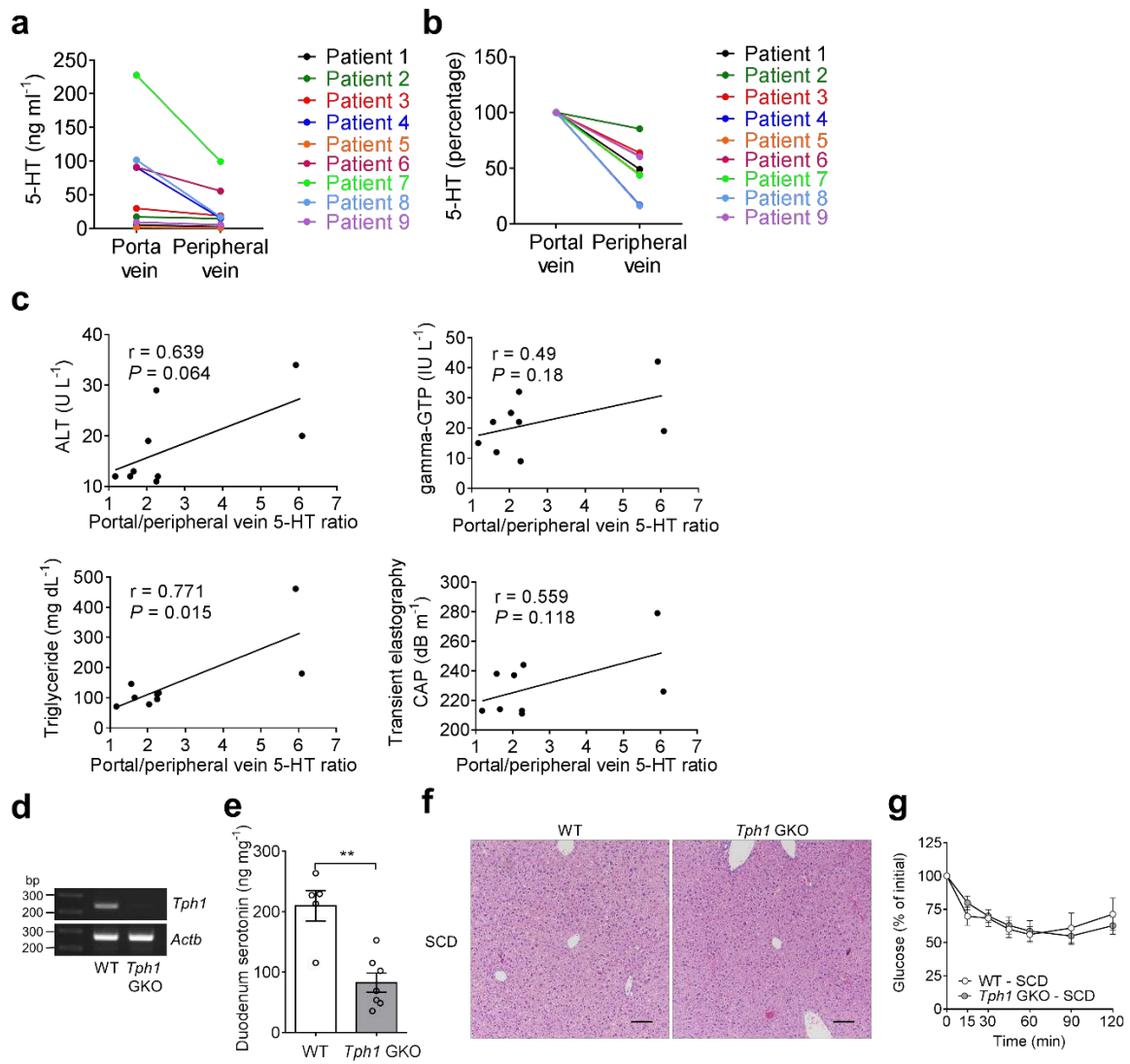


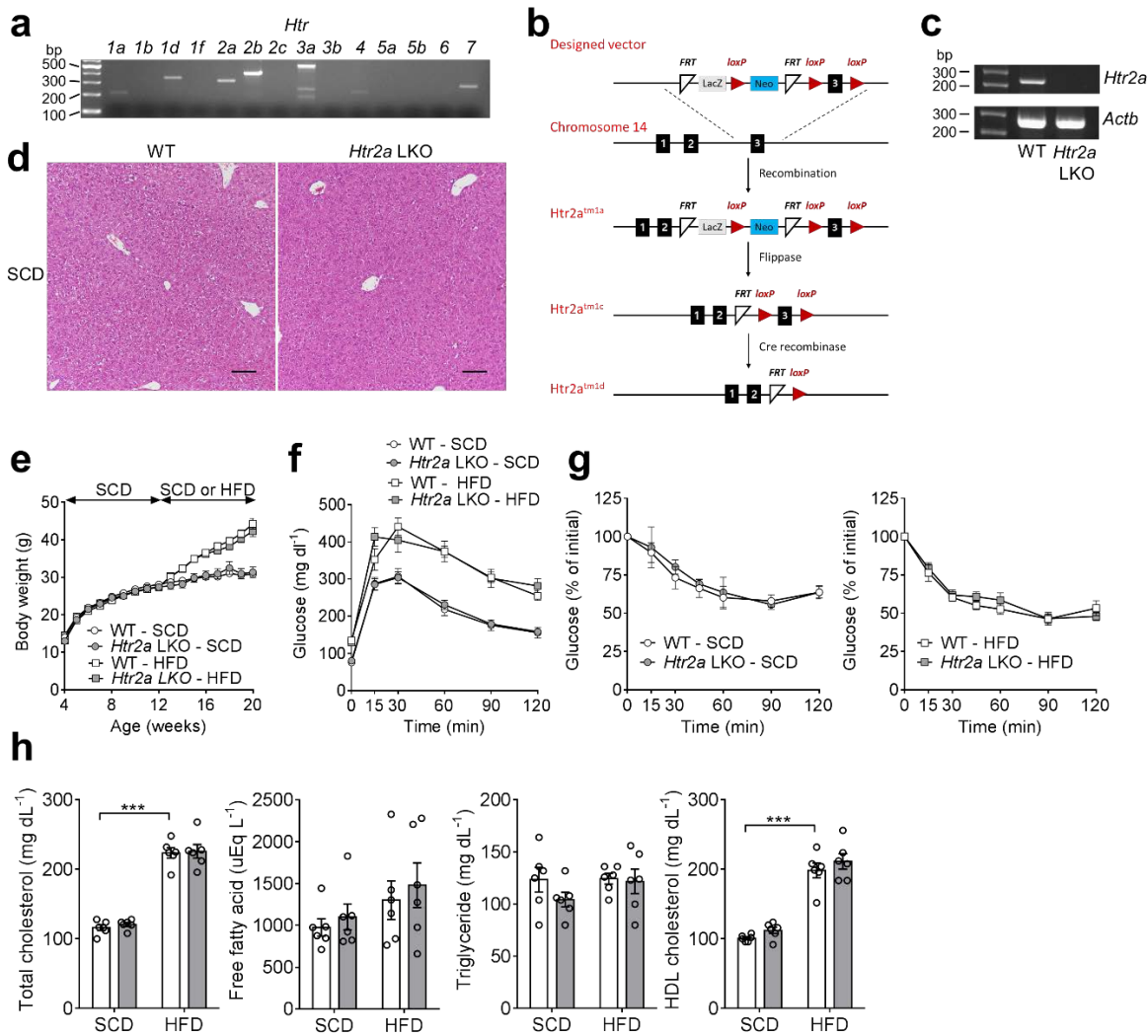
## **Supplementary Information**

**Serotonin signals through a gut-liver axis to regulate hepatic steatosis**

**Choi et al.**

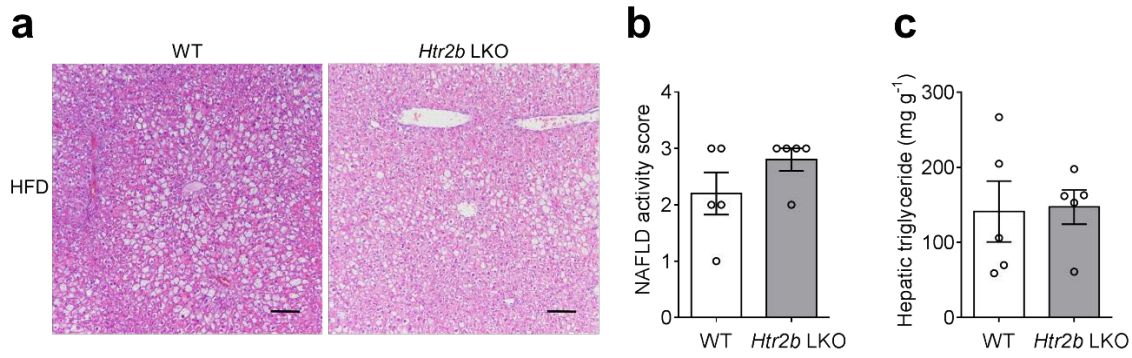


**Supplementary Figure 1. Inhibiting gut-derived 5-HT does not affect hepatic lipid metabolism in SCD-fed mice.** (a) Plasma 5-HT concentrations of portal blood and peripheral blood in humans. n = 9 per group. (b) Plasma 5-HT concentrations of portal blood and peripheral blood in humans. Portal blood concentration set as 100%. n = 9 per group. (c) Scatter plots of values for portal/peripheral blood 5-HT ratio versus ALT, gamma-GTP, TG and transient elastography CAP in humans. n = 9 per group. These data were compared with one another by calculation of the Pearson's *r* correlation coefficient. (d) Representative RT-PCR data of *Tph1* gene expression in duodenum from 20 week old WT and *Tph1* GKO mice. (e) Duodenal 5-HT levels in WT and *Tph1* GKO mice. n = 5-7 per group. (f) Representative liver histology by H&E staining from SCD fed WT and *Tph1* GKO mice. (g) Intraperitoneal insulin tolerance test (IPITT) after 4 h fasting of SCD fed WT and *Tph1* GKO mice. n = 5 per group. Scale bars, 100µm. Data are expressed as the means ± standard error of the mean (SEM). \**P*<0.5, \*\**P*<0.01, \*\*\**P*<0.001, Student's *t*-test (e, g).

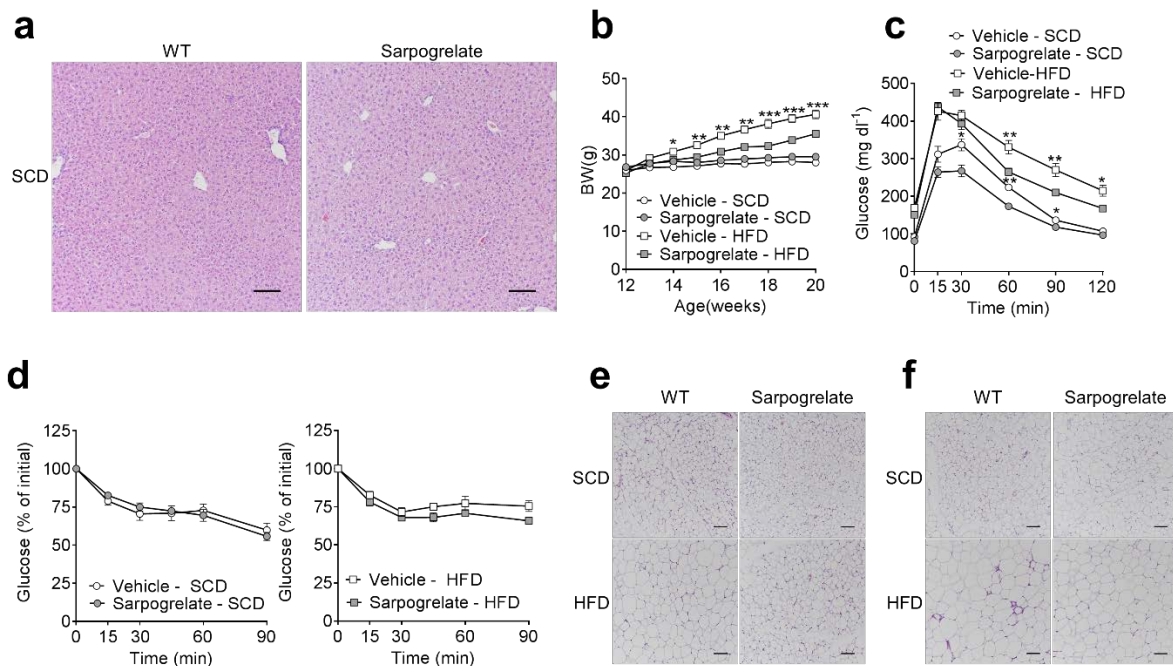


**Supplementary Figure 2. Liver-specific deletion of *Htr2a* does not affect systemic energy metabolism.**

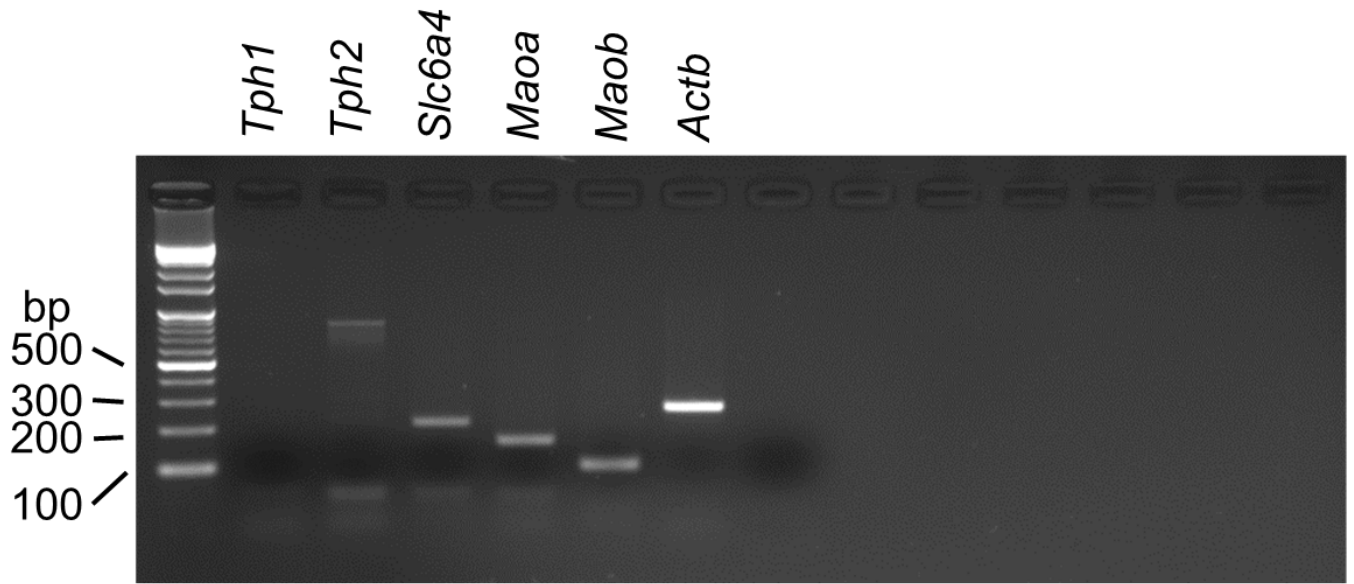
(a) mRNA expression of *Htrs* as assessed by RT-PCR in liver from 8 week old C57BL/6J mice. (b) Schematic of gene targeting strategy to generate the *Htr2a* floxed allele. (c) Representative RT-PCR data of *Htr2a* gene expression in liver from 20 week old WT and *Htr2a* LKO mice. (d-h) 12 week old WT and *Htr2a* LKO mice were fed SCD or HFD for 8 weeks. (d) Representative liver histology by H&E staining from SCD fed WT and *Htr2a* LKO mice. Scale bars, 100 $\mu$ m. (e) Body weight trend. n = 6-10 per group. (f) Intraperitoneal glucose tolerance test (IPGTT) after 16 h fasting. n = 6 per group. (g) Intraperitoneal insulin tolerance test (IPITT) after 4 h fasting. n = 6 per group. (h) Plasma total cholesterol, free fatty acid, triglyceride, and HDL cholesterol levels. n = 6 per group. Data are expressed as the means  $\pm$  standard error of the mean (SEM). \**P*<0.5, \*\**P*<0.01, \*\*\**P*<0.001, Student's *t*-test (g) or one-way ANOVA with *posthoc* Tukey's test (e,f,h).



**Supplementary Figure 3. HTR2B does not mediate the lipogenic action of GDS in the liver.** (a-c) 12 week old mice were fed HFD for 8 weeks. (a) Representative liver histology by H&E staining from HFD fed WT and *Htr2b* LKO mice. Scale bars, 100 $\mu$ m. (b) NAFLD activity scores of HFD fed WT and *Htr2b* LKO mice. n = 5 per group. (c) Hepatic triglyceride levels. n = 5 per group.



**Supplementary Figure 4. HTR2A antagonist ameliorates HFD-induced obesity.** (a-f) 12 week old mice were fed SCD or HFD for 8 weeks and were treated with vehicle or sarpogrelate daily per os. (a) Representative liver histology by H&E staining from SCD fed vehicle and sarpogrelate treated mice. Scale bars, 100 $\mu$ m. (b) Body weight trend. n = 5-6 per group. (c) Intraperitoneal glucose tolerance test (IPGTT) after 16 h fasting. n = 5-6 per group. (d) Intraperitoneal insulin tolerance test (IPITT) after 4 h fasting. n = 5-6 per group. (e, f) Representative iWAT (e) and eWAT (f) histology by H&E staining. Scale bars, 100 $\mu$ m. Data are expressed as the means  $\pm$  standard error of the mean (SEM). \* $P$ <0.5, \*\* $P$ <0.01, \*\*\* $P$ <0.001, Student's  $t$ -test (d) or one-way ANOVA with *posthoc* Tukey's test (b,c).



**Supplementary Figure 5. Uncropped gel photo shown in Figure 1.**

**Supplementary Table 1. List of primers used in this study.**

Gene name	Forward primer (5' to 3')	Reverse primer (5' to 3')
<i>Acaca</i>	CAGTAACCTGGTGAAGCTGGA	GCCAGACATGCTGGATCTCAT
<i>Acat1</i>	CCCCATTGATTTTCCACTTG	AGCACAACCACACTGAATGC
<i>Acox1</i>	CACGGCTATTCTCACAGCAG	CAGGCTGTTAATGTCCACCA
<i>Actb</i>	GGTACCACCATGTACCCAGG	GAAAGGGTGTAACGCAGC
<i>Agpat1</i>	GCGCAATGTGAGAACATGA	TCATTCCAAGCAGGTCGAGG
<i>Apob</i>	TACTTCCACCCACAGTCCCCT	CCTTAGAAGCCTTGGGCACAT
<i>Cd36</i>	TGGCCAAGCTATTGCGACAT	ACACAGCGTAGATAGACCTGC
<i>Coll1a1</i>	ACATGTTTCAGCTTTGTGGACC	TAGGCCATTGTGTATGCAGC
<i>Col3a1</i>	GGAACCTGGTTTCTTCTCACC	TAGGACTGACCAAGGTGGCT
<i>Col4a1</i>	GTCTGGCTTCTGCTGCTCTT	CACATTTTCCACAGCCAGAG
<i>Cpt1a</i>	AGCTCGCACATTACAAGGACA	CCAGCACAAAGTTGCAGGAC
<i>Fabp1</i>	TGAAGGCAATAGGTCTGCCC	GTCATGGTCTCCAGTTCGCA
<i>Fabp2</i>	CTTCGGGAACCACAGGTCTTC	CATAGCAAGGCTGTCCCATAC
<i>Fasn</i>	AAGCGGTCTGGAAAGCTGAA	AGGCTGGGTTGATACCTCCA
<i>Fatp5</i>	TTCGAAAGAACCAACCCTTCTT	GCGTCGTACATTCGCAACAA
<i>F4/80</i>	TCAAGGACACGAGGT TGCTGA	CCAAGGGGCCAATCTGGAA
<i>Gpat1</i>	CCACAGAGCTGGGAAAGGTT	GTGCCTTGTGTGCGTTTCAT
<i>Htr1a</i>	TCAGCTACCAAGTGATCACCTCT	GTCCACTTGTTGAGCACCTG
<i>Htr1b</i>	TGCTCCTCATCGCCCTCTATG	CTAGCGGCCATGAGTTTCTTCTT
<i>Htr1d</i>	CCTCCAACAGATCCCTGAATG	CAGAGCAATGACACAGAGATGCA
<i>Htr1f</i>	TGTGAGAGAGAGCTGGATTATGG	TAGTTCCTTGGTGCCTCCAGAA
<i>Htr2a</i>	AGCTGCAGAATGCCACCAACTAT	GGGATTGGCATGGATATACCTAC
<i>Htr2b</i>	AAATAAGCCACCTCAACGCCT	TCCCGAAATGTCTTATTGAAGAG
<i>Htr2c</i>	TTCTTAATGTCCCTAGCCATTGC	GCAATCTTCATGATGGCCTTAGT
<i>Htr3a</i>	AAATCAGGGCGAGTGGGAGCTG	GACACGATGATGAGGAAGACTG
<i>Htr3b</i>	CGTGTGGTACCGAGAGGTTT	GGATGGGCTTGTGGTTTCTA
<i>Htr4</i>	ATGGACAACTTGATGCTAATGTGA	TCACCAGCACCGAAACCAGCA

<i>Htr5a</i>	GATTGACTTCAGTGGGCTCG	AAAGTCAGGACTAGCACTCG
<i>Htr5b</i>	GGAGCCTTCTACCTGCCTCT	ATGAGCTCCGTCAGGAAGAA
<i>Htr6</i>	CCTCACATGGCTGGGATACT	ATCTGAGTTGGGTGGCAGAG
<i>Htr7</i>	CTCGGTGTGCTTTGTCAAGA	TTGGCCATACATTTCCCATT
<i>Maoa</i>	GCGGTACAAGGGTCTGTTC	CAGCCAATCCTGAGATGCCG
<i>Maob</i>	GGGCGGCATCTCAGGTATGG	AAGTCCTGCCTCCTACACGG
<i>Mcad</i>	GGTTTGGCTTTTGGACAATG	TGACGTGTCCAATCTACCACA
<i>Mcp1</i>	GGCTCAGCCAGATGCAGTTAA	AGCCTACTCATTGGGATCATCTT
<i>Mogat1</i>	TTGACCCATGGTGCCAGTTT	GTGGCAAGGCTACTCCCATT
<i>Mtp</i>	TCTCACAGTACCCGTTCTT	TCTTCTCCGAGAGACATATCC
<i>Ppara</i>	AGAAGTTGCAGGAGGGGATT	TCGGACTCGGTCTTCTTGAT
<i>Pparg</i>	GGTGTGATCTTAACTGCCGGA	GCCCAAACCTGATGGCATTG
<i>Slc6a4</i>	CGCAGTTCCCAGTACAAGC	GAAGGCCCTCCACCATTCTG
<i>Srebp1c</i>	GGAGCCATGGATTGCACATT	GGCCCGGGAAGTCACTGT
<i>Tgfb1</i>	GGGCTACCATGCCAACTTCTG	GAGGGCAAGGACCTTGCTGTA
<i>Timp1</i>	CCCTGCTCAGCAAAGAGC	TCACTCTCCAGTTTGCAAGG
<i>Tph1</i>	ACCATGATTGAAGACAACAAGGAG	TCAACTGTTCTCGGCTGATG
<i>Tph2</i>	GCCATGCAGCCCGCAATGATGATG	CAACTGCTGTCTTGCTGCTC
<i>Ucp1</i>	CTTTGCCTCACTCAGGATTGG	ACTGCCACACCTCCAGTCATT



**Supplementary Table 2. Clinical characteristics of study participants.**

<b>Patient</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>	<b>7</b>	<b>8</b>	<b>9</b>
<b>Age</b>	21	24	37	48	58	37	42	36	53
<b>Sex</b>	Male	Male	Male	Male	Male	Male	Female	Male	Female
<b>Height (m)</b>	1.76	1.74	1.78	1.7	1.62	1.85	1.62	1.68	1.61
<b>Weight (kg)</b>	75	65	75	74	53	81	63	68	53.5
<b>BMI(kg m<sup>-2</sup>)</b>	24.21	21.47	23.67	25.61	20.2	23.67	24.01	24.1	20.64
<b>Hb (g dL<sup>-1</sup>)</b>	16.9	15.5	15.2	15.4	13.3	15.4	10.3	14.1	13.9
<b>WBC (10<sup>3</sup> uL<sup>-1</sup>)</b>	5	5.02	7.37	8.46	6.57	5.72	7.47	5.7	5.69
<b>Seg (%)</b>	46.3	64.9	62.4	55.9	59.6	49.7	74.4	57.1	37.3
<b>Lymph (%)</b>	44.1	27.1	27.4	32.4	30.5	37.6	18.5	33.9	45
<b>PLT (10<sup>3</sup> uL<sup>-1</sup>)</b>	200	255	219	270	239	281	327	276	253
<b>Ca (mg dL<sup>-1</sup>)</b>	9.6	9.7	8.9	9.3	9.5	9.8	9.3	9.4	9.7
<b>P (mg dL<sup>-1</sup>)</b>	3.8	3.6	3.3	2.9	4	3.5	4.4	3.7	3.7
<b>Glucose (mg dL<sup>-1</sup>)</b>	94	95	98	98	117	101	103	105	92
<b>BUN (mg dL<sup>-1</sup>)</b>	16.5	7.6	8.1	12.6	10.2	12.8	12.2	14.1	12.9
<b>Cr (mg dL<sup>-1</sup>)</b>	0.79	0.86	0.83	0.86	0.65	0.94	0.9	0.69	0.67
<b>Uric acid (mg dL<sup>-1</sup>)</b>	6	4.8	5.9	5.4	5	7.2	5.5	5.1	4.7
<b>Cholesterol (mg dL<sup>-1</sup>)</b>	133	154	139	212	211	281	209	180	219
<b>Total protein (g dL<sup>-1</sup>)</b>	7.2	7.3	6.9	6.8	6.4	7.7	7.3	7.4	7.1
<b>Albumin (g dL<sup>-1</sup>)</b>	4.7	4.9	4.5	4.4	4.5	4.6	4.6	4.7	4.6
<b>ALP (IU L<sup>-1</sup>)</b>	79	72	70	81	64	51	36	69	64
<b>AST (IU L<sup>-1</sup>)</b>	16	14	13	17	21	29	14	26	17
<b>ALT (IU L<sup>-1</sup>)</b>	19	12	12	34	11	29	12	20	13
<b>Bilirubin (mg dL<sup>-1</sup>)</b>	1.2	1	0.6	0.9	0.6	0.8	0.9	1.3	0.8
<b>gamma-GTP (IU L<sup>-1</sup>)</b>	25	15	22	42	22	32	9	19	12
<b>TG (mg dL<sup>-1</sup>)</b>	78	71	146	461	95	112	116	180	100
<b>HDL cholesterol (mg dL<sup>-1</sup>)</b>	44	62	36	37	43	68	51	50	75
<b>LDL cholesterol (mg dL<sup>-1</sup>)</b>	78	81	85	89	142	194	137	104	118
<b>Na (mmol L<sup>-1</sup>)</b>	142	142	143	140	142	141	140	143	142
<b>K (mmol L<sup>-1</sup>)</b>	4.4	4	4.9	4.6	4.5	4.7	4.2	4.5	4.7
<b>Cl (mmol L<sup>-1</sup>)</b>	102	100	105	102	102	100	101	103	101

<b>PT (Sec)</b>	11.5	12	11.2	10.7	10.6	10.5	11.6	10.5	10.4
<b>PT (%)</b>	98	92	100	100	100	100	91	100	100
<b>PTT (sec)</b>	37.3	32.7	33.4	29.3	29.8	29.6	32.7	31.2	31.2
<b>Portal vein serotonin (ng ml<sup>-1</sup>)</b>	5.3	17.1	29.6	90.6	1.8	91	227.6	101.6	9.1
<b>Systemic vein serotonin (ng ml<sup>-1</sup>)</b>	2.6	14.6	18.9	15.3	0.8	55.5	99.3	16.7	5.5
<b>Portal vein/systemic vein serotonin</b>	2.04	1.17	1.57	5.92	2.25	1.64	2.29	6.08	1.65
<b>Fibroscan LSM (kPa)</b>	5.3	4.9	3.1	4.6	4.9	3.6	4.6	4.5	8.5
<b>Fibroscan CAP (dB m<sup>-1</sup>)</b>	237	213	238	279	213	211	244	226	214

BMI, body mass index; Hb, hemoglobin; Seg, segmented neutrophil; Lymph, lymphocyte; PLT, platelet; BUN, blood urea nitrogen; ALP, alkaline phosphatase; AST, aspartate transferase; ALT, alanine transferase; TG, triacylglycerol; HDL, high-density lipoprotein; LDL, low-density lipoprotein; PT, prothrombin time; PTT, partial thromboplastin time; LSM, liver stiffness measure; CAP, controlled attenuation parameter.