

1 **Supplementary tables**

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3 **Intestinal inflammation induced by dextran sodium sulphate causes liver inflammation and lipid**
4 **metabolism dysfunction in laying hens**

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20 **Supplementary Table S1** PCR primers used for mRNA expression analysis.

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Target genes	Forward Primer	Revers Primer	Product size	Accession no.
<i>IL-1β</i>	GTTGGGCTCAACATTGCGCTGTA	TGTCCAGGGGTAAGAGATGAAG	214	NM_204524.1
<i>IL-6</i>	AGAAATCCCTCCTGCCAAT	AAATAGCGAACGGCCCTCA	121	NM_204628.1
<i>CXCLi2</i>	CTGTCCTGGCCCTCCTCGTT	TGGCGTCAGCTTACATCTTG	146	NM_205498.1
<i>IL-10</i>	GCTGAGGTGAAGTTGAGGAA	GAAGCCAGCATCTCTGACA	142	NM_001004414.2
<i>TGFβ-2</i>	AGGAATGTGCAGGATAATT	ATTTGGGTGTTGCCAA	269	NM_001031045.3
<i>TGFβ-3</i>	CAGATCCTGGCCTCTACA	GAGGCCCTGGATCATGTCA	141	NM_205454.1
<i>TGFβ-4</i>	ATGAGTATTGGGCCAAAG	ACGTTGAAACACGAAGAAG	109	NM_001318456.1
<i>ACC</i>	CTCAGAATTGCCCTTGTC	CATCTCCTCGTGTCTCC	145	NM_205505.1
<i>FAS</i>	TGAAGGCAATGGGAAACAG	GACATAACAGGATGAGGTTGATT	112	NM_205155.1
<i>SCD-1</i>	ATCTGCACCTCCTTGGTCG	GGGATGCCCTCATGCCATTA	187	DQ645535.1
<i>LDLr</i>	GGAGCAGTCACAGCATCAGCT	CTGTGTCACACTCCGCTGTCTC	109	NM_204452.1
<i>SREBP-1</i>	CTACCGCTCATCCATCAACG	CTGCTTCAGCTCTGGTTGC	145	NM_204126.2
<i>SREBP-2</i>	TCCTTGATGATGCAAAGGTTAAGG	AGGGATGTCAAAGGTTGAAGG	105	XM_015289037.2
<i>ApoB</i>	CTGCAATGCTGGCTGTT	CTGGTTGAGCCATCCAGCTT	106	NM_001044633.1
<i>ApoVLDL-II</i>	AGGGCTGAACGGTACCAACAAAC	GGATGACCAAGCCAGTCACGA	140	NM_205483.2
<i>VTG-II</i>	CAACATATCTCCGTTGAAATTG	TTCACAAACAAAGATTCTCCAGTAGC	147	NM_001031276
<i>ERα</i>	CTTCATCCATCACACAG	TAGACATCCTCTCACGAAT	143	NM_205183.1
<i>RPS17</i>	AAGCTGAGGAGGAGGAGGAGG	GGITGGACAGGCTGCCAAGT	136	NM_204217.1
<i>GLUT-2</i>	CACACTATGGCGCATGCT	ATTGTCCTGGAGGTGTTGGTG	68	NM_207178.1
<i>GLUT-5</i>	TTGCTGGCTTGGGTTGIG	GGAGGTGAGGGCCAAAGTC	60	XM_025142667.1
<i>SGLT-1</i>	GCCGTGGCAGGGCTTA	CAATAACCTGATCTGTGACCCAGTA	71	NM_001293240.1

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24 F, forward; R, reverse, *PCR products were sequenced for verification.

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Supplementary Table S2 Effects of paired feeding on the egg production, egg weight and egg shell weight.

	group	Day 0	Day 1	Day 2	Day 3	Day 4	Day 5
Number of egg laying	Con	4/4	3/4	3/4	3/4	3/4	3/4
	DSS	4/4	4/4	4/4	4/4	4/4	3/4
Egg weight (%)	Con	100.0	97.4 ± 2.4	94.3 ± 1.9	101.6 ± 3.2	104.5 ± 1.6 ^a	97.6 ± 1.8 ^a
	DSS	100.0	100.2 ± 0.8	100.5 ± 1.4	98.1 ± 1.8	96.3 ± 1.4 ^b	90.2 ± 1.5 ^b
Egg-shell weight (%)	Con	100.0	92.4 ± 6.5	81.0 ± 5.0	85.3 ± 8.7	105.0 ± 4.6 ^a	98.4 ± 5.9 ^a
	DSS	100.0	97.7 ± 3.2	88.6 ± 1.2	77.6 ± 1.5	76.2 ± 1.5 ^b	71.4 ± 0.2 ^b

Number of egg laying means produced egg number/total hen number. Values are mean ± SEM of the ratio of egg weight and egg-shell weight against day 0 (%).

Lower case letters (a, b) indicate significant differences between the control and PF groups ($P < 0.05$).

Supplementary Figures

Intestinal inflammation induced by dextran sodium sulphate causes liver inflammation and lipid metabolism dysfunction in laying hens

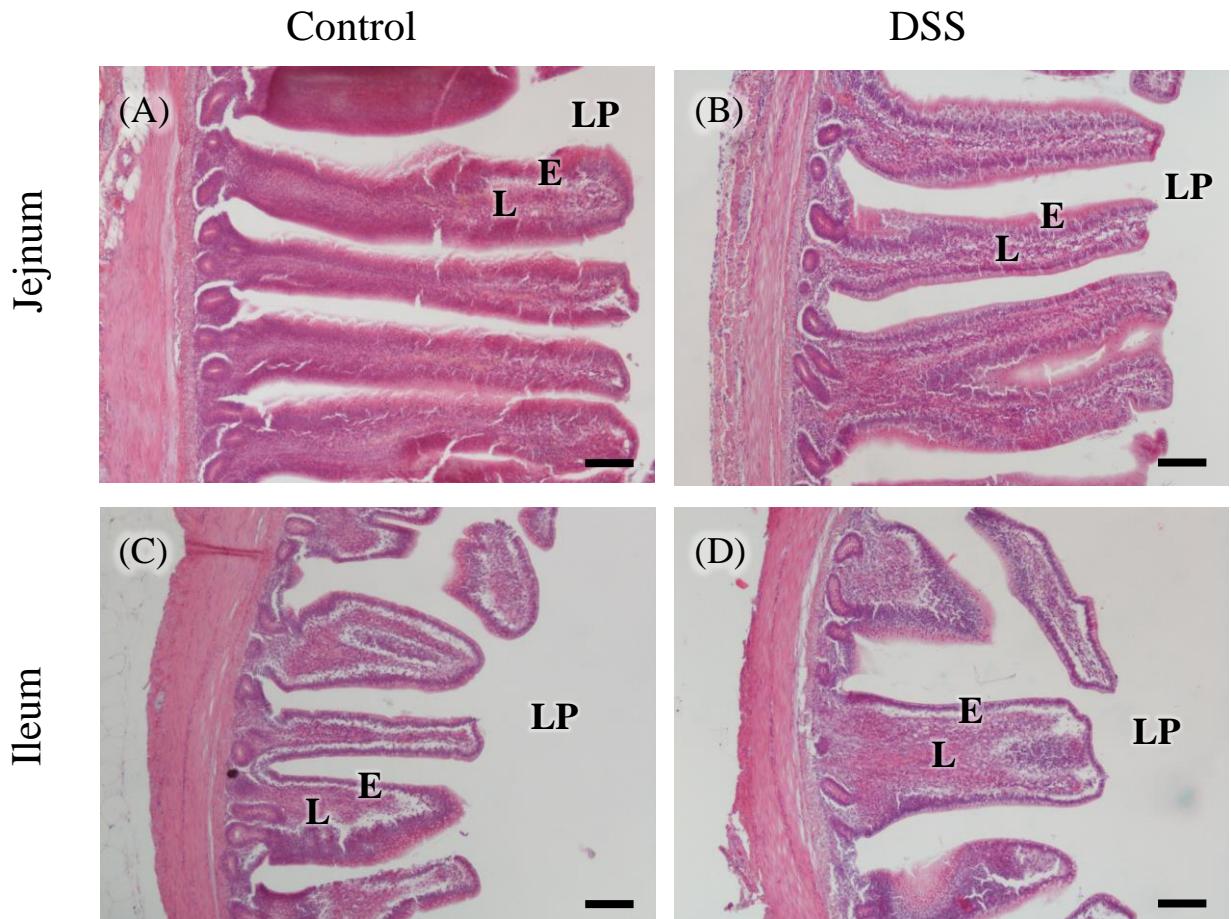
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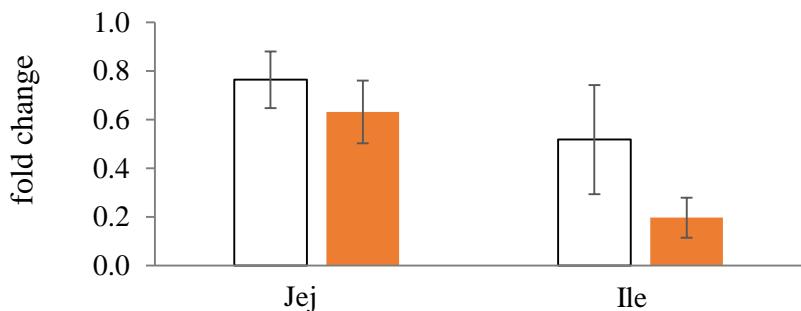
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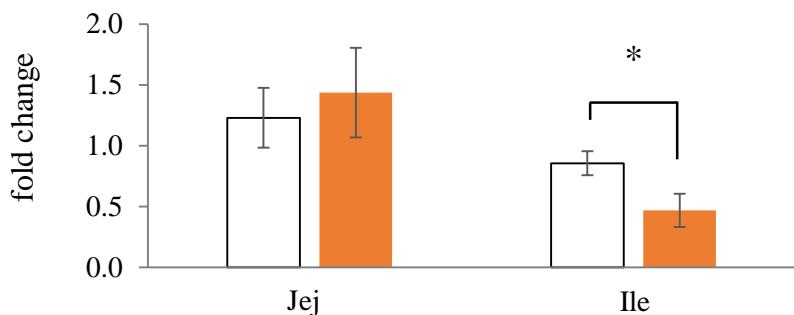


Supplementary Figure S1 Micrographs showing the histology of the jejunum and ileum of hens orally administered DSS (DSS group) or water (control group). HE-stained tissue from the (A) jejunum of the control group, (B) jejunum of the DSS group, (C) ileum of the control group, and (D) ileum of the DSS group. The lamina propria is covered with simple columnar epithelium. E = mucosal epithelium, L = lumen, LP = lamina propria. Scale bars = 100 μ m.

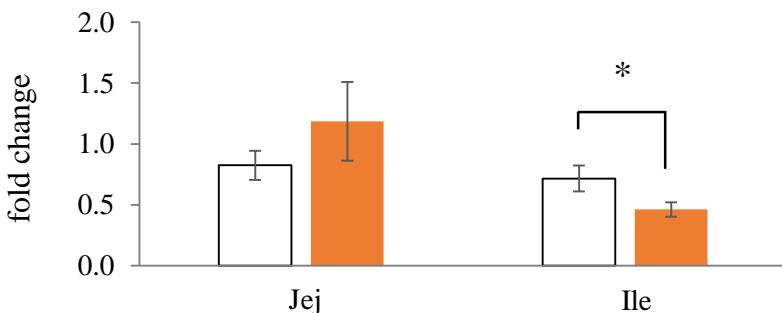
(A) *GLUT2*



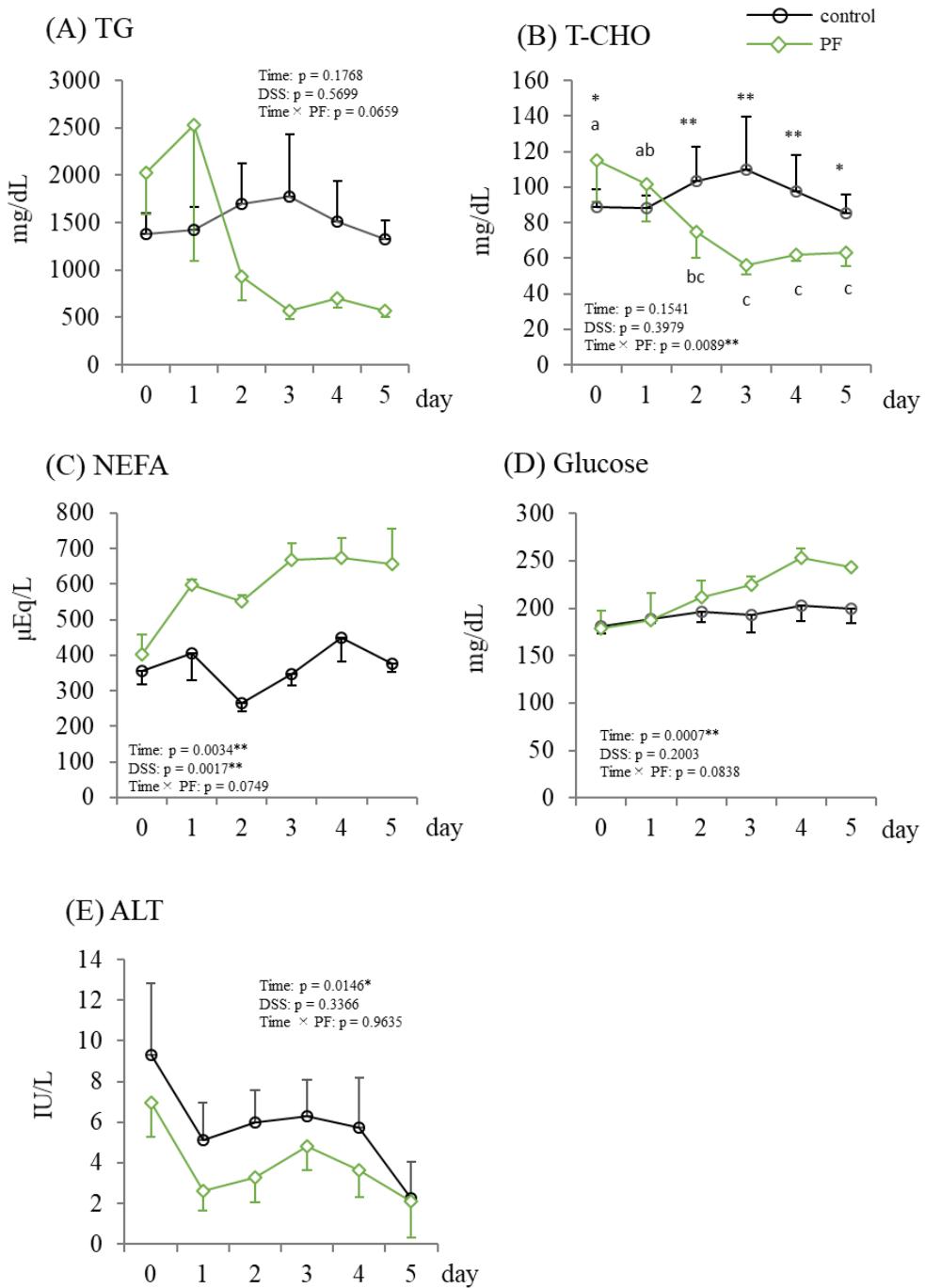
(B) *GLUT5*



(C) *SGLT1*

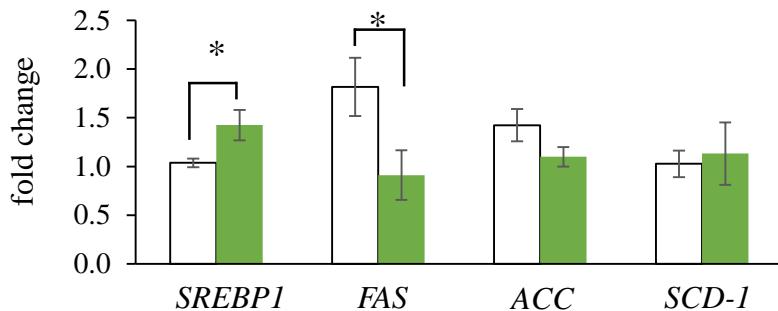


Supplementary Figure S2 Effects of oral administration of DSS on the mRNA expression of the glucose transporters (A) *GLUT2*, (B) *GLUT5*, and (C) *SGLT1* in the jejunum and ileum. Open bars are the control group, and orange filled bars are the DSS group. Values are mean \pm SEM ($n = 8$) of fold change in the target gene expression from a standard sample of each segment in the control group. Target gene expression was normalized to the house-keeping gene RPS17. An asterisk (*) indicates a significant difference between the control and DSS groups ($P < 0.05$).

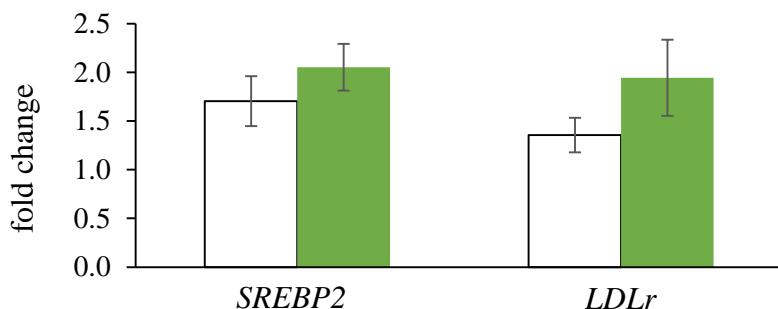


Supplementary Figure S3 Changes in the blood serum parameters TG (A), T-CHO (B), NEFA (C), glucose (D), and ALT (E) of hens during the paired feeding experiment in the restricted feed (PF) and ad libitum feed (control) groups. Open circle plots are the control group, and green open rhombus plots are the PF group. Values are the mean \pm SEM ($n = 4$). Significant differences between the control and PF groups at the same time point are indicated with asterisks (*, ** indicate $P < 0.05$ and $P < 0.01$, respectively). Significant differences at different time points are shown as different letters (a–c; $P < 0.05$).

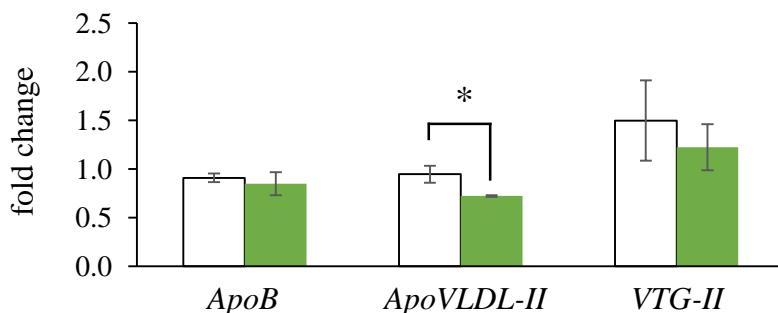
(A) Lipid synthesis-related genes



(B) Lipoprotein uptake-related genes



(C) Yolk precursor synthesis-related gene



Supplementary Figure S4 Effects of paired feeding (PF) on the mRNA expression of lipid synthesis-related genes (A; *SREPB-1*, *FAS*, *ACC*, and *SCD-1*), lipoprotein uptake-related genes (B; *SREPB-2* and *LDLr*), and yolk precursor-related genes (C; *ApoB*, *ApoVLDL-II*, and *VTG-II*) in the liver of hens. Open bars are the control group, and green filled bars are the PF group. Values are mean \pm SEM ($n = 8$) of fold change in the target gene expression from a standard sample of the control group. Target gene expression was normalized to the house-keeping gene *RPS17*. An asterisk (*) indicates a significant difference between the *ad libitum* feed (control) and PF groups ($P < 0.05$).