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Supplemental Information

Differentiation and Functional Comparison of Monocytes and Macrophages from hiPSCs with Peripheral Blood Derivatives

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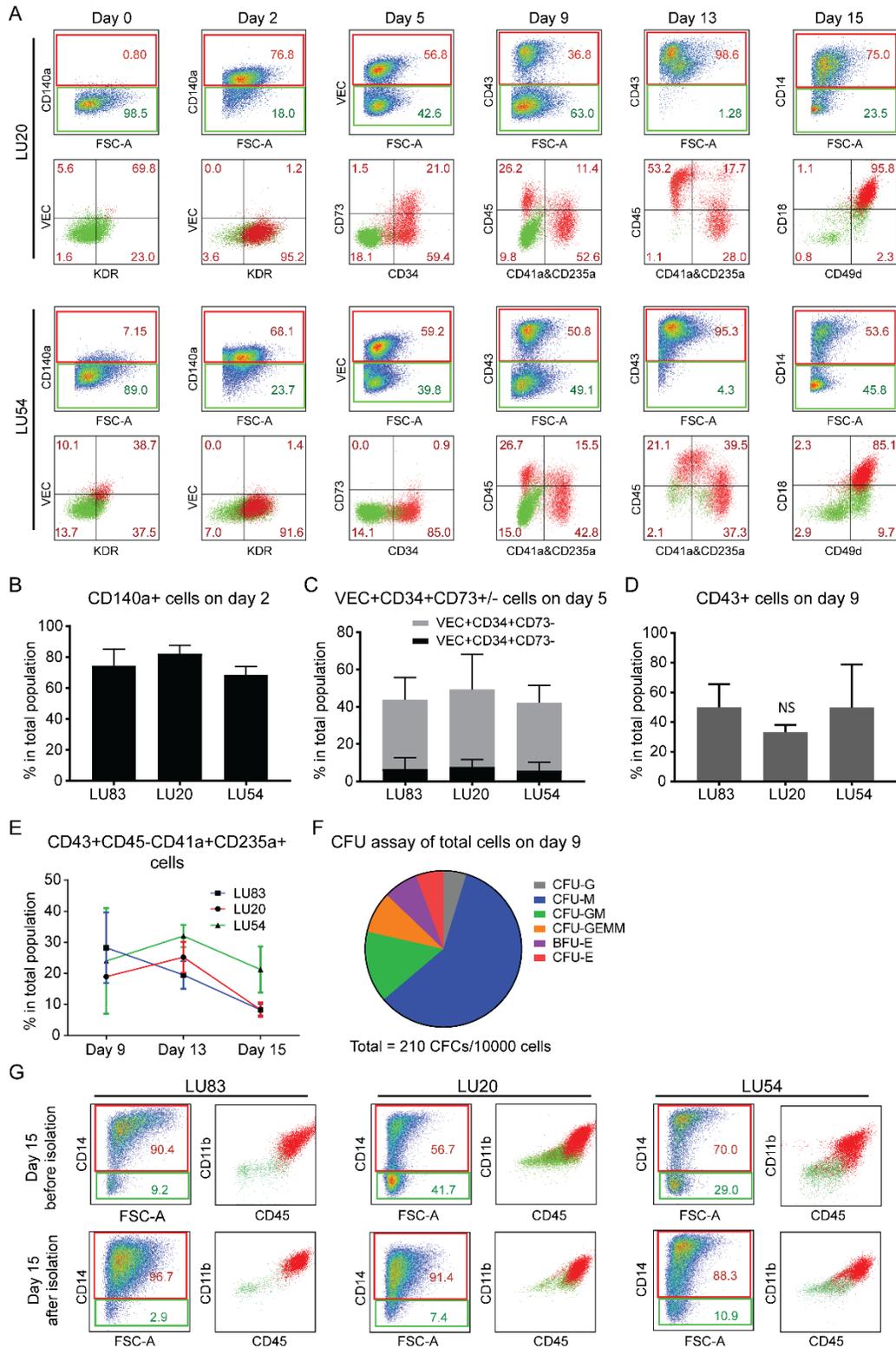
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Supplemental Figures and Legend



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Figure S1. Differentiation of CD14⁺ monocytes from hiPSCs. Related to Figure 1.

(A) FACS analysis of stage-specific markers at day 0, day 2, day 5, day 9, day 13 and day 15 of differentiation from LU20 and LU54. Positive populations are gated in upper panels and their percentages are shown in red in both upper and lower panels. (B) Percentage of early pan-mesodermal cell marker (PDGFR α) on day 2 of differentiation from three hiPSC lines (LU83, LU20 and LU54). (C) Percentage of non-HE (VEC+CD34+CD73⁺) and HE (VEC+CD34+CD73⁻) subsets on day 5 of differentiation from three hiPSC lines (LU83, LU20 and LU54). (D) Percentage of early HPC marker CD43 on day 9 of differentiation from three hiPSC lines (LU83, LU20 and LU54). (E) Percentage of erythro-megakaryocytic lineage cells (CD43+CD45-CD41a+CD235a⁺) in total cell population on day 9, day 13 and day 15 of differentiation from three hiPSC lines (LU83, LU20 and LU54). (F) CFU assay of total cell population on day 9 of differentiation from LU83. (G) Representative FACS analysis of CD14⁺ monocytes before and after MACS isolation on day 15 of differentiation from three hiPSC lines (LU83, LU20 and LU54). Error bars are \pm SD of three independent experiments in (B-E).

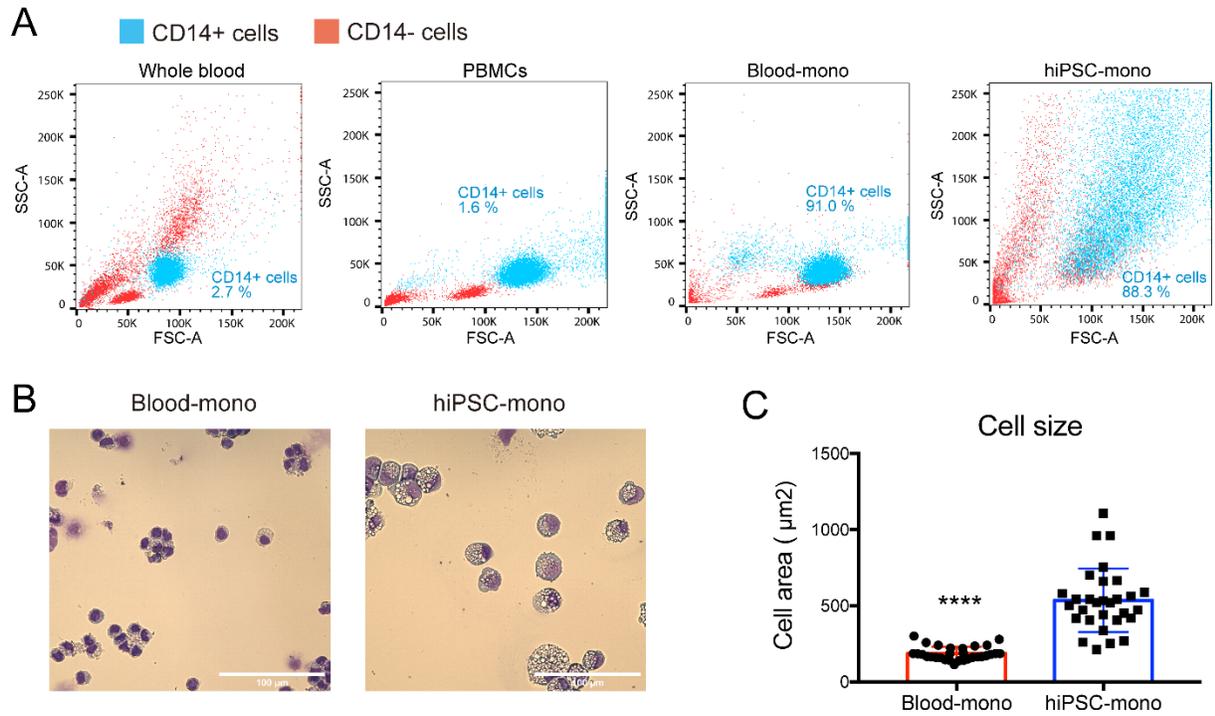


Figure S2. Comparison of cell sizes of whole blood, PBMCs, Blood-mono and hiPSC-mono. Related to Figure 2.

(A) FACS analysis of whole blood, PBMCs, Blood-mono from the same donor and hiPSC-mono on day 15 of differentiation from LU83 hiPSC line. (B) Giemsa staining of blood-mono isolated from human PBMC and hiPSC-mono isolated on differentiation day 15. Scale bar 100 μm . (C) Quantification of cell size of blood-mono and hiPSC-mono using Giemsa staining images. Cell area of 30 intact cells was measured from each cell type. Unpaired t-test. **** $p < 0.0001$.

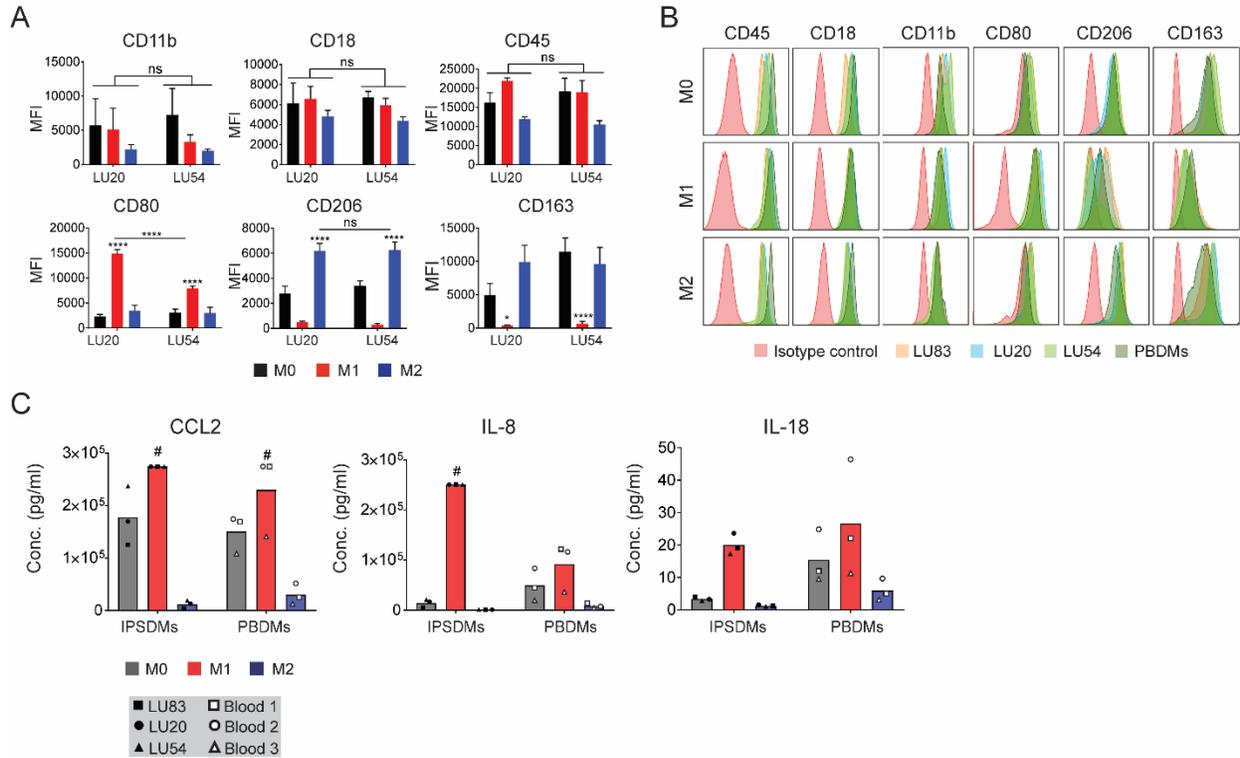


Figure S3. Characterization of IPSDMs and PBDMs. Related to Figure 3.

(A) Quantification of surface expression of pan-specific macrophage markers: CD11b, CD18 and CD45 and subtype-specific markers: CD80 (M1) and CD206 and CD163 (M2) on IPSDMs (differentiated from LU20 and LU54). Error bars are \pm SD of three independent experiments. Uncorrected Fisher's LSD test. ns = non-significant, * $p < 0.05$, **** $p < 0.0001$. (B) Representative FACS plots of pan-specific macrophage markers: CD11b, CD18 and CD45 and subtype-specific markers: CD80 (M1), CD206 and CD163 (M2) on IPSDMs (differentiated from LU20 and LU54) and PBDMs. (C) Quantification of secreted cytokines and chemokines by Multiplex assay using supernatants from IPSDMs and PBDMs after 48hours of polarization. Data are presented as mean of three biological replicates (three hiPSC lines or PBMC samples). # higher than the detection limit of Multiplex.

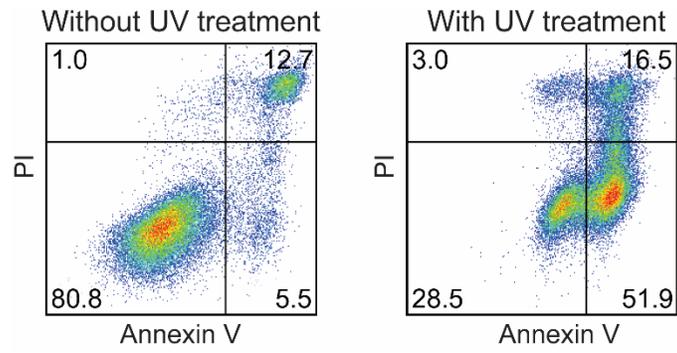


Figure S4. Induction of Apoptosis by UV Radiation. Related to Figure 5.
FACS analysis of apoptotic (Annexin V+ PI-) cells in hiPSCs without and with UV (35 J/cm²) treatment.

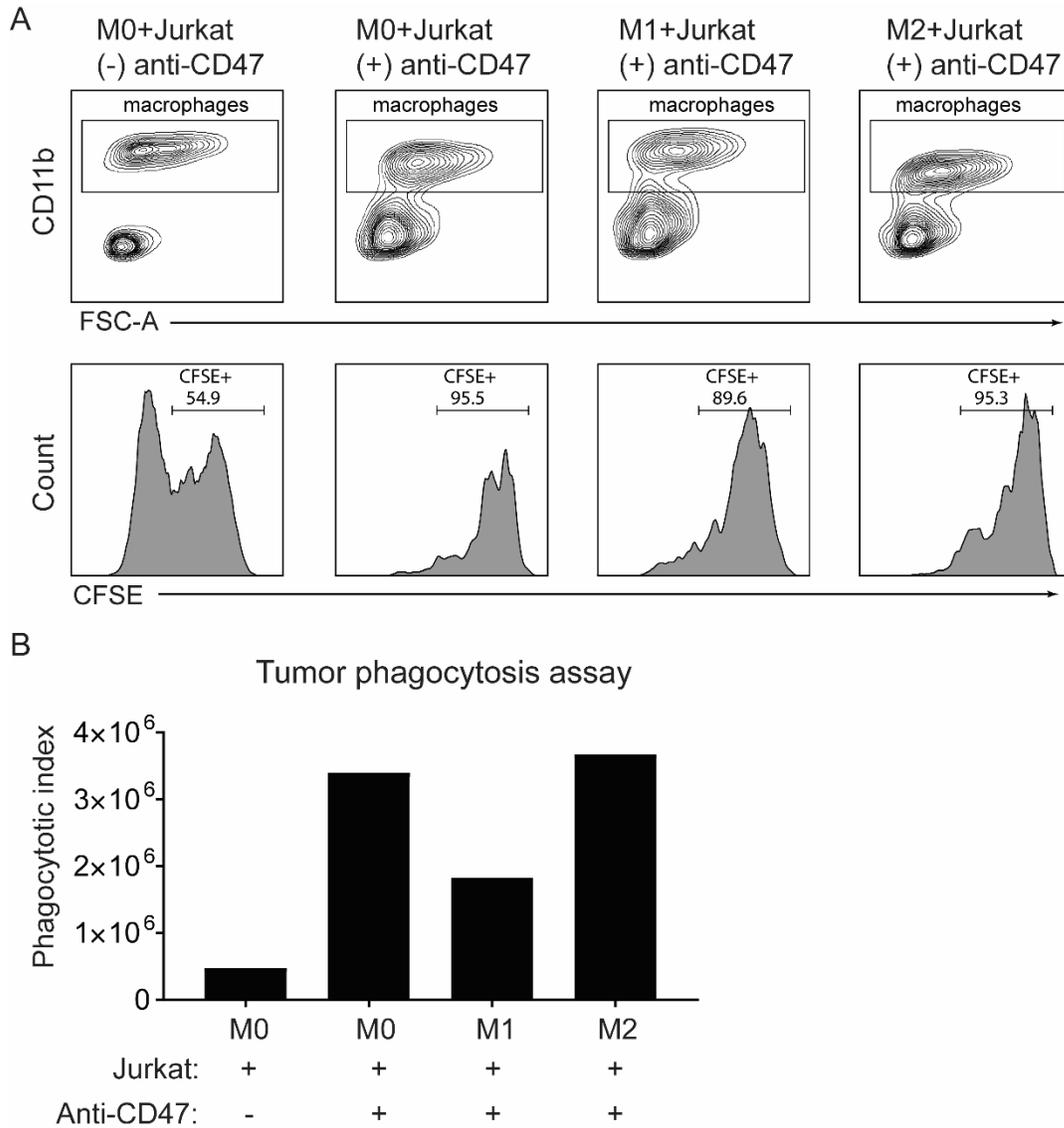


Figure S5. Characterization IPSDMs tumour phagocytosis activity. Related to Figure 6.

(A) FACS analysis of Jurkat cell phagocytosis by different subtypes of IPSDMs in the presence of CD47 blocking antibody. Jurkat cell phagocytosis by IPSDMs (M0) without CD47 blocking antibody is shown as a negative control. CD11b+ IPSDMs are gated (upper panel) and their CFSE intensities are shown as a histogram (lower panel). (B) Phagocytotic index of different subtypes of IPSDMs in the presence of CD47 blocking antibody. Jurkat cell phagocytosis by IPSDMs (M0) without CD47 blocking antibody is shown as a negative control. Percentage of CFSE+ macrophages was multiplied by MFI of CFSE to obtain the phagocytotic index. IPSDMs were differentiated from LU83 in (A-B).

Supplemental Tables

Medium component (stock concentration)	Source	Volume added (250ml final volume)	Final concentration
IMDM	Iscove's modified Dulbecco's medium (IMDM), no phenol red (Gibco, cat. no. 21056-023)	117.25 ml	--
F12	Ham's F-12 nutrient mix, GlutaMAX supplement (Gibco, cat. no. 31765-027)	117.25 ml	--
PVA (5%)	Poly vinyl alcohol (Sigma-Aldrich, cat. no. P8136-250G)	50 ul	10 mg/L
Lipids (100X)	Chemically defined lipid concentrate (Gibco, cat. no. 11905031)	250ul	0.1% (vol%)
ITS-X (100X)	Insulin-transferrin-selenium-ethanolamine (Gibco, cat. no. 51500-056)	5 ml	2% (vol%)
αMTG (1.3% in IMDM)	Mono-thio glycerol (Sigma-Aldrich, cat. no. M6145-25ml)	750 μ l	40 ul/L
AA2P (5 mg/ml)	Sigma-Aldrich, cat. no. A8960	3.2 ml	64 mg/L
GlutaMax (100X)	GlutaMAX-1 supplement (Gibco, cat. no. 35050-038)	2.5 ml	1% (vol%)
NEAA (100X)	MEM Non-Essential Amino Acids Solution (100X) (Gibco, Cat. No. 11140-035)	2.5 ml	1% (vol%)
Pen-strep (5,000 U/ml)	Gibco, cat no. 15070-063	1.25ml	0.5% (vol%)

Table S1. Formulation for IF9S medium. Related to Experiment Procedures.

Antibody	Fluorochrome	Source	Dilution	Catalog #
CD140a	BV421	BD Bioscience	1:100	562799
VE-Cadherin	Alexa488	eBioscience	1:50	53-1449-42
CD34	APC	Miltenyi Biotec	1:20	130-090-954
KDR	PE	R&D	1:20	FAB357P
CD73	PE	BD Pharmingen	1:20	550257
CD43	PE	BD Bioscience	1:20	560199
CD45	FITC	Miltenyi Biotec	1:20	130-080-202
CD41a	Vioblue	Miltenyi Biotec	1:20	130-105-610
CD235a	Vioblue	Miltenyi Biotec	1:20	130-100-273
CD14	PE	Miltenyi Biotec	1:20	130-091-242
CD11b	Vioblue	Miltenyi Biotec	1:20	130-097-336
CD18	FITC	Miltenyi Biotec	1:20	130-101-237
CD49d	PE-Vio770	Miltenyi Biotec	1:20	130-104-326
CD29	PE	eBioscience	1:50	12-0299-71
ICAM1	F	R&D	1:20	BBA20
E-Selectin	F	R&D	1:20	BBA21
VCAM1	PE	R&D	1:20	FAB5649P
CD31	APC	eBioscience	1:50	17-0319
CD105	Vioblue	Miltenyi Biotec	1:20	130-099-666
CD80	PE-Vio770	Miltenyi Biotec	1:20	130-101-218
CD206	FITC	Miltenyi Biotec	1:20	130-095-131
CD163	FITC	Miltenyi Biotec	1:100	130-112-290
CD172a	PE-Vio770	Miltenyi Biotec	1:20	130-099-793
Annexin-V	Pacific Blue	Thermofisher	1:20	A35122

Table S2. List of conjugated antibodies. Related to Experiment Procedures.

Gene	Forward sequence	Reverse sequence	Product size
<i>CD68</i>	GGAAATGCCACGGTTCATCCA	TGGGGTTCAGTACAGAGATGC	247
<i>IL1B</i>	ATGATGGCTTATTACAGTGGCAA	GTCGGAGATTCGTAGCTGGA	132
<i>IL6</i>	ACTCACCTCTTCAGAACGAATTG	CCATCTTTGGAAGGTTTCAGGTTG	149
<i>IL8</i>	AGCACTCCTTGGCAAACTG	CGGAAGGAACCATCTCACTG	116
<i>TNFA</i>	CCTCTCTCTAATCAGCCCTCTG	GAGGACCTGGGAGTAGATGAG	220
<i>CCL2</i>	CAGCCAGATGCAATCAATGCC	TGGAATCCTGAACCCACTTCT	190
<i>CCL5</i>	CCAGCAGTCGTCTTTGTCAC	CTCTGGGTTGGCACACACTT	54
<i>CXCL10</i>	GTGGCATTCAAGGAGTACCTC	TGATGGCCTTCGATTCTGGATT	198
<i>CD64</i>	AGCTGTGAAACAAAGTTGCTCT	GGTCTTGCTGCCCATGTAGA	75
<i>IDO1</i>	GCCAGCTTCGAGAAAGAGTTG	ATCCCAGAACTAGACGTGCAA	96
<i>NOX2</i>	ACCGGGTTTATGATATTCCACCT	GATTCGACAGACTGGCAAGA	135
<i>CD206</i>	TCCGGGTGCTGTTCTCCTA	CCAGTCTGTTTTTGATGGCACT	211
<i>CD163</i>	TTTGTCAACTTGAGTCCCTTCAC	TCCCGCTACACTTGTTTTCAC	127
<i>CD200R</i>	TGGTTGTTGAAAGTCAATGGCT	CTCAGATGCCTTCACCTTGTTT	153
<i>TGM2</i>	GAGGAGCTGGTCTTAGAGAGG	CGGTCACGACACTGAAGGTG	184
<i>IL1RA</i>	CATTGAGCCTCATGCTCTGTT	CGCTGTCTGAGCGGATGAA	167
<i>CCL22</i>	ATCGCCTACAGACTGCACTC	GACGGTAACGGACGTAATCAC	129
<i>CCL24</i>	ACATCATCCCTACGGGCTCT	CTTGGGGTCGCCACAGAAC	176
<i>TLR1</i>	CCACGTTCTAAAGACCTATCCC	CCAAGTGCTTGAGGTTACACAG	248
<i>TLR2</i>	ATCCTCCAATCAGGCTTCTCT	GGACAGGTCAAGGCTTTTTACA	118
<i>TLR4</i>	AGACCTGTCCCTGAACCCTAT	CGATGGACTTCTAAACCAGCCA	147
<i>TLR6</i>	TTCTCCGACGGAAATGAATTTGC	CAGCGGTAGGTCTTTTGGAAC	75
<i>TLR8</i>	ATGTTCCCTCAGTCGTCAATGC	TTGCTGCACTCTGCAATAACT	143
<i>CX3CRI</i>	ACTTTGAGTACGATGATTTGGCT	GGTAAATGTCGGTGACACTCTT	177
<i>S1PR1</i>	TTCCACCGACCCATGFACTAT	GCGAGGAGACTGAACACGG	185
<i>CD36</i>	GGCTGTGACCGGAACTGTG	AGGTCTCCAACCTGGCATTAGAA	92
<i>MERTK</i>	CTCTGGCGTAGAGCTATCACT	AGGCTGGGTTGGTGAAAACA	162
<i>RPL37A</i>	GTGGTTCCTGCATGAAGACAGTG	TTCTGATGGCGGACTTTACCG	84
<i>HARP</i>	CACCATTGAAATCCTGAGTGATGT	TGACCAGCCCAAAGGAGAAG	116

Table S3. Sequence of primes used for qPCR. Related to Experiment Procedures.

Supplemental Videos

Movie S1. Monocyte differentiation day 7 to day 9. Related to Figure 1.

Time-lapse imaging of monocyte differentiation from LU83 hiPSC line. Video was taken from differentiation day 7 to day 9 in a timespan of ~50 hours. The video is 15 frames/second. The interval between each frame is 40 minutes in a real time. Scale bar represents 200 μm .

Movie S2. Monocyte differentiation day 6 to day 8. Related to Figure 1.

Time-lapse imaging of monocyte differentiation from LU83 hiPSC line. Video was taken from differentiation day 6 to day 8 in a timespan of ~48 hours. The video is 15 frames/second. The interval between each frame is 16 minutes in a real time. Scale bar represents 100 μm .

Movie S3. Tumour phagocytosis by IPSDMs. Related to Figure 6.

Tumor cell phagocytosis by M0-IPSDMs differentiated from LU83. Video was taken 30 minutes after co-culture of tumor cells with M0-IPSDMs. Video was made in the same field as Figure 6C. Video is 15 frames/second and interval between each frame is 30 seconds in a real time. Scale bar represents 50 μm .