## Supplemental document

Description	Parameters	Value	References
Henry's constant for oxygen	$K_{H,O_2}$	$1.32\times10^{-3}mol{\cdot}m^{-3}{\cdot}mmHg^{-1}$	[1]
Oxygen partial pressure in atmosphere	$P_{O_2}$	159 mmHg	[1,2]
Oxygen concentration in medium	$C_{\mathrm{in},O_2}$	0.21 mol/m <sup>3</sup>	[3,4]
entering the system			
Oxygen diffusion in medium	$D_{O_2}$	$3 \times 10^{-9} \text{ m}^2/\text{s}$	[3,4,9]
Oxygen permeability in PDMS	$P_{PDMS}$	$3.786 \times 10^{-11} \text{ mol} \cdot \text{m} \cdot \text{m}^{-2} \cdot \text{s}^{-1} \cdot \text{mmHg}^{-1}$	[5,6]
Hepatocyte maximum oxygen	V .	$4.8\times10^{-17}\ mol\cdot cell^{-1}\cdot s^{-1}$	[3,7,8]
consumption rate	• max,O <sub>2</sub>		
Michaelis-Menten constant for	$K_{\mathfrak{m},O_2}$	0.5 mmHg	[3,8]
hepatocyte oxygen consumption			

Table S1. Model parameters for oxygen transport and consumption at 37 °C.

**Table S2.** The primers used for real-time RT-PCR in this study.

Gene Symbol	Primers (forward/reverse; 5' to 3')
CYP1A1	GATGGTCAAGGAGCACTACA/AAAGAGGTCCAAGACGATGT
CYP1A2	TCAATGACATCTTTGGAGCAG/CTCTGTATCTCAGGCTTGGTC
CYP2B6	GGGAGATTGAACAGGTGATTG/GATGATGTACCCTCGGAAGC
CYP2C9	GGATTTGTGTGGGAGAAGC/TGAAGCACAGCTGGTAGAAG
CYP2D6	CGCATCCCTAAGGGAACGACA/CAGGAAGTGTTCGGGGTGGAA
CYP2E1	CCATCAAGGATAGGCAAGAG/TCCAGAGTTGGCACTACGAC
СҮРЗА4	TGTCCTACCATAAGGGCTT/GGCTGTTGACCATCATAAAAG
СҮРЗА5	ATATGGGACCCGTACACATG/CAGAGACCCTGACGATAGGAC
UGT1A1	GAATCAACTGCCTTCACCAA/GACTGTCTGAGGGATTTTGC
UGT2B4	TGTCTACAGCCTCCGCTTCT/GAACTGATCCCACTTCTTCATG
SULT1A1	GAGCCGCACCCACCTGTT/TGAACGACGTGTGCTGAACCAC
SULT2A1	AAAGACGTTAGAACCCGAAGA/TTTCCAGTCCCCAGATACACC
OAT2	GTGATGCTGCTGGCACTGGTT/CTCTTTCACATGGCCTTGGGTC
OCT1	AAGAGGATGTCACCGAAAAGC/GGATGAGCCCCTGATAGAGCA
SLCO1B3	GCCTAACCTTGACCTATGAT/CAGGTAAGTTATTCCATTGTTC
SLCO2B1	GGGAGTCCACGAAGAAGCAG/GACAGGACCACCAGCAGGAA
AHR	GGTTGTGATGCCAAAGGAAGA/TCATTCGGATATGGGACTCG
RXRα	TCGTCCTCTTTAACCCTGACTC/GCTGCTCTGGGTACTTGTGCT
PXR	GGTCCCCAAATCTGCCGTGTA/CCGGGCGTTGCGTTTCATG
CAR	TTGCAGAAGTGCTTAGATGCT/TCAGCTCTTCTTGCTCCTTACT
MRP2	GACAATTCTAATCTAGCCTACTCC/CATCAACTTCCCAGACATCC
BCRP	GTTCTTGGATGAGCCTACA/CTGAGGCCAATAAGGTCA
MDR1	GCTCGTGCCCTTGTTAGAC/GTGCCATGCTCCTTGACTC
BSEP	CCCTCATCCGAAATCCCAAGA/TGCAGTGCCATGTTCAAAACC
Albumin	ACCCCAAGTGTCAACTCCAA/GGTTCAGGACCACGGATAGA



**Figure S1.** FEM model geometry of the 3D-LOC for the simulation analysis of flow and oxygen mass transfer. (A) 3D geometry of the modeled microfluidic channel, which contained  $180 \times 6$  arrays of microwells. (B) A single microwell geometry of 3D-LOC, which contained a microporous membrane and a Ø 200 µm cell spheroid. The microwell had a V-shape cross section with round bottom (Ø 250 µm × 25° Angle). The fluid channel height, top PDMS layer thickness and microwell depth were 200µm, *L* and *D<sub>W</sub>*, respectively. (C) For comparison, a single microwell geometry with open microwell (Conventional perfusion method, 3D-perfusion) was also constructed and the dimension parameters were the same as those in (B).



**Figure S2.** A comparison of wall shear stress in two different perfusion methods (3D-perfusion and 3D-LOC). The wall shear stress distribution along the top red cut lines of the cell spheroid, described in (A), were plotted in different microwell depth configurations including (B)  $D_W$ = 200 µm, (C)  $D_W$ = 300 µm, and (D)  $D_W$ = 400 µm.



**Figure S3.** The percentage of the surface area of the cell spheroid (Oxygen concentration >  $C_{min}$ ) in the 3D-perfusion method with open microwell (OW) and in the 3D-LOC method with membrane (MW) at different microwell depths ( $D_W$  = 200, 300 and 400 µm) and flow rates (Q = 1, 10 and 100 µL/min).



**Figure S4.** A comparision of cell spheroids viability and loss status in three different perfusion methods (3D-perfusion (P1), 3D-perfusion (P2), and 3D-LOC) perfusion cultured for 1-6 days at different flow rates including (A)  $Q = 10 \mu L/min$ , and (B)  $Q = 100 \mu L/min$ . White arrows indicate the direction of fluid flow. Scale bars = 400  $\mu m$ .

## **Supplementary References**

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