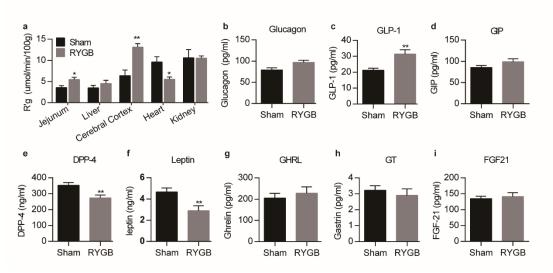
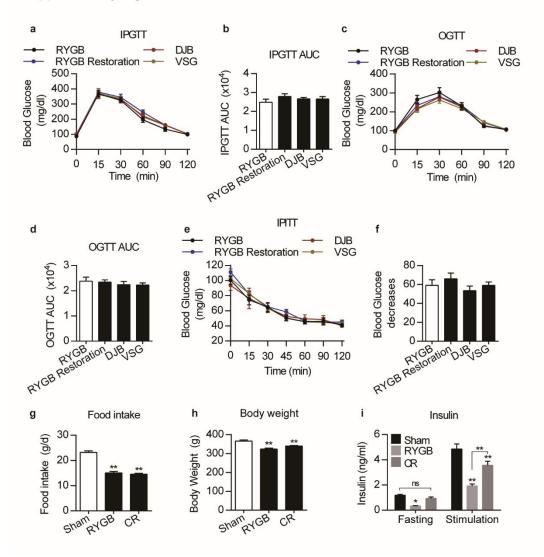


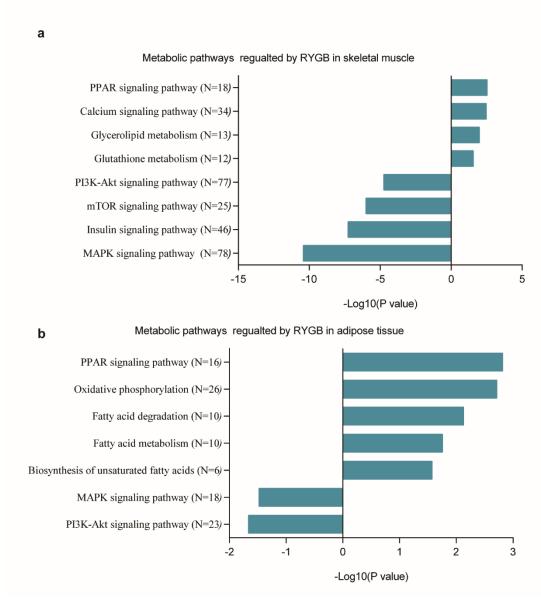
Supplementary Figure S1. Effect of RYGB on physiological parameters in rats. (a) Plasma insulin levels measured by radioimmunoassay in the fasting and glucose-stimulated states are significantly decreased in RYGB rats (n=10). (be) Body weight (b), food intake (c), water intake (d) and urine volume (e) of sham and RYGB rats (n=6-8). Values are shown as the mean \pm s.e.m. Significance was calculated using two-way ANOVA with Tukey's post hoc analysis (a) and repeated-measures analysis (b-e). **P*<0.05 and ***P*<0.01 compared with the sham group.



Supplementary Figure S2. Changes in gastrointestinal hormones and adipokines in plasma after RYGB. (a) Utilization of glucose in jejunum, liver, cerebral cortex, heart and kidney under hyperinsulinemic-euglycemic clamping homeostasis (n=4). (b-i) Plasma levels of gastrointestinal hormones and adipokines (glucagon, GLP-1, GIP, DPP-4, leptin, ghrelin, gastrin and FGF-21) in sham and RYGB rats (n=6-11). Values are shown as the mean \pm s.e.m. Significance was calculated using two-way ANOVA with Tukey's post hoc analysis (a) and two-tailed unpaired t-test (b-i). **P*<0.05 and ***P*<0.01 compared with the sham group.

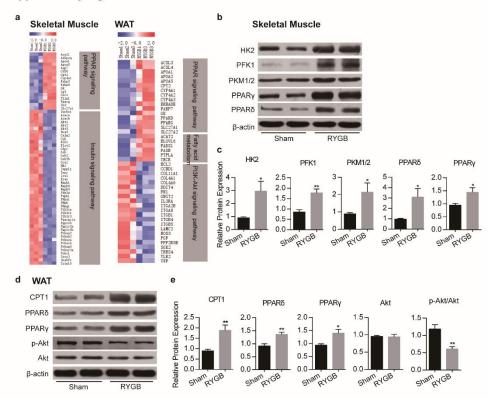


Supplementary Figure S3. Glucose and insulin tolerance tests in rats after different metabolic surgeries. (a-f) IPGTT (2g/kg) (a-b), OGTT(2g/kg) (c-d) and IPITT (e-f) results in rats that underwent RYGB, DJB, VSG or RYGB restoration surgery. (g-h) Food intake, body weight and plasma insulin level in rats that treated with sham, RYGB or caloric restriction (CR). Values are shown as the mean \pm s.e.m. Significance was calculated using two-way ANOVA with Tukey's post hoc analysis (b, d, f, i) and one-way ANOVA with Tukey's post hoc analysis (g, h). **P*<0.05 and ***P*<0.01 compared with the sham group.

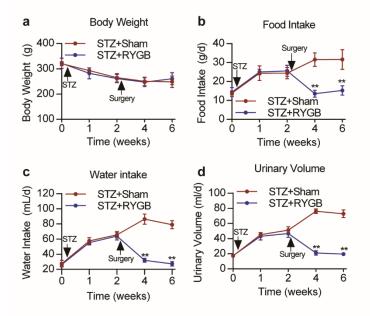


Supplementary Figure S4. Metabolic pathways in rat skeletal muscle and adipose tissue. **(a)** Enriched metabolic pathways in rat skeletal muscle regulated by RYGB (n=3). **(b)** Enriched metabolic pathways in rat adipose tissue regulated by RYGB (n=3).





Supplementary Figure S5. Heatmap and related protein expression in skeletal muscle and WAT after surgery. (a) Expression levels of enriched metabolism pathway genes in skeletal muscle and WAT from sham and RYGB rats. (b-c) Immunoblots of HK2, PFK1, PKM1/2, PPARγ and PPARδ and β-actin in skeletal muscle from sham and RYGB rats (n=6). (d-e) Immunoblots of CPT1, PPARγ, PPARδ, p-Akt and Akt and β-actin in WAT from sham and RYGB rats (n=6). Values are shown as the mean ± s.e.m. Significance was calculated using a two-tailed unpaired t-test (c, e). **P*<0.05 and ***P*<0.01 compared with the sham group.



Supplementary Figure S6. Effect of RYGB on rats with islet disruption. (a-d) Body weight (a), food intake (b), water intake (c) and urine volume (d) of shamor RYGB-treated STZ injected rats (n=6-8). Values are shown as the mean \pm s.e.m. Significance was calculated using repeated-measures analysis (a-d). **P*<0.05 and ***P*<0.01 compared with the sham group.

Supplementary Tables

Table S1

Baseline characteristics of patients with type 2 diabetes underwent metabolic

surgery

	Pre-RYGB (n=87)	Post-RYGB	<i>p</i> value
Age (years)	42±11	_	
Gender (male/female)	49/38	_	—
Duration(year)	5±4	—	_
BMI (kg/m ²)	30.8±2.3	23.4±1.2	0.00
WC (cm)	99.9±2.5	80.6±4.2	0.00
FBG (mmol/L)	8.1±0.6	6.2±0.8	0.00
HbA1c (%)	7.9±0.8	6.2±0.6	0.00
Fasting insulin (mIU/L)	20.1±2.5	12.4±1.6	0.01
SBP (mm Hg)	132±11	122±13	0.00
DBP (mm Hg)	80±11	72±10	0.00
Cholesterol (mmol/l)	4.99±0.25	4.12±0.21	0.00
Triglyceride (mmol/l)	3.06±0.24	1.34±0.17	0.00
HDL-C (mmol/l)	1.28±0.03	1.21±0.05	0.54
LDL-C (mmol/l)	2.70±0.06	2.42±0.07	0.89

The results are mean \pm s.e.m. *p* values are obtained by paired two-tailed Student's t test. BMI, body mass index; FBG, fasting blood glucose; WC, waist circumference; SBP, systolic blood pressure; DBP, diastolic blood pressure.