

Insulin-like peptide 5 is an orexigenic gastrointestinal hormone

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The gut endocrine system is emerging as a central player in the control of appetite and glucose homeostasis, and as a rich source of peptides with therapeutic potential in the field of diabetes and obesity. In this study we have explored the physiology of insulin-like peptide 5 (Insl5), which we identified as a product of colonic enteroendocrine L-cells, better known for their secretion of glucagon-like peptide-1 and peptideYY. i.p. Insl5 increased food intake in wild-type mice but not mice lacking the cognate receptor Rxfp4. Plasma Insl5 levels were elevated by fasting or prolonged calorie restriction, and declined with feeding. We conclude that Insl5 is an orexigenic hormone released from colonic L-cells, which promotes appetite during conditions of energy deprivation.

The success of bariatric surgery in the treatment of morbid obesity and its surprising ability to resolve a high percentage of cases of type 2 diabetes has sparked a renewal of interest in the intestinal endocrine system (1, 2). Gut hormones have a variety of physiological actions outside the intestine, and play a central role in linking food ingestion to peripheral nutrient disposal and appetite (3, 4). Indeed, glucagon-like peptide-1 (GLP-1) analogs and inhibitors of GLP-1 degradation are now widely prescribed for the treatment of type 2 diabetes, offering additional beneficial effects on body weight compared with conventional insulin secretagogues (5, 6). As increasing evidence implicates enteroendocrine L-cells, which secrete GLP-1 together with the anorectic peptides oxyntomodulin and peptideYY (PYY), as substantial players in postbariatric physiology (7), a question that has raised considerable interest is whether L-cells or their close relatives (8, 9) produce additional peptides of therapeutic significance.

Insulin-like peptide 5 (Insl5) is a member of the relaxin family of peptides, similar in tertiary structure to insulin and the insulin-like growth factors (10). It has been identified recently in colonic tissue and neuroendocrine tumors (11, 12), but its function remains unclear. Although some members of the relaxin family have important roles in reproductive physiology and remodeling of connective tissue, the functions of others, including Insl5, have remained elusive (13). One recent report suggested a role of Insl5 in insulin secretion, based on the initial observation of impaired glucose tolerance in aging *Insl5*^{-/-} mice; however, the phenotype was mild and only observed on a 129/Sv, but not on a C57B6, genetic background (14).

Insl5 has been reported to act through the G protein coupled relaxin/insulin-like family peptide receptor-4 (Rxfp4). Human and murine Insl5 have been shown to reduce cAMP through a pertussis toxin sensitive G_{αi} signaling pathway in cells heterologously expressing Rxfp4. By contrast, Insl5 does not stimulate the related Rxfp3, unlike relaxin-3, which is an agonist at both receptors (15, 16). The pairing of receptor and ligand is

further supported by the observation that both *Insl5* and *Rxfp4* are inactivated in tandem by pseudogenization in some species such as rat and dog (17).

We show here that Insl5 is produced by a subset of enteroendocrine L-cells in the colon in close proximity to Rxfp4 positive enteric neurons. However, unlike the other known hormonal products of L-cells, Insl5 acts as an orexigenic signal in mice and is up-regulated under conditions of calorie restriction (CR).

Results

Identification of Insl5 as an Enteroendocrine Hormone in Mouse and Human. *Insl5* was identified as an L-cell transcript in a microarray analysis of mouse L-cell populations purified by their transgenic expression of a fluorescent protein (Venus) driven by the proglucagon (*Gcg*) promoter (GLU-Venus mice) (18). Quantitative RT-PCR (qRT-PCR) analysis of FACS-purified murine L-cells from different regions of the intestine confirmed that L-cells from jejunum, ileum and colon expressed the expected transcripts for *Gcg* and *Pyy*, but that *Insl5* message was largely restricted to the colonic L-cell population (Fig. 1A). *Insl5* message

Significance

Hormonal factors from specialized enteroendocrine cells in the gut epithelium link the availability of dietary nutrients to energy utilization and storage. Many gut hormones also affect behaviors such as appetite and foraging, conveying for example the satiating effects of food consumption. Here we identify insulin-like peptide 5 (Insl5) as a product of colonic endocrine L-cells, and show that levels were elevated in calorie-restricted mice and reduced after feeding. Consistent with this profile Insl5 administration stimulated food intake in mice, indicating it should join ghrelin as only the second identified gut hormone that enhances appetite. Modulating the Insl5 axis presents a new strategy for the treatment of metabolic disease and obesity.

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was ~500-fold lower in non-L-cells compared with colonic L-cells (Fig. 1A). By flow cytometric analysis, *Insl5* immunoreactivity was detected in $56 \pm 2\%$ ($n = 3$) of L-cells isolated from mouse colon/rectum (Fig. 1B). In tissue homogenates from different regions of the mouse large intestine, we detected high *Insl5* expression in the ascending, transverse and descending colon and proximal rectum, with lower levels in the cecum and distal rectum (Fig. 1C). Outside the gastrointestinal tract low levels of *Insl5* mRNA were detected in the pancreas, thymus, and eye (Fig. S1A). No *Insl5* message was detected in pancreatic islet cells (Fig. S1B). Immunostaining of murine primary colonic cultures (Fig. 1D), and of colonic sections from human (Fig. 1E) or mouse (Fig. S1C, controls in Fig. S1D and E), revealed cells costained for *Insl5* together with GLP-1 and PYY.

Sites of Expression of the *Insl5* Receptor, *Rxfp4*. By qRT-PCR, we detected *Rxfp4* expression throughout the length of the mouse colon and cecum, as well as in the nodose ganglion. Expression was low in the distal rectum, and not detected in the hypothalamus (Fig. 2A and B). A similar profile in the intestine and nodose ganglion was found for the GLP-1 receptor (*Glp1r*, Fig. 2B). *Rxfp3*, by contrast, was expressed at higher levels in the hypothalamus than colon (Fig. 2A) (16). Within the mouse colon, in situ hybridization revealed *Rxfp4* signal in a pattern

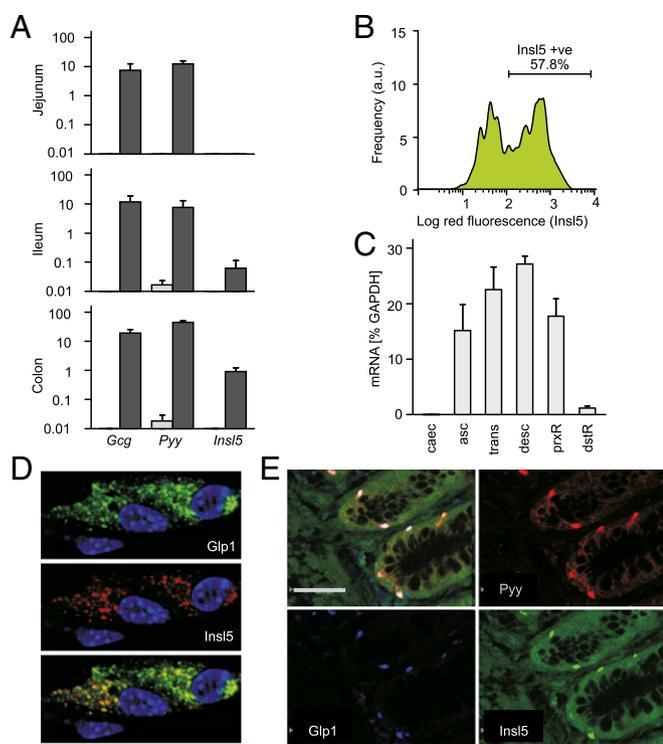


Fig. 1. *Insl5* is expressed in GLP-1-positive enteroendocrine cells. (A) L-cells (dark gray) and control cells (light gray) were isolated by flow cytometry from small and large intestine of transgenic GLU-Venus mice (18). Expression of *Gcg*, *Pyy* and *Insl5* mRNAs were analyzed by qRT-PCR, and presented relative to β -actin ($n = 3$). (B) Primary colon/rectal tissue suspensions from GLU-Venus mice were immunostained for *Insl5* with a red fluorescent secondary, and *Insl5* staining in Venus positive L-cells was recorded by FACS analysis. (C) Expression of *Insl5* in the mouse ascending (asc), transverse (trans) and descending (desc) colon and proximal (prxR) and distal rectum (dstR), but not cecum (Caec), by qRT-PCR. (D) Primary murine colonic cultures immunostained for GLP-1 (Top) and *Insl5* (Middle), revealing colocalization by confocal microscopy (Bottom). Nuclear stain: DAPI. (E) Immunohistochemistry of human colon demonstrated colocalization of *Insl5* with PYY and GLP-1. (Scale bar: 500 μ m).

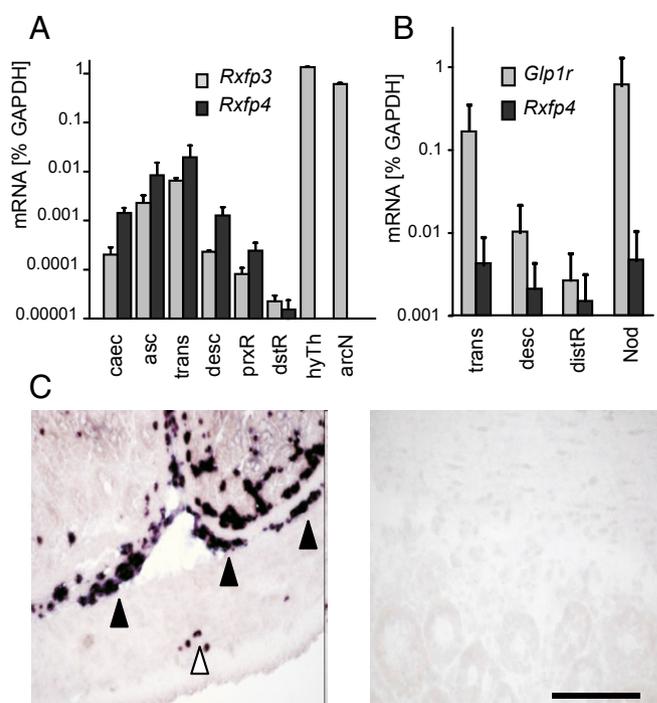


Fig. 2. Patterns of *Rxfp4* expression. (A) mRNAs for *Rxfp4* and *Rxfp3* were detected by qRT-PCR in the different regions of large intestine: asc, ascending colon; caec, cecum; desc, descending colon; distR, distal rectum; prxR, proximal rectum; trans, transverse colon. *Rxfp3* but not *Rxfp4* was detectable in the hypothalamus (hyTh) and the arcuate nucleus (arcN). Data are presented as mean \pm SEM ($n = 3$). (B) *Glp1r* and *Rxfp4* are expressed in the gastrointestinal tract (abbreviations as in A) and in the nodose ganglion (Nod). Data are presented as mean \pm SEM ($n = 3$). (C) In situ hybridization of mouse colon using a probe for *Rxfp4* (Left) revealed a pattern consistent with expression in the myenteric (open arrow heads) and submucosal (filled arrow heads) plexus of the enteric nervous system. No labeling was observed using a sense *Rxfp4* control probe (Right; scale bar: 100 μ m).

consistent with expression in submucosal and myenteric ganglia (Fig. 2C).

Influence of Meal Ingestion and Chronic Energy Balance on *Insl5* Production in the Mouse.

As *Insl5* is a peptide secreted by L-cells, we asked whether its plasma level is dependent on food intake. Mice on an ad libitum chow diet were fasted overnight, and blood samples taken before and at different times after refeeding. *Insl5* concentrations fluctuated initially, but were consistently suppressed at later times of the refeeding phase (Fig. 3A). To validate these results, we repeated the fasting and late refeeding time points on a separate cohort of animals, using an *Insl5* ELISA instead of the RIA. *Insl5* concentrations obtained with the ELISA were generally lower, but a similar pattern was observed with high levels in the fasting state that were suppressed by refeeding (Fig. 3B).

We explored the response to chronic changes in energy balance in groups of male C57BL/6J mice that were either fed on high-fat diet (HFD, 45% fat) or subjected to CR (60% of the ad libitum chow consumption) for 2 wk. Fasting plasma levels were significantly higher in the CR group compared with ad libitum and HFD-fed mice (Fig. 3B). In the CR group, as in the ad libitum fed mice, *Insl5* concentrations fell after refeeding. This pattern was lost in the HFD cohort, and no significant difference was detected between the 10-h-refed concentrations across all three groups. Interestingly, when in an independent experiment mice were subjected to prolonged calorie restriction for 10 wk

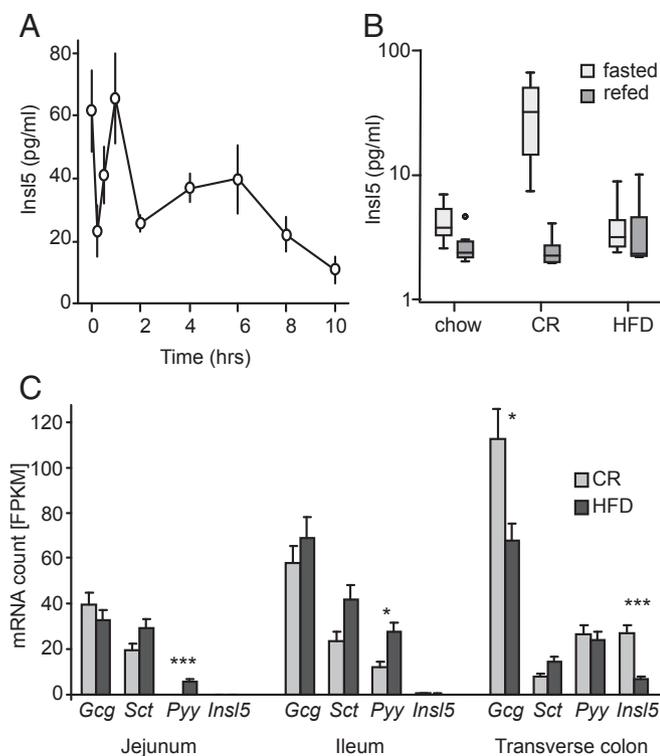


Fig. 3. Insl5 responses to acute and chronic fasting. (A) Plasma levels of Insl5 in mice during refeeding with chow following an overnight fast, determined using an Insl5 RIA (Phoenix). Data are represented as the mean \pm 1 SEM. Concentrations fell significantly after reintroduction of food, as assessed by Kruskal-Wallis test for independent samples ($P < 0.001$; $n \geq 10$). (B) Male C57BL/6 mice were either on chow ad libitum, 60% caloric restriction or HFD for 2 wk. After an overnight fast or after 10hrs refeeding plasma Insl5 were measured by ELISA (Kamiya). Fasting Insl5 levels were higher in the CR cohort than in the HFD or chow cohorts (Kruskal Wallis test, $P < 0.001$). Comparing fasted and refeeding Insl5 levels within each diet group, significant differences were detected for chow and CR, but not the HFD cohort ($P = 0.002$ and $P < 0.001$, respectively. Mann-Whitney u test; $n = 10$, boxplot with median). (C) Expression of *Gcg*, secretin (*Sct*), *Pyy* and *Insl5* mRNAs after CR or HFD in the distal intestine. C57BL/6 mice were either fed a HFD or were restricted to 60% caloric intake of a control group fed chow ad libitum (CR) for 4 wk. mRNA was harvested just before provision of the daily food ration in the CR cohort and at the same time in the ad libitum fed HFD group. Pooled cDNA ($n = 6$) was quantified by sequencing and data are represented as fragments per kilobase exon per million fragments (FPKM + SD). Data were analyzed using Cuffdiff, which provides the following P values adjusted for multiple testing: Colon: *Insl5*: 1.56×10^{-7} ; *Gcg*: 0.0449; Jejunum: *Pyy*: 8.58×10^{-6} ; Ileum: *Pyy*: 0.0119. * $P < 0.05$, *** $P < 0.01$.

instead of 2, Insl5 concentrations remained high at 10-h refeeding (Fig. S2B).

To explore the transcriptional regulation of enteroendocrine hormones in different parts of the gut, another cohort of C57BL/6J males was placed on HFD or 60% CR (chow) for 4 wk. At the end of the study period, intestinal tissue from different regions was harvested and the transcriptome sequenced and quantitatively analyzed. Transcripts of the anorexigenic peptide *Pyy* were elevated in the HFD group in both jejunum and ileum, but were not different in the colon (Fig. 3C). By contrast, in the colon of the CR group we detected a marked increase in *Insl5* as well as an increase in *Gcg* transcripts. The combined findings that CR increased *Insl5* transcripts and plasma Insl5 concentrations, and that circulating levels were lowered by refeeding, suggest that this hormone might help to shape the behavioral response to short and longer term energy balance.

Physiological Roles of Exogenous and Endogenous Insl5. We examined the physiological role of Insl5 by injecting a commercial preparation of the recombinant peptide. i.p. Insl5 dose-dependently increased intake of chow in nonfasted wild-type mice (Fig. 4A). Mice adapted to receive a highly palatable meal daily at a fixed time also showed increased food consumption (Fig. S3A). Similar results were observed with an Insl5 preparation produced in-house (Fig. S3B and C). Consistent with *Rxfp4* being the cognate receptor for Insl5, the orexigenic action of Insl5 was not observed in mice lacking *Rxfp4* (Fig. 4B). To examine whether the action of Insl5 was mediated by a peripheral or central effect, mice were injected intracerebroventricularly (ICV) with Insl5 or relaxin-3, or with ghrelin or neuropeptide Y (NPY) as positive controls. Whereas ICV ghrelin or NPY triggered robust feeding responses, no significant effect was observed with Insl5 (Fig. 4C), presumably reflecting the lack of central expression of *Rxfp4* (Fig. 2A).

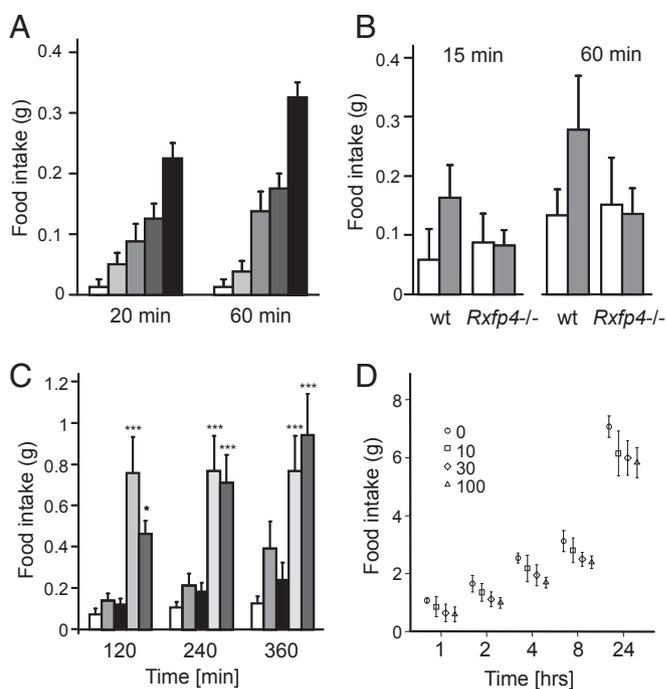


Fig. 4. Dose-dependent stimulation of food intake by Insl5. (A) Effects of a commercial preparation of Insl5 (Phoenix) injected intraperitoneally at concentrations of 0 (white), 8, 40, 200 (light to dark gray), or 1,000 (black) ng per 25 g of body weight, on intake of chow, at 20 and 60 min after administration in free-feeding mice [General linear model (GLM) for repeated measures; $n = 8$; $P < 0.001$]. Significantly increased food intake was observed for the doses of 200 and 1000 ng/25g BW ($P < 0.001$, Tamhane's T2 post hoc test). Data are presented as mean \pm 1 SEM. (B) Insl5 (Phoenix, 1000 ng/25g, black bars), injected as in A, induced food intake in wild-type but not *Rxfp4*^{-/-} mice, compared with saline control (gray bars). (GLM for repeated measures in wild type $P = 0.003$; $n \geq 6$, and in *Rxfp4*^{-/-} $P = 0.7$; $n \geq 8$). Data are presented as mean \pm 95% CI. (C) Twelve-wk-old C57BL/6 mice were administered 4 μ g of peptide or PBS as a sham control, by single injection via an implanted ICV cannula. Bar colors: sham, white; Insl5, midgray; relaxin-3, black; ghrelin, light gray; NPY, dark gray. Food intake was measured at the time points indicated. Data are shown as mean \pm SEM. Significance was tested by two-way ANOVA with post hoc Bonferroni test. * $P < 0.05$, *** $P < 0.001$ compared with sham control at the corresponding time point. (D) Wild-type mice were fasted overnight and injected via the tail vein with rabbit normal serum (open bars), or a polyclonal Insl5 antibody at a dose of 10 (light gray), 30 (dark gray), or 100 (black bars) μ g per mouse. Food was reintroduced and cumulative food intake monitored over the following 24 h. Food intake was reduced in a dose dependent manner ($P < 0.001$, GLM for repeated measures, $n = 5$) at all 3 doses ($P < 0.05$ in Dunnett's T post hoc test vs. control). Data are presented as mean \pm 95% CI.

To examine the role of endogenous *Insl5*, we performed studies involving immunoneutralization of circulating *Insl5* or genetic disruption of *Rxfp4*. i.v. injection of a polyclonal antibody against *Insl5* in overnight-fasted wild-type mice reduced food intake over the following 24-h period (Fig. 4C). The effect of the antibody injection on daily food intake was still evident at 3 d but faded thereafter, in line with common antibody half lives (Fig. 4D). Mice with a homozygous deletion of *Rxfp4* (*Rxfp4*^{-/-}; Fig. S4) were viable and fertile without any obvious differences in a number of morphological, neurological, behavioral assays, clinical chemistry or hematology. Glucose metabolism was assessed in fasted mice (male and female, 3 and 9 mo old, chow diet), in oral glucose tolerance tests, and during a postprandial fast (hourly readings for 9 h) without detecting reproducible differences in either glucose or insulin levels (data not shown). As well as showing no response to i.p. *Insl5* (Fig. 4B), *Rxfp4*^{-/-} mice exhibited no alteration in food intake after injection of anti-*Insl5* antibody (Fig. S3D).

To further investigate the effects of *Rxfp4* knockout on food intake, *Rxfp4*^{-/-} and wild-type animals were fasted overnight and feeding patterns monitored with two different diet options, a HFD and a high-carbohydrate diet (HCD), offered in parallel. *Rxfp4*^{-/-} animals exhibited shorter meal durations, particularly pronounced for the HFD, and appeared to have lost the preference for HFD seen in wild-type animals (Fig. 5A). In a further experiment, mice were fasted overnight before being offered only a HFD for the first 24 h, and then an additional choice of

HCD for a further 2 d. *Rxfp4*^{-/-} mice consumed less HFD than wild-type animals throughout the 3 d observation period (Fig. 5B). When additionally offered HCD, the cumulative combined intake of both diets remained lower in *Rxfp4*^{-/-} animals, although they eventually consumed more HCD than the wild-type mice (Fig. 5C). These data suggest that *Insl5* levels may not only affect hunger after fasting but might also contribute to the normal preference for HFD over HCD, particularly in states of calorie restriction.

Consistent with a physiological role for the *Insl5*/*Rxfp4* axis in the control of food intake, *Rxfp4*^{-/-} mice tended to have a lower body weight than wild-type controls at an age of 3 mo (Fig. 6A). When fed on diets with high caloric density (HFD or HCD) for 22 wk, however, *Rxfp4*^{-/-} mice eventually reached similar body weights to the wild-type controls (Fig. 6A). To examine further the influence of *Rxfp4* on body composition, we analyzed body weight and composition of mice on chow diet at 8 and 11 wk of age (Fig. 6B–E). Although no significant differences in body weight or lean mass were observed in this cohort, *Rxfp4*^{-/-} animals had a trend toward a lower fat mass at 8 wk by NMR spectroscopy (EchoMRI), which reached significance at 11 wk, and had significantly decreased inguinal and gonadal fat pad masses observed *ex vivo*.

Discussion

The data we present here show that *Insl5* is a hormonal product of colonic L-cells that enhances food intake in mice. In line with the expected properties of an orexigenic hormone, *Insl5* plasma levels and colonic mRNA transcripts were elevated by caloric restriction, and circulating *Insl5* concentrations were suppressed by refeeding. Knockout of the receptor *Rxfp4* or treatment with antibodies against endogenous *Insl5* correspondingly reduced refeeding responses after fasting. Taken together our data support the notion that *Insl5* is an orexigenic hormone that plays a physiological role in driving food intake under conditions of energy restriction.

Insl5 was found to be produced by ~50% of L-cells in the colon, but mRNA transcripts in the ileum were much lower, and in the jejunum and duodenum were largely absent. This pattern differs from the other L-cell hormones, GLP-1 and PYY, which show increased production along the proximal-distal gut axis, but are still found in substantial amounts in the small intestine. Indeed the direct stimulation by ingested nutrients of L-cells in the upper small intestine would be sufficient to underlie the rapid rise in plasma concentrations of GLP-1 and smaller elevation in PYY after food intake. It remains debatable, however, whether colonic L-cells contribute substantially to postprandial gut hormone secretion, as they would not normally experience major fluctuations in local nutrient exposure. Although some evidence supports the concept that colonic L-cells can be stimulated by a neural or hormonal loop after food intake, the contribution of this pathway to plasma levels remains unclear. Indeed, our finding that circulating *Insl5* concentrations were high in the fasting state and suppressed after food intake suggests instead that L-cells in the colon are not dramatically stimulated by food ingestion, but may rather signal longer term measures of energy balance.

The observed expression of *Rxfp4* in enteric and nodose ganglia but not the hypothalamus, together with the lack of effect on food intake of ICV *Insl5*, suggests that the orexigenic effects of *Insl5* are initiated peripherally and transmitted to the central nervous system by afferent sensory nerves. The nodose ganglion also expresses receptors for GLP-1, PYY and CCK, supporting the established view that the afferent vagus nerve contributes to the anorexigenic actions of a number of endogenous hormones released locally within the intestinal epithelium (19, 20). Acting on the intrinsic enteric plexus, *Insl5*, like several other metabolically relevant gut hormones, may contribute to the coordination of colon motility and secretion, ensuring the efficient mixing, fermentation

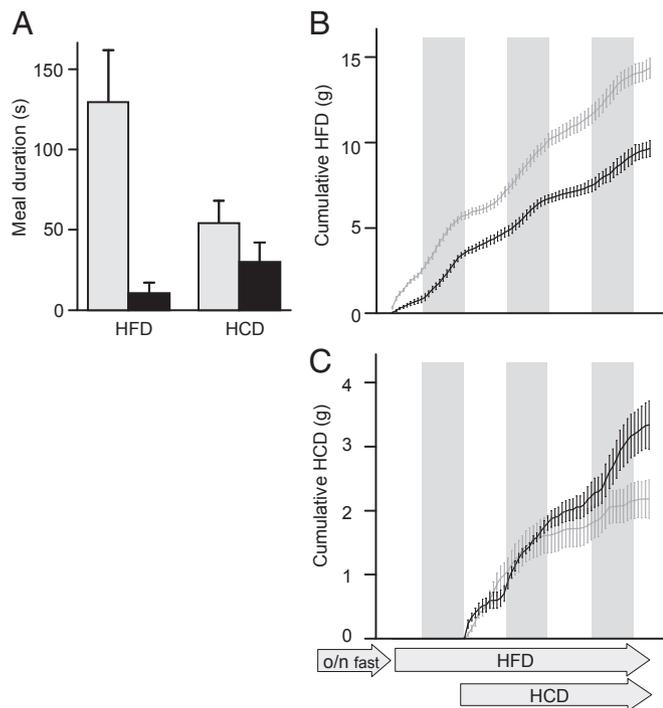


Fig. 5. Deletion of *Rxfp4* changes feeding patterns and macronutrient preference. (A) Single housed wild-type (gray bars) and *Rxfp4*^{-/-} (black bars) mice were fasted overnight and presented with a choice of a HFD and HCD, and consumption of both diets was continuously monitored. Meal durations (exceeding 10s undisturbed feeding at the chosen food hopper) differed between genotype by median test ($P < 0.001$ for HFD, $P = 0.005$ for HCD). Data represent the mean of $n = 4$ and the 95% CI. (B and C) To explore the initial consumption of HFD without a choice after the overnight fast we performed an experiment where the HCD was introduced after a 24 h delay. Continuous monitoring of the consumption of HFD (B) and HCD (C) revealed that *Rxfp4*^{-/-} mice (black lines) consumed less HFD throughout the 3-d observation period than wild-type mice (gray lines). Data are presented as mean \pm 95% CI, $n = 4$; dark phases are indicated by gray boxes.

"Superfrost gold" slides and stored at -80°C . In situ hybridization was performed using DIG RNA Labeling kit (SP6/T7), anti-DIG-AP antibody, and NBT/BCIP according to provider's instructions (Roche Applied Sciences).

Human Tissue Sample Preparation. Tissue was fixed with 10% (wt/vol) buffered formalin, dehydrated, and embedded in paraplast. Sections were cut with a Leica microtome at nominal thicknesses of $3\ \mu\text{m}$ and mounted on superfrost plus (Menzel) slides. Sections were deparaffinized and treated with H_2O_2 /methanol (30 min), and antigens were retrieved by pressure cooking in citrate buffer, 0.001 M, pH 6. After cooling and washing with phosphate buffer (0.1 M, pH 7.4), immunohistochemical detection was performed using primary antibodies against *InsI5* (as described in *InsI5 Antibodies*), PYY (Eud 5201, Acris), and GLP-1 (T4057, Bachem) followed by appropriate secondary antibodies (Alexa-488, Cy3, and Cy5). Triple immunofluorescence staining was recorded with a Leica confocal scanning microscope.

Mouse Diet Study. High fat [45% Atwater Fuel Energy (AFE) fat; 824053], high carbohydrate (70% AFE carbohydrates; 824050), and chow diet (801722) were from Special Diets Services. *InsI5* was prepared as above, and 5% (wt/vol) mouse serum albumin or water constituted sham injection. Peptide was delivered using an insulin syringe (BD Micro-Fine 0.3-mL syringe, Becton Dickinson Medical). All doses were given in a volume of $100\ \mu\text{L}$ i.p. For all i.p. peptide administrations, animals were handled and given sham ip. injections on 3–5 d before the experiment. For refeeding studies shown in Fig. 53C, male C57BL/6 mice aged 10 wk (Charles River) were acclimatized for 1 wk to single housing, appearance of a highly palatable foodstuff [either wet mash (chow diet in water, 1:1.5 wt/wt) or Farley's Breakfast "Peachy Porridge" (HJ Heinz), 1 part dry food to 1.3 parts water for 1 h/d], and handling procedures. In the final 3 d of the acclimatization, mice underwent a sham i.p. injection of $100\ \mu\text{L}$ of PBS. Mice were randomly assigned to receive a different dose on three occasions over a week. At the onset of the 1-h test period, standard chow was removed, and 20 g of highly palatable diet was placed in a clear plastic dish on the floor of the cage. After 1 h, palatable food was removed, standard chow was returned to the hopper, and mice were returned to their holding room.

Intracerebroventricular (ICV) injections were performed on 12-wk-old C57/Bl6 mice (Charles River), mean body weight 27.3 g, using a protocol described previously (23). In brief, a lateral ventricle cannula was placed under isoflurane anesthesia, the animal was given 1 wk to recover, and studies were only performed further if body weight was at or above pre surgery weight. Mice were given ad libitum access to standard chow. Experiments were started at 9:00 AM, following ad libitum feeding the previous night. A single injection of peptide ($4\ \mu\text{g}$) was given, and food intake was measured over the following 6 h. Sham mice were injected with PBS. *InsI5* and Relaxin-3 were obtained from Phoenix Pharmaceuticals (catalog numbers 035–40 and

035–36, respectively), NPY and ghrelin were from Bachem (catalog numbers H6375 and H4862, respectively).

Mouse Plasma Analyses. Mouse *InsI5* was measured using a RIA kit from Phoenix Pharmaceuticals (Fig. 3A and Figs. S2B and S5). The specificity of the assay was confirmed by detection of synthetic *InsI5* (Phoenix Pharmaceuticals) injected intravenously, which resulted in the expected plasma peak followed by a decaying signal (Fig. S5). New cohorts of mice were assessed using an *InsI5* ELISA (Kamiya Biomedical KT-58379), which became available later in the study (Fig. 3B).

Quantitative mRNA Sequencing. Double-stranded cDNA was synthesized from polyadenylated RNA and sheared. The 190- to 210-bp fraction was isolated and PCR amplified to generate the sequencing library, as per the Illumina Genome Analyzer paired end library protocol (Illumina). The resulting libraries were sequenced on an Illumina GA_{II} and paired end reads were mapped to the mm9 mouse genome assembly using TopHat 1.3.1 (24). The resultant alignments were analyzed for differential expression using Cufflinks 1.0.3 (25). Gene definitions were supplied as a gtf file, which was downloaded from the UCSC Table Browser (26) and further processed with cuffcompare (25). Raw read counts were determined using htseq-count (www-huber.embl.de/users/anders/HTSeq/doc/count.html).

Statistics. All tests and post hoc tests are two-sided with $\alpha = 0.05$ unless indicated otherwise. Instead of extensive post hoc tests 95% confidence intervals are indicated where suitable. PASW Statistics 17 and IBM SPSS 19 (IBM) software packages were used.

Note Added in Proof. A recent batch of the mouse *InsI5* ELISA from Kamiya give high unspecific signals. We recommend validating assays with plasma spiked with *InsI5* and plasma from *InsI5*^{-/-} mice. Alternative ELISAs are available from BlueGene and Cusabio.

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