Evidences for KSHV-induced Mesenchymal-to-Endothelial Transition

Transforming Oral Mesenchymal Stem Cell to Kaposi's Sarcoma

Supplementary Data

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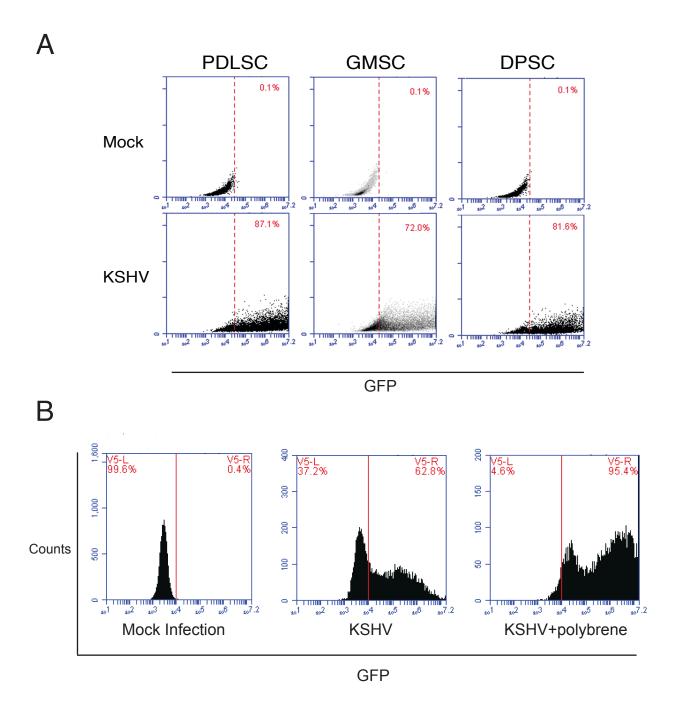


Fig. S1. (A) Primary oral MSCs of different origins (PDLSCs, GMSCs and DPSCs) were infected with GFP-KSHV in an MOI of 50 (KSHV genome equivalent) in the presence of polybrene for 48 hours and analyzed by GFP fluoresces. The rates of KSHV infection were determined by flow cytometric analysis of GFP-expression cells. (B) Comparison of infectivity of KSHV to PDLSCs in the presence and absence of polybrene.

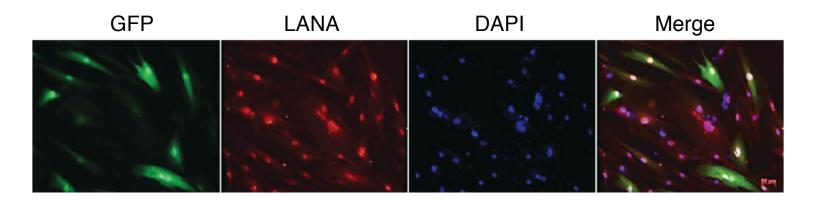


Fig. S2. Persistent KSHV infection in PDLSCs. Primary PDLSCs were infected with GFP-KSHV in an MOI of 50 (KSHV genome equivalent). Cells were cultured for four weeks with out any antibiotics selection. Cells were analyzed by GFP fluoresce and IFA with an anti-LANA antibody.

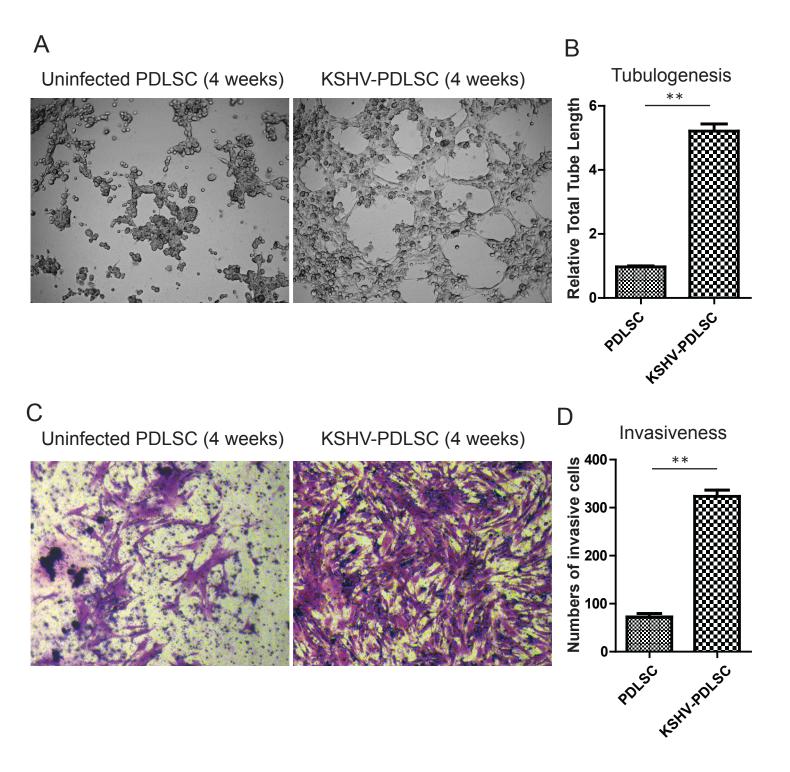


Fig. S3. Long-term cultured KSHV-infected PDLSCs preserve the potential of tubulogenesis and invasiveness. Primary PDLSCs were infected with GFP-KSHV in an MOI of 50 (KSHV genome equivalent). Cells were cultured without any antibiotics selection in cytokine-free medium for four weeks. Mock- and virally infected PDLSCs were loaded on the top of Matrigel and the ability of the cells in formation of capillary-like tubules was analyzed under a microscope (A). Quantfication of tubulogenesis was performed by calculating the average length of the capillary-like tubules (B). Cells were subjected to Transwell migration assay to assess cell invasion ability. PDLSCs or KSHV-PDLSCs were seeded in the upper chamber of Transwell in serum-freea-MEM. a-MEM containing 10% FBS was used as a stimulus for chemotaxis in the lower chamber. Cells migrated to the lower chamber were stained with crystal violet (C). Quantfication of invasiveness was performed by counting the invasive cell numbers from 5 different fields (D).

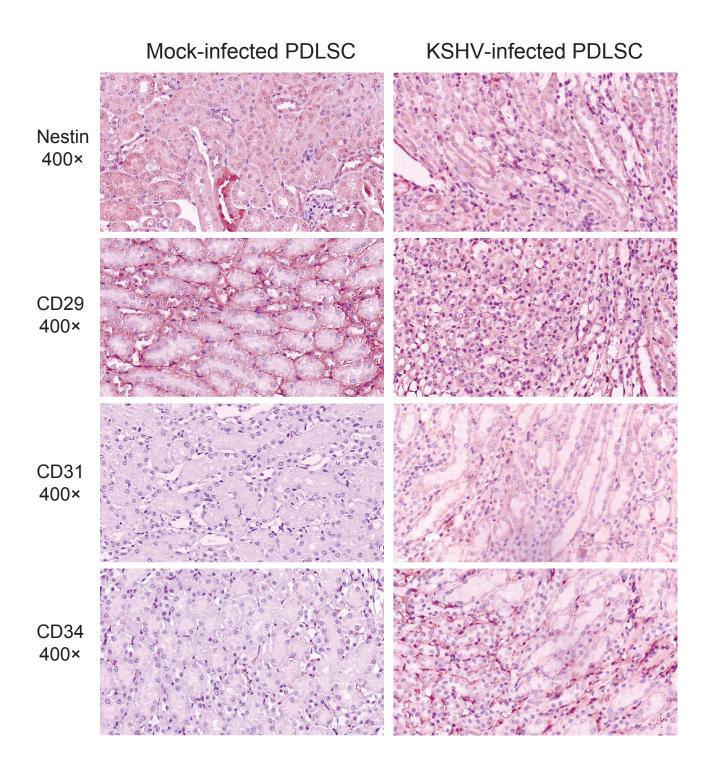


Fig. S4. Expression of oral MSC-specific and endothelial markers in mock- and KSHV-infected MSCs. Sections of kidney capsule implantation of mock- and KSHV-infected PDLSCs were subjected to immunohistochemical analyses with antibodies agianst neural crest-derived stem marker Nestin, oral MSC marker CD29, endothelial marker CD31 and hematopoietic/endothelial progenitor marker CD34.

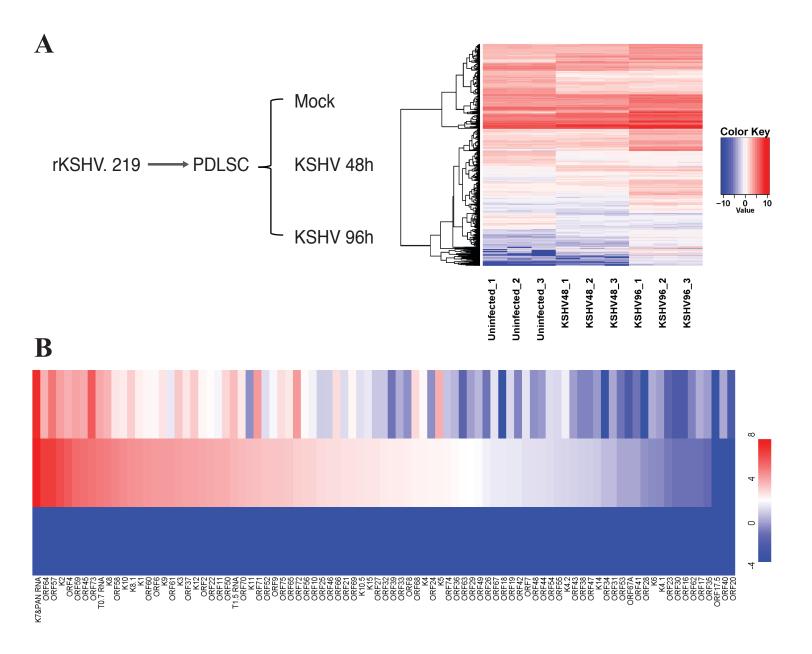


Fig. S5. Gene expression profiling analysis of KSHV-infected oral MSCs. PDLSCs (p2) were infected with rKSHV.219 in an MOI of 50 (viral genomic DNA equivalent). RNA was isolated from mock- and KSHV-infected PDLSCs (48 or 96 hour post-infection) and subjected to RNA-seq analysis. Clean reads were mapped to human reference genome (version hg19/GRCh37) and KSHV reference genome (version CQ994935.1). Differentially expressed cellular (A) and viral genes (B) were shown in heatmap.

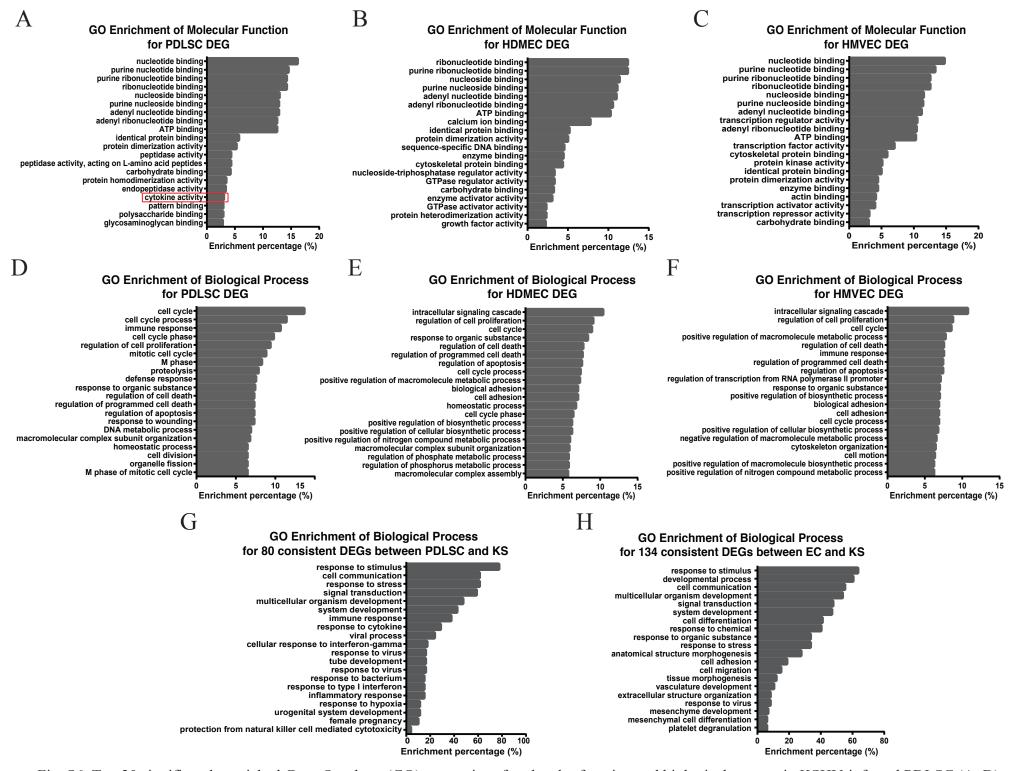


Fig. S6. Top 20 significantly enriched Gene Ontology (GO) categories of molecular function and biological process in KSHV-infected PDLSC (A, D), HDMEC (B, E) and HMVEC (C, F), as well as biological process in consistent DEGs of KS and PDLSC or endothelial cells (G, H).

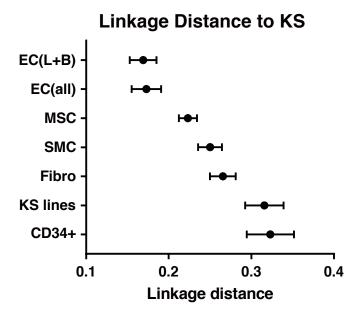


Fig. S7. The average linkage distance between Kaposi's sarcoma (KS) and each cell group was re-calculated from the published data (52) by Pearson correlation coefficient and then subtracted by 1. EC (L+B): endothelial cells (LECs+BECs).

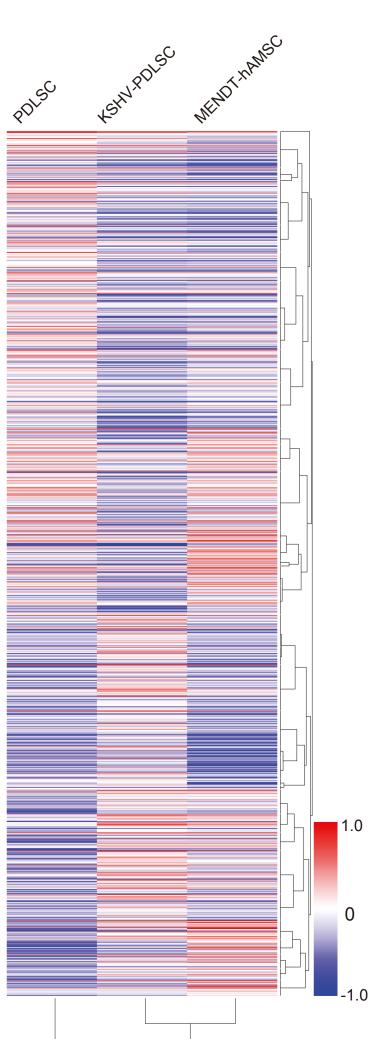


Fig. S8. Relationship of KSHV-infected MSC and an MEndT model (induced hAMSC) in gene expression profile. Mock- and KSHV-infected PDLSCs were normalized and subjected to a comparison with hAMSCs that have been induced for endothelial differentiation. Unsupervised clustering of samples (X-axis) and genes (Y-axis) were performed by average linkage method.

Table S1. Common DEGs in KSHV-PDLSC vs. PDLSC and KS signature

Function	Cluster	Gene	KSHV-	KS vs.	KSHV-	KSHV-	KSHV-	KSHV-
		symbol	MSC vs.	skin	HDMEC vs. HDMEC		KSHV- LEC vs. LEC	BEC vs.
		BGN		↑ ↑	HDMEC			DEC ↓
		DSP	↑ ↑	↓ ↓	_			_
		EDNRA	1 1	1	_	_		_
		FGF2	↑ ·	↑ ↑ ↑ ↑	_	_	_	↓ ↓
	<u>.</u> .	MMP11	' ↑	↑ ↑ ↑ ↑	_	_	J.	1
Mesenchymal to		PDPN	' ↑↑	↑ ↑ ↑ ↑	4 4	^		' ↑
Endothelial 7	Γransition	PROX1	1 1	↑ ↑ ↑ ↑	1			' ↑
		PGF	1 1 1 1 1 1 1 1 1 1 1	↑ ↑ ↑ ↑	• ↓			
		PML	1 1 1 1 1 1 1 1 1 1 1	↑ ↑ ↑ ↑	_	HMVEC vs. L		_
		TGFB3	↑ ↑ ↑ ↑	↑ ↑	_	_	_	_
		TGFBR2	↑ ↑ ↑↑	↑ ↑	_	_	J.	_
		ANGPTL2	<u> </u>	<u> </u>	_	Λ		↑ ↑
Vessel deve	lopment	VCAM1	ሳተ	↑ ↑ ↑↑	_	<u>.</u>	_	↓
	F	WARS	↑ ↑ ↑↑	↑ ↑	1	_	T.	_
		CCL5	<u></u>	<u> </u>		_		↑ ↑
		CCL8	↑↑ ↑↑	↑↑	_	_		_
	Chemotaxis	CXCL10	↑↑	↑↑ ↑↑	_	^	LEC vs. LEC	_
		SHC3	↑↑	↓↓	_	_		_
		STAT1	↑ ↑	ተተ	↑ ↑	_		_
		APOL2	↑ ↑	↑ ↑		_		_
		APOL3	ተተ	ተተ	_	_		4
		CFH	ተተ	ተተ	_	_		_
Immune response		FN1	↑ ↑	↑ ↑	_	$\downarrow \downarrow$	$\downarrow \downarrow$	4
	Inflammation	IFI16	↑ ↑	↑ ↑	_	^	^	_
		IL-1RN	↑ ↑	$\downarrow \downarrow$	_	_	_	_
		LY96	个个	ተተ	_	_	_	↑
		NT5E	个个	ተተ	4	↑ ↓	V	_
		PLA2G4C	ተተ	ተተ	_	_		
	Other genes	FCGRT	ተተ	ተተ	_	_	_	_
	Other genes	GBP1	ተተ	ተተ	↑ ↑	↑ ↑	_	$\downarrow \downarrow$
	involved in	GCH1	ተተ	ተተ	_	^	ተ ተ	_

	immune	HLA-A	↑ ↑	ተተ	_	_	_	_
	response	HLA-B	$\uparrow \uparrow$	ተተ	↑	↑	_	_
	•	HLA-C	$\uparrow \uparrow$	ተተ	↑	↑	_	_
		HLA-E	$\uparrow \uparrow$	ተተ	_	_	_	_
		HLA-F	$\uparrow \uparrow$	$\uparrow\uparrow$	_	_	_	_
		HLA-G	$\uparrow \uparrow$	ተተ	_	_	_	_
		IFI35	$\uparrow \uparrow$	$\uparrow\uparrow$	_	^	_	_
		IFITM1	$\uparrow \uparrow$	ተተ	_	^	↑	_
		IFITM3	$\uparrow \uparrow$	ተተ	_	_	_	_
		ISG15	$\uparrow \uparrow$	ተተ	_	^	$\uparrow \uparrow$	↑
		ISG20	$\uparrow \uparrow$	ተተ	_	_	_	$\downarrow \downarrow$
		NOD2	$\uparrow \uparrow$	$\downarrow \downarrow$	$\uparrow \uparrow$	_	_	_
		PSMB9	$\uparrow\uparrow$	ተተ	↑	^	^	↑
		SP110	$\uparrow \uparrow$	$\uparrow \uparrow$	_	↑ ↑	_	V
		SPON2	$\uparrow \uparrow$	ተተ	_	_	_	_
		TACC3	$\downarrow \downarrow$	$\uparrow \uparrow$	V	V	_	$\downarrow \downarrow$
		TIMP1	$\uparrow \uparrow$	$\uparrow \uparrow$	_	_	↑ ↑	$\uparrow\uparrow$
		TNFSF10	$\uparrow \uparrow$	$\uparrow\uparrow$	↑ ↑	_	$\downarrow \downarrow$	_
		TRIM25	$\uparrow \uparrow$	$\downarrow \downarrow$	_	_	_	_
		TRIM38	↑ ↑	ተተ	_	↑	↑	↑
		CCND1	$\uparrow \uparrow$	$\downarrow \downarrow$	_	_	$\uparrow \uparrow$	$\uparrow \uparrow$
		DHCR24	$\downarrow \downarrow$	$\downarrow \downarrow$	_	_	_	_
		IGF1	$\uparrow \uparrow$	$\uparrow\uparrow$	_	_	$\downarrow \downarrow$	_
Cell prolif	eration	KLF4	$\uparrow \uparrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	4	\	_
		MCM5	$\downarrow \downarrow$	ተተ	4	_	\	V
		MCM6	$\downarrow \downarrow$	ተተ	4	_	\	V
		TOB1	↑ ↑	$\downarrow \downarrow$		<u> </u>	↓ ↓	↑
		CD36	$\uparrow \uparrow$	$\uparrow\uparrow$	↑ ↑	4	$\downarrow \downarrow$	↑
		DAPK2	$\uparrow \uparrow$	$\downarrow \downarrow$	_	_	\	_
		DUSP6	$\uparrow \uparrow$	$\uparrow\uparrow$	_	_	V	V
Signal Tran	sduction	FAM129A	$\downarrow \downarrow$	$\uparrow\uparrow$	_	_	V	_
Signai II an	Suuctivii	GDF5	$\uparrow\uparrow$	$\downarrow \downarrow$	_	_	$\downarrow \downarrow$	_
		GPSM2	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	V	_	_
		LGALS9	$\uparrow\uparrow$	$\uparrow\uparrow$	_	↑ ↑	V	_
		RGS2	$\uparrow \uparrow$	$\downarrow \downarrow$	_	_	_	_

	SRGAP3	↑ ↑	$\downarrow \downarrow$	_	_	_	_
	TBC1D1	$\downarrow \downarrow$	ተተ	_	_	_	_
	BACE2	↑ ↑	↑ ↑	_	_	\	_
	EHF	$\uparrow\uparrow$	$\downarrow \downarrow$	1	_	_	_
	KLHL3	$\uparrow\uparrow$	$\uparrow\uparrow$		V	_	
Metabolism	PBXIP1	$\uparrow \uparrow$	$\uparrow\uparrow$	_	_	V	↑ ↑
	TIMP3	$\uparrow\uparrow$	$\uparrow\uparrow$	$\uparrow \uparrow$	_	$\downarrow \downarrow$	_
	UBE2L6	$\uparrow\uparrow$	$\uparrow \uparrow$	_	$\uparrow \uparrow$	V	_
	UBE2S	$\downarrow \downarrow$	$\uparrow \uparrow$	V	V		4
	C1ORF38	↑ ↑	↑ ↑	_	_	↑	_
Cell adhesion	CLCA2	$\uparrow\uparrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	_	↑	1
Cell adhesion	PCDH17	ተተ	ተተ	↑	↑ ↑	↓	_
	PODXL	ተተ	ተተ	V	↑	ተተ	ተተ
	ADD3	$\uparrow \uparrow$	$\uparrow\uparrow$	1	_	$\downarrow \downarrow$	_
	AKAP12	$\uparrow \uparrow$	$\uparrow\uparrow$	_	1	V	_
	AKR1B1	$\uparrow \uparrow$	$\uparrow\uparrow$	$\downarrow \downarrow$	_	_	_
	ALDH3A1	$\uparrow \uparrow$	$\downarrow \downarrow$	_	_	↑	11
	AVPI1	$\uparrow \uparrow$	$\downarrow \downarrow$	_	_	_	_
	BTN3A3	$\uparrow \uparrow$	$\uparrow\uparrow$	_	_	V	_
	ChGn	$\uparrow \uparrow$	$\uparrow\uparrow$	_	_	V	_
	COTL1	$\downarrow \downarrow$	$\uparrow\uparrow$		_	$\downarrow \downarrow$	•
	CPA4	$\downarrow \downarrow$	$\downarrow \downarrow$	_	_	$\downarrow \downarrow$	_
	CYP2E1	$\uparrow\uparrow$	$\downarrow \downarrow$	_	_	_	_
Others	DTNA	$\uparrow\uparrow$	$\downarrow \downarrow$	_	_	$\downarrow \downarrow$	_
Others	DTYMK	$\downarrow \downarrow$	$\uparrow \uparrow$	V	_	_	_
	EPS8L2	$\uparrow\uparrow$	$\downarrow \downarrow$	_	_	$\downarrow \downarrow$	_
	GABBR2	ተተ	ተተ	$\downarrow \downarrow$	4	\	4
	GCAT	$\downarrow \downarrow$	$\downarrow \downarrow$	_	_	↑	↑
	GOLM1	ተተ	ተተ	_	_	ተተ	_
	HSPA4L	$\uparrow\uparrow$	$\downarrow \downarrow$	_	↑ ↑	_	_
	KCTD12	ተተ	$\uparrow \uparrow$	_	V	$\downarrow \downarrow$	V
	KPNA2	$\downarrow \downarrow$	$\uparrow \uparrow$	_	_	_	V
	KRT10	$\downarrow \downarrow$	$\downarrow \downarrow$	_	_	↑	_
	LAP3	$\uparrow \uparrow$	$\uparrow \uparrow$	_	↑	4	_
	LGMN	ተተ	↑ ↑	_	_	↓	1

```
MDK
                                          \uparrow\uparrow
                                                                    \uparrow\uparrow
 OLFML1
                                                                    \uparrow\uparrow
                                          \uparrow\uparrow
      PAM
                                          \uparrow\uparrow
                                                                    \uparrow\uparrow
                                                                                                                                                                                                     \uparrow
 RPLP0P2
                                          \uparrow\uparrow
                                                                    \downarrow \downarrow
                                                                    \downarrow \downarrow
  SLC9A7
                                          \uparrow\uparrow
                                                                                                                                                                       \downarrow \downarrow
    STOM
                                          \uparrow\uparrow
                                                                    \uparrow\uparrow
                                                                                                  \uparrow\uparrow
 TAGLN2
                                          \downarrow \downarrow
                                                                    \uparrow\uparrow
  TM4SF1
                                          \uparrow\uparrow
                                                                    \uparrow\uparrow
                                                                                                                                        \uparrow\uparrow
                                                                                                                                                                                                     \uparrow
TMEM140
                                          \uparrow\uparrow
                                                                    \uparrow\uparrow
                                                                                                  \uparrow\uparrow
                                                                                                                                        1
                                                                                                                                                                                                   \uparrow\uparrow
      TPP1
                                          \uparrow\uparrow
                                                                    \uparrow\uparrow
```

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↑↑: log<sub>2</sub> (Fold change) > 1;
↑: 0.5 ≤ log<sub>2</sub> (Fold change) ≤1;
-: No significant change;
↓↓: log<sub>2</sub> (Fold change) < -1;</li>
↓: -1 ≤ log<sub>2</sub> (Fold change) ≤ -0.5.
```

Table S2. Consistently regulated genes between KSHV-PDLSC vs. PDLSC and MEndT hAMSC vs. hAMSC

Functi	on Cluster	Gene symbol	KSHV-MSC vs. MSC	MEndT hAMSC vs. hAMSC	KS vs. Skin	Function Cluster	Gene symbol	KSHV-MSC vs. MSC	MEndT hAMSC vs. hAMSC	KS vs. Skin
		ANGPTL1	↑ ↑	↑ ↑	↑		BTN3A1	↑ ↑	↑ ↑	↑ ↑
		EDNRA	$\uparrow \uparrow$	$\uparrow \uparrow$	↑ ↑		BTN3A3	$\uparrow \uparrow$	$\uparrow \uparrow$	$\uparrow\uparrow$
		FGF5	$\uparrow \uparrow$	$\uparrow \uparrow$	↑		CD55	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$
		FGF7	$\uparrow \uparrow$	$\uparrow \uparrow$	↑		C1R	$\uparrow \uparrow$	$\uparrow\uparrow$	_
	F 1 41 12 1	PDPN	$\uparrow \uparrow$	$\uparrow \uparrow$	↑ ↑		C1S	$\uparrow \uparrow$	$\uparrow\uparrow$	_
	Endothelial related	PROX1	$\uparrow \uparrow$	$\uparrow \uparrow$	↑ ↑		CFH	$\uparrow \uparrow$	$\uparrow \uparrow$	$\uparrow\uparrow$
	Telateu	TGFB3	$\uparrow \uparrow$	$\uparrow \uparrow$	↑ ↑		COTL1	$\downarrow \downarrow$	$\downarrow \downarrow$	$\uparrow\uparrow$
		PTGS2	$\uparrow \uparrow$	$\uparrow \uparrow$	↑		EPHA3	$\uparrow \uparrow$	$\uparrow \uparrow$	$\uparrow\uparrow$
		SLC40A1	$\uparrow \uparrow$	$\uparrow \uparrow$	_		IFI44	$\uparrow \uparrow$	$\uparrow\uparrow$	1
		SRPX2	$\uparrow \uparrow$	$\uparrow \uparrow$	↑	Immune	IFI44L	$\uparrow \uparrow$	$\uparrow\uparrow$	$\uparrow\uparrow$
D:664		WARS	$\uparrow \uparrow$	$\uparrow \uparrow$	↑ ↑	response	IFITM1	$\uparrow \uparrow$	$\uparrow \uparrow$	$\uparrow\uparrow$
Different -iation		ACTA2	1	1	$\downarrow \downarrow$		IFITM3	$\uparrow \uparrow$	$\uparrow \uparrow$	$\uparrow\uparrow$
-iation		CMKLR1	$\uparrow \uparrow$	$\uparrow \uparrow$	_		IGDCC4	$\uparrow \uparrow$	ተተ	_
		CNN1	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$		PSMB8	$\uparrow \uparrow$	ተተ	↑
		DACT1	$\downarrow \downarrow$	$\downarrow \downarrow$	4		SAMHD1	$\uparrow \uparrow$	ተተ	$\uparrow\uparrow$
		DDIT3	$\downarrow \downarrow$	$\downarrow \downarrow$	\		TAPBPL	$\uparrow \uparrow$	$\uparrow \uparrow$	1
	Mesenchymal	EFEMP1	$\uparrow \uparrow$	$\uparrow \uparrow$	$\downarrow \downarrow$		TLR3	$\uparrow \uparrow$	ተተ	_
	related	EGFL6	$\uparrow \uparrow$	$\uparrow \uparrow$	$\downarrow \downarrow$		TNFAIP3	$\uparrow\uparrow$	ተተ	_
		ETV1	$\uparrow \uparrow$	$\uparrow \uparrow$	↑		TNFAIP6	$\uparrow \uparrow$	$\uparrow \uparrow$	\
		NBEA	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$		TNFRSF10D	↓ ↓	$\downarrow \downarrow$	$\downarrow \downarrow$
		SVIL	$\uparrow \uparrow$	$\uparrow \uparrow$	$\downarrow \downarrow$		ACSL4	↓ ↓	4 4	_
		TAGLN	$\downarrow \downarrow$	$\downarrow \downarrow$	4		APOD	$\uparrow\uparrow$	↑ ↑	↑
		TPM1	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$		APOE	$\uparrow \uparrow$	↑ ↑	↑

		ADAMTSL4	↑ ↑	↑ ↑	_		APOL1	↑ ↑	↑ ↑	ተተ
		CRABP2	↑ ↑	ተተ	$\downarrow \downarrow$		CHST7	ተተ	$\uparrow \uparrow$	ተተ
	Other genes involved in differentiatio n	DCN	↑ ↑	ተተ	4	Metabolic	ENO1	$\downarrow \downarrow$	$\downarrow \downarrow$	↓
	o .	DIAPH3	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$		PDK4	$\uparrow \uparrow$	$\uparrow \uparrow$	_
		IGF1	↑ ↑	ተተ	ተተ		PLA2G4A	$\uparrow \uparrow$	$\uparrow \uparrow$	$\downarrow \downarrow$
		OXTR	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$		PTGIS	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$
	11	TSHZ2	↑ ↑	ተተ	_		SMOX	$\uparrow \uparrow$	$\uparrow \uparrow$	↑
		TXNIP	↑ ↑	↑ ↑	$\downarrow \downarrow$		TCN2	↑ ↑	$\uparrow \uparrow$	_
							COL21A1	↑ ↑	个个	_
		AKAP12	↑ ↑	个个	ተተ		CORO2B	↑ ↑	↑ ↑	V
		ARHGAP20	$\uparrow \uparrow$	ተተ	_		CPA4	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$
		BST2	↑ ↑	ተተ	ተተ		GADD45A	$\downarrow \downarrow$	$\downarrow \downarrow$	↓
		CHI3L1	$\uparrow\uparrow$	个个	$\downarrow \downarrow$		KCNJ2	ተተ	ተተ	_
		CHI3L2	$\uparrow\uparrow$	$\uparrow \uparrow$	$\downarrow \downarrow$		MAMDC2	$\uparrow\uparrow$	$\uparrow \uparrow$	_
		FAM129A	$\downarrow \downarrow$	$\downarrow \downarrow$	ተተ		MME	$\uparrow\uparrow$	$\uparrow \uparrow$	V
C:14		FAM180A	$\uparrow\uparrow$	$\uparrow \uparrow$	_	Others	PARP14	$\uparrow\uparrow$	$\uparrow \uparrow$	_
Signai t	ransduction	FIBIN	$\uparrow\uparrow$	$\uparrow \uparrow$	_		PLAT	$\uparrow\uparrow$	$\uparrow \uparrow$	↑
		FOS	$\uparrow\uparrow$	$\uparrow \uparrow$	ተተ		SDK1	$\uparrow\uparrow$	$\uparrow \uparrow$	_
		F10	$\uparrow\uparrow$	$\uparrow \uparrow$	$\downarrow \downarrow$		SLC1A3	$\uparrow \uparrow$	$\uparrow \uparrow$	\
		GGH	$\downarrow \downarrow$	$\downarrow \downarrow$	4		SLC38A4	$\downarrow \downarrow$	$\downarrow \downarrow$	V
		HSPA4L	$\uparrow\uparrow$	$\downarrow \downarrow$	$\downarrow \downarrow$		SPON1	$\uparrow \uparrow$	$\uparrow \uparrow$	↑
		PCDHB14	$\uparrow\uparrow$	$\uparrow \uparrow$	_		STOM	$\uparrow \uparrow$	$\uparrow \uparrow$	$\uparrow \uparrow$
		PMEPA1	$\downarrow \downarrow$	$\downarrow \downarrow$	ተተ		TRANK1	$\uparrow\uparrow$	$\uparrow \uparrow$	_

^{↑↑:} log₂ (Fold change) > 1;

DEGs present both in KSHV-infected MSC and KS signature were marked as red.

 $[\]uparrow$: 0.5 ≤ log₂ (Fold change) ≤1;

^{-:} No significant change;

^{↓↓:} log₂ (Fold change) < -1;</pre>

Table S3. Common DEGs in KSHV-HDMEC vs. HDMEC and KS signature

Funct	ion Cluster	Gene symbol	KSHV- HDMEC vs. HDMEC	KSHV- HMVEC vs. HMVEC	KSHV- LEC vs. LEC	KSHV -BEC vs. BEC	KS vs. skin	KSHV- MSC vs. MSC
		COL15A1	ተተ	个个	_	↑	个个	_
		COL5A1	$\downarrow \downarrow$	V	_	4	$\uparrow \uparrow$	V
		EZR	$\downarrow \downarrow$	_	_	V	$\downarrow \downarrow$	V
	ıl-mesenchymal	LMO2	↑ ↑	_	_	_	$\uparrow\uparrow$	_
tra	ansition	S100A4	$\downarrow \downarrow$	_	_	↑	$\uparrow \uparrow$	_
		SEMA3C	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	_	$\downarrow \downarrow$	↑
		TGFA	ተተ	V	V	_	$\downarrow \downarrow$	
		ADORA2B	4 4	_	$\downarrow \downarrow$	1	$\downarrow \downarrow$	_
		ANGPT2	ተተ	↑ ↑	$\uparrow \uparrow$	↑	$\uparrow \uparrow$	↑
Vessel o	development	FGFR2	$\downarrow \downarrow$	_	↑	ተተ	$\downarrow \downarrow$	_
		LYVE1	ተተ	↑ ↑	_	_	$\uparrow \uparrow$	_
		PGF1	↓	↑	V	_	↑ ↑	ተተ
	Chemotaxis	STAT1	ተተ	_	_	_	↑ ↑	ተተ
	Inflammation	A2M	ተተ	↑ ↑	_	ተተ	$\uparrow\uparrow$	_
	Inflammation	MGLL	$\downarrow \downarrow$	_	_	V	$\uparrow\uparrow$	_
		CBLB	ተተ	_	_	ተተ	↑ ↑	_
		GBP1	ተተ	↑ ↑	_	$\downarrow \downarrow$	$\uparrow\uparrow$	ተተ
	Other genes	GIMAP5	ተተ	_	_	↑	$\uparrow \uparrow$	_
Immune	involved in	HLX	ተተ	$\uparrow \uparrow$	$\uparrow \uparrow$	ተተ	$\uparrow \uparrow$	_
response	immune	MICAL2	$\downarrow \downarrow$	↓	_	4	$\uparrow \uparrow$	4
response	response	NOD2	ተተ	_	_	_	$\downarrow \downarrow$	ተተ
		PTGER4	$\downarrow \downarrow$	_	_	$\downarrow \downarrow$	↑ ↑	_
		TNFSF10	ተተ	_	$\downarrow \downarrow$	_	↑ ↑	ተተ
	1	KLF4	↓ ↓	V	V	_	↓ ↓	个个
Cell p	roliferation	MYCN	ተተ	_	V	_	$\downarrow \downarrow$	_
		TBX5	4 4	_	_	_	$\downarrow \downarrow$	_
		CD36	ተተ	V	↓ ↓	↑	↑ ↑	ተተ
		EFS	ተተ	_	_	_	$\downarrow \downarrow$	_
		EGFR	↓ ↓	4 4	V	V	$\downarrow \downarrow$	↑
		EPS8	4 4	_	_	_	↑ ↑	_
Me	tabolism	GPSM2	$\downarrow \downarrow$	V	_	_	$\downarrow \downarrow$	$\downarrow \downarrow$

	RAB38	$\downarrow \downarrow$	_	_	V	$\downarrow \downarrow$	_
	RASGRP3	↑ ↑	ተተ	_	ተተ	$\uparrow \uparrow$	^
	RGS4	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	$\uparrow \uparrow$	↑
	RIPK4	$\downarrow \downarrow$	_	_	V	$\downarrow \downarrow$	_
	STK32B	$\downarrow \downarrow$	↑	_	V	$\uparrow \uparrow$	_
	BTRC	↑ ↑	_	_	个个	↓ ↓	_
Metabolism	CSGALNACT1	$\uparrow \uparrow$	_	_	_	ተተ	ተተ
	MAOA	$\uparrow \uparrow$	V	1	$\downarrow \downarrow$	$\downarrow \downarrow$	_
	TIMP3	$\uparrow \uparrow$	_	$\downarrow \downarrow$	_	$\uparrow \uparrow$	ተተ
	CLCA2	↓ ↓	_	Λ.	^	↓ ↓	ተተ
Cell adhension	COL6A3	$\downarrow \downarrow$	_	_	_	$\uparrow \uparrow$	_
	MPZL2	$\downarrow \downarrow$	↑	_	V	$\downarrow \downarrow$	_
	VCAN	$\downarrow \downarrow$	_	4	\	$\uparrow \uparrow$	1
	ANK3	↓ ↓	_	\	_	↓ ↓	_
Cytoskeleton	EPB41L3	$\downarrow \downarrow$	V	_	_	$\uparrow\uparrow$	^
	KIF1A	$\uparrow \uparrow$	_	ተተ	_	$\downarrow \downarrow$	_
	SEMA6A	$\uparrow \uparrow$	_	1	_	ተተ	_
	AKR1B1	↓ ↓	_	_	_	个个	个个
	ASTN2	$\uparrow \uparrow$	_	1	↑	$\downarrow \downarrow$	_
	BDH1	$\downarrow \downarrow$	_	_	个个	$\downarrow \downarrow$	_
	BIN1	$\downarrow \downarrow$	↓	4	_	$\uparrow \uparrow$	_
	CD200	↑ ↑	_	_	_	↑ ↑	_
	COPS8	$\downarrow \downarrow$	_	_	_	ተተ	_
	ELOVL6	$\downarrow \downarrow$	_	^	^	$\downarrow \downarrow$	_
	ETV5	$\downarrow \downarrow$	↑	_	V	$\downarrow \downarrow$	^
	GABBR2	$\downarrow \downarrow$	V	4	V	$\uparrow\uparrow$	↑ ↑
	GAD2	$\downarrow \downarrow$	_	_	^	$\downarrow \downarrow$	_
	GPR37	$\downarrow \downarrow$	_	_	V	$\downarrow \downarrow$	_
	HPCAL1	1 1	_	↑	_	↑ ↑	_
Others	IRX5	1 1	_	_	_	↓↓	_
	KCNJ13	↑ ↑	_	_	_	$\downarrow\downarrow$	_
	KIAA1045	↑↑ ↑↑	_	_	\	↓ ↓	_
	KLK11	↓ ↓	_	_	_	↓ ↓	_
	LIMD1	1 1	_	→	_	$\downarrow \downarrow$	_
	NAGA	1 1 1	_	_	<u>−</u>	** * *	_
	NAUA	$\Psi\Psi$	_	_	T	TT	_

İ	NES	$\downarrow \downarrow$	_	_	_	$\uparrow \uparrow$	V
	OLFM1	$\uparrow \uparrow$	_	_	↑	$\uparrow\uparrow$	_
	RAMP2	↑ ↑	_	_	$\uparrow\uparrow$	$\uparrow\uparrow$	_
	RAMP3	↑ ↑	4	4	_	$\uparrow\uparrow$	_
	REEP1	↑ ↑	_	4	_	$\uparrow\uparrow$	_
	RTEL1	$\downarrow \downarrow$	_	_	V	$\uparrow\uparrow$	_
	SEPP1	↑ ↑	_	4	_	$\uparrow\uparrow$	ተተ
	SEP6	↑ ↑	_	_	_	$\uparrow\uparrow$	_
	SLC15A1	$\downarrow \downarrow$	_	_	^	$\downarrow \downarrow$	_
	STOM	↑ ↑	_	$\uparrow\uparrow$	^	$\uparrow\uparrow$	ተተ
	TMEM140	↑ ↑	ተተ	_	$\uparrow \uparrow$	$\uparrow \uparrow$	ተተ
	ZNF185	$\downarrow \downarrow$	_	_	_	$\downarrow \downarrow$	4

↑↑: log₂ (Fold change) > 1;

↑: $0.5 \le \log_2$ (Fold change) ≤ 1 ;

-: No significant change;

↓↓: log₂ (Fold change) < -1;</pre>

 \downarrow : -1 ≤ log₂ (Fold change) ≤ -0.5.

Table S4. Common DEGs in KSHV-HMVEC vs. HMVEC and KS signature

Functio	on Cluster	Gene symbol	KSHV- HDMEC vs. HDMEC	KSHV- HMVEC vs. HMVEC	KSHV- LEC vs. LEC	KSHV- BEC vs. BEC	KS vs. skin	KSHV- MSC vs. MSC
		COL15A1	↑ ↑	↑ ↑	_	↑	↑ ↑	_
F 1 - 4112 - 1		GJA4	1	$\downarrow \downarrow$	$\downarrow \downarrow$	_	$\uparrow\uparrow$	_
	-mesenchymal nsition	HEY1	↑	$\downarrow \downarrow$	_	$\uparrow \uparrow$	$\uparrow\uparrow$	_
u ai	iisitioii	NRP1	_	^	$\uparrow \uparrow$	_	$\uparrow\uparrow$	↑
		SEMA3C	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	_	$\downarrow \downarrow$	↑
		ANGPT2	↑ ↑	^	$\uparrow \uparrow$	↑	$\uparrow\uparrow$	↑
		ANGPTL2	_	^	V	$\uparrow \uparrow$	$\uparrow\uparrow$	ተተ
Vessel de	evelopment	LYVE1	↑ ↑	^	_	_	$\uparrow\uparrow$	_
		PGF1	4	^	V	_	$\uparrow\uparrow$	ተተ
		PTPRM	_	↑ ↑			↑ ↑	_
		CXCL10	_	^	_	_	$\uparrow\uparrow$	ተተ
	Chemotaxis	CXCR4	4	^	_	_	$\uparrow\uparrow$	_
	Chemotaxis	GRK5	_	^	_	↑	$\uparrow\uparrow$	↑
		A2M	↑ ↑	↑ ↑	_	个个	↑ ↑	_
	Inflammatio	FN1	_	$\downarrow \downarrow$	$\downarrow \downarrow$	V	$\uparrow\uparrow$	ተተ
	n	IFI16	_	^	^	_	$\uparrow\uparrow$	ተተ
T		TLR2	1	^	_	_	$\uparrow\uparrow$	↑
Immune		GBP1	↑ ↑	↑ ↑	_	1	↑ ↑	ተተ
Response		HLX	^	^	$\uparrow \uparrow$	$\uparrow \uparrow$	$\uparrow\uparrow$	_
	Other genes	HOXB7	_	^	$\uparrow \uparrow$	_	$\uparrow\uparrow$	_
	involved in	IFI35	_	$\uparrow \uparrow$	_	_	$\uparrow\uparrow$	ተተ
	immune	IFITM1	_	^	1	_	$\uparrow\uparrow$	ተተ
	response	ISG15	_	$\uparrow \uparrow$	ተተ	1	$\uparrow\uparrow$	ተተ
		PSMB9	↑	^	↑	↑	$\uparrow \uparrow$	ተተ
		SP110	_	^	_	V	$\uparrow \uparrow$	↑
Cell pr	oliferation	HYAL1	_	↓↓	\	_	1	_
		EGFR	↓↓	↓↓	\	\	4 4	↑
		GUCY1A3	_	$\downarrow \downarrow$	V	_	$\uparrow\uparrow$	_
Signal tr	ansduction	LGALS9	_	↑ ↑	_	_	$\uparrow\uparrow$	ተተ
_	Signal transduction	RASGRP3	↑ ↑	↑ ↑	_	$\uparrow \uparrow$	$\uparrow\uparrow$	↑
		RGS4	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	$\uparrow\uparrow$	↑

	TRIB2	_	$\downarrow \downarrow$	\	_	$\uparrow\uparrow$	_
	NR2F1	_	1 1	\	4 4	↑ ↑	_
Metabolism	PCSK6	↑	↑ ↑	_	$\uparrow \uparrow$	$\uparrow\uparrow$	_
	UBE2L6	_	↑ ↑	V	_	$\uparrow \uparrow$	ተተ
Cell adhension	PCDH12	_	↑ ↑	_	_	个个	_
Cen adnersion	PCDH17	↑	↑ ↑	V	_	$\uparrow \uparrow$	$\uparrow \uparrow$
Cytoskeleton	DIAPH2	_	↑ ↑	个个	↑	个个	_
	CA2	4	↑ ↑	↑ ↑	↑ ↑	↓ ↓	_
	HSPA4L	_	↑ ↑	_	_	$\downarrow \downarrow$	$\uparrow \uparrow$
	LXN	_	↑ ↑	_	_	$\uparrow\uparrow$	4
Others	PPAP2C	_	$\downarrow \downarrow$	_	↑	$\downarrow \downarrow$	_
	TACSTD2	_	$\downarrow \downarrow$	1	↑	$\downarrow \downarrow$	_
	TM4SF1	_	↑ ↑	_	↑	ተተ	$\uparrow \uparrow$
	ZNF117	_	↑ ↑	_	_	$\downarrow \downarrow$	_

↑↑: log₂ (Fold change) > 1;

↑: 0.5 ≤ log₂ (Fold change) ≤1;

-: No significant change;

↓↓: log₂ (Fold change) < -1;</pre>

↓: $-1 \le \log_2$ (Fold change) ≤ -0.5 .

Function Cluster	Gene symbol	KSHV- HDMEC vs. HDMEC	KSHV- HMVEC vs. HMVEC	KSHV- LEC vs. LEC	KSHV- BEC vs. BEC	KS vs. skin	KSHV- MSC vs MSC
	GJA4	↑	4 4	44	_	↑ ↑	_
Endothelial-mesenchymal	NRP1	_	$\uparrow \uparrow$	ተተ	_	$\uparrow \uparrow$	↑
transition	SEMA3C	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	_	$\downarrow \downarrow$	↑
	TGFB1I1	_	_	$\downarrow \downarrow$	V	ተተ	↓
	ADORA2B	$\downarrow \downarrow$	_	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	_
Vessel development	ANGPT2	↑ ↑	ተተ	ተተ	↑	$\uparrow \uparrow$	↑
	LOC100131909	_	_	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	_
	AZGP1	_	_	个个	↑	↓ ↓	_
	AZGP1P1	_	_	$\uparrow \uparrow$	^	$\downarrow \downarrow$	_
	CBLB	$\uparrow \uparrow$	_	$\uparrow \uparrow$	ተተ	$\uparrow \uparrow$	_
	GJA4 NRP1 SEMA3C TGFB111 ADORA2B ANGPT2 LOC100131909 AZGP1 AZGP1P1 CBLB CD47 GCH1 HLX HOXB7 ISG15 TIMP1 CCND1 DERL2 SSR1 ARL4C ATP6V0E1 DGKD ECE1 FLNA IQGAP2 JAK1 RGS4 B3GAT3 CST3	↑	_	ተተ	_	$\uparrow \uparrow$	1
Immune response	GCH1	_	^	ተተ	_	$\uparrow \uparrow$	$\uparrow \uparrow$
_	HLX	^	ተተ	ተተ	ተተ	$\uparrow \uparrow$	_
	HOXB7	_	ተተ	ተተ	_	skin	_
	ISG15	_	ተተ	ተተ	↑	$\uparrow \uparrow$	$\uparrow \uparrow$
	TIMP1	_	_	ተተ	ተተ	skin ↑↑ ↓↓ ↑↑ ↓↓ ↑↑	$\uparrow \uparrow$
	CCND1	_	_	ተተ	ተተ	44	ተ ተ
Cell proliferation	DERL2	_	_	ተተ	_	$\uparrow \uparrow$	_
•	SSR1	_	_	ተተ	_	skin ↑↑ ↓↓ ↑↑ ↓↓ ↑↑ ↑ </td <td>_</td>	_
	ARL4C	4	\	44	4	skin ↑↑ ↓↑↑ ↓↓↑ ↑↑↑ ↑↑↑ ↑↑↑ ↑↑↑ ↑↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑ </td <td>_</td>	_
		_	_	$\uparrow \uparrow$	^	$\uparrow \uparrow$	↑
	DGKD	_	_	$\uparrow \uparrow$	^	$\uparrow \uparrow$	_
C:1	ECE1	_	_	$\uparrow \uparrow$	_	$\uparrow \uparrow$	↑
Signal transduction	FLNA	_	_	$\uparrow \uparrow$	V	$\uparrow \uparrow$	V
	IQGAP2	_	_	$\downarrow \downarrow$	_	$\uparrow \uparrow$	_
	JAK1	_	^	$\uparrow \uparrow$	^	$\uparrow \uparrow$	_
	RGS4	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	skin ↑↑ ↓↑ ↓↓ ↑↑ ↓↓ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑↑ ↑	1
	B3GAT3	_	_	个个	_	ተተ	↑
	CST3	_	_	$\uparrow \uparrow$	^	$\uparrow \uparrow$	↑
Metabolism	IMPA2	_	_	$\downarrow \downarrow$	_	$\downarrow \downarrow$	↑
	ITGB3	↑	^	$\uparrow \uparrow$	^	$\uparrow \uparrow$	_
	UBE2J1	_	_	ተተ	ተተ	ተተ	_

	LOXL2	_	_	↑ ↑	↓ ↓	ተተ	\
Cell adhension	NID1	1	_	$\downarrow \downarrow$	_	ተተ	_
	PODXL	V	↑	ተተ	$\uparrow \uparrow$	↑ ↑	ተተ
Cytoskeleton	DIAPH2	_	ተተ	↑ ↑	↑	↑ ↑	_
Cytoskeleton	KIF1A	↑ ↑	_	↑ ↑	_	$\downarrow \downarrow$	_
	C21ORF7	_	_	↑ ↑	^	↑ ↑	_
	CA2	V	ተተ	↑ ↑	$\uparrow \uparrow$	$\downarrow \downarrow$	_
	CNNM2	_	_	↑ ↑	_	$\downarrow \downarrow$	↑
	GOLM1	_	_	$\uparrow \uparrow$	_	$\uparrow\uparrow$	ተተ
	GPR116	_	_	$\uparrow \uparrow$	↑	$\uparrow\uparrow$	_
	HEG1	_	_	$\downarrow \downarrow$	↑	↑ ↑	_
	IGSF3	_	_	$\downarrow \downarrow$	↑	$\downarrow \downarrow$	_
	KCNIP1	_	_	$\downarrow \downarrow$	_	$\downarrow \downarrow$	_
	KIAA1462	_	_	$\downarrow \downarrow$	_	ተተ	↑
	LOC283398	_	_	$\uparrow \uparrow$	_	$\downarrow \downarrow$	_
04	NCR2	_	_	$\uparrow \uparrow$	↓	$\downarrow \downarrow$	_
Others	PLLP	_	↓	$\downarrow \downarrow$	_	$\downarrow \downarrow$	_
	PLVAP	_	_	$\uparrow \uparrow$	^	ተተ	_
	RNF115	_	_	$\uparrow \uparrow$	^	ተተ	_
	SLC39A14	_	_	$\uparrow \uparrow$	_	ተተ	_
	SNX10	_	_	$\downarrow \downarrow$	_	ተተ	_
	STOM	↑ ↑	_	↑ ↑	↑	ተተ	↑ ↑
	SUCLG2	4	_	$\uparrow \uparrow$	_	$\downarrow \downarrow$	_
	TBCD	_	_	↑ ↑	_	ተተ	_
	TFPI	_	\	$\downarrow \downarrow$	↓	ተተ	↑
	THSD7A	↑	_	$\downarrow \downarrow$	_	ተተ	_
	XPNPEP1	_	_	$\downarrow \downarrow$	_	ተተ	_

↑↑: log₂ (Fold change) > 1;

 \uparrow : 0.5 ≤ log₂ (Fold change) ≤1;

-: No significant change;

↓↓: log₂ (Fold change) < -1;</pre>

Function Cluster		Gene symbol	KSHV- HDMEC vs. HDMEC	KSHV- HMVEC vs. HMVEC	KSHV- LEC vs. LEC	KSHV- BEC vs. BEC	KS vs. skin	KSHV MSC v MSC
Vessel development		ADORA2B	44	_	↓ ↓	↓ ↓	44	_
		ANGPTL2	_	↑	↓	$\uparrow \uparrow$	ተተ	ተተ
		FGFR2	$\downarrow \downarrow$	_	^	$\uparrow \uparrow$	$\downarrow \downarrow$	_
		HEY1	↑	$\downarrow \downarrow$	_	$\uparrow\uparrow$	ተተ	_
		LOC100131909	_	_	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	_
		VCAM1	_	_	_	$\downarrow \downarrow$	$\uparrow\uparrow$	$\uparrow \uparrow$
Immune Response	Chemotaxis	CCL5	_	_	↑	个个	个个	个个
		CYBA	_	_	_	$\uparrow \uparrow$	$\uparrow \uparrow$	1
	Inflammation	A2M	↑ ↑	↑ ↑	_	个个	ተተ	_
		TNFRSF4	_	4	↓	ተተ	ተተ	_
	Other genes	CBLB	↑ ↑	_	_	个个	ተተ	_
		GBP1	↑ ↑	$\uparrow \uparrow$	_	$\downarrow \downarrow$	ተተ	ተተ
		HLX	↑ ↑	$\uparrow \uparrow$	$\uparrow \uparrow$	↑ ↑	ተተ	_
	involved in	ISG20	_	_	_	$\downarrow \downarrow$	$\uparrow \uparrow$	ተተ
	immune	PTGER4	$\downarrow \downarrow$	_	_	$\downarrow \downarrow$	$\uparrow \uparrow$	_
	response	TACC3	V	V	_	$\downarrow \downarrow$	$\uparrow \uparrow$	$\downarrow \downarrow$
	•	TAPBP	_	_	_	$\uparrow\uparrow$	ተተ	1
		TIMP1	_	_	^	ተተ	ተተ	ተተ
Cell nr	oliferation	CCND1	_	_	↑ ↑	个个	↓ ↓	ተተ
Cen promeration		APP	_	_		个个	ተተ	1
		CCDC88A	V	_	_	$\uparrow\uparrow$	ተተ	_
Signal transduction		HPGD	_	_	_	$\downarrow \downarrow$	$\downarrow \downarrow$	_
		RASGRP3	↑ ↑	$\uparrow \uparrow$	_	$\uparrow \uparrow$	ተተ	↑
		RGS4	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	$\downarrow \downarrow$	ተተ	1
		SIK1	_	_	^	$\downarrow \downarrow$	$\downarrow \downarrow$	_
		SPRY1	↑	1	↓	$\uparrow\uparrow$	ተተ	1
		BTRC	↑ ↑	_	_	个个	↓ ↓	_
Metabolism		MAOA	$\uparrow \uparrow$	V	^	$\downarrow \downarrow$	$\downarrow \downarrow$	_
		MEF2C	_	1	_	$\uparrow\uparrow$	$\uparrow\uparrow$	_
		MGAT2	4	_	_	$\uparrow \uparrow$	$\uparrow \uparrow$	_
		NR2F1	_	$\downarrow \downarrow$	4	$\downarrow \downarrow$	ተተ	_

	PBXIP1	_	_	\	$\uparrow \uparrow$	↑ ↑	$\uparrow \uparrow$
	PCSK6	1	↑ ↑	_	$\uparrow \uparrow$	$\uparrow \uparrow$	_
	PTPRN2	_	_	\	$\uparrow\uparrow$	$\uparrow \uparrow$	↑
	UBE2J1	_	_	↑ ↑	$\uparrow \uparrow$	ተተ	_
	COL6A1	_	↑	1	个个	个个	↑
	LOXL2	_	_	↑ ↑	$\downarrow \downarrow$	ተተ	V
	PARVB	_	_	4	$\downarrow \downarrow$	$\uparrow \uparrow$	V
	PCDHGA12	_	_	_	$\uparrow \uparrow$	ተተ	_
Cell adhension	PCDHGC3	_	_	_	$\uparrow \uparrow$	ተተ	_
	PCDHGC4	_	_	_	$\uparrow\uparrow$	ተተ	_
	PCDHGC5	_	_	_	$\uparrow\uparrow$	ተተ	↑
	PODXL	V	↑	↑ ↑	$\uparrow\uparrow$	ተተ	$\uparrow \uparrow$
	ZYX	_	V	_	$\downarrow \downarrow$	ተተ	_
	ALDH3A1	_	_	1	个个	44	个个
	ASPA	_	_	_	$\uparrow \uparrow$	$\downarrow \downarrow$	^
	BDH1	$\downarrow \downarrow$	_	_	$\uparrow\uparrow$	$\downarrow \downarrow$	_
	C21ORF7	_	_	↑ ↑	$\uparrow\uparrow$	ተተ	_
	CA2	V	↑ ↑	↑ ↑	$\uparrow\uparrow$	$\downarrow \downarrow$	_
	CYB5R2	_	_	_	$\uparrow \uparrow$	$\downarrow \downarrow$	_
	CYP3A43	_	_	↑	$\downarrow \downarrow$	$\downarrow \downarrow$	_
	GOLGA8A	_	_	_	$\uparrow\uparrow$	$\downarrow \downarrow$	↑
	GOLGA8B	_	_	_	$\uparrow\uparrow$	$\downarrow \downarrow$	_
	GPR126	_	V	_	$\downarrow \downarrow$	$\downarrow \downarrow$	_
	HIP1R	_	_	_	$\downarrow \downarrow$	$\downarrow \downarrow$	_
Others	LZTS1	_	_	↑	$\uparrow \uparrow$	$\downarrow \downarrow$	_
	METRN	V	_	_	$\uparrow\uparrow$	ተተ	↑
	MTERF	_	_	_	$\downarrow \downarrow$	$\downarrow \downarrow$	_
	NOL3	_	_	↑	$\uparrow \uparrow$	$\downarrow \downarrow$	_
	PLVAP	_	_	ተተ	$\uparrow \uparrow$	ተተ	_
	RAMP2	↑ ↑	_	_	$\uparrow \uparrow$	ተተ	_
	RTN1	_	_	_	$\uparrow\uparrow$	ተተ	_
	SDPR	_	_	_	$\downarrow \downarrow$	ተተ	^
	SEP6	↑ ↑	_	_	ተተ	ተተ	_
	SLC38A7	_	_	V	ተተ	ተተ	_
	TLL2	_	_	_	$\downarrow \downarrow$	$\downarrow \downarrow$	_
	TMEM140	_	↑ ↑	_	$\uparrow\uparrow$	$\uparrow \uparrow$	$\uparrow \uparrow$

```
↑↑: log<sub>2</sub> (Fold change) > 1;

↑: 0.5 ≤ log<sub>2</sub> (Fold change) ≤1;

-: No significant change;

↓↓: log<sub>2</sub> (Fold change) < -1;

↓: -1 ≤ log<sub>2</sub> (Fold change) ≤ -0.5.
```