Insulin internalizes GLUT2 in the enterocytes of healthy but not insulin-resistant mice.

Vanessa Tobin\textsuperscript{1}, Maude Le Gall\textsuperscript{1}, Xavier Fioramonti\textsuperscript{2}, Emilie Stolarczyk\textsuperscript{1}, Alba G. Blazquez\textsuperscript{5}, Christophe Klein\textsuperscript{3}, Magali Prigent\textsuperscript{4}, Patricia Serradas\textsuperscript{1}, Marie-Hélène Cuif\textsuperscript{4}, Christophe Magnan\textsuperscript{2}, Armelle Leturque\textsuperscript{1}, Edith Brot-Laroche\textsuperscript{5}

\textsuperscript{1}Université Pierre et Marie Curie-Paris6, UMR S 872, Les Cordeliers, Paris, F-75006 France; INSERM, UMR S 872, Paris, F-75006 France;
\textsuperscript{2}Université Paris 7, CNRS, Paris cedex 05, France;
\textsuperscript{3}IFR58, Centre de Recherche des Cordeliers, 15 rue de l’école de Médecine, 75006 Paris, France;
\textsuperscript{4}Institut de Génétique et Microbiologie, Centre Scientifique d’Orsay, Université Paris 11-Orsay, 91405 Orsay Cedex France
\textsuperscript{5}Laboratory of experimental pathology and drug targeting (HEVEFARM) University of Salamanca, CIBERehd, Spain

Running title: Insulin regulates intestinal sugar absorption

Corresponding Author:
Edith Brot-Laroche
CRC, Team 9
15 rue de l’Ecole de Médecine
75006 Paris, France
edith.brot-laroche@crc.jussieu.fr

Received for publication 16 July 2007 and accepted in revised form 20 November 2007.

Additional information for this article can be found in an online appendix at http://diabetes.diabetesjournals.org.
**ABSTRACT**

**Objectives:** A physiological adaptation to a sugar-rich meal is achieved by increased sugar uptake to match dietary load, resulting from a rapid transient translocation of the fructose/glucose GLUT2 transporter to the brush border membrane (BBM) of enterocytes. The aim of this study was to define contributors and physiological mechanism controlling intestinal sugar absorption focusing on the action of insulin and contribution of GLUT2 mediated transport.

**Research design:** The studies were performed in the human enterocytic Caco-2/TC7 cells, and in vivo during hyperinsulinemic-euglycemic (HI-EG) clamp experiments in conscious mice. Chronic high-fructose or high-fat diets were used to induce glucose intolerance and insulin resistance in mice.

**Results & Conclusions:** In Caco-2/TC7 cells, insulin action diminished the transepithelial transfer of sugar and reduced BBM and basolateral (BLM) GLUT2 levels, demonstrating that insulin can target sugar absorption by controlling the membrane localization of GLUT2 in enterocytes. Similarly, in HI-EG clamp experiments in sensitive mice, insulin abolished the GLUT2 (i.e. cytochalasin B-sensitive component of fructose absorption), decreased BBM GLUT2 and concomitantly increased intracellular GLUT2. Acute insulin treatment prior to sugar intake prevented the insertion of GLUT2 into the BBM. Insulin resistance in mice provoked a loss of GLUT2 trafficking and GLUT2 levels remained permanently high in the BBM and low in the BLM. We propose that, in addition to its peripheral effects, insulin inhibits intestinal sugar absorption to prevent excessive blood glucose excursion after a sugar meal. This protective mechanism is lost in the insulin-resistant state induced by high-fat or high-fructose feeding.

**ABBREVIATIONS.** BBM (brush border membrane), BLM (basolateral membrane), Caco-2/TC7 (colon carcinoma TC7 subclone), HI-EG (Hyperinsulinemic Euglycaemic Clamp), GIR (glucose infusion rate), PepT1 (Peptide transporter-1), PIP\(_3\) (Phosphatidylinositol (3,4,5)-trisphosphate)
Intestinal sugar transport constantly adapts to the dietary environment. At low levels, the end products of carbohydrate digestion are absorbed by a two-step membrane-transport processes involving the sodium-dependent glucose cotransporter (SGLT1) and the facilitative fructose transporter (GLUT5) in the brush border membrane (BBM) (1) lining the lumen. GLUT2 in the basolateral membrane (BLM) (2), ensures sugar exit into the blood stream (3). The level of sugar absorption is also regulated by a rapid and transient recruitment of GLUT2 into enterocyte BBM (4; 5). A high sugar intake is a physiological regulator of this process, increasing monosaccharide uptake 3-fold in vivo (6). The recruitment of GLUT2 in BBM was also observed in conditions of increased calorie demand and Glucagon-Like-Peptide 2 treatment (7-11).

The mechanisms by which GLUT2 leaves the BBM in the absence of luminal sugar (interprandial periods) are unknown. Of the possible physiological stimuli occurring during feeding, insulin was thought to be a candidate because it exerts systemic hypoglycemic effects by stimulating the translocation of GLUT4 into the plasma membrane of skeletal muscle and adipose cells (reviewed in (12)) and decreasing liver glucose output. Furthermore, the presence of GLUT2 in BBM was initially found under conditions of experimental diabetes i.e. lack of insulin and consequent hyperglycaemia (13). The underlying mechanisms for the loss of BBM GLUT2, i.e. degradation, internalization and transcytosis, need to be determined.

An acute inhibition of sugar absorption by insulin was interpreted as being an indirect consequence of metabolic flow and entirely attributed to SGLT1 (14). The aim of this study was therefore to analyze the impact of insulin on intestinal sugar absorption, focusing on the localization of GLUT2 in enterocyte membranes, intestinal sugar absorption, focusing on the localization of GLUT2 in enterocyte membranes.

RESEARCH DESIGN AND METHODS
Cell culture. Caco-2/TC7 on cells filters (3µm HD, BD Biosciences, France) were grown in DMEM (Gibco, Paisley, UK) supplemented with 25mmol.L⁻¹ glucose and 10% decomponented fetal calf serum (FCS) (AbCys, France) (15). Insulin stimulation (1µM) in basolateral compartment, 1h, 37°C was performed in serum-free media. Differentiated cells were washed with ice-cold PBS/Ca²⁺/Mg²⁺ buffer before processing.

Fluorescence microscopy, image acquisition and treatment. Caco-2/TC7 cells were fixed with 4% paraformaldehyde (W/V) at 4°C and antibody labeling was made on permeabilized cells with 1% TritonX100. Antibodies were added for 1h and increased washing with PBS/glycine buffer performed between steps. Secondary antibodies were labeled with cyanin. Images were taken using a confocal microscope (Zeiss, LSM 510) and Leica DMRXA microscope with CCD camera (5MHz Micromax 1300Y Roper Instruments) with the Metamorph software (Universal Imaging Corp.) to deconvolute Z-series. Colocalizations were analyzed with the ImageJ1.37 software (http://rsb.info.nih.gov).

Cell surface biotinylation. Cells were labeled from the BLM or BBM side with 1mg.ml⁻¹ membrane-impermeant sulfo-NHSS-Biotin and biotinylated proteins purified on NeutrAvidin™ gel columns (Pierce, Rockford, IL) as described in (11).

Insulin signaling pathway. Total AKT and phospho-AKT contents were evaluated on frozen cells. Western blots on cleared lysates (mmol.L⁻¹: 20 Tris, 150 NaCl, 5 EDTA, 1% v/v TritonX100, 0.5 % deoxycholate) were revealed with specific AKT and Ser473P-AKT antibodies (Cell Signaling Technologies).

Animals. C57bl/6J mice were fed with isocaloric 65% glucose-, fructose- or protein-rich diets (6) or 72% fat, low sugar (18) or standard chow (M25, Safe, France). Body weights of standard chow and fructose-fed age-matched mice after 30 days (23.5±0.5 g vs. 23.1±0.4; n=24) as well as age-matched fed standard chow and high-fat diet after 4 month (27.4±0.3 g vs. 24.6±0.4; n=20) were similar. Gavages were performed after an 18h-fast using saline or 4 mg.g⁻¹ bwt glucose or fructose solutions. Intraperitoneal injections of saline (sham) or 1mU.kg⁻¹ insulin (Actrapid, Novo Nordisk) were performed 20 min prior to gavage. Tail blood glucose (Accu-check, Roche Diagnostic) and plasma insulin (EIA, Linco Research, Missouri) were measured. The hyperinsulminemic-euglycemic (HI-EG) clamp experiments were performed in conscious mice after a 5h-fast
Insulin regulates intestinal sugar absorption

(19). Insulin perfusions in the jugular vein were at 0.6 mU/kg·h and glucose (15 g/dl) at variable rate to maintain euglycaemia (mg/dl). Fructose diet: clamp 122±17 n=9, sham 135±10 n=9; high-fat diet: clamp 148±37 n=4; sham 152±15; n=4. Glucose infusion rates (GIR) were calculated at the end of a 40 min clamp. Membrane preparations. Fresh jejunal mucosa scrapings were processed for membrane preparations. Membrane preparations in iodixanol (Optiprep™ Abscys) gradients were performed as described in (20) with 6 mg protein (BCA protein kit, Pierce) were laid on top of a 7.5%-30% gradient and step fractions (500µl) collected for density fractions and Western blot analysis. BBMs were purified using the MgCl₂/EGTA precipitation method (21).

Western blotting. Western blots were performed as described in (6). Primary antibodies against rGLUT5, hGLUT5-Cterm (22), sucrase-isomaltase (23), NaKATPase-alpha1 (Abcam, Cambridge, UK), Insulin receptor-ß1 (Santa Cruz, Ca) and PepT1 were used. For GLUT2 labels, rhGLUT2 (extracellular TM1-TM2 loop) (6), hm GLUT2 (Santa Cruz Biotech, Ca), hGLUT2-Nter peptide gave similar results.

Sugar transport assays. The transepithelial transfer of [³H]-3-O-methyl-glucose (3-OMG) in 25 mmol.L⁻¹ glucose-DMEM was measured in control and insulin-treated Caco-2/TC7 cells. The radioactivity was added at the BBM side of cells and 50µl triplicate BLM medium samples were counted after 30 min. Results: pmoles ± SEM from 9 filters. Fructose uptake in jejunal tissue-rings was measured as described in (6). Uptakes are in nmol. mg protein⁻¹·s⁻¹.

Statistics: Results are given as (means ± SEM; n), and ANOVA tests were performed.

RESULTS
BBM GLUT2 is decreased by insulin in Caco-2/TC7 cells. First of all, the effects of insulin on enterocytes were studied in Caco-2/TC7 cells (15; 24), which enabled a separation between the effects of insulin and glucose in vitro. Differentiated Caco-2/TC7 cells segregate sucrase-isomaltase in BBM and NaKATPase in BLM (Figure 1A, On-line appendix [available at http://diabetes.diabetesjournals.org]). The β1-insulin receptor was found in BBM and BLM and in intracellular stores (XZ sections). In this experimental setting, glucose was added to the apical side and insulin to the basolateral side of the cells in order to mimic in vivo sugar absorption. Culture media were supplemented with 10% FCS containing insulin that could induce internalization of the insulin-receptor. Glucose concentrations had not changed significantly at harvest (not shown). Insulin receptors were functional, as shown by AKT phosphorylation kinetics (Figure 1B, Online appendix).

Confocal microscopy analysis of the colocalization of GLUT2 with sucrase-isomaltase (Figure 2A) indicated that GLUT2 was associated with BBM. GLUT2 was also abundant in sub-apical compartments where sucrase-isomaltase was undetectable. The quantification of GLUT2 and sucrase-isomaltase colocalization (merge, line 1) indicated that insulin (1mU.ml⁻¹) reduced BBM GLUT2 by 60% (Figure 2D, Fluoro).

To distinguish GLUT2 in intracellular membranes from BBM GLUT2, we performed surface biotinylation of proteins with impermeant NH-SS-biotin. When biotin was added to the BBM side of cells, western blotting of biotinylated proteins revealed GLUT5 and GLUT2 in BLM (Figure 2C). NaKATPase, could be biotinylated from the BLM (data not shown), but not from the BBM side of cells (Figure 2C), indicating integrity of intercellular tight junctions. After exposure to insulin for 1h, GLUT5, GLUT2 and NaKATPase levels in total membranes were unchanged (Figure 2C) but decreased BBM GLUT2 by 80% (Figure 2D, Biotin) whereas GLUT5 remained unchanged.

A precise deconvoluted image of a single microscopy section at mid height of enterocytes shows association of GLUT2 with BLM identified by NaKATPase labeling (Figure 2E). The addition of 25mM glucose increased BLM GLUT2 2-fold after 15 min and 4-fold after 30 min as compared to sugar-deprived cells (Figure 2F). BLM GLUT2 at 30 min contained 12% of total GLUT2 in whole cell. Insulin treatment decreased BLM GLUT2 by 45% and 34% after 15 and 30 min respectively (P<0.05) (Figure 2F).

To determine the functional consequences of insulin action, we measured the transfer of the non-metabolizable glucose analogue 3-OMG from the apical to the basolateral compartment. The results revealed a 30% (P<0.01) inhibition after a 30 min insulin-treatment (Figure 2B), which reflected the inhibition of GLUT2-mediated transport in Caco-2/TC7 cells. Thus insulin decreases BBM and BLM GLUT2 levels in enterocytes and limits the transepithelial transport of sugars by a specific internalization of GLUT2 from plasma membranes.

High-fructose or high-fat feeding on intestinal GLUT2 expression and glucose homeostasis in mice in vivo. Mice fed a fructose-rich diet have a 2-fold (2.25±0.9; n=14) higher expression of GLUT2 in the jejunum as compared to chow fed mice. The maximum increase was attained after 5 days and
remained stable thereafter. On the contrary, mice fed a high-fat diet expressed basal levels of GLUT2 (1.04±0.3; n=7) like chow fed mice.

The consumption of fructose-rich and high-fat diets perturbed mice glucose homeostasis (Table I, On-line appendix). Glucose intolerance (ΔG) was found as soon as 5 days after the beginning of the fructose diet and after 3 months of fat feeding (not shown). Increased ΔG/ΔI ratios indicated that mice became insulin-resistant and fasting blood glucose levels were slightly increased.

To investigate insulin action in the intestine, we performed hyperinsulinemic-euglycemic (HI-EG) clamps in conscious mice. Moderate hyperinsulinemia (2.36±0.34 vs. 0.49±0.05 ng.mL⁻¹; n=4) was obtained. Glucose infusion rates (GIR) were similar in chow- and fructose-fed mice for 15 days (67±1 and 65±4 mg.kg⁻¹.min⁻¹; n=4) indicating similar insulin sensitivity. However GIR were decreased at day 30 in fructose fed mice, indicating they were insulin-resistant (36±6 mg.kg⁻¹.min⁻¹; n=6). The GIR of mice fed a high-fat diet for 4 months was reduced even more (7±4 mg.kg⁻¹.min⁻¹; n=6) indicating very high insulin resistance. We were then able to study insulin effects on GLUT2 trafficking by comparing sensitive and resistant mice.

**Insulin action on the membrane distribution of GLUT2 in insulin-sensitive mice.** Enterocyte membranes were separated according to densities (1.057 to 1.150 g.L⁻¹) in iodixanol gradients (20). GLUT2 was found in BBM fractions containing the PepT1 marker and in BLM containing the NaKATPase marker (Figure 3A,B). Intracellular (non BBM, non BLM membranes) also contained GLUT2. These results confirmed the presence of GLUT2 in BBM and intracellular GLUT2 stores in vivo.

Insulin lowered GLUT2 by 60% in PepT1 fractions indicating removal from BBM (Figure 3B,C) in absence of major protein degradation since total GLUT2 levels in sham and HI-EG clamp conditions were unchanged (Figure 3B insert). Accordingly, GLUT2 increased in intracellular membranes (Figure 3C). Thus, insulin promoted GLUT2 internalization from BBM to intracellular membrane stores.

The functional significance of the insulin-dependent internalization of BBM GLUT2 with respect to sugar absorption was measured in everted jejunal rings of sham and HI-EG mice (Figure 3D). Insulin reduced fructose uptake by 40% (P<0.001), indicating that sugar transport correlated to BBM transporter abundance. We used cytochalasin B (CB) to determine the contributions of GLUT2 and GLUT5 (insensitive to CB) to fructose uptake. Fructose uptakes were identical in HI-EG and sham mice exposed to CB, thus confirming that insulin did not change localization and activity of GLUT5. Insulin specifically decreases GLUT2-dependent fructose transport.

**Impairment of GLUT2 trafficking by insulin injection prior to a sugar bolus.** The effect of insulin in HI-EG mice reduced GLUT2-dependent fructose uptake in conditions of fast. During a sugar meal, luminal sugars are high triggering GLUT2 translocation to the BBM, then blood glucose and plasma insulin rise. To document the impact of luminal sugar and insulin on the distribution of GLUT2 in enterocyte membranes, we force-fed insulin-sensitive mice with fructose and elicit GLUT2 trafficking into BBM (Figure 3 E,F,G).

Intraperitoneal insulin injections performed 20 min prior to the fructose test meal produced a 3-fold elevation of plasma insulin (range 1.2 to 1.7 ng.mL⁻¹) to a level similar to that produced during HI-EG clamp. The recruitment of GLUT2 into BBM in response to luminal fructose was abolished by insulin injection, and the distribution of GLUT2 was similar to that in enterocytes of fasted mice (Figure 3E). Interestingly, insulin effects tended to be weaker in purified BBM from mice fed 5 days with the fructose-rich as compared to glucose-rich diet (Figure 3G). Prior insulin treatment counteracted the insertion of GLUT2 into BBM induced by luminal dietary sugar (Figure 3E).

**GLUT2 membrane distribution and trafficking in insulin-resistant mice.** The membrane distribution of GLUT2 in insulin-resistant mice obtained by prolonged fructose (Figure 4 A,B,C) or fat feeding (Figure 4 D,E,F) was markedly perturbed. High levels of GLUT2 were found in BBM-PepT1 fractions (Figure 4A,D), which represented a 40 % of total GLUT2 in the enterocyte membranes of mice fed fructose-rich diet for 30 days (Figure 4C). This proportion increased to 80% in high-fat conditions (Figure 4E). BLM GLUT2 levels were 10% (high-fructose; Figure 4C) and 20% (high-fat; Figure 4F) of total GLUT2 as compared to 40% in insulin-sensitive mice (Figure 3C). The functional impact of this redistribution of GLUT2 was revealed by the 2-fold increase in the initial slope (0-15min) of blood glucose after an OGTT in resistant mice (high-fat 0.30 ± 0.02; high-fructose 0.28 ± 0.01 mg.dl⁻¹.min⁻¹; n=8) as compared to sensitive mice (0.13 ± 0.01 mg.dl⁻¹.min⁻¹; n=8; p< 0.001). Thus in insulin resistant states, BBM GLUT2 was high and increased blood glucose entry.

In insulin resistant mice, BBM localizations were unaffected by insulin in HI-EG clamp conditions. BLM GLUT2 remained below 20% of total GLUT2 (Figure 4 C,F). In addition BBM GLUT2 levels were correlated with GIR. Insulin
resistant animals, a permanent localization of GLUT2 in enterocyte BBM, a pathological consequence of long-term fructose or fat feeding.

DISCUSSION

We had previously reported the rapid and transient recruitment of GLUT2 into enterocyte BBM after a bolus of simple sugars, generating a 3-fold enhancement of uptake (6). The present study demonstrates and quantifies in vitro and in vivo, the inhibition by insulin of intestinal sugar uptake, as a result of the internalization of GLUT2 from plasma membranes back into intracellular pools. This important physiological process mediated by insulin at the level of enterocytes probably reveals another mechanism by which insulin limits sugar excursion in the blood during a sugar-rich meal. Furthermore, insulin resistance provoked a loss of control by insulin of GLUT2 membrane trafficking, a pathological consequence of the long-term consumption of a fructose-rich or high-fat diet.

Insulin sensitivity and glucose tolerance are important factors in the regulation of intestinal GLUT2 trafficking. In insulin-resistant mice obtained by prolonged fructose-rich or high-fat diets, the distribution of GLUT2 was drastically altered; BBM GLUT2 was permanently high and insulin was unable to promote GLUT2 internalization and enrichment in intracellular membranes. Insulin resistance resulted in increased intestinal sugar delivery, as reflected by higher initial rise in blood glucose after oral glucose load.

Insulin action is transduced via receptors in enterocyte plasma membranes in vivo (25; 26) and in vitro in Caco-2/TC7 cells. The functional significance of BLM insulin receptors is obvious and we speculate that enterocytes akin epithelial renal cells undergo a rapid PiP3 diffusion after insulin to trigger trafficking of a BBM protein (27). The role of insulin receptors in the BBM is unclear but it creates an opportunity for oral insulin treatment (28) that might target BBM GLUT2 localization and function. Conflicting results concerning acute insulin treatment report inhibition (14) and activation (29) of SGLT1-dependent intestinal sugar absorption ex vivo. In the present work, we describe an insulin action on the high capacity GLUT2 component of fructose transport that leads to a 50% inhibition of total transport, down to GLUT5 basal transport levels. We speculate that this regulation applies to intestinal glucose absorption. In insulin resistant animals, a permanent BBM GLUT2 might create glucose efflux from enterocytes into the lumen. This is unlikely since permanent SGLT1 recapture would operate. These results highlight the complexity of the regulation of intestinal sugar absorption by insulin to modulate but not block intestinal sugar absorption.

In adipocytes and muscle cells, insulin provokes a massive translocation of GLUT4 to the plasma membrane (for a review (12; 30)). In sharp contrast, insulin internalizes BBM GLUT2 into intracellular pools. Insulin, by regulating GLUT2 and GLUT4 traffic in opposite direction, controls glucose homeostasis and limits glucose excursion in the course of digestion by increasing peripheral GLUT4-dependent glucose uptake and slowing down intestinal GLUT2-dependent sugar delivery. We therefore propose that the small intestine constitutes an early checkpoint to limit postprandial glucose excursion. We also anticipate that insulin treatment will remove GLUT2 from its permanent BBM localization in type I diabetic patients that are insulin sensitive.

Modern westernized diets and eating habits have changed, increasing the amounts of dietary fructose and fat, probably a nutritional basis for the obesity and type II diabetes pandemics (31). Intestinal adaptation to these diets may be an early event to the onset of metabolic disorders due to the rapid increase in sugar transport capacities and the alteration of insulin action in enterocytes. With time insulin resistance and metabolic disorders (32) might be worsen by uncontrolled sugar absorption in a small intestine that is no longer responsive to attenuation by insulin. A vicious circle thus develops that equips the intestine for high transport of dietary sugar in organisms already suffering from excessive blood glucose levels. Strategies to ameliorate insulin action on intestinal function therefore constitute another target for therapeutic intervention and control of post-prandial glycaemia.

ACKNOWLEDGMENTS

This work received support from INSERM, University Paris6, ATC Nutrition ASEO22129DSA, ALFEDIAM/Merk-Lipha grants. VT and ES were recipients of MRT PhD fellowships, MLG of a Benjamin Delessert prize, and AGB from FPI Spanish Ministerio de Educacion y Ciencia (BES-2004-4685). We wish to thank E. Petridi and V. Ondet for their contribution, C. Lasne and E. Dussaux for technical assistance. We are grateful to J. Chambaz for support, to GW Gould (Glasgow, UK) and GL.Kellett (York, UK) for antibodies and GLK for fruitful discussions.
REFERENCES

2. Thorens B: Molecular and cellular physiology of GLUT-2, a high-Km facilitated diffusion glucose transporter. *Int Rev Cytol* 137:209-238, 1992


Insulin regulates intestinal sugar absorption

FIGURE LEGENDS

Figure 1 (On-line appendix): Caco-2/TC7 cells express a functional insulin receptor at the plasma membrane.
Confocal microscopy analysis of differentiated Caco-2/TC7 cells cultured on a porous support in 25 mmol.L$^{-1}$ glucose-DMEM supplemented with 10% FCS. Arrows across enterocyte schemes show the planes of confocal imaging. (A) Images of insulin receptor ß1, sucrase-isomaltase, NaKATPase taken at BBM (XY, column 1) and BLM plans (XY, column 2) and XZ sections (column 3). Secondary antibodies were labeled with CY3. (B) Representative experiment of AKT phosphorylation following the addition of 1mU.ml$^{-1}$ insulin for 10, 20, 40 and 60 min to 25 mmol.L$^{-1}$ glucose DMEM without FCS.

Figure 2: Impact of insulin on BBM and BLM GLUT2 in Caco-2/TC7 cells
Caco-2/TC7 cells were cultured on a porous support, in 25 mmol.L$^{-1}$ glucose-DMEM for 21 days and exposed to 1 mU.ml$^{-1}$ insulin for 1h in serum free media. (A) Immunofluorescence study of GLUT2 (CY2, green) and sucrase-isomaltase (CY3, red) membrane colocalization (merge yellow) in control and insulin-treated cells. Confocal images were taken through the BBM plane or just below BBM in a subapical compartment or through nuclei. (B) Transepithelial transport after 30 min across control and insulin-treated cells was measured using 3-OMG as a tracer. Results nmoles ± SEM from 9 filters are shown. (C) Surface biotinylation with membrane-impermeant NH-S-S-biotin was performed from the BBM side of cells. Western blots of GLUT2, GLUT5 and NaKATPase in 10% of total protein or biotin-proteins purified on Neutavidin$^{TM}$ columns from 90% total proteins are shown. (D) Quantification of insulin effects by colocalization of GLUT2 and sucrate-isomaltase over total sucrate-isomaltase positive cells (Fluo) counted from 4 independent cultures with 10 fields and 200 control and 200 insulin-treated cells. Scan density analysis of biotin-GLUT2 in insulin-treated cells (Biotin) were quantified in 8 monolayers from 4 experiments and expressed relative to biotin-GLUT2 in parallel control cells. (E,F) Deconvoluted images taken at a single plane are shown. (E) BLM GLUT2 (CY2) and NaKATPase (CY3) in a representative experiment out of 3, showing cells cultured in glucose-free (G0), in glucose (G25) or in glucose with 1mU.L$^{-1}$ insulin. (F) The colocalization of GLUT2 with NaKATPase was quantified in all the planes imaging the whole cell and expressed as percent of total GLUT2 in glucose and/or insulin treated cells. Statistical significance *** $P<0.001$, **$P<0.01$, *$P<0.05$

Figure 3: GLUT2 membrane trafficking in insulin-sensitive mice
(A,B,C) Hyperinsulinenic-euglycaemic (HI-EG) clamp experiment in mice. (A) Western blot of GLUT2, PepT1 and NaKATPase abundance in fractions of iodoxanol density-gradient of total membranes prepared from a single intestine in either sham or HI-EG clamp conditions. HI-EG clamp experiments were performed after a 5h-fast in mice fed a high-fructose diet for 15 days. The intestines of sham and HI-EG mice were taken at the end of a 2h clamp. Gradient fractions are sorted from left to right by increasing densities ranging from 1.06 to 1.16 g/ml. (B) Quantification of NaKATPase (open circles), PepT1 (open triangles) and GLUT2 in fractions from 4 paired sham (open squares) and HI-EG (black squares) mice. Scan densities are shown as ratios of protein levels in a fraction to its maximum level in the gradient (arbitrary unit ± SEM). Insert: GLUT2 in the total membranes loaded on gradients (sham: white bar; HI-EG: black bar). (C) GLUT2 membrane distribution calculated from B in BBM (sum in fractions containing PepT1), BLM (sum in fractions containing NaKATPase) and non plasma membranes (intracellular membranes; all other fractions). (D) Fructose uptake (10 mmole.L$^{-1}$ for 2 min at 37°C) in jejunal everted rings from sham (white bar; n=3) and HI-EG (black bar; n=3) and sham after 100 µmole.L$^{-1}$ cytochalasin B preincubation for 10 min (hatched bar; n=3). Uptakes were measured in individual mice intestines (12 rings per mouse). (E,F,G) Effect of an insulin injection prior to a fructose test meal. Mice were fed a fructose diet for 5 days and fasted overnight. Mice were then injected either intraperitoneal PBS or insulin (1 mU.kg$^{-1}$), 20 min prior to a fructose (90 mg.kg$^{-1}$) test meal. Intestines were taken 30 min after the test meal. (E) Representative western blot (12 mice in 4 experiments) of the distribution of GLUT2 in iodoxanol fractions. Fractions containing BLM (NaKATPase) and BBM (PepT1) have been boxed for clarity. (F) Comparison of the effect of a 5-day glucose-rich or fructose-rich diet on BBM GLUT2 levels in the response an insulin injection prior a fructose test meal (protocol similar to E). Representative western blot of purified jejunal BBM using the Mg$^{2+}$/EGTA precipitation method. (G) Scan density quantification (means ± SEM; n=6) of BBM GLUT2 in conditions of PBS (open bars) or 1mU/Kg insulin (black bars). Statistical significance *** $P<0.001$; ** $P<0.01$. 

9
Figure 4: Insulin resistance remodels the distribution of GLUT2 in enterocyte membranes

Insulin-resistance was induced in mice by feeding a fructose-rich diet for 30 days (A,B,C) or a high-fat diet for 4 months (D,E,F) and intestines were processed as described in Figure 3. (A,D) Western blot (out of 4) of GLUT2, PepT1 and NaKATPase in sham and HI-EG clamp in mice. (B,E) Average values from 4 mice of GLUT2, NaKATPase and PepT1 protein abundance in gradient fractions (same symbols as in Figure 3B). Data are expressed as the ratio of the level of protein in a given fraction to the maximum level of protein in the gradient (arbitrary units ± SEM). Insert: GLUT2 in the total membranes loaded on gradients (sham white bar; HI-EG black bar). (C,F) GLUT2 membrane distribution calculated from B and E in BBM (sum in fractions containing PepT1), BLM (sum in fractions containing NaKATPase) and non plasma membranes (intracellular membranes; all other fractions).
Figure 2
Insulin regulates intestinal sugar absorption

Figure 3
Insulin regulates intestinal sugar absorption

Figure 4