

## SUPPLEMENTAL TABLES AND FIGURES

### Placenta-specific *Slc38a2*/SNAT2 knockdown causes fetal growth restriction in mice.

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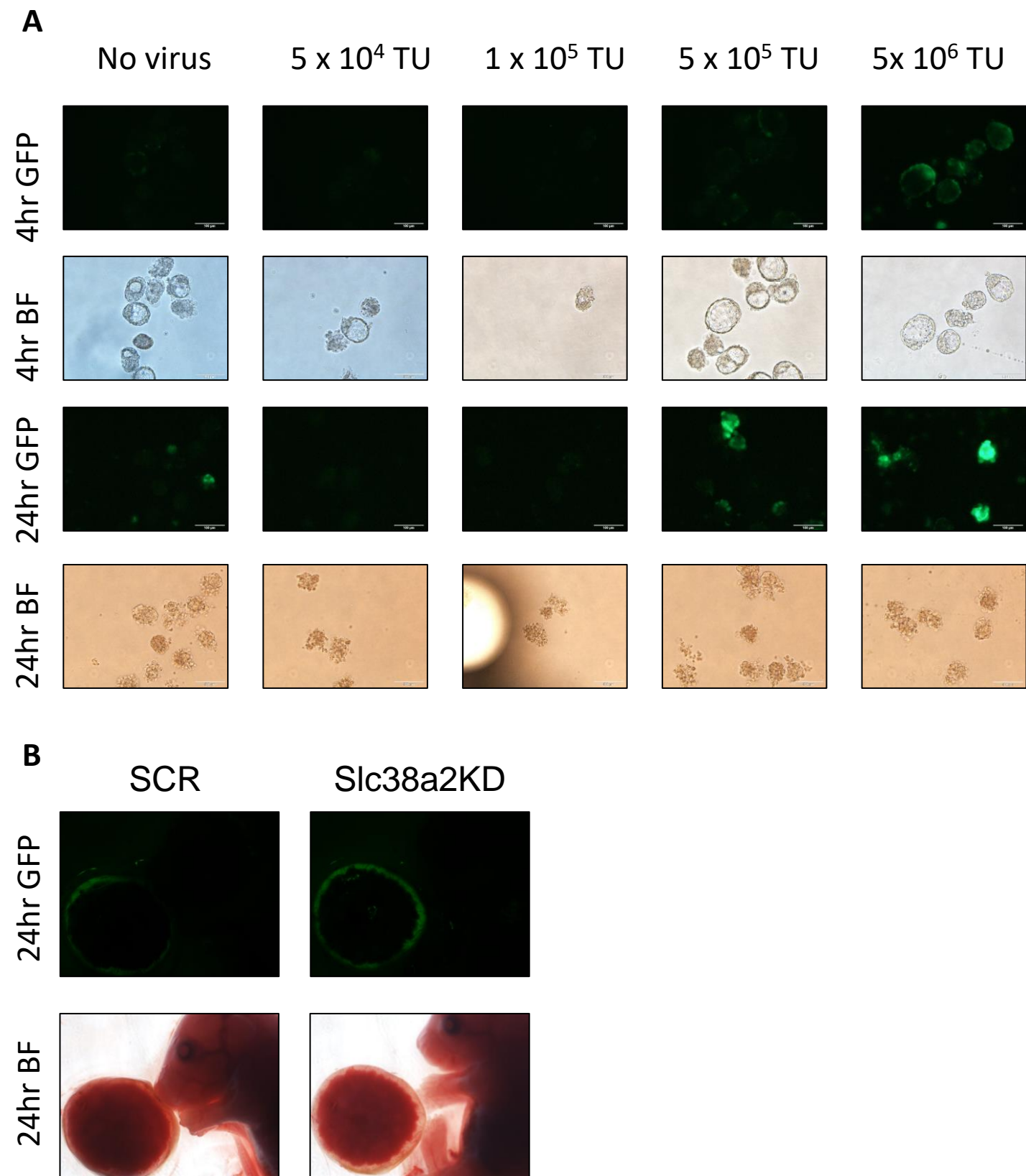
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#### Contact information:

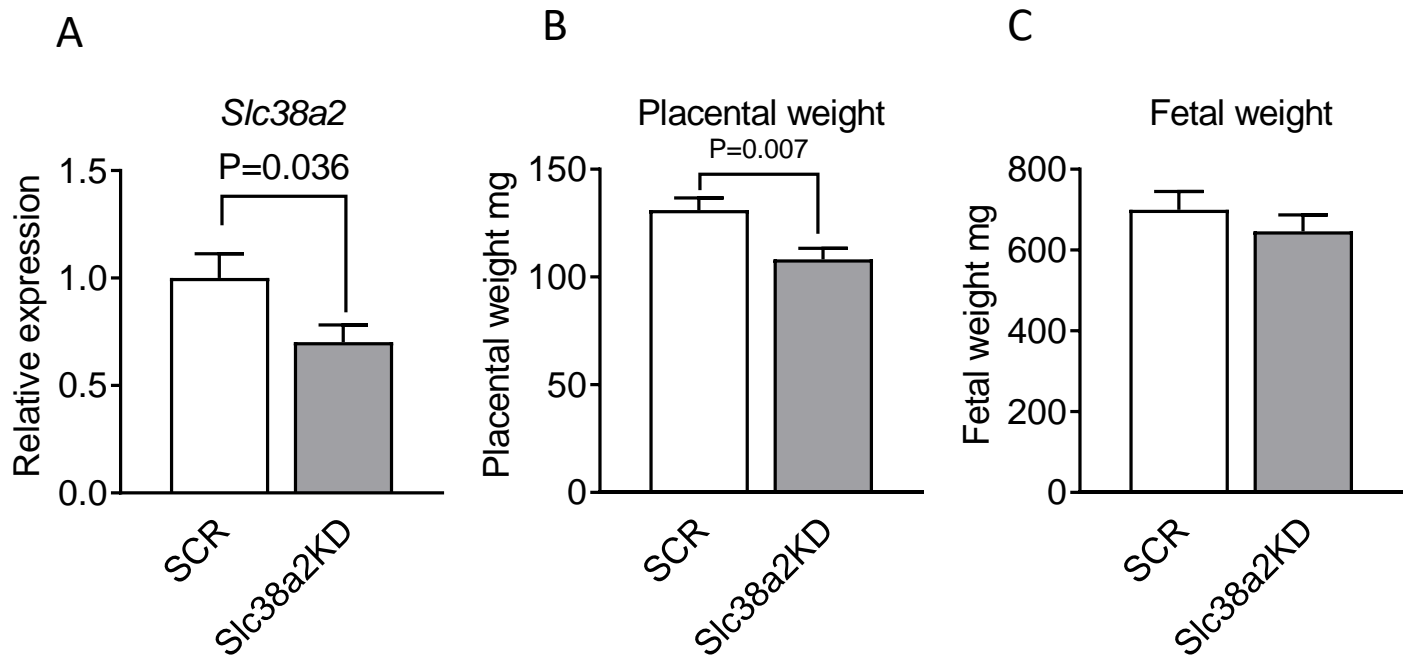
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#### Conflict of interest statement.

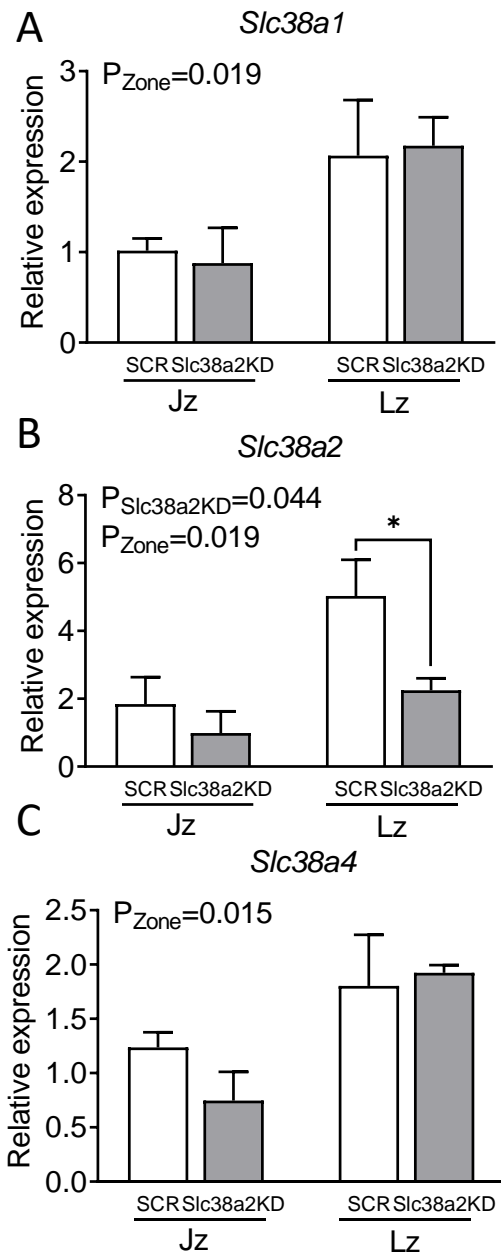
The authors have declared that no conflict of interest exists.



**Fig. S1 GFP reporter expression in lentivirus transduced embryos.** (A) Fluorescence and bright field (BF) images of blastocysts transduced with varying titers (TU, transforming units) of Slc38a2KD lentivirus, immediately after 4 hr transduction (E3.5) or 24 later (E4.5). (B) Fluorescent and bright field images of embryos collected near term (E17.5) following transduction with SCR or Slc38a2KD lentivirus.



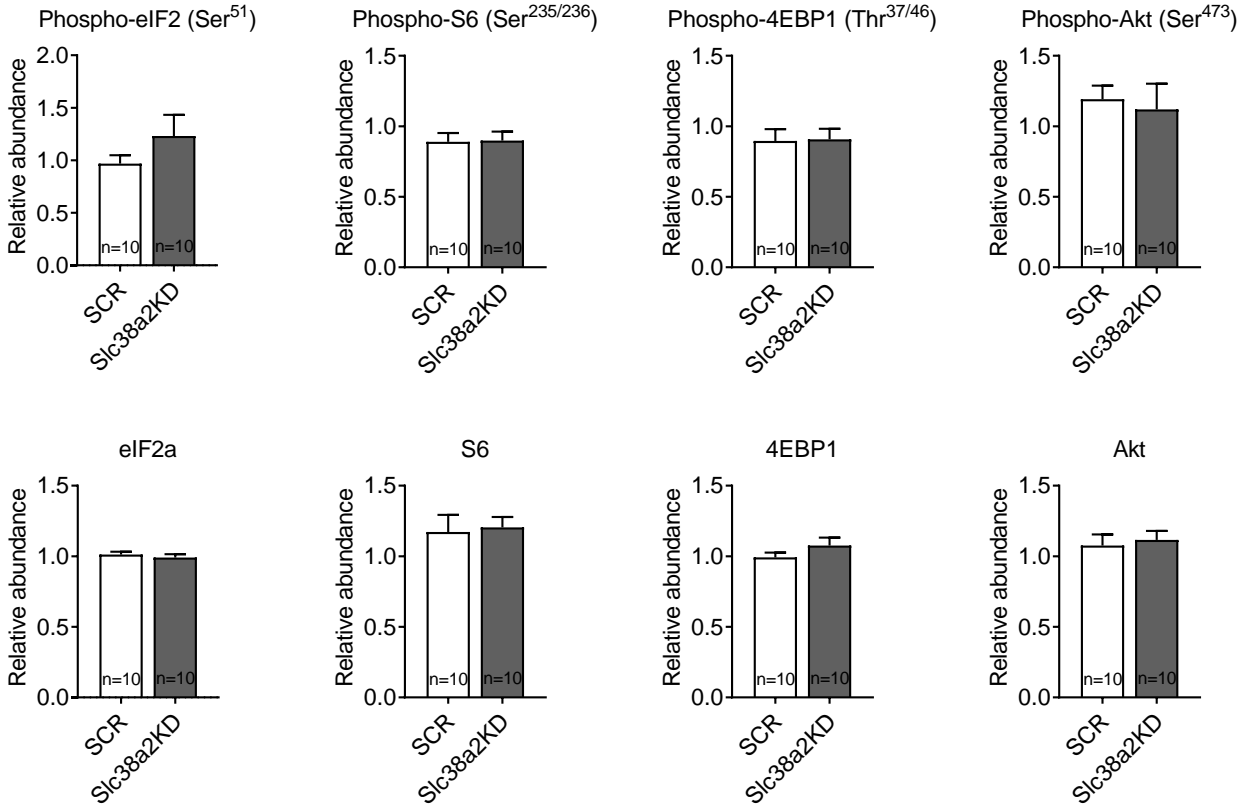
**Fig. S2 *Slc38a2* gene expression (A), placental (B) and fetal weights (C) in SCR and *Slc38a2* transduced embryos at E17.5.** SCR, n=11 conceptuses and *Slc38a2*, n=14 conceptuses from four litters, each with at least 1 conceptus transduced with each virus. Mean + SEM. P values for statistically significant differences ( $P<0.05$  by Student's t-test) between SCR and *Slc38a2* given in figure.



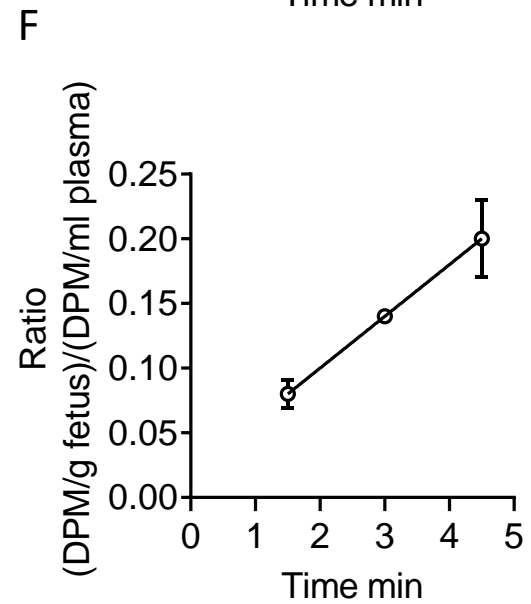
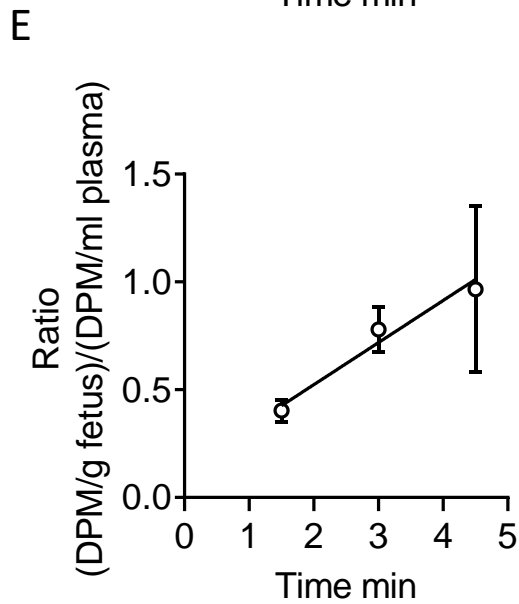
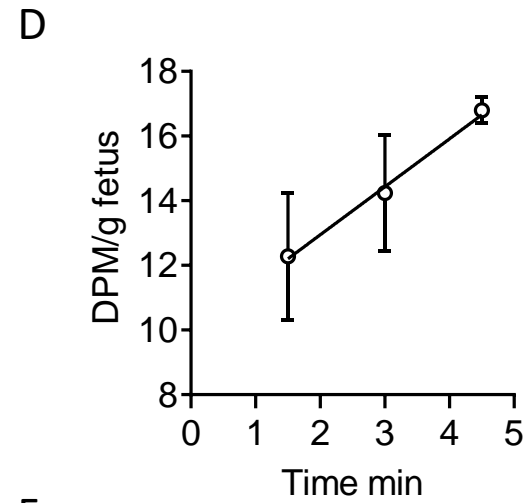
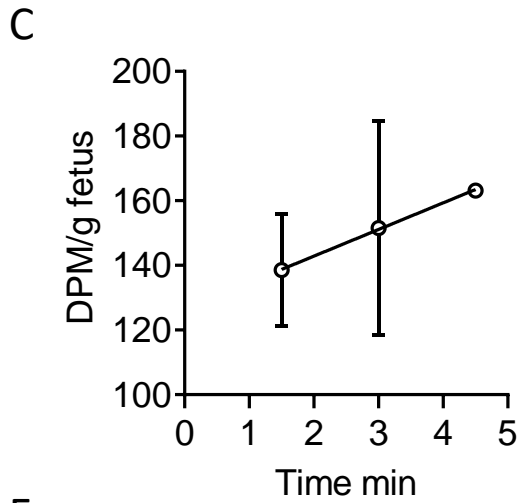
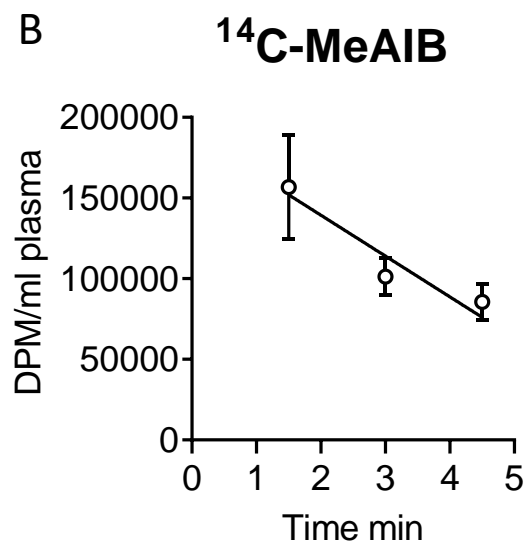
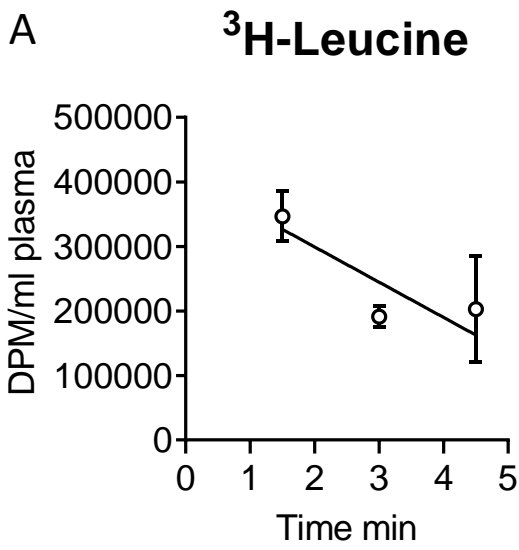
**Fig. S3** *Slc38a1* (A), *Slc38a2* (B) and *Slc38a4* (C) gene expression in dissected junctional (Jz) and labyrinthine (Lz) zones of SCR and *Slc38a2KD* placentae. n=3 litters (15 SCR placentas, 9 *Slc38a2KD* placentas). Main effects of *Slc38a2KD* ( $P_{\text{Slc38a2KD}}$ ) and zone ( $P_{\text{zone}}$ ) determined by two-way ANOVA. P values for significant overall effects ( $P < 0.05$ ) given in figure. \*,  $P < 0.05$  Fisher's LSD post-hoc test. Mean + SEM

**Amino acid response signaling**

**mTOR signaling**



**Fig. S4 Amino acid response and mTOR signalling pathway activity in SCR and Slc38a2KD placentae.** Phosphorylated and total protein abundance determined by western blot in pooled placental homogenates. SCR and Slc38a2KD pools compared by paired Student's t –test. No significant differences at  $P < 0.05$ . Mean + SEM



**Fig. S5 Time courses of maternal and fetal radioactivity up to 4.5 min after bolus infusion of  $^{14}\text{C}$ -MeAIB and  $^3\text{H}$ -leucine.**  $^3\text{H}$  (A, C, E) and  $^{14}\text{C}$  radioactivity (B, D, F), in decays per minute (DPM) determined by liquid scintillation counting. (A, B) Maternal plasma radioactivity per ml, (C, D) fetal radioactivity per gram bodyweight, (E, F) ratio of maternal and fetal radioactivity. Points are mean  $\pm$  SEM. 1.5 min, n=4; 3 min, n=3; 4.5 min, n=2. Dependence of radioactivity on time from infusion determined by least squares linear regression. (A-D) Slope does not differ significantly from zero ( $P > 0.05$ ). (E)  $P = 0.03$ ,  $R^2 = 0.53$ ,  $y = (0.20 \pm 0.07)x + (0.13 \pm 0.20)$ . (F)  $P < 0.01$ ,  $R^2 = 0.86$ ,  $y = (0.04 \pm 0.01)x + (0.02 \pm 0.02)$ .

**Table S1 Implantation and viability rates for SCR and Slc38a2KD transduced embryos at E17.5**

	<b>SCR</b>	<b>Slc38a2KD</b>	<b>P value</b>
<b>Total embryos transferred (n)</b>	49	53	
<b>Implanted (n)</b>	11	17	
<b>%</b>	22%	32%	0.375 <sup>a</sup>
<b>Viable fetus at E18.5</b>	11	17	
<b>%</b>	22%	32%	0.375 <sup>a</sup>
<b>Number of fetuses per uterine horn (median, min-max)</b>	3 (1-4)	3 (1-7)	0.971 <sup>b</sup>

Embryos were transferred to 6 recipient dams, of which 5 were pregnant, with any number of implantations, and 1 was non-pregnant on E17.5. SCR and Slc38a2 were compared by <sup>a</sup>Fisher's exact test or <sup>b</sup>Mann-Whitney U-test.

**Table S2 Relationship between human placental *SLC38A2* and clinical characteristics of AGA and FGR subjects delivered at term.**

		<b>Placental <i>SLC38A2</i></b>	<b>Maternal age</b>	<b>Maternal BMI</b>	<b>Gestation al age</b>	<b>Placental weight</b>	<b>Birth weight</b>
<b>Placental <i>SLC38A2</i></b>	<b>r</b>	<b>1.000</b>	<b>0.120</b>	<b>-0.050</b>	<b>0.501</b>	<b>0.664</b>	<b>0.256</b>
	Confidence interval		-0.272 to 0.478	-0.429 to 0.344	0.150 to 0.740	0.251 to 0.873	-0.137 to 0.580
	<i>P</i>		0.550	0.810	0.008	0.005	0.198
	n	27	27	26	27	16	27
<b>Maternal age</b>	<b>r</b>		<b>1.000</b>	<b>0.144</b>	<b>0.195</b>	<b>0.215</b>	<b>0.343</b>
	Confidence interval			-0.258 to 0.503	-0.200 to 0.535	-0.314 to 0.642	-0.0426 to 0.640
	<i>P</i>			0.484	0.330	0.424	0.080
	n		27	26	27	16	27
<b>Maternal BMI</b>	<b>r</b>			<b>1.000</b>	<b>-0.098</b>	<b>-0.071</b>	<b>-0.001</b>
	Confidence interval				-0.468 to 0.301	-0.563 to 0.458	-0.388 to 0.386
	<i>P</i>				0.634	0.800	0.996
	n			26	26	15	26
<b>Gestational age</b>	<b>r</b>				<b>1.000</b>	<b>0.806</b>	<b>0.709</b>
	Confidence interval					0.516 to 0.930	0.451 to 0.858
	<i>P</i>					<0.001	<0.001
	n					16	27
<b>Placental weight</b>	<b>r</b>						<b>0.877</b>
	Confidence interval						0.674 to 0.957
	<i>P</i>						<0.001
	n						16
<b>Birthweight</b>	<b>r</b>						<b>1.000</b>
	Confidence interval						
	<i>P</i>						
	n						27

Interdependence of variables determined by Pearson's correlation.



**Table S3 Estimated parameters for multiple linear regression model to predict effect of FGR status on term placental *SLC38A2* expression, adjusted for confounding variables.**

<b>Variable</b>	<b><math>\beta</math></b>	<b>95% Confidence interval</b>	<b>P value</b>
<b>Intercept</b>	-4.349	-9.87 to 1.17	0.116
<b>Late-onset FGR</b>	-0.389	-0.793 to 0.0150	0.058
<b>Early-onset FGR</b>	0.869	0.503 to 1.23	<0.001
<b>Ethnicity (Asian, black or mixed)</b>	-0.072	-0.413 to 0.270	0.667
<b>Gestational age</b>	0.139	-0.00232 to 0.279	0.054
<b>Hypertension in pregnancy</b>	-0.044	-0.584 to 0.496	0.866

Regression model: Placental *SLC38A2* ~ Intercept + FGR + Ethnicity + Gestational age + Hypertension in pregnancy. Reference levels for FGR status, ethnicity and hypertension were AGA, white and no hypertension, respectively. n=27 subjects

**Table S4 Primer sequences used for qRT-PCR in mouse tissues.**

<b>Target mRNA</b>	<b>Forward primer</b>	<b>Reverse Primer</b>	<b>Reference</b>
<i>Slc38a1</i>	CTTCTCCAGATTTCGTGCCG	AGGGAGAGAGAGAGAAGCCA	(Matoba et al., 2019)
<i>Slc38a2</i>	ACATAAGGCGTATGGTCTG	TACCACAACCCATTTGTATC	(Uno et al., 2015)
<i>Slc38a4</i>	GGCTTCTTCTGCCACTATGC	AAGACCAAAGCCCCAATCTT	(Matoba et al., 2019)
<i>RNA28S</i>	AAGTCCTTCTGACTGAGGCC	ATTCCCAAGCAACCCGACTC	(Uno et al., 2015)