SUPPLEMENTARY FIGURES AND TABLE

Chronic Repression of mTOR Complex 2 Induces Changes in the Gut Microbiota of Diet-induced Obese Mice

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Supplementary Figure S1. Resveratrol improves, whereas rapamycin impairs, glucose homeostasis. (A) HOMA2-%S index and (B) QUICKI were calculated from fasting glucose and insulin levels (n = 5 per group). Data are expressed as the mean \pm SEM. *F*- and *p*-values are from two-way ANOVA after Bonferroni's post hoc test. **P* < .05, ***P* < .005.



Supplementary Figure S2. Resveratrol ameliorates, whereas rapamycin exacerbates, intestinal inflammation. (A) The concentration of faecal LCN-2 (ng/g faeces) and (B) colon length (cm) were measured after 8 weeks of treatment (n = 5 per group). (C) The correlation between faecal LCN-2 concentrations and colon length was determined using two-tailed Pearson's *r* correlation analysis. Data are expressed as the mean \pm SEM. *F*- and *p*-values are from two-way ANOVA after Bonferroni's post hoc test. **P* < .05, ***P* < .005.



Supplementary Figure S3. Effects of resveratrol or rapamycin treatment on the mTOR signaling pathway in NCD- or HFD-fed mice. (A) p-S6 (Ser235/236)/total S6 (n = 5 per group), (B) p-Akt1 (Ser473)/total Akt1 (n = 3 per group) and (C) cAMP concentrations (n = 5 per group) in liver tissue samples. To examine Akt1 phosphorylation, mice were injected with 0.5U of insulin per kg body weight prior to sacrifice. Data are expressed as the mean \pm SEM. *F*- and *p*-values are from two-way ANOVA after Bonferroni's post hoc test. **P* < .05, ***P* < .005.



Supplementary Figure S4. Changes in the faecal bacterial community following resveratrol or rapamycin treatment. Pie charts showing the relative abundance (%) of different bacterial phyla in the different diet and treatment groups (n = 5 per group).



Supplementary Figure S5. Summary of the effects of resveratrol or rapamycin treatment on the interaction between host physiology and gut microbiota under normal-chow diet or high-fat diet conditions.

Supplementary Table S1. Alpha diversity of the gut microbial communities in NCD or HFD-fed mice after resveratrol or rapamycin treatment. 16S rRNA V1-2 sequences obtained by 454 pyrosquencing were used to analyze alpha diversity at the OTU level. *F*- and *p*-values are from two-way ANOVA after Bonferroni's post hoc test. *P < .05; **P < .005 compared with NCD-CT, #P < .05 compared with HFD-CT.

Group	# Observed OTUs	Chao1	Shannon	Phylogenetic Diversity	Simpson
NCD-CT	301±14	405±16	5.81±0.23	25.13±0.38	0.96±0.01
NCD-Res	306±24	432±30	5.90±0.11	24.86±0.85	0.96±0.01
NCD-Rapa	251±25	335±20	5.00±0.32	21.55±1.07*	0.92±0.01
HFD-CT	229±12*	318±29	5.07±0.13	20.49±0.82**	0.92±0.01
HFD-Res	171±6**	246±20**	4.48±0.06**	17.18±0.77** ^{,#}	0.91±0.01*
HFD-Rapa	215±19*	305±27*	5.15±0.37	18.89±1.21**	0.93±0.02
Interaction	F=3.72, P<.05	F=5.05, P<.05	F=6.06, P<.05	F=4.05, P<.05	F=5.14, P<.05
Treatment	F=1.69, P=0.21	F=1.39, P=0.27	F=1.36, P=0.28	F=4.44, P<.05	F=1.10, P=0.35
Diet	F=29.07, P<.005	F=24.60, P<.005	F=12.30, P<.005	F=47.12, P<.005	F=8.36, P<.05

Source of Variation	Lactococcus	Clostridium XI	Oscillibacter	Pseudoflavonifractor	Flavonifractor	Hydrogeno- anaerobacterium	Howardella
Interaction	<i>F</i> =9.50, <i>P</i> <.005	<i>F</i> =17.61, <i>P</i> <.005	<i>F</i> =13.11, <i>P</i> <.005	<i>F</i> =23.30, <i>P</i> <.005	<i>F</i> =27.30, <i>P</i> <.005	<i>F</i> =10.98, <i>P</i> <.005	<i>F</i> =5.97, <i>P</i> <.05
Treatment	<i>F</i> =10.01, <i>P</i> <.005	<i>F</i> =15.84, <i>P</i> <.005	F=23.19, <i>P</i> <.005	<i>F</i> =11.71, <i>P</i> <.005	<i>F</i> =16.35, <i>P</i> <.005	<i>F</i> =3.88, <i>P</i> =0.07	<i>F</i> =8.92, <i>P</i> <.05
Diet	F=205.9, <i>P</i> <.005	<i>F</i> =85.69, <i>P</i> <.005	<i>F</i> =23.54, <i>P</i> <.005	<i>F</i> =13.51, <i>P</i> <.005	<i>F</i> =3.76, <i>P</i> =0.07	<i>F</i> =8.42, <i>P</i> <.05	<i>F</i> =26.27, <i>P</i> <.005
Source of Variation	Turicibacter	UC_Marinilabiliaceae	Alloprevotella	UC_Porphyro- monadaceae	Ruminococcus	Turicibacter	UC_Marinilabiliaceae
Interaction	<i>F</i> =17.61, <i>P</i> <.005	<i>F</i> =2.75, <i>P</i> =0.12	<i>F</i> =14.49, <i>P</i> <.005	F=7.37, P<.05	<i>F</i> =27.30, <i>P</i> <.005	<i>F</i> =10.53, <i>P</i> <.05	<i>F</i> =12.39, <i>P</i> <.005
Treatment	<i>F</i> =15.84, <i>P</i> <.005	<i>F</i> =3.0, <i>P</i> =0.10	<i>F</i> =19.87, <i>P</i> <.005	<i>F</i> =0.51, <i>P</i> =0.49	<i>F</i> =16.35, <i>P</i> <.005	<i>F</i> =10.44, <i>P</i> <.05	<i>F</i> =14.09, <i>P</i> <.005
Diet	F=85.69, <i>P</i> <.005	<i>F</i> =78.11, <i>P</i> <.005	<i>F</i> =20.84, <i>P</i> <.005	<i>F</i> =47.94, <i>P</i> <.005	<i>F</i> =3.76, <i>P</i> =0.07	<i>F</i> =10.50, <i>P</i> <.05	<i>F</i> =9.01, <i>P</i> <.05
Source of Variation	Alloprevotella	UC_Porphyro- monadaceae	Ruminococcus				
Interaction	<i>F</i> =1.95, <i>P</i> =0.18	<i>F</i> =13.11, <i>P</i> <.005	<i>F</i> =3.36, <i>P</i> =0.09				
Treatment	<i>F</i> =0.49, <i>P</i> =0.50	<i>F</i> =13.11, <i>P</i> <.005	<i>F</i> =2.84, <i>P</i> =0.11				
Diet	<i>F</i> =21.40, <i>P</i> <.005	<i>F</i> =13.11, <i>P</i> <.005	<i>F</i> =7.60, <i>P</i> <.05				

Supplementary Table S2. F- and p-values of Figures 4A and 4B are from two-way ANOVA after Bonferroni's post hoc test.

Supplementary Table S3. Formula of normal chow diet (NCD).

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Nutrients			Minerals			
Protein	20	g%	Ash	7.25	g%	
Arginine	Arginine 1.26 g%		Calcium	1.2	g%	
Cystine	0.37	g%	Phosphorus	0.62	g%	
Glycine	0.87	g%	Phosphorus (non-phytate)	0.4	g%	
Histidine	0.5	g%	Potassium	0.82	g%	
Isoleucine	0.82	g%	Magnesium	0.16	g%	
Leucine	1.47	g%	Sulfur	0.22	g%	
Lysine	1.01	g%	Sodium	0.34	g%	
Methionine	0.33	g%	Chlorine	0.47	g%	
Phenylalanine	0.98	g%	Fluorine	21.38	ppm	
Tyrosine	0.63	g%	Iron	112.93	ppm	
Threonine	0.72	g%	Zinc	128.85	ppm	
Tryptophan	0.25	g%	Manganese	95.49	ppm	
Valine	0.91 g% Copper		Copper	22.74	ppm	
			Cobalt	0.76	ppm	
Fat (ether extract)	4.5	g%	Iodine	1.42	ppm	
Linoleic Acid	1.1	g%	Chromium	0	ppm	
Linolenic Acid	0.12	g%	Selenium	0.32	ppm	
Arachidonic Acid	0.02	g%	Vitamins			
Omega-3 Fatty Acids	1.11	g%	Vitamin K	6.69	ppm	
			Thiamin Hydrochloride	11.02	ppm	
Fiber (Crude)	3.7	g%	Riboflavin	11.57	ppm	
			Niacin	217.7	ppm	
			Pantothenic Acid	88.72	ppm	
			Choline Chloride	3447.9 6	ppm	
			Folic Acid	13.6	ppm	
			Pyridoxine	11	ppm	
Calories provided by :			Biotin	0.15	ppm	
Protein 24.52 %		B12	41	ppm		
Fat (ether extract)	12.41	%	Vitamin A	28.03	IU/kg	
Carbohydrates 63.07 %		%	Vitamin D3 (added)	4	IU/kg	
Total 3,940 k		kcal/kg	Vitamin E	100	IU/kg	

Purina Lab. Rodent Chow (38057)

Supplementary Table S4. Formula of high-fat diet (HFD).

Research Diet #D12492	g%	kcal%
Protein	26.2	20
Carbohydrate	26.3	20
Fat	34.9	60
Total		100
kcal/g	5.24	
Ingredient	g	kcal
Casein, 30 Mesh	200	800
L-Cystine	3	12
Corn Starch	0	0
Maltodextrin 10	125	500
	68.8	275.2
Cellulose, BW200	50	0
Soybean Oli	20	225
Lalu Minoral Mix S10026**	240	2205
DiCalcium Phosphate	10	0
Calcium Carbonate	55	0
Potassium Citrate 1 H2O	16.5	0
Vitamin Mix V10001***	10.5	40
Choline Bitartrate	2	-0 0
ED&C Blue Dye #1	0.05	0
Total	773.85	4057
*Typical analysis of cholesterol in lard	= 0.72 mg/g.	
Cholesterol (mg)/4057 kcal = 216.4	00	
Cholesterol (mg)/kg = 279.6		
**Mineral Mix S10026		g
Sodium Chloride (39.3% Na 60.7% C	l)	2.59
Magnesium Oxide, Heavy, DC USP (6	60.3% Mg)	0.419
Magnesium Sulfate, Heptanydrate (9.	87% Mg 13.0% S)	2.576
Chromium Potassium Sulfate (10.4%)	Cr)	0.0005
Copper Carbonate (57.5% Cu)	,	0.0105
Ferric Citrate (17.4% Fe)	•• •	0.21
Manganese Carbonate Hydrate (47.8	Mn)	0.1225
Sodium Eluoride (45.2%% El)		0.00035
Sodium Selenite (45.7% Se)		0.00035
Zinc Carbonate (52.1% Zn)		0.056
Sucrose		3.99105
***Vitamin Mix V10001		a
Vitamin A Palmitate (500,000 IU/gm)		0.008
Vitamin D3 (100,000 IU/gm)		0.01
Vitamin E Acetate (500 IU/gm)	(anadiana)	0.1
Biotin 1.0%	lenadione)	0.0008
Cyanocobalamin, 0.1%		0.02
Folic Acid		0.002
Nicotinic Acid		0.03
Calcium Pantothenate		0.016
гунцохше-пон Riboflavin		0.007
Thiamin HCI		0.006
Sucrose		9.78