

An energy supply network of nutrient absorption coordinated by calcium and T1R taste receptors in rat small intestine

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T1R taste receptors are present throughout the gastrointestinal tract. Glucose absorption comprises active absorption via SGLT1 and facilitated absorption via GLUT2 in the apical membrane. Trafficking of apical GLUT2 is rapidly up-regulated by glucose and artificial sweeteners, which act through T1R2 + T1R3/ α -gustducin to activate PLC β 2 and PKC β II. We therefore investigated whether non-sugar nutrients are regulated by taste receptors using perfused rat jejunum *in vivo*. Under different conditions, we observed a Ca²⁺-dependent reciprocal relationship between the H⁺/oligopeptide transporter PepT1 and apical GLUT2, reflecting the fact that trafficking of PepT1 and GLUT2 to the apical membrane is inhibited and activated by PKC β II, respectively. Addition of L-glutamate or sucralose to a perfusate containing low glucose (20 mM) each activated PKC β II and decreased apical PepT1 levels and absorption of the hydrolysis-resistant dipeptide L-Phe(Ψ S)-L-Ala (1 mM), while increasing apical GLUT2 and glucose absorption within minutes. Switching perfusion from mannitol to glucose (75 mM) exerted similar effects. L-Glutamate induced rapid GPCR internalization of T1R1, T1R3 and transducin, whereas sucralose internalized T1R2, T1R3 and α -gustducin. We conclude that L-glutamate acts via amino acid and glucose via sweet taste receptors to coordinate regulation of PepT1 and apical GLUT2 reciprocally through a common enterocytic pool of PKC β II. These data suggest the existence of a wider Ca²⁺ and taste receptor-coordinated transport network incorporating other nutrients and/or other stimuli capable of activating PKC β II and additional transporters, such as the aspartate/glutamate transporter, EAAC1, whose level was doubled by L-glutamate. The network may control energy supply.

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The T1R taste receptor family comprises three G-protein coupled receptors (GPCRs), which act through α -gustducin and/or transducin to activate a PLC β 2-dependent pathway (McLaughlin *et al.* 1992; Adler *et al.* 2000; Montmayeur *et al.* 2001; Li *et al.* 2002; Nelson *et al.* 2002). T1R receptors may also activate a cAMP-dependent pathway (Margolskee, 2002). The T1R2 + T1R3 heterodimer senses sweet taste, with simple sugars (glucose, fructose and sucrose) acting

at concentrations in the hundred millimolar range and artificial sweeteners (acesulfame-K, sucralose and saccharin) at a few millimolar (Li *et al.* 2002). The T1R1 + T1R3 heterodimer senses amino acids, notably L-aspartate and L-glutamate (umami). In 2007, four papers identified roles for T1R2 + T1R3 sweet taste receptors in the regulation of glucose absorption and enteroendocrine hormone secretion, prompting new lines of research (Jang *et al.* 2007; Le Gall *et al.* 2007; Margolskee *et al.* 2007b; Mace *et al.* 2007b).

The first paper concerned regulation of the facilitated component of glucose absorption. There are two pathways of intestinal glucose absorption. At low concentrations,

We dedicate this paper to the memory of Professor J. R. Bronk.

This article has online supplemental material.

the predominant pathway is classical active absorption mediated by the Na⁺–glucose cotransporter. However, SGLT1 becomes saturated at concentrations below those generated at the apical membrane after a meal (30–100 mM; Mace *et al.* 2007b). SGLT1 therefore displays an important regulatory role by providing the first essential signal to generate additional transport capacity through rapid insertion of GLUT2 into the apical membrane (Helliwell *et al.* 2000a,b; Kellett & Helliwell, 2000; Kellett, 2001; Kellett & Brot-Laroche, 2005; Kellett *et al.* 2008). SGLT1 does so by depolarizing the apical membrane to induce rapid influx of Ca²⁺ through the neuroendocrine L-type channel Ca_v1.3, so that the rate of Ca²⁺ absorption is increased 3-fold at 10–20 mM glucose (Morgan *et al.* 2003, 2007). The ensuing phosphorylation of myosin II in the terminal web and the peri-junctional actomyosin ring is associated with the enterocyte cytoskeletal rearrangement necessary for apical GLUT2 insertion (Madara & Pappenheimer, 1987; Berglund *et al.* 2001; Mace *et al.* 2007a).

However, no insertion over basal levels of GLUT2 occurs until the second essential signal is provided by the activation of sweet taste receptors in the apical membrane: the activation range of T1R2 + T1R3 by glucose in a heterologous expression system coincides with that for apical GLUT2 insertion in intestine (30–100 mM glucose; Li *et al.* 2002; Mace *et al.* 2007b). At these high concentrations, apical GLUT2 then provides the major pathway of absorption. Alternatively, apical GLUT2 insertion at low glucose concentrations (20 mM) may be rapidly induced by artificial sweeteners (Mace *et al.* 2007b).

Simple sugars and artificial sweeteners act synergistically through a T1R2 + T1R3– α -gustducin–PLC β 2 pathway to stimulate PKC β II activation, which is essential for apical GLUT2 insertion (Helliwell *et al.* 2000b, 2003). The necessary taste reception signalling components have been detected in the enterocytes of glucose-perfused jejunum from a fed rat and in solitary chemosensory cells (SCC) labelling with α -gustducin ($G_{\alpha, \text{gust}}$) or transducin ($G_{\alpha, \text{t1}}$), which therefore include enteroendocrine cells, brush cells and bipolar SCCs (Sbarbati & Osculati, 2005; Bezencon *et al.* 2007; Mace *et al.* 2007b). Rapid agonist-induced receptor internalization, a characteristic of GPCRs (Tan *et al.* 2004), has been observed for T1R2, T1R3 and α -gustducin and occurs simultaneously with externalization of T1R1, transducin and PLC β 2 (Mace *et al.* 2007b).

Margolskee and colleagues further reported that enteroendocrine cells in the duodenum of starved mouse secrete incretins GIP and GLP-1 from K- and L-cells, respectively, in response to sweet taste receptor stimulation by high concentrations (550 mM) of glucose (Jang *et al.* 2007). In addition, sucralose increased SGLT1 mRNA, SGLT1 protein and active glucose absorption of mice on a low carbohydrate diet for 4 weeks. Increased SGLT1

up-regulation and incretin secretion were both attenuated in T1R3 knockout and in α -gustducin knockout mice (Margolskee *et al.* 2007a).

Long-term, taste receptor-mediated up-regulation of SGLT1 was also observed in the clonal enterocytic cell line, Caco-2. Under sugar-replete conditions, Caco-2 cells contain transcripts for T1R3 and α -gustducin and also T1R2 and T1R3 protein in the plasma membrane (Le Gall *et al.* 2007), so that fructose-induced increases in SGLT1 mRNA and protein were blocked by the T1R3 inhibitor lactisole. GLP-2, secreted from L-cells, up-regulates apical GLUT2 (Au *et al.* 2002). Work from several laboratories therefore demonstrates clearly that there are both enteroendocrine and enterocyte-based mechanisms for controlling sugar absorption (for a review, see Kellett *et al.* 2008).

The observations in the four papers raise several interesting questions: What other nutrients, if any, do sweet taste receptors regulate? What is the function of amino acid taste receptors – do they also regulate absorption of any nutrients? If so, is there any cross-talk between sweet and amino acid taste reception pathways, that is, can nutrient absorption be coordinated by different taste receptors?

The opportunity to provide positive answers to these questions was prompted by a preliminary observation that levels of the oligopeptide transporter, PepT1, appeared to decrease under conditions that increased those of apical GLUT2. PepT1 is proton dependent (Ganapathy & Leibach, 1985; Daniel, 2004; Thwaites & Anderson, 2007). It transports di- and tri-peptides and a variety of pharmacological agents (for reviews see Meredith & Boyd, 2000; Daniel, 2004). Of interest in the present context, there is evidence that PepT1 in Caco-2 cells is inhibited by Ca²⁺ and PKC and that rapid regulation of PepT1 involves trafficking to the apical membrane from an intracellular pool (Brandsch *et al.* 1994; Thamotharan *et al.* 1999b; Buyse *et al.* 2001; D'Souza *et al.* 2003; Watanabe *et al.* 2004). Investigation of the parallels between PepT1 and apical GLUT2 regulation has now led to the first functional demonstration of amino acid taste receptors in nutrient absorption and to the discovery of a Ca²⁺ and taste-receptor mediated network of nutrient absorption.

Methods

Animals

Male Wistar rats (240–270 g), fed *ad libitum* on a standard Bantin and Kingman (Hull, UK) rat and mouse diet, had free access to water and were kept under a 12 h day–night cycle. All procedures used conformed to the UK Animals (Scientific Procedures) Act 1986 and had the approval of the Ethical Review Process Committee of the Department of Biology at the University of York. The number of animals used specifically for this paper was 36. In addition, data were obtained from 65 other animals using archived

vesicle and immunocytochemical samples that were prepared for and retained after previously published work.

Procedures

The following procedures have been previously described (Helliwell *et al.* 2000b; Mace *et al.* 2007b): single pass perfusion of jejunum from a fed rat *in vivo* and *in vivo/in vitro*, preparation of apical membrane vesicles, and Western blotting and immunocytochemistry, including the antibodies used. We therefore emphasize just some key points here. Rats were anaesthetized by an intra-peritoneal injection of a mixture of 1.0 ml Hypnorm (Janssen Animal Health, High Wycombe, UK) and 0.5 ml Hypnovel (Roche Diagnostics, Welwyn Garden City, UK) per kg body weight. For *in vivo* perfusions, tail pinch, foot pinch and corneal reflexes were carefully monitored throughout the duration of the perfusion. Additional anaesthetic was administered by intramuscular injection of a mixture of 0.4 ml Hypnorm and 0.2 ml Hypnovel per kg body weight when required. Rats were humanely killed by exsanguination under anaesthetic at the conclusion of the experiment. The *in vivo* single-pass perfusion technique uses two perfusate reservoirs to permit a paired comparison between a control (0–40 min) and experimental (40–90 min) period; for each set of conditions, data were collected from four perfusions. [^3H]inulin (0.7 kBq ml^{-1}) was added to correct for changes in water transport. When nutrient concentrations were less than 75 mM, their total concentration was made up to 75 mM by the addition of mannitol to obviate any potential osmotic effects. Each membrane vesicle preparation was made from two rats and three preparations were used for each condition. Immunocytochemistry employed spectral unmixing techniques to subtract all non-specific background contributions, including autofluorescence, reflection and non-assigned residuals. Where antigen was available for a given antibody, neutralization of antibody with excess peptide therefore resulted in a totally black image. Accordingly, we show only individual instances to avoid large areas of black print. Dual luminal and vascular perfusion ($n = 4$) for characterization of dipeptide transport was performed as described by Shepherd *et al.* (2002). Expression of PepT1 in oocytes was performed as described in Pieri *et al.* (2008).

Antibodies

Antibody raised in rabbit to the C-terminal region AEIEAQFDEDEKKK of PepT1 was commissioned from Invitrogen. Goat IgG to EAAC1 (EAAT3) was sc-7761 from Santa Cruz (Autogen Bioclear, UK). Other antibodies are identified in Mace *et al.* (2007b).

L-Phenylalanine(Ψ)-L-alanine was synthesized as described by Bailey (2005) and Bailey *et al.* (2006). This

dipeptide is hydrolysis resistant, is transported by PepT1 and undergoes transepithelial transport at a high rate (Fig. 3). Perfusate samples were analysed for peptide by HPLC at 210 nm on a $5 \mu\text{m}$ octadecyl silane silica C18 column (Jones Chromatography, Hengoed, Glamorgan, UK). The mobile phase was 20% methanol–80% 21 mM KH_2PO_4 (pH 5.0).

Statistical analysis

Values are presented as means \pm S.E.M. and were tested for significance using paired or unpaired Student's *t* test as appropriate.

Results

A reciprocal relationship between the levels of PepT1 and GLUT2 in the apical membrane

We have previously published many Western blots revealing the regulation of apical GLUT2, PKC β II and taste reception signalling components under different conditions. In Fig. 1 we now reproduce the main bands of interest for each set of conditions, in order that the reader can compare them immediately in one place with new data showing the regulation of PepT1 in each condition.

Rat jejunum was perfused *in vitro* with modified Krebs–Henseleit buffer (KHB) containing 5 mM fructose in the absence or presence of 200 nM PMA and/or $2 \mu\text{M}$ chelerythrin, which activate and inhibit conventional PKC isoforms, respectively. Western blots of GLUT2 and PKC β II in apical membrane vesicles prepared immediately after perfusion for 30 min are reproduced in Fig. 1A from Helliwell *et al.* (2000b). They show that PMA increased apical GLUT2 3.8 ± 0.6 -fold compared with the control of 5 mM fructose alone. The increase was blocked by chelerythrin, which had no effect alone, and correlated with PKC β II activation, as indicated by its increased level at the apical membrane. When blots of vesicles were subsequently probed with PepT1 antibody to obtain new information, a single strong band was detected at $\sim 70 \text{ kDa}$, consistent with its reported molecular mass of 75 kDa for PepT1 (Saito *et al.* 1995; Ogihara *et al.* 1999; Hussain *et al.* 2002). In contrast to apical GLUT2, PMA almost halved the level of PepT1 to $55 \pm 12\%$ ($n = 3$, $P < 0.01$) of the fructose control value; the decrease was blocked with chelerythrin, which alone had no effect on PepT1 levels. PMA has no effect on PepT1 synthesis (Shiraga *et al.* 1999). These results demonstrate that PKC β II activation down-regulated apical PepT1 levels.

Apical GLUT2 insertion is dependent on enterocyte cytoskeletal rearrangement involving myosin contraction and on the activity of PKC β II; both these processes are Ca^{2+} dependent (Morgan *et al.* 2003; Morgan *et al.* 2007; Mace *et al.* 2007a). The blots in Fig. 1B are reproduced from Mace *et al.* (2007a) and show the regulation of

apical GLUT2 and PKC β II by Ca^{2+} . They reveal that when jejunum is perfused *in vivo* with 75 mM glucose, apical GLUT2 and PKC β II are high; however, when 10 μM nifedipine is included in the perfusate to block the glucose-dependent increase in Ca^{2+} absorption through $\text{Ca}_v1.3$, apical GLUT2 and PKC β II levels are halved to 44 ± 4 and $52 \pm 4\%$, respectively, compared with control.

We now find that PepT1 is doubled (2.2 ± 0.1 -fold, $P < 0.001$). Very similar results were obtained when Ca^{2+} absorption was abolished by removing Ca^{2+} from the perfusate (Ca^{2+} -deplete conditions, Fig. 1B). The ability of Ca^{2+} to induce cytoskeletal rearrangement is prevented by 20 μM ML-7, at which concentration it is a relatively selective, cell permeant inhibitor of MLCK;

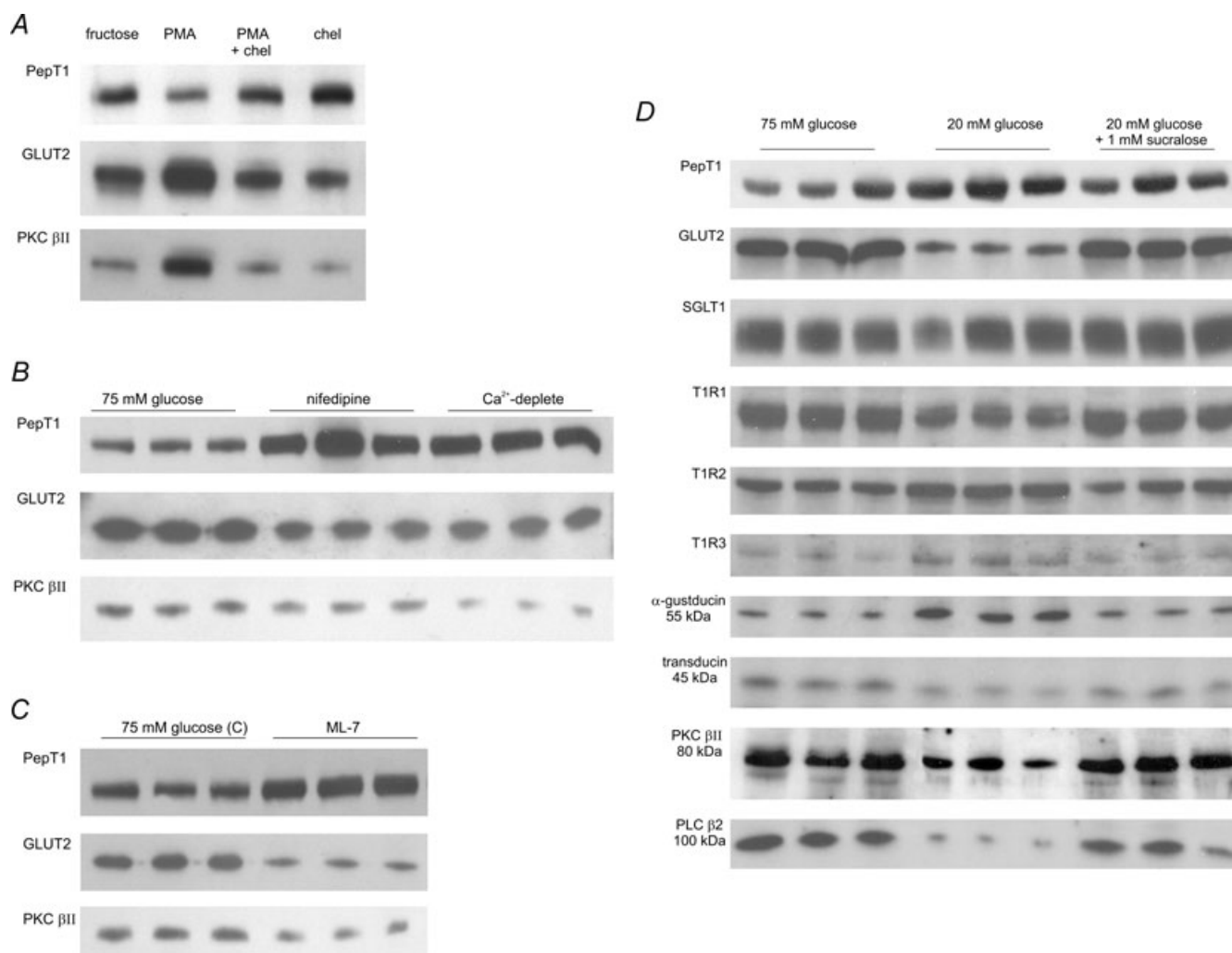


Figure 1. The levels of GLUT2 and PKC β II in the apical membrane show a reciprocal relationship with those of PepT1

Rat jejunum was perfused in KHB for 30 min as described and apical membrane vesicles prepared. Proteins (20 μg) were then separated on 10% SDS-PAGE, transblotted onto PVDF membrane and immunoblotted for GLUT2, PKC β II and PepT1. With the exception of panel A, three separate preparations from two rats each are shown for each condition. All controls, in which antibody has been neutralized by preincubation with excess immunogenic peptide, are described in the references given. The blots for apical GLUT2 and PKC β II are reproduced with permission from the *Biochemical Journal* (A, Helliwell *et al.* 2000b), and *The Journal of Physiology* (B, C and D, Morgan *et al.* 2007; Mace *et al.* 2007a,b). The PepT1 blots are new; PepT1 controls were negative (data not shown). A, effect of PMA. Rat jejunum was perfused with 5 mM D-fructose in the presence or absence of 200 nM PMA and/or 2 μM chelerythrin. B, effects of nifedipine and luminal Ca^{2+} depletion. Jejunum was perfused with 75 mM glucose alone, 75 mM glucose and 10 μM nifedipine or 75 mM glucose in KHB from which Ca^{2+} was omitted (Ca^{2+} -deplete conditions). C, effect of MLCK inhibition by ML-7. Rat jejunum was perfused with either 75 mM glucose alone or 75 mM glucose and 5 μM ML-7. D, effects glucose and artificial sweetener. Jejunum was perfused with either 75 mM glucose, 20 mM glucose or 20 mM glucose and 1 mM sucralose. Note that the first three samples of the PKC β II were inadvertently placed at the opposite end of the gel; they have been digitally transposed solely for presentational purposes; these PKC β II blots are new.

ML-7 therefore blocks the Ca^{2+} -dependent increase in myosin II phosphorylation and subsequent cytoskeletal rearrangement in response to glucose. We now observe that ML-7 increases PepT1 by 1.6 ± 0.1 -fold (Fig. 1C), whereas the blots for apical GLUT2 and PKC β II reproduced from Mace *et al.* (2007a) demonstrate that both are sharply diminished to 37 ± 4 and $37 \pm 7\%$ of the value for the glucose control, respectively.

Our early work revealed that perfusion of rat jejunum *in vivo* with 75–100 mM glucose doubled apical GLUT2 and increased PKC β II 1.6-fold compared with mannitol alone (Kellett & Helliwell, 2000). We have since found that PepT1 was concomitantly halved in the presence of glucose compared with mannitol; furthermore, phloridzin at high glucose concentrations strongly diminishes PepT1 (data not shown).

When we investigated the role of sweet taste receptors in regulating glucose absorption in perfused rat jejunum *in vivo*, we monitored changes in apical membrane levels of GLUT2, T1R1, T1R2, T1R3, α -gustducin, transducin and PLC β 2 using three conditions with respect to apical GLUT2 (Mace *et al.* 2007b). The first was 20 mM glucose alone, at which there is a basal level of GLUT2 in the apical membrane. Since rapid glucose-induced insertion of apical GLUT2 is first detectable at 30 mM glucose, use of 20 mM glucose permitted detection of apical GLUT2 regulation by 1 mM sucralose. The results were compared with those for 75 mM glucose alone, at which there is substantial apical GLUT2 insertion above the basal level. Ca^{2+} absorption was maximal in each of the three conditions. We have previously presented the full Western blots for these proteins, which include bands deriving from rapid turnover products (Mace *et al.* 2007b). We therefore reproduce in Fig. 1D only the main bands of immediate interest, to which we now add PepT1 and PKC β II (not done previously). Apical GLUT2 is strongly enhanced at both 75 mM glucose and 20 mM glucose + 1 mM sucralose compared with 20 mM glucose alone, demonstrating a sweet taste receptor response. We now find that PKC β II is correspondingly increased to an average of 1.5 ± 0.2 -fold ($P < 0.05$, $n = 3$), but, in contrast, PepT1 is decreased by an average of $48 \pm 17\%$ ($P < 0.01$, $n = 3$). There is no effect of sugar on the level of SGLT1, which appears as a broad band, but is in fact a closely spaced triplet (Balén *et al.* 2008). Note that 75 mM glucose or 20 mM glucose + 1 mM sucralose induce rapid internalization of T1R2, T1R3 and α -gustducin, whereas T1R1, transducin and PLC β 2 are inserted into the apical membrane. For GLUT2, SGLT1, T1R2, α -gustducin, transducin, PLC β 2 and PKC β II, preincubation of antibody with excess immunogenic peptide abolished all bands for that protein, confirming the specificity of detection (data not shown). For T1R1 and T1R3, no peptide was available, since the sequence was deemed 'commercially sensitive'. However, the fact that blots and

immunocytochemistry (see below) gave consistent results for the apical membrane indicates that labelling is specific.

The changes revealed by Western blotting in the levels of apical GLUT2 under the three conditions have been confirmed by immunocytochemistry previously (Mace *et al.* 2007b). Additional immunocytochemistry in Fig. 2 now confirms the changes seen for PepT1 and PLC β 2, showing how each traffics in the opposite direction, with the former being high and the latter low at 20 mM glucose compared with 75 mM glucose or 20 mM glucose + 1 mM sucralose. PepT1, an established marker for the brush-border enterocyte, is readily seen at the apical membrane over the full length of the villus, in agreement with previous reports (Ogihara *et al.* 1996, 1999; Hussain *et al.* 2002). PLC β 2 was also located over the full length of the villus at the apical membrane; however, it is essential to observe its enterocyte location under optimal conditions, when there is a strong sweet taste receptor signal (Figs 1D and 2). Note that for both proteins the peptide control for antibody labelling is completely negative. This is because all images were obtained by spectral unmixing techniques, which automatically subtract non-specific background contributions and unassigned residuals (Mace *et al.* 2007b); all images shown in this and the previous paper therefore display specific labelling only. Interestingly, the PLC β 2 images reveal three distinct types of SCCs, which are indicated by the arrowheads in Fig. 2. The left image most likely shows an enteroendocrine cell and the middle a typical bipolar SCC (Sbarbati & Osculati, 2005); the right hand image shows a triangular cell with the apex at the basolateral membrane and very strong labelling of the base at the apical membrane. Such PLC β 2-labelled triangular cells are relatively common.

We have used the term 'internalization' as shorthand to denote loss of proteins from the membrane without specifying their ultimate fate. Transporters appear to reside in intracellular compartments or vesicles from which they can recycle (Mace *et al.* 2007b; Tobin *et al.* 2008). However, signalling components such as GPCRs, G-proteins, PKC β II and PLC β 2 also undergo rapid turnover and proteolytic degradation (Helliwell *et al.* 2003; Mace *et al.* 2007b).

The regulation of dipeptide transport by glucose and artificial sweetener

The viability of PepT1-mediated transport in perfused rat jejunum *in vivo* was assessed by determining the luminal disappearance of the novel hydrolysis-resistant dipeptide L-Phe(Ψ S)-L-Ala (Bailey *et al.* 2005). With 1 mM dipeptide in the presence of 20 mM glucose, the high rate of dipeptide transport was maintained at a steady-state rate of $1.22 \pm 0.06 \mu\text{mole min}^{-1} (\text{g dry weight})^{-1}$ ($n = 6$) from 10 to 90 min, the full duration of perfusions described below

(Fig. 3A); the corresponding rate of glucose absorption was $10.31 \pm 0.81 \mu\text{mol min}^{-1} (\text{g dry weight})^{-1}$, similar to that reported previously (Kellett & Helliwell, 2000; Mace *et al.* 2007b). Tests of the kinetic characteristics of the dipeptide transport at pH 6.8 in a luminally and vascularly perfused preparation of jejunum revealed that $\sim 80\%$ of the L-Phe(Ψ S)-L-Ala absorbed across the apical membrane appeared in the vascular circuit, demonstrating effective transepithelial transport (Fig. 3B). When PepT1 was expressed in oocytes, the transport of $0.4 \mu\text{M}$ D-[^3H]phenylalanine-L-glutamine, a known non-hydrolysable substrate of PepT1, was competitively inhibited by L-Phe(Ψ S)-L-Ala with a K_i of $0.32 \pm 0.08 \text{ mM}$ (Fig. 3C). The characteristics of L-Phe(Ψ S)-L-Ala therefore make it an excellent tool with which to monitor the regulation of PepT1.

In order to investigate the effect of glucose on dipeptide transport, so-called 'paired' perfusions were performed in which the sugar conditions were changed half way through to provide a direct comparison between two sets of conditions within a single perfusion; this approach minimizes interanimal variations in data. Thus jejunum was perfused initially *in vivo* with 1 mM L-Phe(Ψ S)-L-Ala and

75 mM mannitol for 40 min (arrow, Fig. 4A). A steady-state rate of peptide transport of $1.56 \pm 0.12 \mu\text{mole min}^{-1} (\text{g dry weight})^{-1}$ was achieved after a marked up-regulation over the first 15–20 min. After 40 min, the perfusate was switched to an identical one containing 75 mM glucose instead of mannitol, when the rate of peptide transport decreased within 20 min to $0.43 \pm 0.06 \mu\text{mole min}^{-1} (\text{g dry weight})^{-1}$ ($n = 4$; $P < 0.001$). The $72 \pm 4\%$ decrease in peptide transport compares with a decrease in PepT1 levels of $57 \pm 16\%$ determined by Western blotting of apical membrane vesicles (data not shown).

When 20 mM glucose was perfused *in vivo* with 1 mM sucralose and 1 mM L-Phe(Ψ S)-L-Ala (Fig. 4B), there was a large increase in the rate of glucose absorption over the first 15 min to give a steady-state rate of glucose absorption of $23.95 \pm 1.94 \mu\text{mol min}^{-1} (\text{g dry wt})^{-1}$, similar to that reported by Mace *et al.* (2007b); the corresponding rate of dipeptide absorption was $0.73 \pm 0.15 \mu\text{mol min}^{-1} (\text{g dry wt})^{-1}$. When sucralose was omitted from the perfusate at 40 min, the rate of glucose absorption rapidly fell to $9.08 \pm 0.67 \mu\text{mol min}^{-1} \text{ g dry wt}^{-1}$ ($n = 4$; $P < 0.001$), caused entirely by trafficking of GLUT2 away

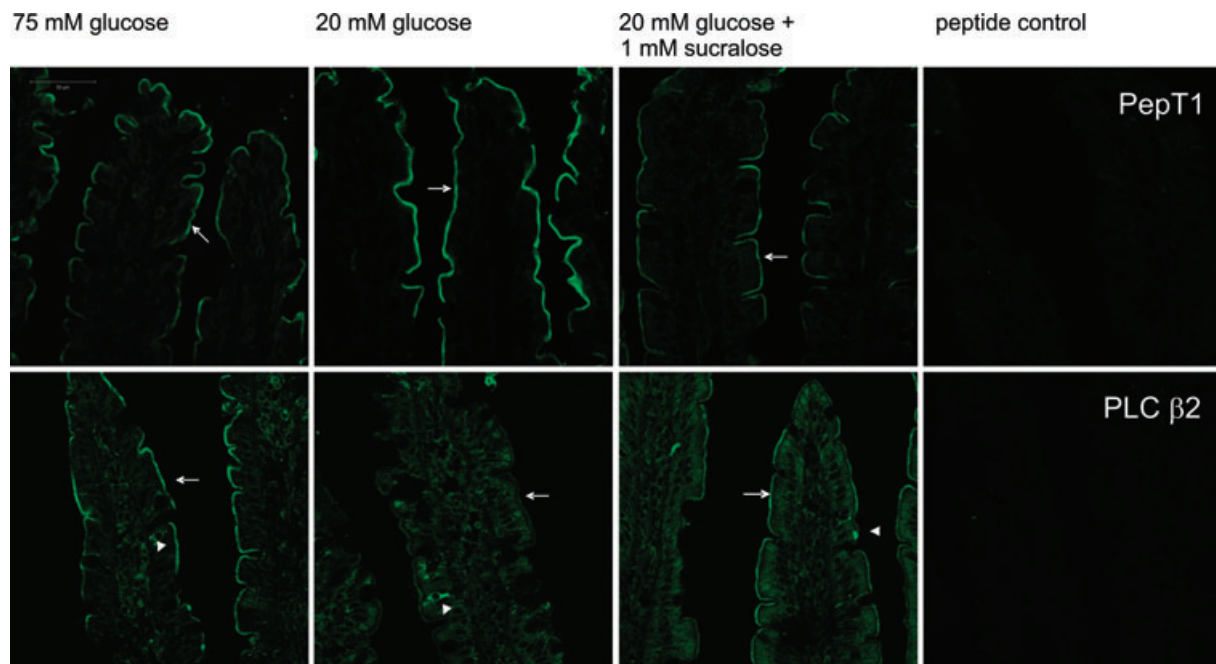


Figure 2. Immunocytochemistry of PepT1 and PLC $\beta 2$ regulation by glucose and artificial sweeteners in rat jejunum

Rat jejunum was perfused with 75 mM glucose, 20 mM glucose or 20 mM glucose and 1 mM sucralose for 30 min. Sections ($7 \mu\text{m}$) were labelled with a rabbit primary antibody detecting either PepT1 or PLC $\beta 2$ followed by a goat anti-rabbit secondary antibody conjugated to Alexa 488 (green). Arrows: apical membrane; arrowheads: three different types of SCC containing PLC $\beta 2$. The peptide controls were obtained by neutralization of primary antibody with excess immunogenic peptide. Images were processed using spectral unmixing techniques (see Methods), which automatically subtract all background contributions to leave only specific labelling as demonstrated by the peptide controls, which are totally black.

from the apical membrane (Mace *et al.* 2007b): in the presence of sucralose, apical GLUT2 is some 3-fold greater than in its absence (Fig. 1D). In contrast, omission of sucralose resulted in a concomitant 1.8 ± 0.2 -fold increase in peptide transport to $1.33 \pm 0.16 \mu\text{mol min}^{-1} (\text{g dry wt})^{-1}$ ($n = 4$; $P < 0.001$), compared with the 1.9 ± 0.2 -fold increase in PepT1 level ($n = 3$, $P < 0.001$; Fig. 1D). The rates of glucose absorption in the presence and absence of sucralose in the present experiments were

not significantly different from those reported previously in experiments without dipeptide, showing that 1 mM L-Phe(Ψ S)-L-Ala had no effect on glucose absorption.

The regulation of dipeptide and glucose absorption by L-glutamate

The experiments above show that glucose and artificial sweetener inhibit PepT1-mediated transport by activation

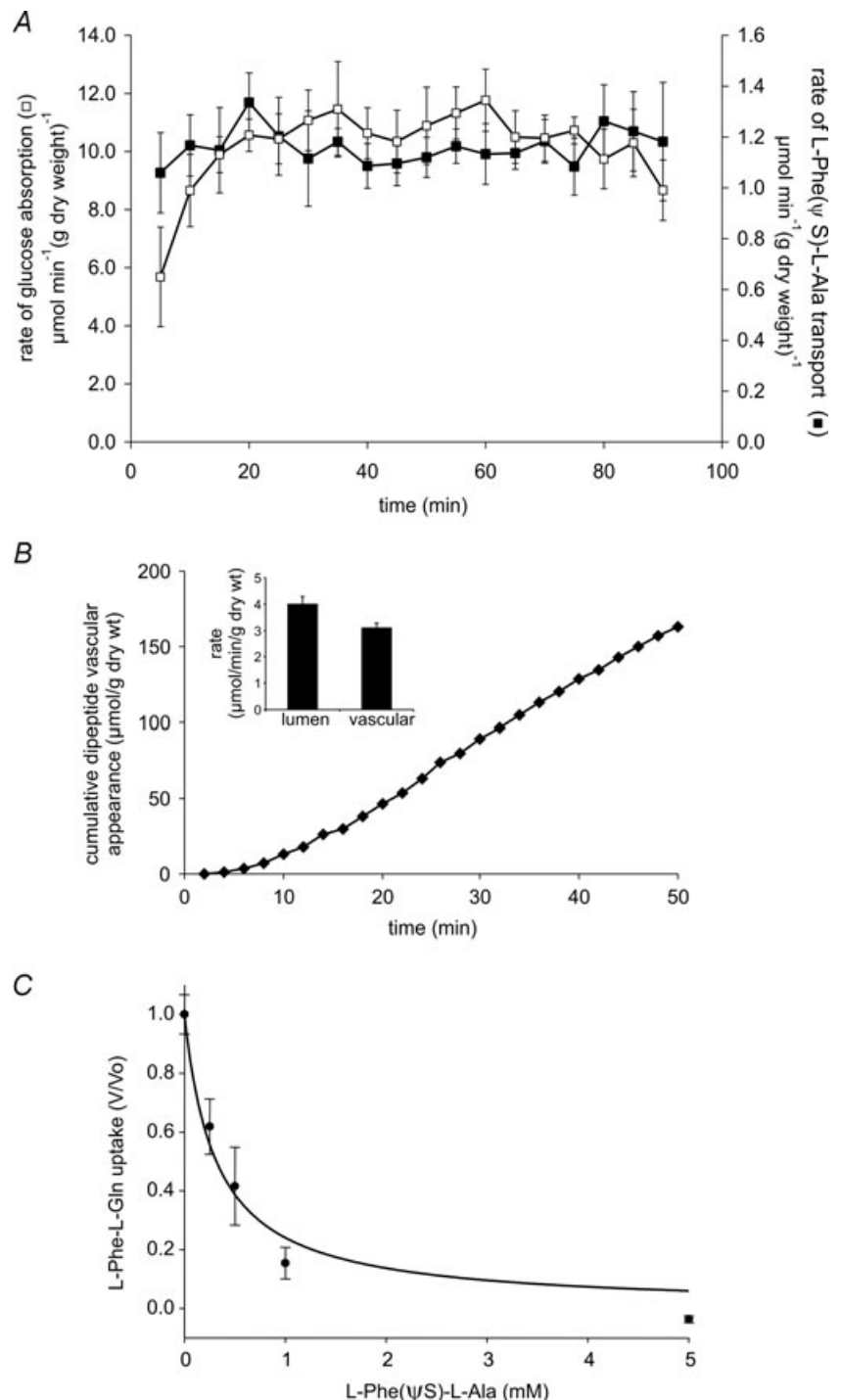


Figure 3. Transport characteristics of the novel hydrolysis-resistant dipeptide, L-Phe(Ψ S)-L-Ala, in rat jejunum

A, rat jejunum was perfused *in vivo* for 90 min with KHB containing 20 mM glucose and 1 mM L-Phe(Ψ S)-L-Ala. The time course depicts the rate of glucose absorption (□, left-hand axis) and rate of L-Phe(Ψ S)-L-Ala transport (■, right hand axis). Rates are measured in $\mu\text{mol min}^{-1} (\text{g dry wt})^{-1}$. *B*, transport characteristics of L-Phe(Ψ S)-L-Ala at pH 6.8. The time course of the cumulative appearance of 1 mM L-Phe(Ψ S)-L-Ala in the vascular circuit of a luminal and vascularly perfused preparation of rat jejunum *in situ*. The inset shows the rate of vascular appearance compared with the rate of luminal disappearance. *C*, specificity of L-Phe(Ψ S)-L-Ala for PepT1. Competitive inhibition of the transport of 0.4 μM D-phenylalanine-L-glutamine by L-Phe(Ψ S)-L-Ala, when PepT1 was expressed in oocytes.

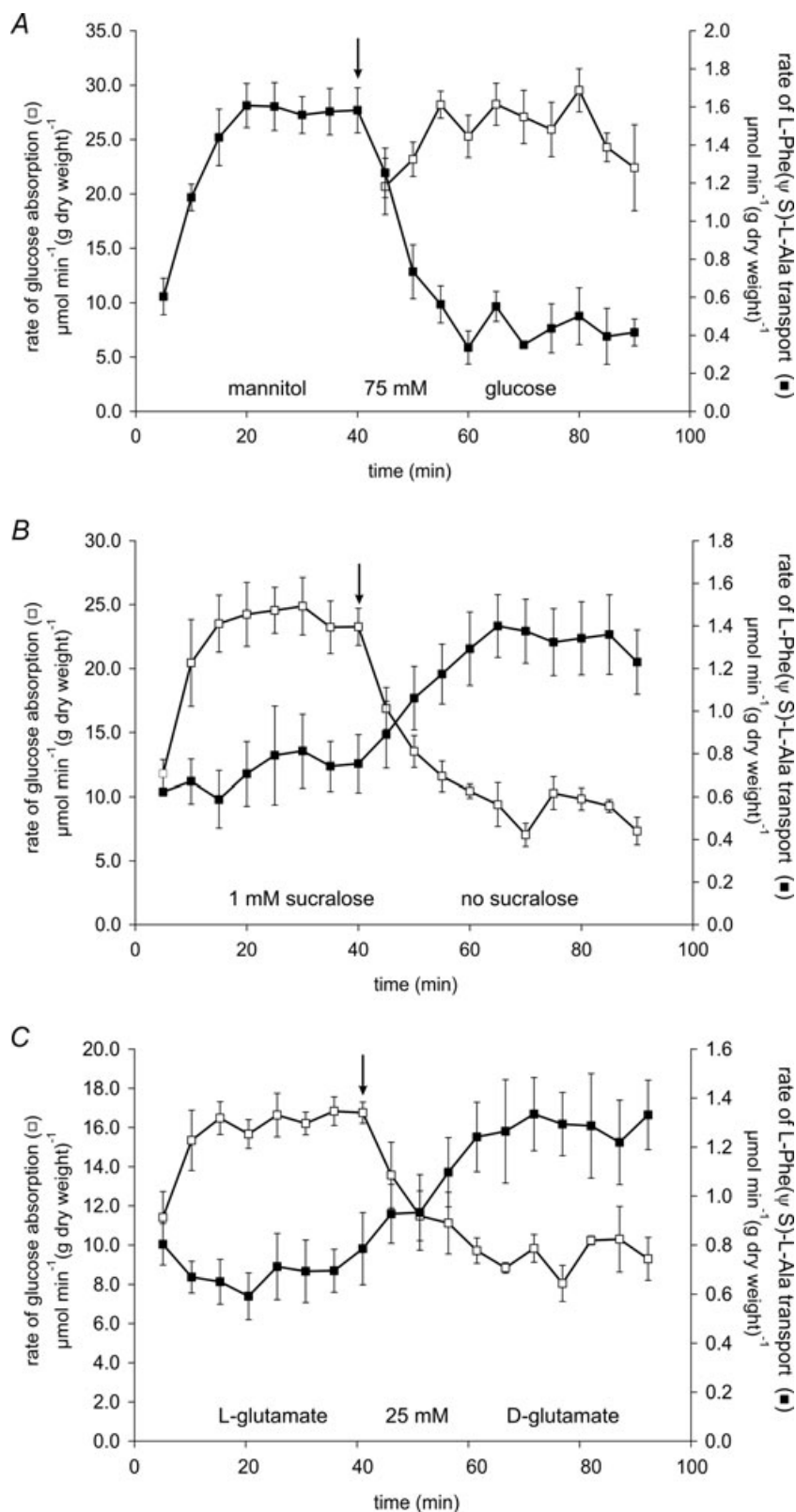


Figure 4. Regulation of glucose and peptide absorption by intestinal taste receptors

A, regulation by glucose. Jejunum was perfused *in vivo* with 75 mM mannitol (0 mM glucose) and 1 mM L-Phe(Ψ S)-L-Ala for 40 min, when the perfusate was switched (arrow) to an identical one containing 75 mM glucose (0 mM mannitol). The time course depicts the rate of glucose absorption (\square , left-hand axis) and the rate of L-Phe(Ψ S)-L-Ala transport (\blacksquare , right-hand axis). **B**, regulation by sucralose. Jejunum was perfused *in vivo* with 20 mM glucose, 1 mM D-Phe(Ψ S)-L-Ala and 1 mM sucralose for 40 min, when the perfusate was switched to an otherwise identical perfusate without sucralose. **C**, regulation by L-glutamate. Jejunum was perfused *in vivo* with 20 mM glucose, 1 mM L-Phe(Ψ S)-L-Ala and 25 mM L-glutamate for 40 min, when the perfusate was changed to an otherwise identical perfusate in which L-glutamate was replaced by D-glutamate.

of PKC β II through a sweet taste receptor- and PLC β 2-dependent pathway. Moreover, enterocytes contain both T1R1 and T1R3 (Mace *et al.* 2007b) and the T1R1 + T1R3 amino acid receptor has the potential to regulate PKC β II. We therefore investigated whether L-glutamate, one of the two cognate amino acids for the rat T1R1 + T1R3 amino acid taste receptor (Li *et al.* 2002), might also regulate PepT1 and apical GLUT2.

A concentration of 25 mM was chosen, because it produces a near-maximal increase in intracellular Ca^{2+} when rat T1R1 and T1R3 are coexpressed in a heterologous system (Li *et al.* 2002). Rat jejunum was perfused *in vivo* with KHB containing 25 mM L-glutamate, 1 mM D-Phe(Ψ)-L-Ala and 20 mM glucose. At 40 min, the perfusate was changed for an otherwise identical one in which L-glutamate was replaced by its inactive analogue, D-glutamate (arrow, Fig. 4C). The rate of L-Phe(Ψ)-L-Ala transport in the presence of L-glutamate reached a steady-state rate of $0.69 \pm 0.011 \mu\text{mol min}^{-1}$ (g dry weight) $^{-1}$ ($n = 4$), which was increased by 1.9-fold to $1.28 \pm 0.16 \mu\text{mol min}^{-1}$ (g dry weight) $^{-1}$ ($n = 4$; $P < 0.01$; Fig. 4C) in the absence of L-glutamate. This was not significantly different from the rate of L-Phe(Ψ)-L-Ala transport in the presence of 20 mM glucose only ($P = 0.13$; Fig. 4B vs. 4C). The rate of glucose absorption in the presence of L-glutamate reached a steady-state rate of $16.43 \pm 0.59 \mu\text{mol min}^{-1}$ (g dry weight) $^{-1}$ ($n = 4$; Fig. 4C), which rapidly diminished some 42% to $9.47 \pm 0.43 \mu\text{mol min}^{-1}$ (g dry weight) $^{-1}$ in its absence ($n = 4$; $P < 0.01$ compared to L-glutamate).

In order to correlate these changes in absorption rates with those of transporter levels, rat jejunum was perfused *in vivo* with either 25 mM L-glutamate or D-glutamate in the presence of 20 mM glucose for 30 min, when apical membrane vesicles were prepared for Western blotting. Figure 5 shows that D-glutamate increased the level of PepT1 by 1.8 ± 0.2 -fold compared with L-glutamate ($n = 3$; $P < 0.001$) to correlate with the change in peptide transport. In contrast, L-glutamate increased PKC β II 1.6 ± 0.1 -fold ($n = 3$; $P < 0.01$) and apical GLUT2 2.0 ± 0.1 -fold compared with D-glutamate ($n = 3$; $P < 0.001$). Given these marked effects of L-glutamate, we also probed the Western blot for EAAC1, which has been identified as the L-glutamate/aspartate transporter in intestine (Iwanaga *et al.* 2005). When viewed on a light box, EAAC1 appeared as a tight doublet at ~ 70 kDa; there were no other bands. However, as shown in Fig. 5, the resolution of the scanner used to scan the image for publication was not sufficient to resolve the two bands, which were merged to appear as a single band. For L-glutamate, the apical levels of EAAC1 were increased to 1.7 ± 0.1 -fold ($n = 3$; $P < 0.01$) compared with those in its absence.

Western blotting also revealed that L-glutamate induced profound changes in the pattern of rapid trafficking of taste receptor and signalling components compared

with D-glutamate. All taste receptor components (T1R1, T1R2 and T1R3) show some evidence of proteolysis in Western blots; moreover, the signalling components, α -gustducin, transducin, PLC β 2 and PKC β II, undergo some combination of cleavage, turnover and ubiquitylation to produce multiple species on activation. We have described the changes induced by sugars in some detail (Helliwell *et al.* 2003; Mace *et al.* 2007b) and find now that L-glutamate induces similar changes. It is apparent that, compared with D-glutamate, L-glutamate causes the rapid internalization of T1R1, T1R3 and transducin, whereas it

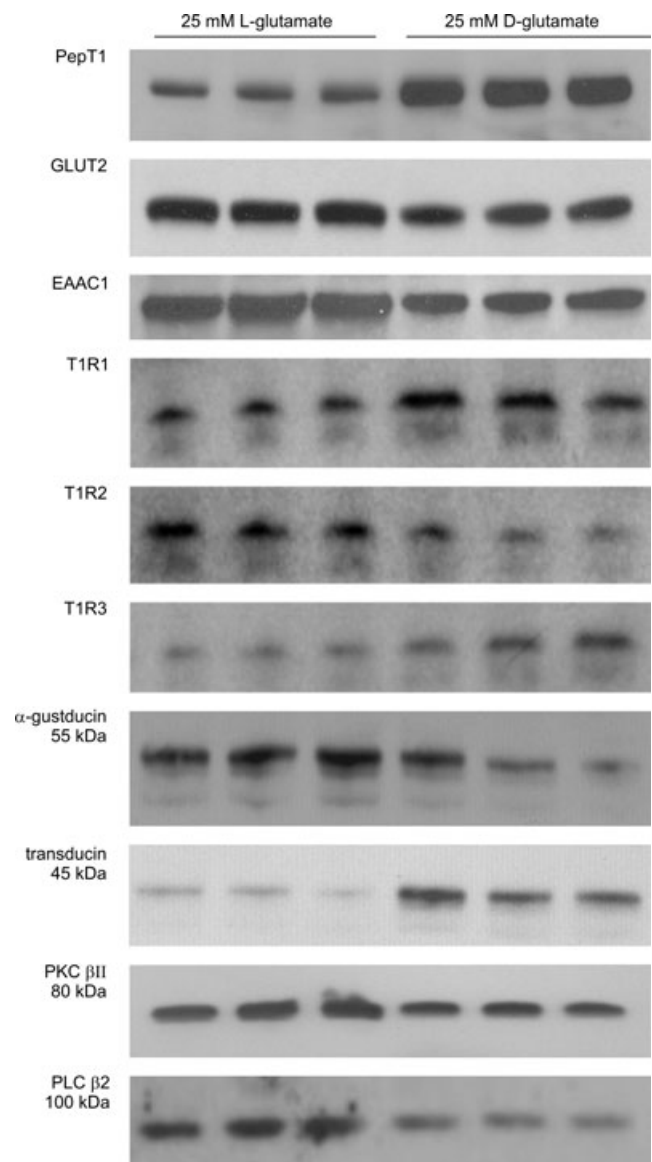


Figure 5. Regulation of transporters and taste reception signalling components by L-glutamate

Rat jejunum was perfused for 30 min with 20 mM glucose and either 25 mM L-glutamate or 25 mM D-glutamate. Apical membrane vesicles (20 μg) were immunoblotted for the proteins shown.

causes apical membrane insertion of T1R2, α -gustducin, PLC β 2 and PKC β II.

In the jejunum of a fed rat, T1R1 and T1R3 are localized in enterocytes, SCCs and in Paneth cells (Mace *et al.* 2007b). Figure 6 shows T1R1 and T1R3 colocalized in the apical membrane (arrow) over large parts of the villus and in the enterocyte cytoplasm. Note particularly the absence of T1R1 and T1R3 in the basolateral membrane in the villus as indicated by the membrane's appearance as a dark line between neighbouring enterocytes (arrowhead). In the crypts, T1R1 and T1R3 are again to be found in the Paneth cells and cytosolic vesicles. However, in contrast to the villus, the basolateral boundary of the cell is clearly seen (arrowheads). Supplementary Figs 1A and B confirm T1R1, T1R3 and transducin expression in Paneth cell granules by colocalization of each with lysozyme, an established Paneth cell marker. Supplementary Fig. 1C shows that T1R3 appears to be secreted from Paneth cell granules in fed and glucose-perfused conditions.

Discussion

A general model for the coordination of nutrient absorption by Ca^{2+} and taste receptors

In a remarkable display of coordinated regulation, L-glutamate, D-glucose, sucralose and Ca^{2+} regulate peptide, L-glutamate and glucose absorption. The fundamental concept underlying the working model for regulation in Fig. 7 is that sweet and amino acid taste receptors located in enterocytes target a common pool of PKC β II in an SGLT1- and Ca^{2+} -dependent manner (left

hand enterocyte); PKC β II in turn coordinates transporter trafficking (right hand enterocyte).

The first signal for PKC β II activation arises from glucose absorption through SGLT1, which depolarizes the apical membrane to stimulate $\text{Ca}_v1.3$ and increase Ca^{2+} absorption 3-fold (Morgan *et al.* 2007; Mace *et al.* 2007a). The resulting increase in intracellular Ca^{2+} induces terminal web and cytoskeletal rearrangement, which changes the pattern of protein trafficking to the apical membrane (see ML-7 data, Fig. 1C). Increased Ca^{2+} further promotes the translocation of inactive, but competent (phosphorylated) PKC β II from the cytosol to the apical membrane by increasing its affinity for phosphatidylserine, but PKC β II is not yet fully activated (Newton, 1997). For the latter, a taste reception signal is now required. While Ca^{2+} absorption is maximal at 20 mM glucose, the sweet taste receptor signal is provided by T1R2 + T1R3, which operates over the range 30–100 mM glucose, identical to that for apical GLUT2 trafficking (Kellett & Helliwell, 2000; Li *et al.* 2002). The sweet taste receptor is also maximally stimulated by 1 mM sucralose to double glucose absorption at 20 mM glucose by a 3-fold increase in apical GLUT2 (Mace *et al.* 2007b). Stimulation of T1R2 + T1R3 activates a heterotrimeric G-protein containing α -gustducin to which it is preferentially coupled (see below), resulting in dissociation of the $G_{\alpha, \text{gust}}$ subunit from the membrane into the cytosol. The $\beta\gamma$ subunits remain anchored in the membrane, where they bind and activate PLC β 2, thereby promoting the translocation of inactive PLC β 2 from the cytosol to the membrane (Rhee, 2001). PLC β 2 now generates diacylglycerol (DAG), which completes the activation of PKC β II by promoting removal of the N-terminal pseudosubstrate region from the active site (Newton, 1997). Additional translocation of PKC β II may occur as the affinity of PKC β II for membrane phospholipid increases further on full activation; see the effect of sucralose in Fig. 1D.

The right hand enterocyte of Fig. 7 now shows that active PKC β II inhibits trafficking of PepT1 to the apical membrane and promotes that of apical GLUT2, accounting for their reciprocal relationship in all blots presented above. Dipeptide undergoes hydrolysis within the enterocyte or exits across the as yet unidentified basolateral peptide transporter (BLPT) (Saito & Inui, 1993; Thwaites *et al.* 1993; Shepherd *et al.* 2002). The left hand enterocyte shows that the amino acid taste receptor, T1R1 + T1R3, is also coupled through a G-protein, in this case transducin ($G_{\alpha, \text{tl}}$), to activate PLC β 2 and hence PKC β II, thereby inducing changes in transporter regulation similar to those of T1R2 + T1R3.

The mechanism in Fig. 7 illustrates how the separate components can be distinguished experimentally by the use of five specific conditions in rat jejunum perfused *in vivo*: (i) 20 mM glucose alone, where sweet taste receptors

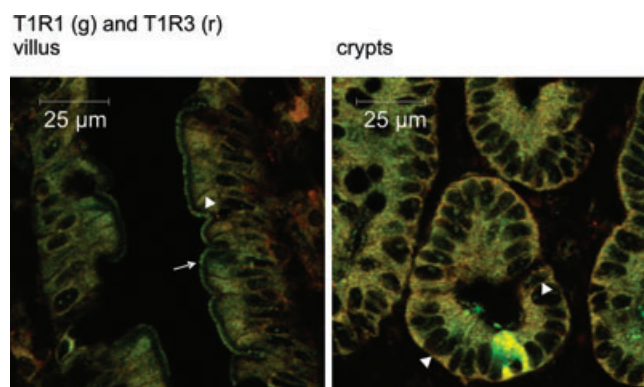


Figure 6. Colocalization of T1R1 and T1R3 in villus and crypts
Sections (7 μm) were dual-labelled with primary antibodies detecting T1R1 (green) and T1R3 (red) using Alexa 488- and 568-conjugated secondary antibodies. In the merged images for villus, T1R1 and T1R3 are colocalized in the apical membrane (arrow), but not in the basolateral membranes, which appear as dark lines (no labelling, arrowhead) where the membrane separates enterocytes. Note also the contrast with crypts, in which T1R1 and T1R3 are colocalized in the clearly visible basolateral membrane (arrowheads).

are not effectively stimulated, so that PepT1 and apical GLUT2 are at maximal and basal level levels, respectively; (ii) 20 mM glucose plus 1 mM sucralose, where sweet taste receptors are maximally stimulated by sucralose, which is neither absorbed nor metabolized; (iii) 75 mM glucose alone, so that sweet taste receptors are approaching maximal stimulation by glucose; 20 mM glucose plus either (iv) 25 mM L-glutamate or (v) 25 mM D-glutamate, so that amino acid taste receptors are maximally or not effectively stimulated, respectively. In addition, the Ca^{2+} and taste receptor components of the mechanism in Fig. 7 can be bypassed completely by perfusion with PMA to activate PKC β II directly.

The inclusion of glucose at a concentration of 20 mM or more in each of the conditions ensures that $\text{Ca}_v1.3$ is always maximal (Morgan *et al.* 2003, 2007). The upstream signal from SGLT1/ Ca^{2+} can be blocked by phloridzin, nifedipine or use of Ca^{2+} -deplete perfusate. In each case, the ensuing downstream taste reception signals are prevented, so that at 75 mM glucose, for example, both apical GLUT2 and PKC β II are diminished as PepT1 is simultaneously increased

(Fig. 1B). Such reciprocal control of intracellular free Ca^{2+} and PepT1 has been reported by Daniel and colleagues in Caco-2 cells (Wenzel *et al.* 2002); in particular, their use of nifedipine and other antagonists is in agreement with our finding that L-type channels play an important role in epithelial Ca^{2+} entry in rat jejunum (Morgan *et al.* 2003, 2007; Mace *et al.* 2007a).

Inhibition of PepT1 by Ca^{2+} in Caco-2 cells or the mutual inhibition of amino acid and active sugar transport is membrane potential dependent (Hindmarsh *et al.* 1966; Murer *et al.* 1975). Several lines of evidence, however, indicate that the changes in dipeptide and sugar absorption described above are caused by PepT1 and apical GLUT2 trafficking. Thus changes in dipeptide absorption correlate with changes in PepT1 levels and changes in glucose absorption are caused solely by change in apical GLUT2 without change in active transport (Mace *et al.* 2007b). All solutions contain 20 mM glucose at which concentration Ca^{2+} absorption through $\text{Ca}_v1.3$ is maximal; nevertheless, sucralose and L-glutamate independently and rapidly induce changes in PepT1, apical GLUT2 and EAAC1 levels at 20 mM glucose (Figs 1D

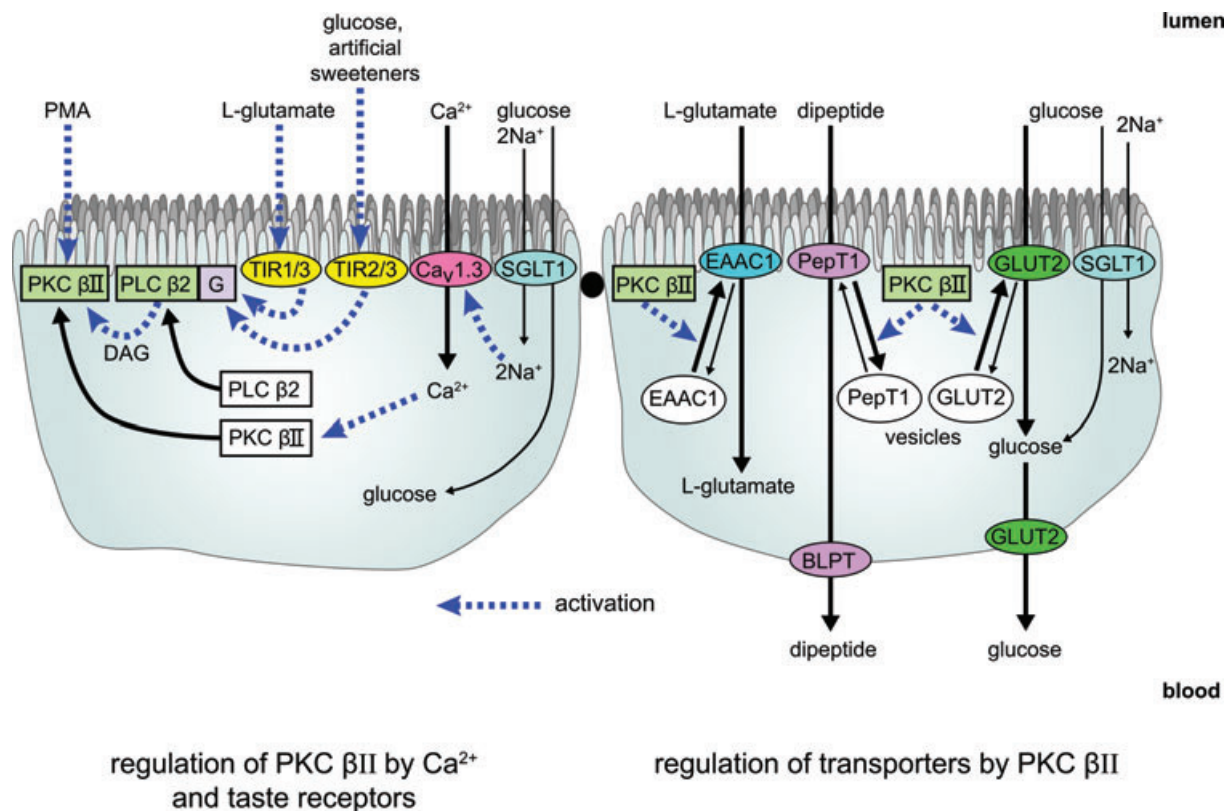


Figure 7. An energy supply network of nutrient absorption coordinated by calcium and T1R taste receptors in rat small intestine

The left hand enterocyte depicts the signalling mechanisms for the activation of a common PKC β II pool by Ca^{2+} absorption and by sweet and amino acid taste receptors; the right hand enterocyte depicts the regulation of transporters by PKC β II.

and 5). Related changes induced by ML-7 (Fig. 1C) have no effect on the upstream signal of Ca^{2+} absorption (Mace *et al.* 2007a). PMA promotes PepT1 trafficking independently of membrane potential in Caco-2 cells (Brandsch *et al.* 1994). PepT1 contains either two (human) or one (rat) consensus sites for PKC (Liang *et al.* 1995; Saito *et al.* 1995); phosphorylation of the latter site might provide a basis for direct effects on PepT1 trafficking.

Taste receptor regulation of PepT1, apical GLUT2 and EAAC-1

At 20 mM glucose, apical GLUT2 is at basal and PepT1 at maximal levels. Increasing the glucose concentration to 75 mM or the addition of 1 mM sucralose each provides a potent sweet taste receptor signal, which strongly increases apical GLUT2 through T1R2 + T1R3 (Fig. 1D). We have now shown that both signals activate PKC β II, while at the same time halving PepT1. Of note, new steady-state rates of dipeptide transport are attained within 15 min and are reciprocal to changes in the rates of glucose absorption mediated by apical GLUT2. As argued in detail elsewhere, sucralose and other artificial sweeteners of different potencies act through sweet taste receptors (Mace *et al.* 2007b).

We had previously established that the amino acid taste receptor, T1R1 + T1R3, is present in the apical membrane. Since T1R1 + T1R3 also targets PLC β 2, we investigated the possibility that T1R1 + T1R3 might also regulate peptide and glucose absorption. In the presence of 20 mM glucose, 25 mM L-glutamate rapidly up-regulated glucose absorption while dipeptide absorption remained low, but the situation was reversed within 15 min of switching to 25 mM D-glutamate (Fig. 4C).

On this basis, we predicted all the changes in transporters and signalling proteins induced by L-glutamate (Fig. 5). The changes in steady state rates of glucose and dipeptide absorption were again reflected in changes in apical GLUT2 and PepT1 levels. Confirmation that L-glutamate acts through the amino acid taste receptor is provided by its effect on receptor trafficking. A recognized characteristic of GPCRs is that activation by agonist results in rapid internalization from the target membrane to cytosolic vesicles (Tan *et al.* 2004; Scherrer *et al.* 2006). L-Glutamate simultaneously internalizes T1R1, T1R3 and transducin and induces trafficking of T1R2 and α -gustducin to the apical membrane, as well as activating and externalizing the downstream signalling components of amino acid taste reception, namely PLC β 2 and PKC β II (Fig. 5). These changes contrast with those induced by high glucose or low glucose plus sucralose, which internalize T1R2, T1R3 and α -gustducin while externalizing T1R1 and transducin together with PLC β 2 and PKC β II. The two sets of results are consistent with the view that T1R3 is a required partner for both T1R1

and T1R2 to be functional as amino acid and sweet taste receptors, respectively. The concerted trafficking of PLC β 2 and PKC β II to the apical membrane is consistent with the regulation of the latter by the former. The trafficking patterns indicate that transducin is the preferred partner for functional T1R1 + T1R3 and α -gustducin that for T1R2 + T1R3. This pattern is repeated in the secretory granules of the crypts, where T1R1 and T1R3 are routinely colocalized with transducin (Fig. 6, Supplementary Fig. 1A and B).

Apical GLUT2, PepT1 and PLC β 2 are distributed throughout the full length of the jejunal villus; in addition T1R1 and T1R3, which target PLC β 2, were detected over the whole of the villus, as were T1R2, α -gustducin, transducin and $\text{Ca}_v1.3$ (Fig. 6; see also Mace *et al.* 2007b). The changes in transporter levels and in glucose and dipeptide absorption in the L-glutamate perfusions correlate with those observed in the glucose and sucralose perfusions. We therefore conclude that L-glutamate can act via amino acid taste receptors and sugars via sweet taste receptors to coordinate the regulation PepT1 and apical GLUT2 reciprocally and to achieve a similar distribution of transporters through a common enterocytic pool of PKC β II, which is activated by receptor-mediated activation of PLC β 2.

In principle, any transporter with access to PKC β II in enterocytes could be part of a coordinated network with PepT1 and apical GLUT2. Thus the network might be larger than the core revealed so far. In order to test this possibility, we looked to L-glutamate, which is transported by EAAC1 (system X_{AG}^- or EAAT3; Kanai & Hediger, 1992). Interestingly, EAAC1 specificity for aspartate and glutamate appears to be the same as that for T1R1 + T1R3 (Li *et al.* 2002). Moreover, PKC-dependent EAAC1 trafficking results in a doubling of L-glutamate transport in C6 glioma cells (Fournier *et al.* 2004). Figure 5 reveals that the level of EAAC-1 in the intestinal apical membrane was also doubled by L-glutamate compared with D-glutamate. Trafficking correlated with increased PKC β II and increased apical GLUT2 and glucose absorption, whereas PepT1 and dipeptide absorption were decreased. Although EAAC-1 is present in the villus, it is predominant in lower villus and crypts and in ileum compared with jejunum, in contrast to apical GLUT2 and PepT1 (Fan *et al.* 2004; Iwanaga *et al.* 2005). The balance of signalling pathways for different transporters within the network will therefore depend on regional differences.

A network of nutrient absorption for the control of energy supply

What is the physiological significance of the coordinated regulation of PepT1, EAAC-1 and apical GLUT2? Part of the answer, at least, seems to be control of energy supply, for the substrates of apical GLUT2 (glucose) and

EEAC-1 (the non-essential amino acids L-glutamate and L-aspartate) are important energy sources for enterocytes, while PepT1 provides an alternative to amino acid transport. Aspartate and glutamate are also readily formed within the cell from their immediate precursors, glutamine and asparagine. Glutamine is taken up by intestine so efficiently as an energy source that it is extracted from the circulation in starvation and converted to glutamate.

Intestinal mucosa is a rapidly dividing tissue requiring amino acids and energy to sustain a high rate of amino acid synthesis. At the same time, an important function of mucosal metabolism is to deliver the carbon skeleton of glucose across the intestine in the form of alanine and lactate (the phenomenon of aerobic lactate production). This is achieved by transamination of pyruvate by glutamate to alanine and 2-oxoglutarate, a key TCA cycle intermediate. 2-Oxoglutarate is converted ultimately to oxaloacetate, which is derived also from transamination of 2-oxoglutarate by aspartate. Replenishment of TCA cycle intermediates from aspartate and glutamate not only generates ATP by oxidative phosphorylation, but also diverts pyruvate away from oxaloacetate to lactate or alanine, which enter the circulation. Glucose and glutamate/aspartate absorption therefore go hand in hand, assisted by complementary signals for mutual up-regulation by PKC β II activation through sweet and amino acid taste receptors. Energy, of course, is also required to power Na^+ - and H^+ -dependent nutrient absorption. Up-regulation of apical GLUT2 rather than SGLT1 has the advantage that absorption by apical GLUT2 is facilitated and therefore energetically cost-free, which is desirable when energy demand is high during postprandial absorption (Walker *et al.* 2005).

Significant absorption of amino acids occurs in the form of dipeptides; once in the enterocyte, they are rapidly cleaved by cytosolic peptidases, so that dipeptide absorption occurs down the concentration gradient and avoids amino acid absorption against the concentration gradient. However, PepT1 transports primarily dipeptides and, to a lesser extent, tripeptides; longer peptides are not effectively transported. In the fed state therefore the presence of elevated membrane peptidases at the luminal surface ensures significant free amino acid production, especially when there is a large dietary load during a meal and as longer peptides are slowly cleaved on their way to the ileum. The generation of glutamate in the lumen can again provide a signal to both up-regulate its own absorption and stimulate glucose uptake to assist in transamination. As the proportion of amino acids and therefore glutamate absorbed by Na^+ -dependent transporters increases, PKC β II activation signals diminished requirement for peptide absorption and therefore PepT1. When sodium-dependent transporters are working near their maximum capacity, the AMP/ATP ratio is high, which may activate AMPK to result in stimulation of

apical GLUT2 (Walker *et al.* 2005) and inhibition PepT1 (E. L. Morgan and M. Pieri, unpublished observations). A relation between AMPK and PKC β II remains to be established.

In overnight starvation, apical GLUT2 is decreased (P. A. Helliwell, unpublished observation), as absorption switches to the scavenging role of SGLT1 at luminal glucose concentrations less than those of plasma. Figures 2 and 5 show that PepT1 and apical GLUT2 levels in the apical membrane of jejunum from fed rats increase and decrease, respectively, within minutes as the concentrations of glucose or L-glutamate in the lumen decrease. These changes appear to prepare the intestine for a 3-fold increase in apical membrane PepT1 following a one-day fast (Thamotharan *et al.* 1999a), which is attenuated somewhat by longer term starvation for 4 days (Ogihara *et al.* 1999; Ihara *et al.* 2000). Indeed, in phase 3 starvation (4 days, protein catabolism), where there is marked villus atrophy, apical GLUT2 is abolished and SGLT1 tripled (Habold *et al.* 2005). However, on refeeding, large amounts of apical GLUT2 appear almost exclusively at the apical membrane within just 2 h and SGLT1 is normalized. In starvation, recycling of peptides from villus desquamation is favoured by decreased membrane peptidase activity and increased uptake by PepT1, which is inherently more energy efficient than Na^+ -dependent amino acid uptake. As in starvation, the reciprocal relationship of apical GLUT2 and PepT1 is preserved in streptozotocin-diabetes, which results in strong activation of PKC β II, persistently high GLUT2 and diminished PepT1 in the apical membrane (Corpe *et al.* 1996; Bikhazi *et al.* 2004).

Consistent with these findings, insulin rapidly doubles dipeptide transport and PepT1 trafficking to the apical membrane of Caco-2 cells (Thamotharan *et al.* 1999b; Watanabe *et al.* 2004), while in mice, circulating insulin promotes rapid trafficking of both basolateral and apical GLUT2 to an intracellular pool (Tobin *et al.* 2008). Since insulin secretion is augmented by incretins, these findings implicate taste receptor-dependent GLP-1 and GIP secretion from L- and K-cells, respectively, in attenuation of glucose absorption by insulin to limit postprandial glucose excursions (Jang *et al.* 2007; Kellett *et al.* 2008).

Interestingly, vesicular L-glutamate is colocalized with GLP-1 in mouse L-cells and the GLUTag cell line, from which release of both is induced by glucose (Uehara *et al.* 2006); GLP-2 from L-cells also promotes apical GLUT2 (Au *et al.* 2002). L-Glutamate from L-cells might therefore play a role in regulating PepT1, apical GLUT2 and EAAC1, through T1R1 + T1R3 in the basolateral membrane of crypts (Fig. 6), especially in starvation when villus atrophy occurs. It is further conceivable that T1R1, T1R3 and transducin are secreted into the crypt lumen for insertion into the apical membrane during the starved to fed transition (Supplementary Fig. 1C).

Conclusion

Studies in intestinal absorption have in the past been largely advanced separately by specialists in the fields of individual nutrients. Our data reveal, however, that absorption of major energy nutrients is in fact mediated by an integrated network – systems biology. The absorption of sugars, peptides and amino acids is coordinated by newly discovered pathways of Ca^{2+} absorption and the signalling of intestinal sweet and amino acid taste receptors to target a common enterocytic pool of PKC βII ; enteroendocrine hormones and metabolites also have important roles in some conditions. Any transporter that can be regulated by PKC βII or, indeed, any other nutrient that can regulate PKC βII , has the potential to be part of this network.

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Supplemental material

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