of the transcellular Ca transport. A reduction of dietary N led to decreased expression of the apical Ca channel *TRPV6* revealed by two-way ANOVA and in combination with a reduced dietary

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Fig. 3. Semi quantification of (a) transient receptor potential vanilloid channel type 6 (*TRPV6*), (b) Ca-binding protein D9K (*CaBP*_{D9K}), (c) plasma membrane Ca²⁺ ATPase (*PMCA*), (d) Na⁺-dependent P_i transporter IIb (*NaPillb*), (e) Na⁺-dependent P_i transporter 1 (*PiT1*), (f) Na^+/K^+ -*ATPase*, (g) vitamin D receptor (*VDR*), (h) Ca-sensing receptor (*CaR*) and (i) insulin-like growth factor 1 receptor (*IGF1-R*) protein expression in mid-jejunum of goats receiving a dietary nitrogen and/or calcium reduction. Values are means, with their standard errors represented by vertical bars. * Significant effects between the single groups revealed by Tukey's multiple comparisons test after two-way ANOVA (*P*<0.05). *GAPDH*, glyceraldehyde-3-phosphate dehydrogenase (\Box , Ca+; \blacksquare , Ca–).

Table 12. Intestinal Na^+/K^+ *ATPase* activity in young goats as affected by different dietary nitrogen and calcium supply (Mean values with their pooled standard errors; n 6 - 7 animals)

Items	N+/Ca+	N-/Ca+	N+/Ca-	N–/Ca–	SEM	P: two-way ANOVA		
						N reduction	Ca reduction	Interaction
Proximal jejunum	49.80	42.60	53.90	42.70	11.19	0.11	0.71	0.73
Mid-jejunum	54.60	42.90	54·10	43.0	9.12	0.02	0.97	0.95
lleum	24.90	27.40	26.80	25.60	5.15	0.80	0.99	0.47

Ca supply led to significantly decreased expression compared with solitary Ca reduction, whereas the latter stimulated TRPV6 expression in all the intestinal segments. These findings were in accordance with results from studies with young goats fed a simultaneous N- and Ca-reduced diet, showing reduced TRPV6 expression in mid-jejunum⁽¹¹⁾, and adult goats on a Ca-restricted diet having increased expression levels of $TRPV6^{(12)}$. Expression of the intracellular $CaBP_{D9K}$ diminished in all the intestinal segments due to the reduced dietary N content. The almost missing effect of a dietary Ca reduction on CaBP_{D9K} expression in the anterior parts of the small intestine had been shown in adult goats fed a Ca-restricted diet too⁽¹²⁾. This, therefore, indicates that $CaBP_{D9K}$ is not the rate-limiting step of transcellular Ca transport or that increased expression of TRPV6 was not associated with an exceeding Ca influx into the enterocyte, requiring an increased intracellular buffer capacity. The expression of the basolateral PMCA was reduced due to N-reduced feeding exclusively in the mid-jejunum, which can partly explain the reduced Ca absorption in this intestinal segment. A single dietary Ca reduction tended to increase PMCA expression in the mid-jejunum, comparable with results of adult goats fed Ca-restricted diets, showing no change in PMCA expression in the jejunum⁽¹²⁾. It might be that changes in pump activity or an activation of additional extrusion systems, such as the Na⁺ Ca exchanger 1, which was shown to be expressed in sheep intestinal epithelia⁽³¹⁾, contribute to basolateral Ca extrusion.

All the described changes of expression patterns can be explained by reduced plasma calcitriol concentrations in goats fed N-reduced diets, assuming calcitriol-mediated regulation of expression via the *VDR* and vitamin D responsive elements (VDRE) similar to monogastric species^(13,38,39). In addition, changes in the expression of *VDR* are an important mechanism to modulate the responsiveness of target tissues of calcitriol⁽⁴⁰⁾. Therefore, the slightly (not statistically significant) increased *VDR* mRNA expression in ileal tissues of N+/Ca– goats in the present study, probably based on increased calcitriol concentrations as reported for ileal tissues of rats after treatment with calcitriol⁽⁴¹⁾, might be one reason for the increased expression of Ca-transporting proteins and the transporting capacity in the ileum in these animals.

Even the concomitant reduction of N and Ca led to a decrease in calcitriol content (not statistically significant) compared with solitary dietary Ca reduction and additionally to increased calcidiol concentrations. This is in line with our former studies^(9–11). The decrease in plasma calcitriol concentrations in goats fed N-reduced diets could be based on diminished plasma IGF1 concentrations (Table 6). IGF1 is a potential modulator of renal 1*a*-hydroxylase expression and/or activity^(42,43), and therefore conversion of calcidiol to calcitriol. By binding to the renal *IGF1-R*, which had been detected in bovine and ovine renal tissues^(44,45), IGF1 increased 1*a*-hydroxylase activity in a Ca-dependent manner in cell cultures⁽⁴⁶⁾.

A possible link between mineral homoeostasis and protein metabolism is CaR, expressed in the basolateral membrane of the small intestine of monogastric species^(47,48). In this study, expression of CaR mRNA was increased in the proximal and mid-jejunum during dietary Ca reduction, and protein expression in the mid-jejunum was decreased due to dietary N reduction. Thus far it is not known whether the intestinal CaR is able to modulate intestinal Ca absorption as CaR in the kidneys⁽⁴⁹⁾. In rats, an up-regulation of *CaR* could be shown in the parathyroid glands and kidneys through a supraphysiological calcitriol application⁽⁵⁰⁾. Furthermore, a *VDR*E was identified in the promoter region of the human $CaR^{(51)}$. Assuming the presence of VDRE in caprine CaR too, this could be an explanation for the changes in expression of intestinal CaR in the present study and potential evidence of a modulating effect of CaR on intestinal Ca absorption.

Due to the unchanged *IGF1-R* expression, it is an unlikely candidate linking protein and Ca metabolism in young goats. More likely, decreased plasma IGF1 concentrations could be associated with decreased intestinal Ca absorption in the goats fed the N-reduced diets in the present study, assuming a positive correlation between these two parameters as it was shown in men⁽¹⁹⁾ with an unknown underlying molecular mechanism.

Reduced intestinal Jnet Pi flux rates in the proximal and midjejunum during dietary Ca reduction were based on a decrease in active transcellular and paracellular Pi absorption. This was verified by a positive correlation between $J_{ms} P_i$ and J_{ms} mannitol in both the intestinal segments (Fig. 2(a) and (b)). The highest P_i absorption, which was mainly transcellular, indicated by the missing correlation between $J_{ms} P_i$ and mannitol (Fig. 2(c)), could be measured in the ileum independently from the feeding regimen, which was shown for adult sheep too⁽⁵²⁾. This might be explained by an ileal pH of 8.0, which led to a shift in the equilibrium constant of P_i to a more divalent P_i (HPO₄^{2–}), which is preferably transported by NaPiIIb. In our previous study, Jnet Pi were increased in the mid-jejunum of young goats due to a simultaneous reduction of dietary N and Ca⁽¹¹⁾. The discrepancy between these former results and the present data could be based on about 50 % lower dietary P feed content, and therefore lower daily P intake of the goats in the previous study, which might have stimulated intestinal P_i absorption. This was further supported by low but potentially adapted equal plasma P_i concentrations in all goats in the previous study, which may indicate that in the

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N–/Ca– group the stimulated P_i absorption was a compensatory mechanism to restore previously lower P_i plasma concentrations.

In the present study, goats fed Ca-reduced diets showed no change in plasma P_i concentrations (Table 6), indicating that the remaining intestinal P_i absorption probably was sufficient or compensatory mechanisms such as releasing P_i and Ca from bones were induced (K Elfers, A Liesegang, MR Wilkens, G Breves and AS Muscher-Banse, unpublished results), and therefore high intestinal P_i absorption was not required. Decreased intestinal P_i absorption based on sufficient P_i mobilisation from bone to maintain physiological P_i plasma concentrations has already been shown in goats, whose Ca homoeostasis was challenged by lactation (J Richter, B Schröder and MR Wilkens, unpublished results).

Transcellular P_i transport in the small intestine of goats was mainly mediated by a Na⁺-dependent P_i co-transport mechanism^(27,34,52), which was confirmed by successful inhibition with Na⁺-arsenate. Decreased P_i flux rates in the proximal and mid-jejunum during dietary Ca reduction were not based on changes in NaPiIIb expression. An additional likewise Na⁺-coupled P_i transporter is *PiT1*, which was expressed in rats' duodenal and jejunal BBM⁽¹⁵⁾ and has been shown for the first time in caprine intestinal epithelia in the present study. The expression of *PiT1* was not changed in any intestinal segment investigated, and therefore seemed not to be responsible for reduced Jnet Pi during Ca reduction. It could be speculated that in the proximal and mid-jejunum decreased P; net flux rates were potentially based on changes in activity of NaPiIIb and/or PiT1. Both NaPiIIb and PiT1 depend on an intracellular-directed Na⁺-gradient that is generated by the basolateral Na^+/K^+ -ATPase. However, Na⁺/K⁺-ATPase mRNA expression remained constant in all the three intestinal segments independently from dietary changes (Tables 9-11), and although Na⁺/K⁺-ATPase protein expression was decreased due to dietary Ca reduction in the mid-jejunum (Fig. 3(f)) no effects on activity were observed. Therefore, reduced intestinal P_i absorption in the proximal and mid-jejunum during dietary Ca reduction was probably not based on modulation of Na^+/K^+ -ATPase. In goats maintained on dietary N reduction, Na⁺/K⁺-ATPase activity diminished in the midjejunum. This was previously shown in rats fed low-protein diets and was supposed to be connected with lower amounts of apical Na⁺-coupled transport processes⁽⁵³⁾. In the present study, no changes in intestinal Pi flux rates during N reduction could be observed, indicating that changes in pump activity did not result in a smaller Na⁺-gradient, and thus a lower driving force for apical, Na⁺-coupled transport mechanisms. However, the decreased pump activity might also be explained by a shift to more Na⁺-independent intestinal amino acid absorption⁽⁵⁴⁾ during dietary N reduction, and therefore a decreased necessity for ATP-driven Na⁺-extrusion from the cell. This is supported by the fact that during dietary N reduction changes in plasma amino acid composition were observed in young goats (AS Muscher-Banse and K Huber, unpublished results)⁽⁵⁵⁾, potentially indicating altered ruminal microbial protein production, and therefore altered intestinal amino acid composition and uptake.

In contrast to monogastric species, where protein abundance of *NaPiIIb* was up-regulated by calcitriol^(13,56), in the anterior parts of the small intestine of goats in this study, *NaPiIIb* expression was not stimulated by calcitriol, which was in accordance with our previous studies⁽³⁴⁾. Even more contrary to monogastric species, increased plasma calcitriol concentrations of the goats fed Ca-reduced diets in the present study were associated with reduced intestinal Pi absorption in the proximal and mid-jejunum. This was confirmed by the negative correlation between J_{net} P_i and plasma calcitriol concentrations, indicating a regulatory relationship between these parameters. Regarding regulation of PiT1 expression in monogastric species, unchanged expression of PiT1 in the intestine during a Pi-restricted diet and high plasma calcitriol concentrations was shown in rats⁽¹⁵⁾. Decreased protein expression of the Na⁺/K⁺-ATPase in the mid-jejunum of goats fed Ca-reduced diets could be explained by increased plasma calcitriol concentrations in these animals, as it was shown in murine intestinal epithelia after calcitriol administration⁽⁵⁷⁾. Taking into account the involvement of the Na^+/K^+ -ATPase in the formation of tight-junction proteins, an inhibition of the Na^+/K^+ -ATPase might have led to increased epithelial permeability, comparable with studies in cell culture⁽⁵⁸⁾. A greater permeability of the epithelium might have increased paracellular Ca absorption, and therefore the overall amount of intestinal Ca absorption in the goats fed Ca-reduced diets in the present study.

In summary, the results of the present study showed that feeding an N-reduced diet to young goats diminished intestinal transcellular Ca absorption due to reduced expression of Ca-transporting structures. Modulation of expression levels were at least in part based on a decrease in calcitriol plasma concentrations during this feeding regimen. An extension of Ca absorption capacity into the ileum during Ca-reduced feeding could be determined. Effects of dietary N reduction became particularly obvious during a concomitant Ca reduction, which did not stimulate intestinal Ca absorption to a level which would normalise plasma Ca concentrations. Involvement of CaR, VDR as well as IGF1-R seemed not to play a role in mediating the effects of an N-reduced diet or an N- and Ca-reduced diet in young goats. Reduced intestinal Pi absorption during dietary Ca reduction in proximal and mid-jejunum did not affect plasma P_i concentrations and was not based on the altered expression of apical Pi transporters NaPiIIb or PiT1 or modulation of Na^+/K^+ -ATPase.

Therefore, it can be concluded that, although goats are able to recycle N efficiently, and therefore able to cope with a reduced dietary N supply, this dietary intervention, especially in combination with a reduced dietary Ca content, impaired intestinal Ca absorption in a calcitriol-dependent manner leading to decreased plasma Ca concentrations. Therefore, during the life period investigated in the present study, which is characterised by intensive growth, and therefore a special need for Ca and P of the animals, a sufficient dietary N supply has to be ensured. Further investigations are needed to clarify whether, for example, alterations in the binding affinity of CaR or VDR or modulated synthesis of the Vitamin D-binding protein could be involved in the described functional and molecular changes. In addition, Pi uptake into BBMV could give information about modulation of NaPiIIb and/or PiT1 activity. Furthermore, the impact on the paracellular transport of intestinal Ca and Pi absorption has to be considered, and

examination of the involved tight-junction proteins could provide an additional explanation for the changes in intestinal Ca absorption during dietary N and/or Ca reduction in young goats.

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A. S. M.-B. designed the experiments; A. S. M.-B., M. R. W. and K. E. conducted the research; A. S. M.-B., K. E. and M. R. W. analysed the data and K. E. and A. S. M.-B. wrote the paper. All authors discussed the results and commented on them in the manuscript.

There are no conflicts of interest.

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