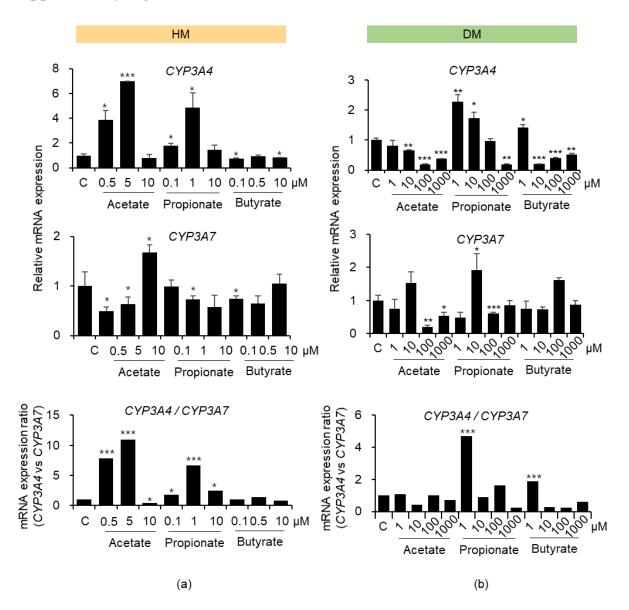
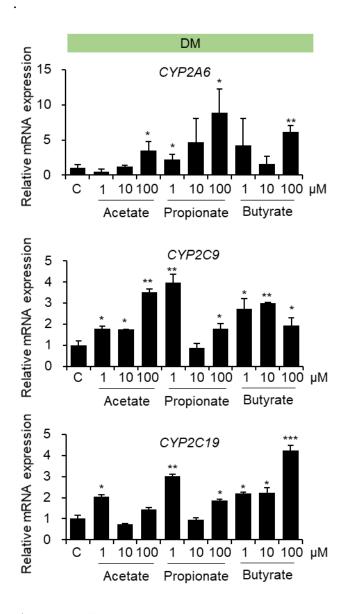
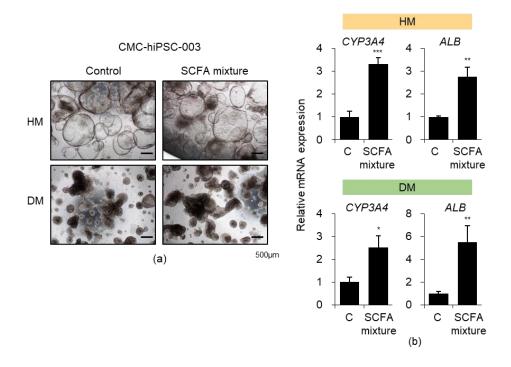
## **Supplementary Figures**



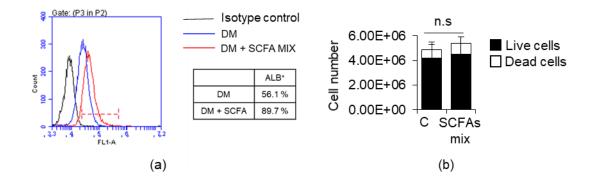
**Figure 1.** Effects of acetate, propionate, and butyrate at various concentrations on *CYP3A4* and *CYP3A7* expression in iPSCs-derived liver organoids. (**a**) mRNA expression levels of *CYP3A4* (*top*), *CYP3A7* (*middle*), and *CYP3A4/CYP3A7* ratio (*bottom*) in SCFAs-treated HM-cultured organoids at each indicated concentration. (**b**) mRNA expression levels of *CYP3A4* (*top*), *CYP3A7* (*middle*), and *CYP3A4/CYP3A7* ratio (*bottom*) in SCFAs-treated HM-cultured organoids at each indicated concentration. (**b**) mRNA expression levels of *CYP3A4* (*top*), *CYP3A7* (*middle*), and *CYP3A4/CYP3A7* ratio (*bottom*) in SCFAs-treated DM-cultured organoids at each indicated concentration. Data are presented as the mean  $\pm$  SEM (*n* = 3) and analyzed by Student's *t* test. \**p* < 0.05, \*\**p* < 0.01, and \*\*\**p* < 0.001.



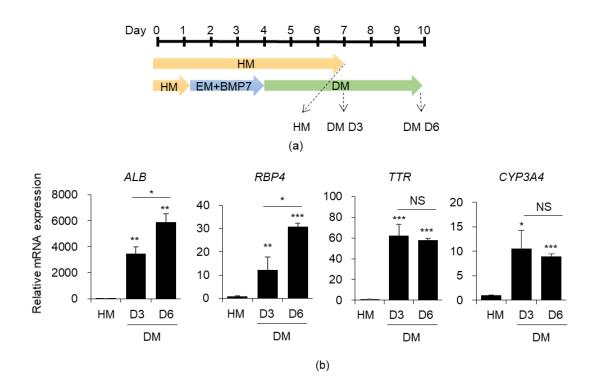
**Figure S2**. Effects of acetate, propionate, and butyrate at various concentrations on *CYPs* expression in iPSC-derived liver organoids. *CYP2A6 (top)*, *CYP2C9 (middle)*, and *CYP2C19 (bottom)* mRNA expression levels in SCFAs-treated DM-cultured organoids at each indicated concentration. Data are presented as the mean  $\pm$  SEM (*n* = 3) and analyzed by Student's *t* test. \**p* < 0.05, \*\**p* < 0.01, and \*\*\**p* < 0.001.



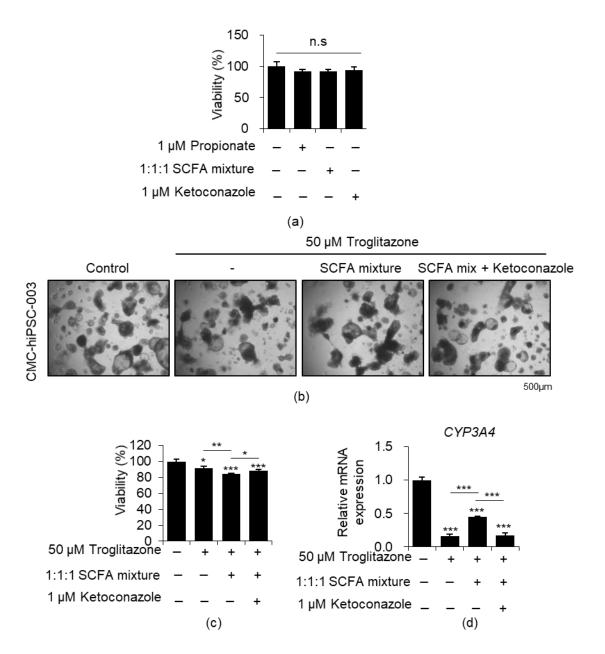
**Figure S3**. Effects of SCFA mixture on CMC-hiPSC-003-derived liver organoids. (**a**) Representative morphology of untreated control and SCFA mixture-treated HM (*upper*)- and DM (*lower*)-cultured organoids. (**b**) *CYP3A4* and *ALB* mRNA expression levels in control and SCFA mixture-treated HM (*upper*)- and DM (*lower*)-cultured organoids. Data are presented as the mean  $\pm$  SEM (*n* = 3) and analyzed by Student's *t* test. \**p* < 0.05, \*\**p* < 0.01, and \*\*\**p* < 0.001.



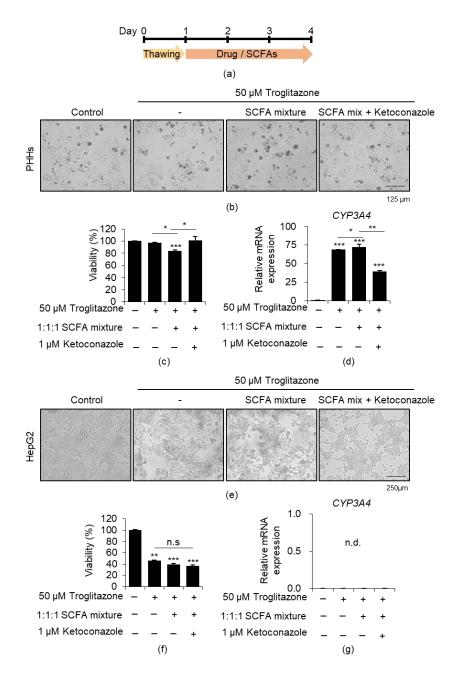
**Figure S4**. Effects of SCFA mixture on ALB expression and cell viability. (**a**) Representative flow cytometry analysis of ALB-stained organoids in DM control and SCFA mixture treated groups. (**b**) Live/Dead cells in DM control and SCFA mixture treated organoids were counted after Trypan Blue staining.



**Figure S5**. Schedule of the optimization of liver organoid differentiation. (**a**) Scheme of liver organoid differentiation. Hepatocyte marker-expression levels were examined at each indicated time point. (**b**) *ALB, RBP4, TTR,* and *CYP3A4* mRNA expression levels in HM-cultured organoids at D7, and DM-cultured organoids at D3 and D6 after differentiation. Data are presented as the mean  $\pm$  SEM (*n* = 3) and analyzed by Student's *t* test. \**p* < 0.05, \*\**p* < 0.01, and \*\*\**p* < 0.001.



**Figure S6**. Effects of SCFAs on cell viability of CRL-2097-derived liver organoids and CYP3A4mediated drug toxicity in CMC-hiPSC-003-derived liver organoids. (**a**) Cell viability was determined by cell counting in each treated DM-cultured organoids. (**b**) Representative morphology of DMcultured CMC-hiPSC-003-derived liver organoids under each indicated treatment condition. (**c**) Relative cell viability under each indicated condition. (**d**) *CYP3A4* mRNA expression levels under each indicated condition. Data are presented as the mean  $\pm$  SEM (*n* = 3) and analyzed by Student's *t* test. \**p* < 0.05, \*\**p* < 0.01, and \*\*\**p* < 0.001.



**Figure S7**. Effects of SCFA mixture on PHHs and HepG2 cells. (**a**) Scheme of drug and SCFAs treatment. (**b**) Representative morphology of PHHs under each indicated treatment condition. (**c**) Relative cell viability under each indicated condition. (**d**) *CYP3A4* mRNA expression levels under each indicated condition. (**e**) Representative morphology of HepG2 cells under each indicated treatment condition. (**f**) Relative cell viability under each indicated condition. (**g**) *CYP3A4* mRNA expression levels under each indicated reatment condition. (**f**) Relative cell viability under each indicated condition. (**g**) *CYP3A4* mRNA expression levels under each indicated reatment condition. (**f**) Relative cell viability under each indicated condition. (**g**) *CYP3A4* mRNA expression levels under each indicated condition. Data are presented as the mean ± SEM (*n* = 3) and analyzed by Student's *t* test. \**p* < 0.05, \*\**p* < 0.01, and \*\*\**p* < 0.001.

Gene	Primer (Forward)	Primer (Reverse)	
AFP	AGCTTGGTGGTGGATGAAAC	CCCTCTTCAGCAAAGCAGAC	
ALB	TTTATGCCCCGGAACTCCTTT	AGTCTCTGTTTGGCAGACGAA	
β-ACTIN	GGACTTCGAGCAAGAGATGG	AGCACTGTGTTGGCGTACAG	
СК19	CGCGGCGTATCCGTGTCCTC	AGCCTGTTCCGTCTCAAACTTGGT	
CYP2A6	CAGCACTTCCTGAATGAG	AGGTGACTGGGAGGACTTGAGGC	
СҮР2С9	CTACAGATAGGTATTAAGGACA	GCTTCATATCCATGCAGCACCAC	
CYP2C19	ACAAGGGCAATCTGACTGGA	AGTGTTTCAGGTGGCTGGTA	
CYP2D6	TGAAGGATGAGGCCGTCTGGGA	CAGTGGGCACCGAGAAGCTGAAG	
	GA	Т	
CYP3A4	CTTCATCCAATGGACTGCATAAAT	TCCCAAGTATAACACTCTACACAG	
		АСАА	
CYP3A7	AAACTTGGCCGTGGAAACCT	CAGCATAGGCTGTTGACAGTC	
HNF4A	GGCCAAGTACATCCCAGCTTT	CAGCACCAGCTCGTCAAGG	
RBP4	GAGTTCTCCGTGGACGAGAC	TCCAGTGGTCATCATTTCCTTTC	
TTR	TGGGAGCCATTTGCCTCTG	AGCCGTGGTGGAATAGGAGTA	
L		I	

**Table S1.** Primer sequences used in this study

Table S2. Antibodies used in this study	,
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Antibodies	Company	Catalog No.	Dilution
anti-ALB	Bethyl Laboratories	A80-129a	1:100
anti-E-cadherin	BD Biosciences	610181	1:200
anti-HNF4A	Cell Signaling Technology	3113s	1:200
Anti-Ki67	Abcam	ab15580	1:200