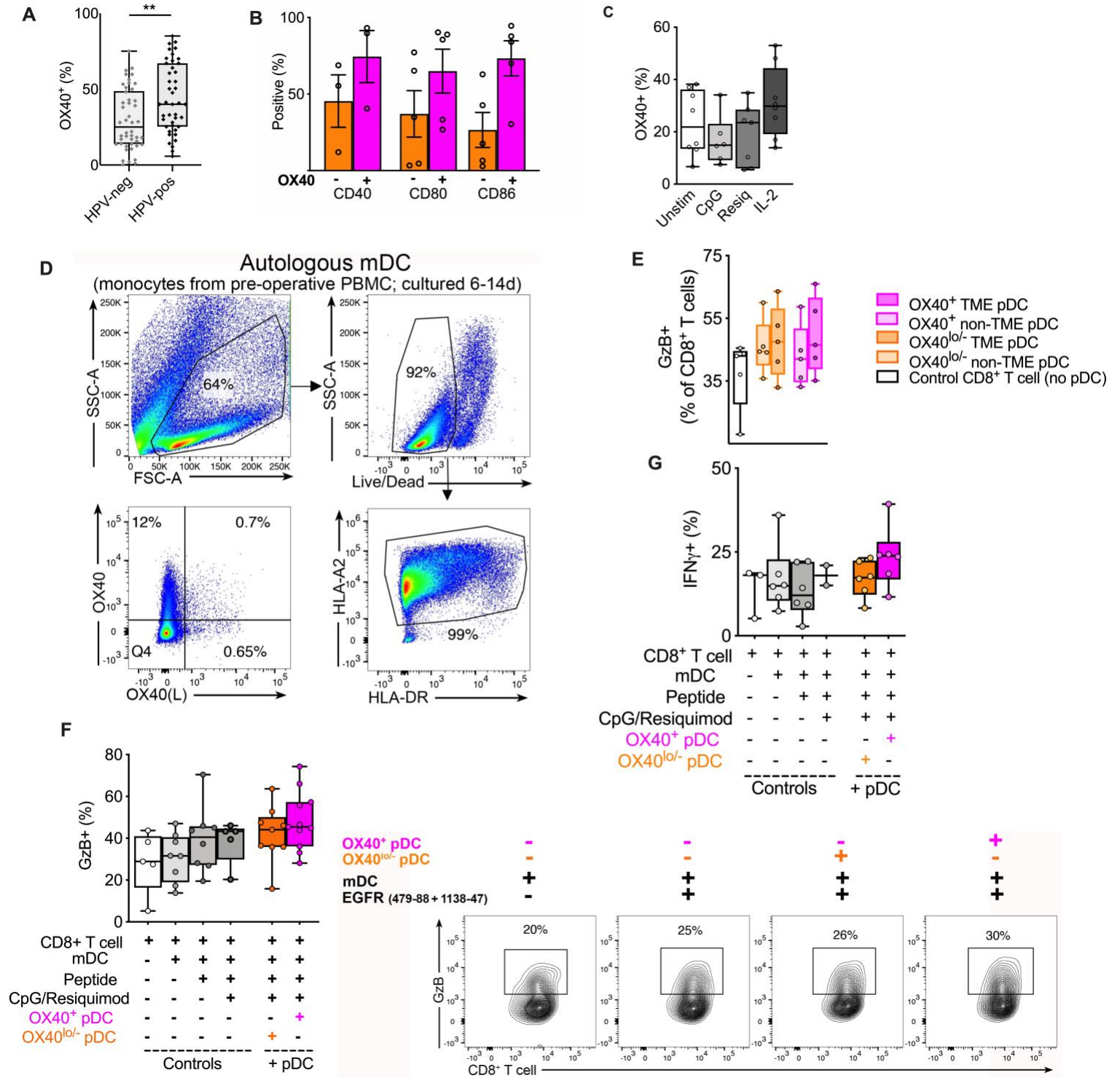


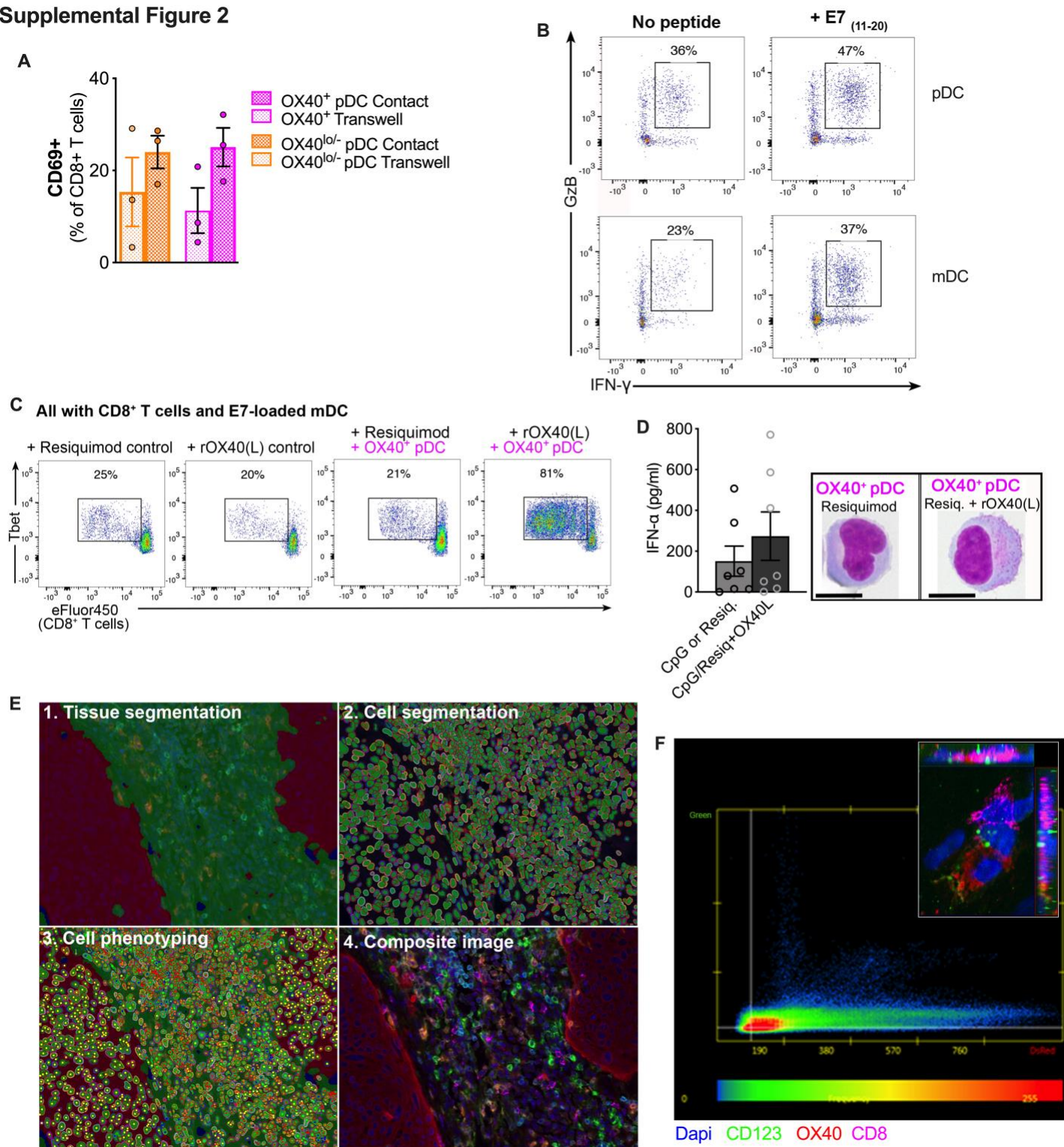
Supplemental Figure 1



OX40-status defines a mature pDC population that drives antigen-specific autologous CD8⁺ T cell responses. (a) OX40 expression on intratumoral pDC from HPV-positive (n= 43) and HPV-negative (n= 46) HNSCC patients. (b) Sorted pDC from dLN_{neg} from HNSCC patients (n= 5) were incubated overnight and measured by flow cytometry for expression of CD40, CD80 and CD86 on pDC subsets. (c) Expression of OX40 measured on whole pDC from dLN_{neg} from HNSCC patients (n=8) after overnight incubation in culture media alone (unstimulated) or in presence of IL-2, CpG or Resiquimod. (d) FACS gating strategy for sorting harvested autologous HLA-DR⁺HLA-A2⁺ monocyte-derived dendritic cells (mDC) for use in autologous antigen-specific co-culture assays (Methods). OX40 and OX40(L) expression levels were also measured. (e) CD8⁺ T cell positivity for granzyme B (GzB) after co-culture with peptide-loaded mDC without pDC (control) or with

OX40⁺ or OX40^{lo/-} pDC from the TME vs. non-TME (dLN_{neg}) (n= 5). **(f-g)** Percentage of CD8⁺ T cells positive for GzB and IFN- γ after co-culture with tumor-associated antigen peptide-loaded mDC and with or without autologous OX40⁺ or OX40^{lo/-} pDC from tumor or dLN_{neg} from HNSCC patients (n= 11). One-way ANOVA followed by Tukey's post hoc test (b-c, e-g). Two-tailed paired T test (a). Bar graph data are mean \pm s.e.m. Middle line of box-and-whisker plot indicates the median, box limits indicate the first and third quartiles, and 'whiskers' indicate 'extreme' for all data points (a, c, e-f). *****P* < 0.01**. Representative flow plots are shown (f).

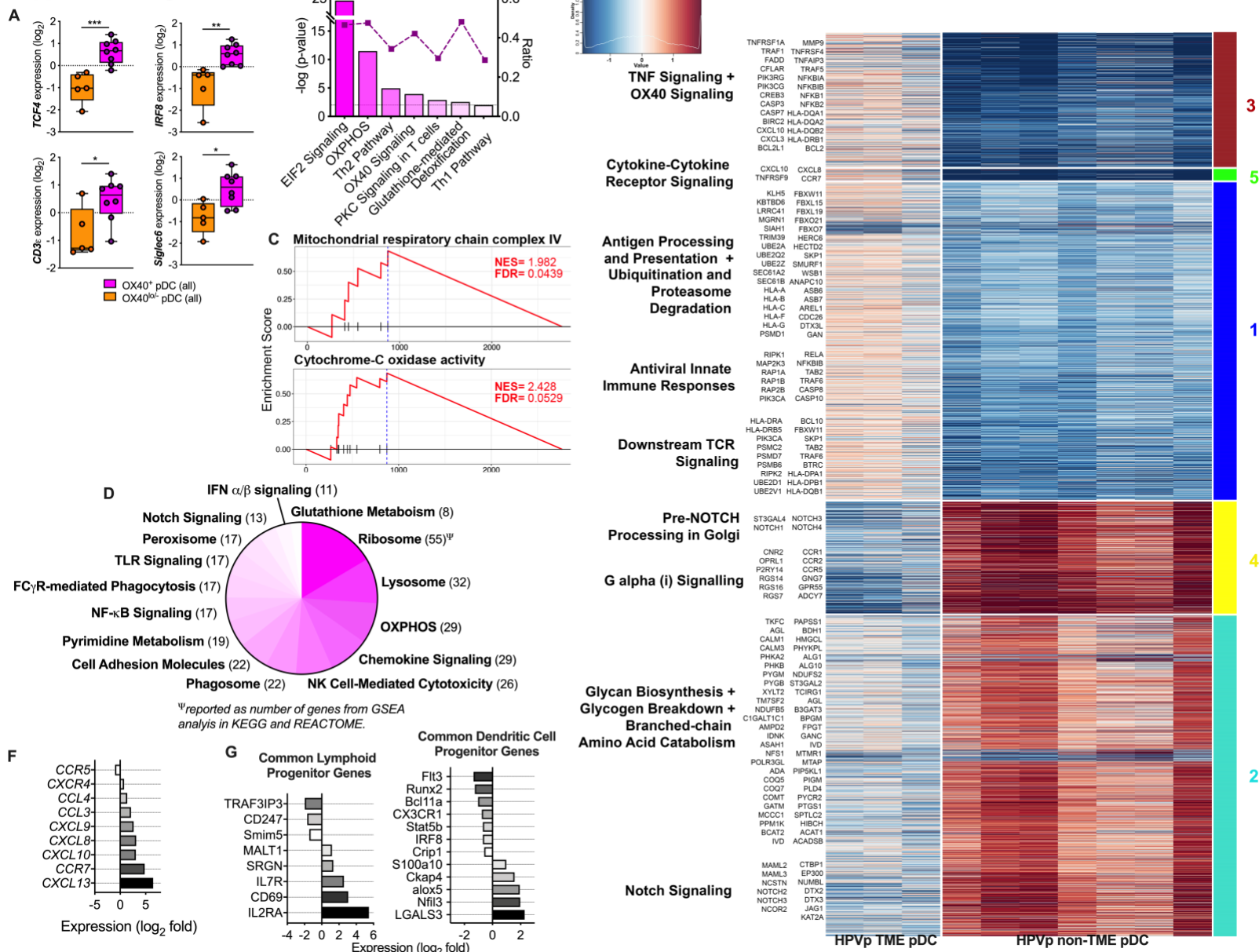
Supplemental Figure 2



The effect of OX40-ligation on pDC morphology, IFN- α secretion and immunostimulatory capacity. (a) Percentage of activated (CD69⁺) CD8⁺ T cells in transwell vs. contact co-culture with pDC subsets (n=3). (b) The production of granzyme B (GzB) and IFN- γ by E7-specific CD8⁺ T cells co-cultured with autologous pDC and mDC without or with prior loading with HPV E7 peptide. (c) Tbet expression on proliferating antigen-specific (eFluor450-low) CD8⁺ T cells from autologous co-culture with peptide-loaded monocyte-derived dendritic cells (mDC) and with OX40⁺ pDC pre-stimulated with either Resiquimod or rOX40(L). (d) The concentration (pg/ml) of IFN- α in the supernatant from OX40⁺ pDC from HNSCC patients (n=7) stimulated with either CpG or Resiquimod or combination treatment of either CpG or Resiquimod with recombinant OX40(L) [rOX40(L)]. Data normalized to 2×10^3 pDC per sample. May-Grunwald staining of FACS-sorted OX40⁺ pDC from the TME of a patient, stimulated with either Resiquimod alone or Resiquimod with

rOX40(L). n= 3; three experimental repeats. 100× magnification, scale bar, 5 μm. (e) Example of image analysis workflow in inForm 2.1 (Akoya Biosciences) software. Following spectral unmixing, single images were segmented according to tissue type (i.e. tumor and stroma) followed by cell segmentation based on a proprietary inForm active learning phenotyping algorithm. Cells were then phenotyped in inForm into one of three different types according to the markers of interest: tumor cells (panCK⁺), CD8⁺ T cells (CD3⁺CD8⁺), pDC (CD3⁺CD123⁺), macrophages (CD68⁺) and 'other cells' (panCK⁻CD68⁻CD8⁻CD123⁻). (f) Colocalization of OX40 (red) with CD123 (green) signals shown as a two-dimensional scatter plot of signal intensities (red on X-axis, green on Y-axis) from a CD123⁺ cell adjacent to a CD8⁺ cell (inset: three-dimensional x-y-z plot), from the tissue section of the TME from a HNSCC patient. One-way ANOVA followed by Tukey's post hoc test (a) and Two-tailed paired T-test (d). Bar graph data are mean ± s.e.m

Supplemental Figure 3



Transcriptomes of pDC in HNSCC based on TME- and OX40-status.

a-g: Bulk RNA sequencing was performed on cell-sorted pDC, including OX40⁺ and OX40^{lo/-} pDC, from the TME and non-TME in HNSCC patients, including analysis restricted to HPV-positive (HPVp) patients (n= 4). Differentially expressed genes (DEG) were determined using the criterion of fold-changes (FC) ≥ 1.5 . **(a)** Expression of *TCF4*, *IRF8*, *CD3e* and *Siglec6* in OX40⁺ and OX40^{lo/-} pDC from HNSCC patients (n= 7) (**Supplemental Table 2**). **(b)** Ingenuity pathway analysis (IPA)-generated canonical pathway analysis of top enriched pathways ($-\log(p\text{-value}) > 2.0$) in HPVp OX40⁺ pDC. **(c)** Gene set enrichment analysis (GSEA) plots of oxidative phosphorylation-related gene pathways enriched in HPVp OX40⁺ pDC with normalized enrichment scores (NES) and false discovery rates (FDR). **(d)** Pie chart of the top enriched-pathways ($p < 0.05$) generated by enriched gene set analysis (KEGG and Reactome) of DEG from OX40⁺ and OX40^{lo/-} pDC from all patients (n= 7). **(e)** Heat map reporting relative expression (\log_2 values) in rows of DEG (**Supplemental Table 4**) in pDC across primary tumors from HPVp patients, labeled as HPVp TME (n= 3) and matched non-TME tissue (dLN_{neg}/PBMC, n= 7 including some patient duplicates). Heat map color scheme is based on \log_2 FC expression levels from -2 (blue) to 2 (red). Color bars in right margin highlight five main gene clusters, with their corresponding labels 1-5, according to their gene expression states. Labeling in left margin highlights annotation of clusters with notable pathways (KEGG). **(f)** Expression of chemokine receptor/ligand genes

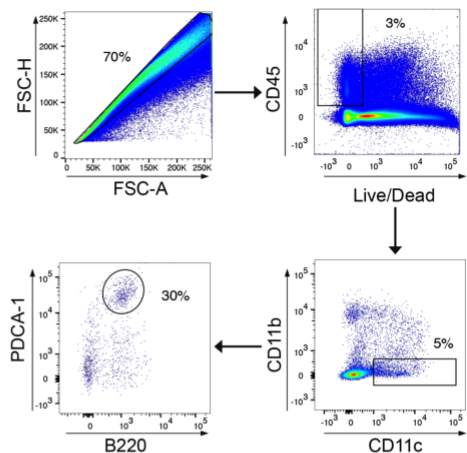
upregulated in TME pDC from HNSCC patients. (g) Expression of common lymphoid progenitor and common dendritic cell progenitor lineage genes upregulated in TME pDC.

Unpaired Student's T test (a). Middle line of box-and-whisker plot indicates the median, box limits indicate the first and third quartiles, and 'whiskers' indicate 'extreme' for all data points. IPA ratios/Z-scores in Source Data 2.

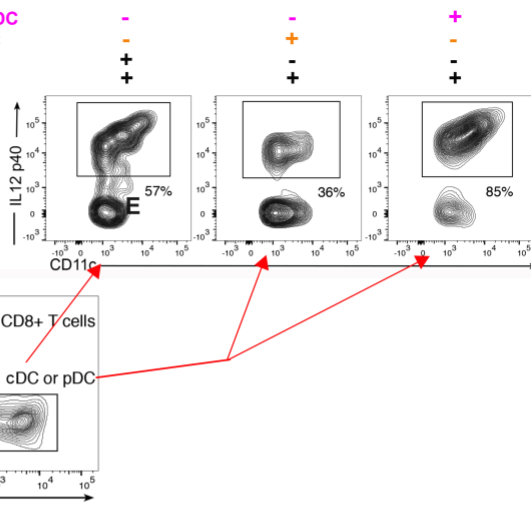
OX40+ pDC correlate to survival and slower tumor growth.

Supplemental Figure 4

A Mouse pDC gating strategy

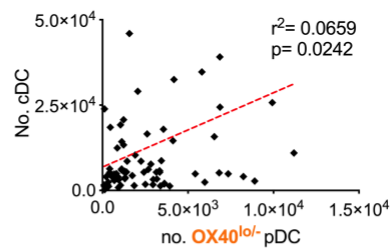
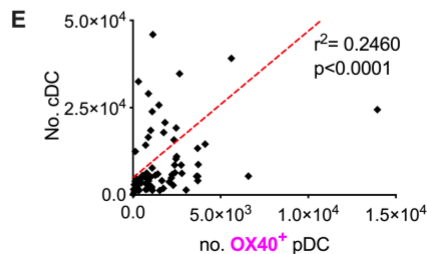
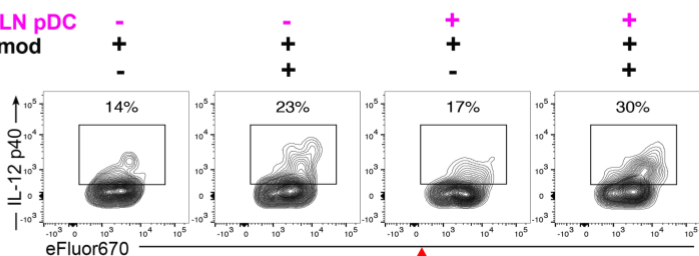


B B16CCR7 dLN pDC B16F10 dLN pDC mDC (WT) gp100

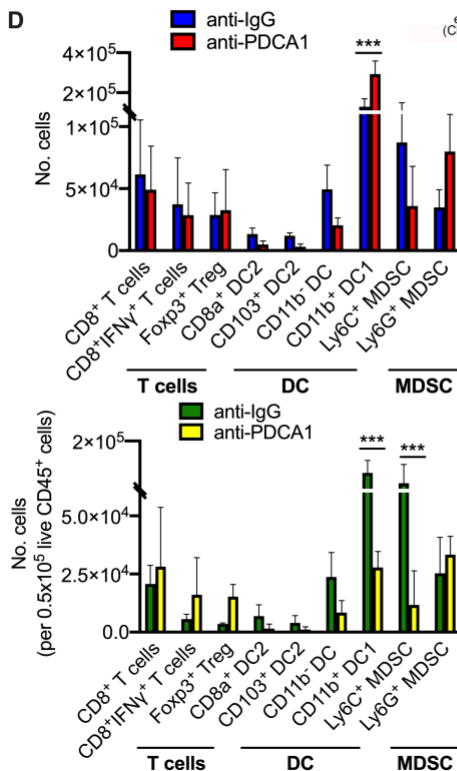


C All with CD8+ T cell and gp100-loaded

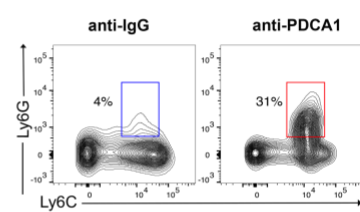
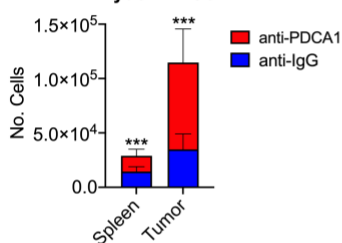
B16CCR7 dLN pDC Resiquimod OX86



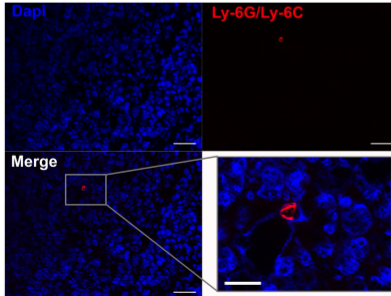
D



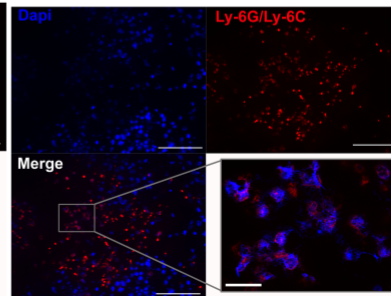
F Ly6G+ MDSC



G B16-CCR7 + anti-IgG



B16-CCR7 + anti-PDCA1



(a) Murine tumor pDC gating strategy. Live singlets were first selected using FSC-H and FSC-A and CD45⁺ live/dead viability stain. pDC were then gated from CD11^{cm}CD11^{bneg} cells as PDCA1⁺B220⁺. **(b)** Flow plots of IL-12 p40 production by either cDC (from WT B6 mice) and pDC (from dLN of B16-F10- or B16CCR7-bearing mice) co-cultured with Pmel-1 CD8⁺ T cells. n= 2; two experimental repeats. **(c)** Flow plots of IL-12 p40 production by cDC when co-cultured with Pmel-1 CD8⁺ T cells and pDC from B16CCR7-bearing mice pre-stimulated with Resiquimod and OX86. **(d)** Number of different immune cell populations in TME of either B16-F10- or B16CCR7-bearing mice treated with either anti-PDCA1 or anti-polyclonal IgG. Data are pooled from individual experiments and normalized to 5×10⁵ live cells. **(e)** Correlation (Pearson) of cDC and either OX40⁺ or OX40^{lo}- pDC numbers (all normalized to 1×10⁶ live cells) in the TME of HNSCC patients (n= 77). A line of best fit is shown. **(f)** Quantification of granulocytic myeloid-derived suppressor cells (MDSC) (CD11b⁺Ly6G⁺Ly6C⁺) in the tumor and spleen from B16CCR7-bearing mice treated with either pDC-depleting antibody (anti-PDCA1) or anti-polyclonal IgG (control). Corresponding flow plots show Ly6G⁺ (granulocytic) and Ly6C⁺ (monocytic) MDSC in pDC-depleted (red gate) and control (blue gate) mice. **(g)** Immunofluorescence microscopy of Ly-6G/Ly-6C MDSC in B16CCR7-bearing mice treated with either anti-polyclonal IgG antibody (control) or anti-PDCA1 pDC depleting antibody. Top (anti-IgG): 10×, scale bar 50 μm and 40×, scale bar 20 μm (inset); Bottom (anti-PDCA1): 20×, scale bar 50 μm and 40×, scale bar 20 μm (inset). Red, Ly-6G/Ly-6C; blue, Dapi. One-way ANOVA followed by Tukey's post hoc test (d, f). Bar graph data are mean ± s.e.m. ****P* < 0.001.

Patient	Age/ Sex	TNM* Stage	HPV	1° Site	1° Site Collected	LN Collected	PBMC Collected	Monocyte Expansion†	Flow Cytometric Phenotyping	FACS sort	ELISA/ Luminex	Co- culture	RNA -seq	IF	IHC‡
NMH67	66/M	T4N0	-	Larynx	Y	L neck contents	N		×	×		×			
NMH68	36/F	T2N2	-	Oral tongue	Y	N/A	Y	×		×	×	×			
NMH69	56/M	T2N0	-	Oral tongue	Y	N/A	Y			×		×	×		
NMH70	54/M	T3N1	-	Oral tongue	Y	R level 2A	Y			×	×	×		×	
NMH71	70/M	T3N2	-	BOT	Y	N/A	Y		×	×	×	×			
NMH72	83/F	T4N0	-	FOM	Y	N/A	Y			×	×				
NMH73	56/M	T2N2	-	Oral tongue	Y	L level 2	Y		×	×					
NMH74	35/F	T2N0	-	Oral tongue	N	N/A	Y			×	×				
NMH75	57/M	T2N0	-	FOM	Y	N/A	Y	×		×		×			
NMH76	67/M	T3N0	-	Oral tongue	Y	R neck contents	Y	×		×		×			
NMH77	56/M	T4N3	-	Larynx	Y	L level 2A L level 3	Y	×	×	×	×	×			
NMH78	29/F	T1N3	-	Oral tongue	N	R level 2A,3	Y	×		×		×			
NMH79	49/F	T3N1	-	Oral tongue	Y	L level 4	N		×	×					
NMH80	63/M	T4N0	-	Maxilla	Y	N/A	Y		×	×					
NMH81	65/M	T4N0	-	Larynx	Y	R level 4	Y		×	×			×	×	
NMH82	50/M	T3N2	-	Oral tongue	Y	R level 4	N		×	×					
NMH83	58/F	T2N3	-	Oral tongue	N	R level 2A,3	N		×	×	×				
NMH84	73/M	T1N1	-	FOM	N	L level 1B	N			×	×				
NMH85	84/M	T3N0	-	Oral tongue	Y	N/A	Y		×	×					
NMH86	60/M	T4N3	-	Oral tongue	N	L level 3 R level 4	N		×	×					
NMH87	57/M	T4N1	-	Hypopharynx	N	R level 3,4	N		×	×					
NMH88	59/M	T4N3	-	Mandible	N	R level 3 L level 3			×	×	×				
NMH89	77/M	T1N0	-	Oral tongue	Y	N/A	Y			×		×			
NMH90	72/M	T1N1	-	Mandible	N	R level 1b			×	×					
NMH91	69/M	T4N0	-	Larynx	Y	R level 2	N		×	×					
NMH92	77/M	T1N0	-	Oral mucosa	Y	N/A	Y		×	×					
NMH93	25/M	T4N3	-	Oral tongue	N	L level 2A R level 3			×	×					
NMH94	59/M	T4N3	-	Lip	N	L level 1b R level 1b	Y		×	×	×				
NMH95	77/M	T2N0	-	Scalp	N	Left neck contents	N		×						
NMH96	65/M	T3N0	-	Oral tongue	Y	Left neck contents	N		×	×					
NMH97	47/M	T2N2	-	BOT	Y	R level 2A,4	Y		×		×				
NMH98	78/M	T1N0	-	Mandible	Y	N/A	Y		×		×				
NMH99	58/M	T3N3	-	Larynx	N	N/A	Y		×	×	×				
NMH100	56/M	T4N2	-	Mandible	N	R level 3	Y		×	×	×		×		
NMH101	76/M	T4N2	-	Larynx	N	R level 2A,3	N		×						
NMH102	62/M	T3N0	-	Larynx	Y	L level 3	N		×						

*Tumor-node-metastasis (TNM) staging according to the AJCC Staging Manual (Eighth edition); †Monocytes obtained from patient pre-operative blood and differentiated into monocyte-derived dendritic cell (mDC) (Methods); ‡Seven-plex immunohistochemistry (IHC) using the Vectra Polaris platform (Akoya Biosciences). §Primary tumor cannot be staged (unknown primary). HPV = Human Papillomavirus; 1° = primary tumor; LN = lymph node; PBMC = peripheral blood mononuclear cells; IF = immunofluorescence; M = male; F = female; BOT = base of tongue; FOM = floor of mouth; R = right; L = left; N/A = not applicable.

Supplemental Table 2: HNSCC patient RNA-seq pDC samples

Sample ID	Pt. ID	HPV	Sample details	TME-status	OX40-status*
pDC-1	NMH19	+	dLN _{pos}	TME	High
pDC-2			dLN _{neg}	non-TME	High
pDC-3	NMH61	-	1° [†] (gingiva)	TME	High
pDC-4			PBMC	non-TME	High
pDC-5	NMH69	-	1° (tongue)	TME	Low
pDC-6			dLN _{neg}	non-TME	High
pDC-7			dLN _{neg}	non-TME	N/A
pDC-8	NMH81	-	dLN _{neg}	non-TME	High
pDC-9			dLN _{neg}	non-TME	N/A [‡]
pDC-10	NMH29	+	dLN _{neg}	non-TME	High
pDC-11			dLN _{neg}	non-TME	N/A
pDC-12			dLN _{pos}	TME	Low
pDC-13	NMH27	+	dLN _{neg}	non-TME	High
pDC-14			1° (tonsil)	TME	N/A
pDC-15			1° (tonsil)	TME	Low
pDC-16	NMH31	+	dLN _{neg}	non-TME	N/A
pDC-17			dLN _{neg}	non-TME	Low
pDC-18			1° (tonsil)	TME	Low
pDC-19			PBMC	non-TME	N/A

*Determined at the time of sorting and based on the gating schemata outlined in Figure S1a; †Primary tumor site; ‡Not-applicable because it represents the whole pDC population and not an OX40 subset. dLN_{neg} = draining cervical lymph node, tumor-negative; dLN_{pos} = draining cervical lymph node, tumor-positive; PBMC = peripheral blood mononuclear cells; TME = tumor microenvironment.

Supplemental Table 3: Upregulated differentially expressed genes in OX40+ pDC grouped according to enriched pathways (KEGG). Genes in each category are listed in order of highest expression levels.

TNFR2 NF-κB pathway*	I. Ribosome/ EIF2 Signaling	II. Oxidative Phosphorylation	III. Detoxification of ROS [†]	IV. Lysosome	Phagosome	V. FcγR-mediated Phagocytosis	VI. Type I IFN Signaling	TLR [‡] Pathway	VII. NK Cell Mediated Cytotoxicity
LTBR	RPS6KA4 RPS15A	NDUFB3 ATP5I	ABCD1 IDH2	AP1M1 PSAP	ACTB ACTG1	LYN PLCG2	NUP93 USP18	TOLLIP	ITGAL TYROBP
LTB	MRPL41 RPL38	ATP5E COX17	PRDX5 HSD17B4	AP1S1 NAPSA	RILP CORO1A	PIK3CD MAPK3	PIN1 NLRC5	LY96	ITGB2 PTPN6
PSME1	RPL19 MRPL11	NDUFB7 ATP5H	MPV17 ACSL5	ARSB CTSW	NCF2 CD36	ARPC4 INPP5D	UBE2E1 OAS1	IRAK1	CD48 NCR2
PSMB10	RPL15 RPS25	CYC1 TCIRG1	CRAT CYBA	LGMM DNASE2	NCF1 TAP2	PLPP1 SYK	UBE2L6 BST2	MAPK9	PTK2B HCST
PSMB8	RPS3 RPL30	COX7C NDUFA4	PMVK GSTP1	PPT1 GNS	NCF4 SCARB1	PRKCD VAMP8	UBA7 IFI35	MAP2K6	PPP3CB FYN
PSMB9	MRPL36 RPS27	ATP5G2 ATP5J2	ACOX3 GPX7	ASAH1 NPC1		ARPC1B NIPSNAP1		TAB1	FCER1G SH3BP2
TNFSF12	RPLP1 RPS28	COX7A2L NDUFA3	GSTK1 TXN2	TPP1 NPC2		RAC2 TRAP1		IRF5	NFATC2 LCP2
TNFSF13B	RPS14 RPL32	ATP5G1 COX8A	CAT GSS	AP1S2 ARSA		GSN FLOT1			SELL
DRAP1	RPL11 RPL7	COX5B NDUFA1	MGST2	NAGA SUMF1					
	RPL12 RPS29	NDUFB1 NDUFV1		MAN2B1 CTSH					
	RPS27A RPL9	NDUFB2 ATP5C1		AGA GGA2					
	MRPL4 RPL8	NDUFS6 IDH3G							
	RPS18 RPL10A	NDUFS4 IDH3A							
	MRPS9 RPS21	NDUFS8 ECH1							
	RPS15 RPS23	COX6B1 MDH2							
	RPS4Y1 RPS24								
	RPS11 RPL27								
	UBA52 MRPS21								
	RPL35 RPS8								
	RPL36 RPS7								
	RPL37 RPL29								
	RPL27A MRPL24								
	MRPL23 RPL23								
	RPL21								

*TNFR2 non-canonical NF-κB pathway/TNFSF members mediating non-canonical NF-κB pathway; †Detoxification of ROS/Peroxisome/Glutathione metabolism; ‡TLR signaling pathway. ROS = reactive oxygen species; IFN = interferon; TLR = Toll-like receptor; NK = natural killer.

Supplemental Table 4: 4,197 differentially-expressed genes (2,181 upregulated, 2,016 downregulated) for HPV non-TME compared to TME pDC, mapped as Clusters 1-5.

CLUSTER 1													
DEG	FC (log ₂)*	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)
LRP5L	-1.07097	CASKIN2	-0.9989	PLEKHD1	-0.9368	FAM129A	-0.8956	DSTN	-0.84393	TOR1AIP2	-0.7996	KCTD17	-0.76603
FAM214B	-1.06955	FAS	-0.9984	JUP	-0.9367	VPS33B	-0.894	TOP1	-0.84354	DDAH2	-0.7991	ZFYVE1	-0.76577
ABCF3	-1.06796	VCPKMT	-0.9981	PTRH2	-0.9362	PTBP2	-0.8932	YY1AP1	-0.84299	UBR7	-0.7991	ZNF354A	-0.76527
PNPLA8	-1.06745	PTTG1	-0.9973	FAM177A1	-0.9357	SLC9A8	-0.8919	HN1	-0.84296	C10orf10	-0.7973	ERMN	-0.7651
TK1	-1.06561	RIMKLB	-0.9958	NDUFS5	-0.9356	ZNF8	-0.8912	ATP6V1F	-0.84205	MMAA	-0.7972	ZNF649	-0.76507
SESTD1	-1.06531	ADCY3	-0.9956	ZSCAN26	-0.9356	FGFR1OP	-0.8905	RAB29	-0.83944	FAM45A	-0.797	PNRC2	-0.76464
TMEM57	-1.06494	PHF1	-0.9939	TRPM4	-0.9356	SLC7A1	-0.8905	FRG1	-0.83784	ZNF195	-0.797	TRAF6	-0.7642
AMZ1	-1.0643	WARS	-0.9924	CSNK1A1	-0.9351	ITPKC	-0.8873	NPH4	-0.83742	CANT1	-0.7961	FTL	-0.76333
FAM118A	-1.06408	FRMD8	-0.9922	ILF2	-0.9344	INPP1	-0.8871	TERF1	-0.83706	RPS6KB1	-0.7957	GTPBP4	-0.76331
SDCBP	-1.05937	BAG2	-0.992	HARS2	-0.9341	USP11	-0.8869	MRPL50	-0.83698	KLF5	-0.7947	PDIK1L	-0.76323
TVP23A	-1.05906	ZNF687	-0.9891	STAB1	-0.9334	PDCL3	-0.8869	MPHOSPH10	-0.83669	AXIN1	-0.7946	FBRS	-0.76189
TCAF2	-1.05766	RNF10	-0.9876	MAN1A1	-0.9328	CYTH2	-0.8869	RPP38	-0.83623	CDC42BPB	-0.7941	SSSCA1	-0.76168
SOCS5	-1.05662	TRAPPC4	-0.987	TMEM188	-0.9318	MBIP	-0.8866	HGSNAT	-0.83539	MRPL13	-0.7939	FAM131A	-0.76114
SLC35F6	-1.05575	PEX14	-0.9868	PNMA1	-0.9311	LLGL2	-0.885	ZNF100	-0.83531	GRAMD4	-0.7938	ZNF836	-0.76107
ANXA2	-1.05333	GPKOW	-0.9866	HGS	-0.9308	ENY2	-0.8846	THBS1	-0.83471	SYVN1	-0.7935	HBP1	-0.76097
TNFAIP8	-1.05168	ZUFSP	-0.9863	EGFL8	-0.9301	ARSG	-0.8844	ZNF555	-0.83412	C11orf84	-0.7934	RNF24	-0.75978
SWAP70	-1.05158	SLC1A3	-0.9863	ALCAM	-0.9299	PISD	-0.8833	MTF2	-0.8331	TYW5	-0.793	TMF1	-0.75978
ZNF675	-1.05147	SIK3	-0.9846	LGALS1	-0.9292	BCL10	-0.8831	USP31	-0.83307	MICALL2	-0.7914	DNAJA2	-0.75882
TDG	-1.04968	PILRB	-0.9844	RAB34	-0.9287	MAP1S	-0.8803	ZSCAN12	-0.83247	HS3ST1	-0.7906	EIF6	-0.75877
ZER1	-1.0489	PHLDB3	-0.9833	ALDH1L2	-0.9284	SLAMF7	-0.8795	LDOC1L	-0.83244	CARD19	-0.7901	UBE2Z	-0.7584
MAPK6	-1.04613	SLFN5	-0.9828	CNG1	-0.9272	MED8	-0.8794	LRRC41	-0.83232	HERPUD1	-0.7894	ZNF436	-0.758
DGCR14	-1.04527	CLIP4	-0.9816	WIPI1	-0.9247	ING2	-0.879	UPP1	-0.83158	IRF2BP1	-0.7863	CGRRF1	-0.75789
PUS3	-1.04434	LRRC8C	-0.9805	EIF5A2	-0.9237	SEC61B	-0.8789	BASP1	-0.83157	PPP1CB	-0.7863	CCNC	-0.75687
MTMR6	-1.04401	MOB1B	-0.9802	C1orf216	-0.9232	TUBA1A	-0.8784	SSH1	-0.83153	DYNC1L1	-0.7863	RPL22L1	-0.75652
STX3	-1.04399	FAM107B	-0.9791	ZNF879	-0.9232	ZNF681	-0.8784	ARL6IP1	-0.83145	NBPF19	-0.7859	DDX24	-0.75618
ANKLE2	-1.04332	C1orf56	-0.9789	ZNF211	-0.922	CD82	-0.8764	MFSD5	-0.82864	TICAM1	-0.7853	BMP8B	-0.75469
ANKRD28	-1.04317	ITGAE	-0.9771	CDK14	-0.9215	TFEC	-0.8757	IPPK	-0.82737	NRP1	-0.7853	NPIPA5	-0.75457
ZNF773	-1.04315	TIMM22	-0.9754	METTL22	-0.9194	ELOF1	-0.8744	MFSD2A	-0.82711	PPP4R2	-0.7847	CNNND1	-0.75456
ZC3H7A	-1.04275	TBC1D17	-0.9753	EXO5	-0.9174	PGBD1	-0.8738	CIITA	-0.8268	RNF2	-0.7846	CWC15	-0.75363
SEC24A	-1.04216	C1orf50	-0.9748	CTNNA1	-0.9174	ASAP1	-0.8722	RDX	-0.8266	AMER1	-0.7833	SART3	-0.75357
ACTN1	-1.04194	SPAG5	-0.9745	VCP	-0.9173	ZNF155	-0.8697	RAB3IP	-0.82639	DUSP12	-0.7826	NDUFA11	-0.75275
CHMP4B	-1.04117	ZFP1	-0.9743	CLIP2	-0.9165	CFAP45	-0.8695	YES1	-0.8262	INTS5	-0.7826	TPD52L2	-0.7523
BACH2	-1.03981	FGD6	-0.9742	CORO1C	-0.9157	SMCHD1	-0.8692	ENO2	-0.82573	TMEM41B	-0.7817	RNF121	-0.751
GIN54	-1.03966	ZNF639	-0.9726	FAM108C1	-0.9156	SCOC	-0.8682	PQBP1	-0.82559	ZNF260	-0.7814	GNA15	-0.74986
NAA50	-1.03962	MAP3K11	-0.9716	GPX4	-0.9151	TAF1D	-0.8672	MYO1D	-0.82527	CYSTM1	-0.7811	PALB2	-0.74979
EDEM1	-1.03769	CCPG1	-0.9689	SLC9B2	-0.9145	SLC16A3	-0.8652	HAT1	-0.82526	SSBP2	-0.7794	GOLPH3L	-0.74968
CCNB1	-1.03712	C2orf88	-0.9664	ADAP2	-0.9144	TMEM136	-0.8649	SLC36A4	-0.82448	TAB2	-0.7788	EIF5B	-0.74961
WEE1	-1.03515	INF2	-0.9656	POMGNT2	-0.9135	MTMR9	-0.8646	MANF	-0.82446	MRFAP1	-0.7777	TTC33	-0.7491
SAMD12	-1.0338	PAM	-0.9633	MED17	-0.913	CCP110	-0.8641	ZFP36L1	-0.82418	PHYH	-0.777	BAZ1A	-0.74828
CCDC71L	-1.03214	SFXN3	-0.9617	GRAMD1B	-0.9129	SLC15A4	-0.8637	WHAMM	-0.82233	ZC3H12A	-0.7768	INSIG1	-0.74767
SLC39A8	-1.03078	CASP2	-0.9611	UBA5	-0.9124	IRGQ	-0.8634	RPS27L	-0.8223	PDE7A	-0.7767	HLA-DRA	-0.74741
TMEM189	-1.02979	G3BP2	-0.9605	LAPTM4A	-0.9122	MCM7	-0.8634	NUP58	-0.82126	TMEM60	-0.7767	ADPRHL2	-0.74733
ZNFX1	-1.02911	KDM5B	-0.9604	RACGAP1	-0.9121	STXBP3	-0.8632	VPS37C	-0.82044	MIER2	-0.7759	MTERF3	-0.74662
ZNF35	-1.02768	HOMER2	-0.9601	HMGXB4	-0.9117	PRKAR2B	-0.8631	BUD13	-0.81843	SRRT	-0.7755	ZNF581	-0.74592
CD180	-1.02729	ORC6	-0.9575	MED31	-0.9103	C18orf21	-0.8625	TGIF2	-0.81828	SOCS6	-0.7749	TAF5L	-0.74547
MPC2	-1.02679	RHOH	-0.9566	SUOX	-0.9093	GNG5	-0.8624	S100A10	-0.81792	DOLK	-0.7746	SMARCE1	-0.74524
LYST	-1.0253	ACVR2B	-0.9551	PRKCI	-0.9091	PIK3CA	-0.8621	UBXN1	-0.81677	ASL	-0.774	PAK1	-0.74472
THAP3	-1.02483	RPS6KC1	-0.9534	CEBPD	-0.9085	BAK1	-0.8614	HLA-A	-0.81667	GALNT2	-0.7733	LNX2	-0.74386
PHAX	-1.02458	BACH1	-0.9531	SLC3A2	-0.9079	RGS12	-0.8601	TMED7	-0.8157	SOAT1	-0.7729	GAN	-0.74357
ATAT1	-1.02192	LGALS9	-0.9516	ZNF17	-0.9069	TNFAIP1	-0.8599	KIF7	-0.81123	N4BP1	-0.7722	SMG8	-0.74256
USP12	-1.01825	NOP58	-0.9513	GCNA	-0.906	DFNB31	-0.8598	HECTD2	-0.8112	CASZ1	-0.772	HOXB2	-0.74181
HIST1H1C	-1.0148	DAPP1	-0.9503	ZNF189	-0.9043	RCC2	-0.8593	LRRC8A	-0.81115	ZSWIM1	-0.7715	KRT5	-0.74121
CASP10	-1.01469	SPAG4	-0.9495	MORF4L2	-0.904	DNAJC12	-0.8589	ACP6	-0.81075	NAP1L3	-0.7713	ZFP3	-0.74114
STARD3NL	-1.01393	TMEM176B	-0.9494	CYB561	-0.9036	RAP1A	-0.8586	SESN2	-0.81032	LMLN	-0.7703	CHSY1	-0.74088
COMMD5	-1.01344	HIST2H2AC	-0.9481	GNLY	-0.9031	KBTBD6	-0.8579	NDUFA6	-0.80923	FOXP1	-0.7702	PDCD10	-0.74058
HPS6	-1.01324	PDGFB	-0.9473	RRAS2	-0.9031	TCTN3	-0.8572	PSMC6	-0.80843	MCMBP	-0.7701	PPP1R8	-0.7402
HOMER2	-1.01323	MSN	-0.9464	SMC4	-0.9027	NUP62	-0.8568	ZNF229	-0.80738	TBC1D23	-0.7698	EED	-0.73986
ARF6	-1.01221	PATL1	-0.9461	DOCK4	-0.9021	PSMA2	-0.8559	FDX1	-0.80556	DNAH17	-0.7695	OTUD5	-0.73953
MAP4K3	-1.01172	RASGRP3	-0.946	CNOT6L	-0.902	WSB1	-0.8556	AKAP6	-0.80489	LIN7B	-0.7695	SIAH1	-0.73932
LUC7L	-1.01159	HLA-F	-0.9452	RNF145	-0.9017	TRAPPC2B	-0.8549	CDC37	-0.80466	RC3H1	-0.7694	ZSCAN22	-0.73916
AK6	-1.01144	CTD-3088G3.8	-0.9446	C2orf47	-0.901	LIMS1	-0.8543	YWHAH	-0.80431	STAG3	-0.7692	MRPL14	-0.7382
SPAG9	-1.01062	FAM102B	-0.9446	HMCES	-0.9004	ZFYVE16	-0.8542	TVP23C	-0.80404	CCDC120	-0.769	FBN1	-0.73792
ELOA	-1.01051	YDJC	-0.9445	DYNLT3	-0.9004	MAPK8	-0.8529	CTNNB1	-0.80394	WDR48	-0.7681	UBQLN4	-0.73781
PI4K2A	-1.00882	GSKIP	-0.944	SIDT1	-0.9	FTH1	-0.8509	ACTR1B	-0.80308	PPP1R3D	-0.768	DDX52	-0.73711
HBS1L	-1.00666	SMAP2	-0.9422	CRY2	-0.8996	TOR3A	-0.8505	GEN1	-0.80304	RPAIN	-0.7678	KIAA1841	-0.73639
INPP5F	-1.00637	DENND5A	-0.9405	SGTB	-0.8995	STRIP1	-0.8505	CEP89	-0.8028	RFFL	-0.7678	SEC61A2	-0.73552
ZNF48	-1.00518	TRO	-0.9402	BTG3	-0.8991	ZNF281	-0.8495	MFAP1	-0.80265	MRPL15	-0.7672	RUNX3	-0.73416
IFT57	-1.0002	ATAD3B	-0.9377	ZNF672	-0.8989	PSMD11	-0.8469	PAK4	-0.80192	KIF20B	-0.7672	SAV1	-0.73402
GPBP1	-0.99984	NOCT	-0.9374	ACTR1A	-0.8989	ENTPD7	-0.8469	PPTC7	-0.80178	ZBTB34	-0.767	ARPP19	-0.73242
CORO7	-0.99957	RAB12	-0.9372	CCNJ	-0.8976	CCDC126	-0.8464	SEL1L	-0.80121	NLRP3	-0.767	SLC10A3	-0.73154
FXYD2	-0.99929	TSPYL1	-0.9368	FLNA	-0.8957	EI24	-0.8445	SEC24D	-0.80071	ANKRD50	-0.7661	ZNF283	-0.73125

CLUSTER 1													
DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)
PKM	-0.72903	TMEM199	-0.695	ZNF823	-0.6594	TUBE1	-0.6251	WDR73	-0.59935	VDAC3	-0.5732	SEC13	-0.54787
ZNF543	-0.72893	GLTP	-0.6944	FH	-0.6593	ATP6V0D1	-0.6249	MAML1	-0.59861	ZBTB45	-0.5725	PSMD5	-0.54781
ENSA	-0.72811	ABL1	-0.6943	EIF2S2	-0.6592	TMEM127	-0.6248	KHDRBS1	-0.59837	NOB1	-0.5725	LAG3	-0.54779
ZNF562	-0.72782	CMC2	-0.6935	ZNF749	-0.6574	COPS2	-0.6247	RBMXL1	-0.5981	CCDC127	-0.5718	HSF2	-0.54755
ZBTB5	-0.7276	GK	-0.693	CINP	-0.6568	MORF4L1	-0.6246	UBQLN2	-0.59787	GOLT1B	-0.5713	OAZ2	-0.54755
PRNP	-0.72734	ZNF430	-0.6924	MYH9	-0.6563	VAV2	-0.6245	FZD2	-0.59744	AGPAT6	-0.5713	ZC4H2	-0.54714
SERINC2	-0.72706	MAP1LC3A	-0.692	TXNDC17	-0.6562	TSEN54	-0.6241	SLC25A19	-0.5973	CTD-2568A17.1	-0.5706	ENDOG	-0.54692
FAM110A	-0.72696	HNRNPUL1	-0.6912	SPRYD4	-0.6552	MGRN1	-0.6235	CDC42	-0.59728	PSMD6	-0.5698	ARFIP2	-0.54687
ZC3HC1	-0.72644	PDK1	-0.6896	CXorf40B	-0.6552	MIEF1	-0.6231	KIAA1191	-0.59703	TMUB1	-0.5697	FAM234A	-0.54674
EBLN2	-0.72534	IST1	-0.6894	POP7	-0.6546	POLR1C	-0.6217	POC1B	-0.59699	KLF3	-0.5694	UTP3	-0.54669
ZBTB49	-0.72505	CYLD	-0.6881	WDFY1	-0.6545	CA5B	-0.6216	NECAP2	-0.59664	CMTM4	-0.568	WIPI2	-0.54649
COPRS	-0.72455	OXSRI	-0.6873	RNF114	-0.6534	STX4	-0.621	QSOX1	-0.59649	AEBP2	-0.5676	SS18L2	-0.54635
SDHAF2	-0.72452	HLA-DRB5	-0.6872	TBC1D25	-0.6532	ZNF674	-0.6201	ZNF552	-0.59622	IMMT	-0.5674	KAT2B	-0.54605
NANS	-0.7242	TEC	-0.6869	GORAB	-0.6525	PSMD4	-0.6197	MPST	-0.59609	ZEB2	-0.567	MPV17L2	-0.54552
CENPP	-0.72294	YTHDF3	-0.6866	CD86	-0.6518	CCM2	-0.6195	RNF25	-0.59571	SMARCD1	-0.5663	C17orf49	-0.54521
ALDH8A1	-0.7222	LRRC40	-0.6861	ZNF550	-0.6517	NCOA5	-0.6192	MSTO1	-0.59479	POP1	-0.5658	PGRMC2	-0.54517
GALE	-0.72137	MYOM2	-0.6854	PITPNM1	-0.6512	PI4K2B	-0.6191	ATG4B	-0.59343	TP53BP2	-0.5658	DNAJC25	-0.545
ISCA1	-0.72122	DYNC1I2	-0.6853	MAP2K3	-0.6511	ATP2A2	-0.6185	GEMIN2	-0.59335	HMGXB3	-0.5657	DDX39A	-0.54499
MCOLN2	-0.72026	ZNF451	-0.684	ENTPD1	-0.65	HLA-DQB1	-0.6176	INO80D	-0.59335	FAM189B	-0.5653	RAB22A	-0.54454
PCGF5	-0.71964	SAMD4B	-0.6828	FICD	-0.6491	CEP76	-0.6171	PLAG1	-0.59334	BET1L	-0.5652	CEP57	-0.54426
MRPL44	-0.71964	MFN1	-0.6817	DOT1L	-0.649	ZNF136	-0.6169	TIPRL	-0.59314	RELA	-0.5651	PML	-0.54411
FAM92A1	-0.71603	STAU1	-0.681	FBXO33	-0.649	WDR25	-0.6168	MTFR1	-0.59292	NET1	-0.565	OCLR	-0.54389
TRIP6	-0.71556	MPRIIP	-0.6804	ASNSD1	-0.6483	SLC30A7	-0.6165	B3GNT2	-0.5929	BAG5	-0.5646	PDXK	-0.5438
HLA-G	-0.71474	USP42	-0.6799	TUBB	-0.6473	CLTA	-0.6165	SETD9	-0.59268	CNOT2	-0.5644	FAM615	-0.54348
E2F6	-0.7147	ZNF606	-0.6794	SLC17A5	-0.647	RANBP10	-0.6165	TBC1D15	-0.59199	SERAC1	-0.5646	PTPN9	-0.54333
PDE3B	-0.71465	PJA1	-0.6792	UBTD2	-0.6461	MGLL	-0.6165	SAP30	-0.59188	FAM127B	-0.564	CNOT6	-0.54308
BRD8	-0.71448	BNIP3L	-0.679	SCML1	-0.6456	TULP3	-0.6161	PICALM	-0.59183	DDX20	-0.5625	RSPH3	-0.54233
LCAT	-0.71444	DHCR7	-0.6786	SAR1B	-0.6455	FAM76B	-0.6159	TXNL1	-0.5918	KLHL36	-0.5625	RAP1B	-0.5423
NRARP	-0.71417	UQCRCF51	-0.6782	CLPX	-0.6455	LANCL3	-0.6158	MRPL22	-0.59159	ZNF91	-0.5617	PPP1R18	-0.54221
HMGNS	-0.71389	TRIP10	-0.6774	SGPP1	-0.6453	PSMD2	-0.6141	SUPT4H1	-0.59122	SPRY1	-0.5616	HIST4H4	-0.54213
MPHOSPH6	-0.71372	ZNF326	-0.677	AREL1	-0.6453	CCZ1	-0.6136	NT5C3A	-0.591	CCDC12	-0.5616	ERLEC1	-0.54212
UBE2V1	-0.71332	UBE2D1	-0.676	DSTYK	-0.6446	ACTR3B	-0.6135	UPRT	-0.59096	TSSC4	-0.5613	LILRB4	-0.54204
AKT1S1	-0.71309	GXYLT1	-0.6759	ZC3H10	-0.6442	ELOVL5	-0.6133	SNX22	-0.58954	RAP2C	-0.5608	MTA2	-0.54194
DERL2	-0.71233	RARS	-0.6758	COX7B	-0.643	RTP5	-0.6132	C11orf57	-0.58929	HECA	-0.5606	MESDC1	-0.54187
RABGGTB	-0.71211	PUM1	-0.6755	CTNS	-0.6429	GPS2	-0.6127	TTF1	-0.58907	SRP54	-0.5598	DHX8	-0.54128
CNOT8	-0.71192	GORASP2	-0.6755	MPP1	-0.6417	C1orf52	-0.6125	TMEM208	-0.58898	PSMB4	-0.5595	FZD6	-0.54102
SNX33	-0.71161	ZNF737	-0.6749	BCKDK	-0.6408	MMD	-0.6117	DHDDS	-0.5885	FAM83G	-0.5592	PTMA	-0.54096
FBXO45	-0.71119	SPATA2	-0.6748	DAGLB	-0.6405	TXNL4B	-0.6115	PSMB5	-0.58833	FBXL3	-0.559	DFFA	-0.54015
KIAA1522	-0.71109	GAS8	-0.6745	MGST3	-0.6404	PITRM1	-0.6114	XRN2	-0.58818	ELF1	-0.559	ARIH2	-0.54001
RSPRY1	-0.71105	PRUNE	-0.6741	ABT1	-0.64	RAB18	-0.6112	ZNF180	-0.58791	XAB2	-0.5588	ZNF219	-0.53993
RBMX	-0.71073	TXNDC9	-0.6732	RAD21	-0.6397	FKBP14	-0.6111	GRPEL2	-0.58772	AP5Z1	-0.5586	SEC61G	-0.53972
TIMM17A	-0.71066	ABTB2	-0.673	DYNLT1	-0.6389	PITHD1	-0.6106	RAB13	-0.58736	RIOK2	-0.5585	HLA-B	-0.53939
TMC6	-0.71017	ARID4B	-0.6729	SFMBT1	-0.6388	PAIP2	-0.6102	IL18RAP	-0.58712	KLHL21	-0.5578	IL2RG	-0.53902
SCD	-0.70955	ADGRD1	-0.6726	EMC8	-0.6381	EIF1AX	-0.6102	MANBAL	-0.58623	SLC38A7	-0.5578	CCDC71	-0.53763
ZNF266	-0.70944	43164	-0.6725	ARL14EP	-0.638	STARDA4	-0.6102	TXNDC15	-0.58608	PRELID3B	-0.5564	FBXL15	-0.53737
KLHL5	-0.70941	SF3B4	-0.6717	CFAP36	-0.638	TOX4	-0.6102	SFR1	-0.58534	AP3M2	-0.5559	DHX40	-0.53669
MAN2A1	-0.70869	BLOC1S3	-0.6713	PANX1	-0.637	MED4	-0.6098	TTC9C	-0.58425	TET2	-0.5548	BIN3	-0.53603
TSPAN14	-0.70789	ATG4A	-0.6708	SF3B5	-0.6363	RPP40	-0.608	DCP2	-0.58392	ZNF154	-0.5547	TRMT10C	-0.53552
RQCD1	-0.70757	PSMC4	-0.6707	FNIP1	-0.6363	ZNF202	-0.6079	PIEZO2	-0.58322	CCDC82	-0.5543	NDUFA5	-0.53513
SPTSSA	-0.70727	VPS26A	-0.6704	WRAP53	-0.6351	HLA-C	-0.6078	SEC62	-0.58317	PSMA1	-0.554	PLAA	-0.53487
PSMC2	-0.70671	CLIC1	-0.6702	FBXO8	-0.6348	ABCF2	-0.606	UNG	-0.5831	NLE1	-0.5529	ISG15	-0.53317
MYO1C	-0.7067	MED21	-0.6688	RP11-437B10.1	-0.6344	KPNA5	-0.6059	POR	-0.58234	RNF8	-0.5527	ADNP2	-0.53313
DNAJC2	-0.70587	CEP85	-0.6678	ATP1B1	-0.6337	KCTD11	-0.6053	FAM104A	-0.58228	GPBP1L1	-0.5521	EFTUD2	-0.53288
ISCU	-0.7053	SMAD4	-0.6676	UBN1	-0.6332	GPR183	-0.6053	ANK3	-0.58216	MRPS2	-0.552	GMPS	-0.53191
SETMAR	-0.70328	USP30	-0.6675	TIMM23	-0.6328	ZMYM5	-0.6049	ARHGAP21	-0.58036	LEO1	-0.5518	VAMP3	-0.53188
TP53RK	-0.70293	BMP2K	-0.6667	SAMD8	-0.6326	KPNA6	-0.6047	PAWR	-0.58036	SLC25A34	-0.5513	HPCAL1	-0.53167
RRBP1	-0.70288	MAP2K1	-0.6662	ZNF557	-0.6326	TCP11L1	-0.6045	RIOX1	-0.57979	TXNDC11	-0.5511	CAPZA1	-0.53124
CDK7	-0.70243	TCEAL3	-0.666	DYRK1B	-0.6325	CLK4	-0.6038	HSD17B12	-0.57944	PLPP5	-0.551	CMIP	-0.53102
AGFG1	-0.70168	SURF4	-0.6648	UBAP2	-0.6321	JAGN1	-0.6034	PPP1R16B	-0.57931	RNF185	-0.5505	COPB2	-0.53087
PDCL	-0.70059	C6orf47	-0.6647	LRRC8B	-0.6312	MGAT4B	-0.6026	FOXP1	-0.57912	SCYL2	-0.5501	PLA2G4A	-0.53052
KIN	-0.70018	GAK	-0.6642	CHD4	-0.6311	ZNF300	-0.6016	PSMD7	-0.57894	ZNF225	-0.55	MRPL12	-0.53045
RABIF	-0.6993	TAGLN	-0.6641	SH3BP5L	-0.6308	FAM65C	-0.601	FLAD1	-0.57871	MSANTD3	-0.5497	SAT2	-0.53043
EMD	-0.69846	CBX1	-0.6641	SKP1	-0.6307	CASP6	-0.6008	ANAPC10	-0.57853	RPF1	-0.5497	ORC1	-0.52974
PTP4A2	-0.69802	KIF3A	-0.6637	DCLRE1B	-0.6306	TP53INP1	-0.6002	DTD1	-0.57704	TAOF9B	-0.5496	C8orf41	-0.52962
KBTBD2	-0.69784	FAM103A1	-0.6624	ZNF711	-0.6296	COPA	-0.6002	ARHGAP26	-0.57661	IFO7	-0.5492	IQSEC3	-0.52946
MED7	-0.69774	PRR14	-0.662	BMPR2	-0.6292	AKAP8L	-0.6002	NBPF1	-0.57543	DNAJC6	-0.5491	FEM1B	-0.529
RIPK2	-0.69757	PRPF39	-0.6608	LYPLA2	-0.6269	CARS	-0.6001	DOCK1	-0.57509	PEX26	-0.5491	SAFB	-0.52889
ASB7	-0.69727	TFG	-0.6607	CCDC174	-0.6266	LSM8	-0.5996	ZNF707	-0.57393	UBE2Q2	-0.549	GNAI3	-0.5287
UBA6	-0.69636	TARS	-0.6603	TSFM	-0.6263	TERF2IP	-0.5996	RNF181	-0.57392	VKORC1	-0.5489	EHD1	-0.5283
ZC3H8	-0.69547	KIF3C	-0.6602	STAP1	-0.6257	RBM15B	-0.5995	SELT	-0.57389	CCDC26	-0.5486	HNRNP80	-0.52789
FBXO21	-0.69539	MED9	-0.6598	KLHL24	-0.6251	NFKBIB	-0.5994	CCDC130	-0.57328	PCSK7	-0.5479	ALG9	-0.5278

CLUSTER 1													
DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)
UPK3A	-0.52721	PIK3R6	-0.5084	FBXO42	-0.484	INIP	-0.3657	ACTR10	-0.4289	PPP2R1A	-0.4328	ESPL1	-0.25635
POLD3	-0.52719	PKIG	-0.5083	NDE1	-0.4839	PATJ	-0.3655	SMARCD3	-0.42845	TLE1	-0.431	HIST1H2AL	-0.26759
PDRG1	-0.52675	MFSD12	-0.5078	HIST1H2BN	-0.4832	ATMIN	-0.3653	FAM160A2	-0.42821	TTL4	-0.4306	MLLT6	-0.26666
SEC22A	-0.52669	MICAL3	-0.5066	SEPSECS	-0.4822	PPP4C	-0.4614	IL10RA	-0.42806	SHOC2	-0.4304	DOCK3	-0.2598
NACC1	-0.52663	TRIM62	-0.5063	CFL2	-0.4813	WARS2	-0.4609	TBC1D10B	-0.4275	GFOD2	-0.4299	CYP2R1	-0.25931
EIF3J	-0.52663	EPT1	-0.5059	LY6G5B	-0.4812	ZMAT2	-0.4609	ICA1L	-0.42748	ZCCHC17	-0.4297	KLHL2	-0.25906
YEATS4	-0.52657	CTDP1	-0.504	TIMP2	-0.481	CEBPG	-0.4608	MYO9B	-0.42703	PSMD1	-0.4291	CCDC66	-0.2556
ZNF85	-0.52645	TOR1B	-0.504	IMPA1	-0.4805	RBM5	-0.4605	SPIN1	-0.42631	TJAP1	-0.3701	CDC42SE2	-0.25404
ZNF486	-0.5261	FAM63A	-0.5036	OTUD4	-0.4804	MTCH1	-0.4603	NDUFB8	-0.42611	RDH10	-0.370	DUS4L	-0.25193
JMJD1C	-0.52604	NSMCE2	-0.5033	NUP88	-0.4803	PRDX6	-0.4593	MINOS1	-0.42516	MFSD6	-0.3695	CHRNA10	-0.24993
TRAFD1	-0.52598	GPATCH3	-0.5031	BCAT1	-0.4801	TOR2A	-0.4592	KPNA1	-0.42457	FUBP1	-0.3658	MYBL2	-0.24931
TRIM13	-0.52587	LINS1	-0.5031	PARL	-0.4799	PROSER2	-0.4584	BTBD10	-0.42405	PGM2L1	-0.3652	ZFP69	-0.24468
RBSN1	-0.52585	EIF5A	-0.5022	ZNF586	-0.4798	GMFB	-0.4583	ABHD6	-0.42204	GPATCH2L	-0.365	SPNS2	-0.24026
HARS	-0.52585	GTF2E1	-0.5017	GSPT2	-0.4795	GUCD1	-0.4581	CAPN2	-0.42171	WASL	-0.3623	ARID5A	-0.23896
C11orf95	-0.52527	HIF1AN	-0.501	TFRC	-0.4794	N4BP2L1	-0.4577	RREB1	-0.42069	PRMT1	-0.361	DCUN1D4	-0.23803
MAPK7	-0.5251	WDR13	-0.5007	RELN	-0.4793	IMP3	-0.4575	ILKAP	-0.41947	KLHL20	-0.3587	TG	-0.2365
TIGAR	-0.52427	YOD1	-0.5006	EMSY	-0.4791	TGS1	-0.4574	PCDH9	-0.41871	CCNG2	-0.3583	PPFIBP2	-0.22931
BAP1	-0.52419	MUL1	-0.5002	PSME4	-0.479	IGHMBP2	-0.456	ZNF16	-0.41821	NEDD1	-0.3569	PLCH2	-0.22722
MFSD9	-0.5241	CDH1	-0.5	NFIC	-0.4789	UBE2R2	-0.4553	ESCO1	-0.4182	PUF60	-0.3567	HSPA1B	-0.2253
VDAC1	-0.52348	HINFP	-0.5	RAPGEF2	-0.4788	RHOG	-0.4552	EGLN3	-0.41809	SDHAF1	-0.3561	ERBB2	-0.22314
C1orf122	-0.52291	THAP4	-0.4999	CLMN	-0.4787	RP3-329A5.4	-0.4549	UBXN7	-0.41672	TRAK2	-0.3559	CSTB	-0.22103
LARPA4	-0.52283	MYL6	-0.4998	DONSON	-0.4786	BLOC1S2	-0.4545	ZNF7	-0.4166	UBR4	-0.3547	ZNF217	-0.21986
TEX30	-0.52215	NUBP1	-0.4998	SOS2	-0.4776	ARL13B	-0.454	KIAA0895L	-0.41646	ZNF253	-0.3545	CDC6	-0.21468
CACNA1F	-0.52183	RAB11FIP4	-0.4997	TMPPE	-0.4769	MYEF2	-0.4538	NSMF	-0.41514	PRKAA1	-0.3542	C17orf107	-0.19783
ABCE1	-0.52176	DIP2C	-0.4995	CCZ1B	-0.4769	ZFP91	-0.4536	COG1	-0.41246	TMEM214	-0.3533	ITSN1	-0.18778
ARID3B	-0.52035	COPB1	-0.4994	PPP2CB	-0.4762	MBD4	-0.4534	CSNK2B	-0.41123	AKT3	-0.3533	CSRP1	-0.18564
ABCA5	-0.51985	EBP	-0.4984	EYA3	-0.476	C12orf29	-0.4527	ELP5	-0.40984	PANK3	-0.3514	CACNB4	-0.18125
LIN9	-0.5198	TINAGL1	-0.4976	NCAPD3	-0.4755	NRXN2	-0.4515	BAG4	-0.4083	NOL4L	-0.3498	MRPL54	-0.17919
TOE1	-0.51978	C12orf65	-0.4971	SIGLEC10	-0.4753	C14orf119	-0.4514	UNC45A	-0.40788	SACM1L	-0.3488	RC3H2	-0.1779
RALGAPA1	-0.51949	BFAR	-0.4971	NPLOC4	-0.4751	ZNF805	-0.4512	ATP5G3	-0.40518	ODF3B	-0.3457	RNF144A	-0.17497
OTUD3	-0.51941	PHTF1	-0.497	MATR3	-0.4744	FBXL19	-0.4501	ANXA11	-0.40458	C8orf76	-0.3452	ZNF671	-0.17326
FBXW11	-0.51935	IFI16	-0.4963	STK35	-0.4738	ZNF706	-0.4491	G2E3	-0.40405	HLA-DMA	-0.344	PLXDC2	-0.16996
GNB4	-0.5192	RABAC1	-0.4961	SLC35E4	-0.4737	SLC26A2	-0.4489	SREBF2	-0.40342	RORA	-0.3432	RNF34	-0.16379
PIIG	-0.51912	FBXO46	-0.4955	HDAC2	-0.4735	DPF2	-0.4487	PCNX4	-0.40322	STARD9	-0.3398	MAP109326.1	-0.16361
RNF144B	-0.51897	CERS5	-0.4953	RAB4B	-0.4723	MAGT1	-0.4469	ZNF853	-0.40193	JADE3	-0.3388	AIG1	-0.15333
ZKSCAN8	-0.51844	QPR1	-0.4951	CEP97	-0.4718	AAMP	-0.4463	NAPB	-0.40095	SLC36A1	-0.3376	LRRC3	-0.15235
MAP4K5	-0.5181	RPL36AL	-0.495	PHF10	-0.4713	RAN	-0.4455	TUBA1C	-0.40038	REV3L	-0.3368	CDT1	-0.15141
LUZP1	-0.51803	TTC13	-0.4949	TRIM23	-0.4709	PWP1	-0.4454	POMP	-0.39817	NEU3	-0.3353	RARRES2	-0.14527
DEDD	-0.51747	USP14	-0.4944	ZNF622	-0.4708	FBXO7	-0.4447	CAMKK1	-0.39745	TTN	-0.3352	TRIM39	-0.14523
ST3GAL1	-0.51735	BTG1	-0.4942	ZNF747	-0.4705	CCL3	-0.4437	C14orf156	-0.3972	TMEM44	-0.3315	POLR3B	-0.14168
ZNF616	-0.5172	SGPL1	-0.4941	CHRNB1	-0.4703	ZFAND6	-0.4435	TMEM64	-0.39698	KRTCAP2	-0.3303	NSF	-0.13056
ARPC5L	-0.51652	LDB1	-0.494	ISY1	-0.4701	TRIM68	-0.4431	TTL	-0.39531	SUDS3	-0.328	ABCA9	-0.1271
SLC25A13	-0.51641	L2HGDH	-0.4939	CTS2	-0.4694	DCAF8	-0.4425	APMAP	-0.39185	PIGV	-0.3278	KIF11	-0.12663
PHF20L1	-0.51584	SNRNP27	-0.493	MESDC2	-0.4692	DCUN1D1	-0.4425	FAM175B	-0.39009	DHRS7B	-0.3248	SCN3A	-0.12578
GBF1	-0.51541	ASXL1	-0.493	NIPSNAP3B	-0.4692	SMURF1	-0.4425	NARS2	-0.38943	TECPR2	-0.3214	PLCB1	-0.12544
INO80C	-0.51529	NEK7	-0.4927	ZKSCAN4	-0.469	TMED1	-0.4417	C6orf1	-0.38917	MAP3K6	-0.3201	CENPK	-0.1242
AAED1	-0.5149	FAM160B1	-0.4927	TMSB10	-0.4685	KNSTRN	-0.4415	PLAGL1	-0.38908	BTRC	-0.3184	EHHADH	-0.12048
RAP2B	-0.51439	DNAJC5	-0.4921	RIPK1	-0.4684	BAG3	-0.4411	ARID5B	-0.38832	SNX27	-0.3175	IRF2	-0.11588
ZNF345	-0.51394	FAM134B	-0.4919	ZNF197	-0.4682	SNRPB	-0.44	HMG20A	-0.38746	FCHO2	-0.3161	SH3GL1	-0.11522
DCAF16	-0.51384	MIER1	-0.4918	STK26	-0.4681	CASP8	-0.4397	ZNF107	-0.38554	POMK	-0.312	FSTL1	-0.11063
RBM7	-0.51373	LILRA6	-0.491	C19orf79	-0.4676	DCTN6	-0.4396	PFKL	-0.38546	FAH	-0.3101	KIAA1328	-0.09798
TP11	-0.512	MEI1	-0.4909	SLC44A1	-0.4673	ZNF653	-0.4391	TSTD2	-0.38345	PUS7	-0.3097	THBD	-0.09386
PLEKHA3	-0.51192	PHF19	-0.4906	CDK9	-0.4665	ATL3	-0.4389	MYO1B	-0.38285	USP9Y	-0.3067	MT2A	-0.09279
ASB6	-0.51192	SLC35B2	-0.4902	DLGAP4	-0.4661	ABCB10	-0.4388	RSL24D1	-0.38251	ZNF841	-0.3051	MAP3K20	-0.08221
TP53BP1	-0.5117	IGBP1	-0.4891	AGO1	-0.4659	ZNF772	-0.4385	SRPK1	-0.38029	E2F3	-0.3035	C5orf30	-0.07377
TMEM120A	-0.51167	UPF3B	-0.4889	BDH2	-0.4658	ZFP82	-0.4382	RHBDD2	-0.37887	RPS10	-0.3027	ZNF568	-0.07339
ARPI1	-0.51133	GNP1	-0.4889	PRRC1	-0.4657	PSMB6	-0.4382	NAP1L4	-0.37786	HERC6	-0.299	ALDH7A1	-0.06744
FAM89B	-0.5111	NOD2	-0.4884	KANSL3	-0.4649	SEL1L3	-0.4372	MTSS1	-0.37678	SLC25A28	-0.2981	RP11-40A7.2	-0.06399
MRPL17	-0.51034	FAM133B	-0.488	DTX3L	-0.4649	GTPBP2	-0.4369	LY6E	-0.37637	IL1RAP	-0.2957	CATSPERG	-0.05932
CDK2	-0.51023	RBPJ	-0.4878	C7orf43	-0.4648	GPR137	-0.4367	BMP1	-0.37633	FAM69A	-0.292	ZNF324B	-0.04673
UBE2A	-0.51013	AC017099.3	-0.4874	C5orf24	-0.4644	ZNF431	-0.4364	FPGS	-0.37476	FNDC3A	-0.2899	SP140	-0.04104
CRTC2	-0.51008	WIZ	-0.4863	CNPY2	-0.4642	43165	-0.4354	ACIN1	-0.37356	SECISBP2L	-0.289	ID3	-0.03814
WBP11	-0.50989	ARFGAP1	-0.4858	TMTCC3	-0.4641	TAF9	-0.4353	HLA-DPA1	-0.37338	ZBTB41	-0.289	SSPO	-0.03723
ARL8A	-0.5096	AKR1C3	-0.4857	STAT6	-0.4628	ZFP28	-0.4352	ARHGAP31	-0.37334	ZW10	-0.283	RAB20	-0.02956
APLF	-0.50921	WIPF2	-0.4849	MED28	-0.4628	MED10	-0.4346	ZNF169	-0.37234	MFHAS1	-0.2827	TEX14	-0.01612
MYPOP	-0.50899	REST	-0.4849	NAPG	-0.4619	URB2	-0.4333	ZNF660	-0.37197	MYH11	-0.2801	APOE	-0.00905
CLN5	-0.50866	ACSL4	-0.4848	NACC2	-0.4617	TRPC4AP	-0.4332	FAM221A	-0.37188	SGO1	-0.2775	PHLDB2	0.003597
HLA-DPB1	-0.50848	ACADVL	-0.484	TSPYL4	-0.4615	MTR	-0.4329	SVIL	-0.37146	SCAND1	-0.2755	GPR18	0.014173
								MSH6	-0.37027	WDHD1	-0.2726	AK1	0.026351
										MED12	-0.2588	MYH7B	0.029058

CLUSTER 2													
DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)
SLC27A1	1.542972	ZNF618	1.4769	ZC3H12D	1.42515	MICU1	1.36019	FBP1	1.31514	C1RL	1.25991	SUPT3H	1.213714
ADD3	1.54288	ORAI2	1.47679	TFAP4	1.42473	NEURL1	1.35994	ZFP64	1.31482	ITGB2	1.25871	PHF2	1.213164
ZNF652	1.542596	NMRAL1	1.47662	TRAM2	1.42381	GF11	1.35961	GLTSCR1L	1.3148	POLR3K	1.25817	AKR7A2	1.211585
NUDT17	1.540387	PHACTR1	1.47482	ANKRD13D	1.4223	P2RX7	1.35939	L3MBTL2	1.31424	SH3KBP1	1.25806	RALGAP2	1.209408
ZNF395	1.53956	HIST1H2BE	1.47457	SNX29	1.42219	ZMYM1	1.35889	TMEM168	1.31421	DIP2A	1.25791	SNTB1	1.208882
TMEM170B	1.539339	RBL2	1.47431	PTGS1	1.42178	LACTB2	1.35848	RECQL5	1.31297	ERI3	1.25713	LTBR	1.208391
PYGM	1.538316	CH507-42P11.8	1.4735	IQGAP2	1.42175	CRTC1	1.35725	C1orf162	1.31218	SUSD1	1.2569	TMEM19	1.208378
GLRX	1.537061	ATP2B4	1.4726	VAMP8	1.42143	MAML3	1.35719	MMP15	1.31207	INTS3	1.25646	MSH5	1.207787
CCDC186	1.535772	MDN1	1.47227	HIC1	1.42061	SCAMP5	1.35684	C6orf203	1.31112	FPGT	1.2549	PSMB9	1.206977
KLHL15	1.535289	MAP2K6	1.47226	SEMA5A	1.419	ST3GAL2	1.35615	FAM46A	1.31097	POU2F2	1.25429	RTKN	1.205673
JUNB	1.533821	EPHX1	1.47212	MEF2A	1.41841	CNTNAP1	1.35571	PRIM2	1.30865	PRMT3	1.2537	CLDN23	1.205326
APOBEC3C	1.533547	DCLK2	1.47177	ADRKB1	1.41796	ADAM11	1.35454	CUX1	1.30738	CYP2D6	1.2534	MYO15B	1.204918
IQSEC1	1.529911	ATG4C	1.47044	MYCBP	1.41737	PLCXD2	1.35448	IQCE	1.30573	CD36	1.25295	POLR3GL	1.203699
IGSF6	1.527765	FBXO32	1.4687	MGME1	1.41562	LMBRD2	1.35397	ACSS1	1.30534	ANAPC16	1.25288	NSMCE3	1.203255
OBFC1	1.52756	HSPG2	1.46636	PRR5L	1.41548	HERC3	1.353	NR1D1	1.30517	SHISA5	1.25218	IL11RA	1.202948
STMN3	1.526867	SHTN1	1.46636	NLRP7	1.41453	PRKCA	1.35213	DNAJC13	1.30434	ARID1B	1.25185	RHPN1	1.201772
SDHAF3	1.5264	USP18	1.46404	ZNF573	1.40898	MBTPS2	1.35206	RNF139	1.30433	KPTN	1.25182	LMO4	1.201392
XYLT2	1.525807	SP110	1.46367	NHLRC3	1.40868	TMEM129	1.35194	TET3	1.30388	METTL25	1.25167	RCN3	1.201335
FAM168A	1.524063	PIK3AP1	1.46267	CCDC191	1.40694	XXYL1	1.35162	BTK	1.30387	KLHDC1	1.24978	UBE2E1	1.200175
STAC3	1.522519	PAIP2B	1.46227	CXorf23	1.40687	MGST2	1.35085	NCOR2	1.30351	THEM6	1.24943	MAP3K4	1.199999
TTC24	1.521241	DNASE1	1.46088	TXNIP	1.40567	MSRB3	1.35066	HIGD1A	1.30335	KIAA2013	1.24744	MXI1	1.199169
ANKRD39	1.518195	COQ7	1.45921	PLCB4	1.40449	RFT1	1.35027	CECR5	1.30131	MAP2K5	1.24608	DOCK2	1.199159
CDKN2AIPNL	1.517658	EPS8L1	1.45875	HIBADH	1.40085	RP11-167P11.3	1.34995	CDK5	1.3007	AMDHD2	1.24583	OBSCN	1.198616
PQLC3	1.515242	SLC12A3	1.45771	RPP21	1.39971	TNFSF13B	1.34974	RELL2	1.30051	SMYD2	1.24581	SYNE1	1.198589
HSPA2	1.51518	SMARCAL1	1.45766	SAMD14	1.39885	ALDOC	1.34827	BICD2	1.29945	LMNTD2	1.2448	CARNS1	1.198353
SLC25A20	1.514876	MZB1	1.45737	B3GNTL1	1.3986	KCNK10	1.34761	CASP1	1.29865	ARHGAP18	1.2443	FAM173A	1.197638
TDRD1	1.514178	SHANK1	1.45698	LPCAT4	1.39803	C19orf38	1.34756	WDR92	1.29754	SLC35B3	1.24322	ZNF692	1.196634
NOD1	1.512964	GPR162	1.45691	ABHD8	1.39762	SRGAP2	1.34647	EVC2	1.2968	SLC23A2	1.24258	DENND5B	1.194003
GALNT6	1.512702	TTC39A	1.45662	FEZ2	1.39634	ACSL5	1.34609	PTK2B	1.29421	SERPINF1	1.24242	CCDC50	1.192517
CEACAM21	1.512044	ZMAT5	1.45431	GNPDA1	1.39309	ADA	1.34558	C17orf67	1.29414	C9orf142	1.23805	DXO	1.190048
TGFBR2	1.509682	DDX60	1.45392	TCFL5	1.39307	FARS2	1.34536	LRRRC8D	1.29285	UTRN	1.23745	CISD3	1.189748
SNAPC5	1.509119	TPK1	1.45383	DHRS4L2	1.39276	QRICH2	1.34517	ALG8	1.29271	TMBIM4	1.23721	JMY	1.18845
DGKD	1.508862	CYP4V2	1.45209	TASP1	1.39092	PRX	1.34353	PRAG1	1.29079	INPP5D	1.23565	GTF2IRD2	1.18765
EXOSC5	1.508852	FAM53A	1.45149	CBX8	1.39038	FAM111A	1.34273	GAB3	1.2906	TRMT2B	1.23444	DTWD2	1.186848
ACADSB	1.507561	MCCC1	1.45081	CCDC146	1.38976	HERC5	1.34201	FYCO1	1.29033	BCL7A	1.23442	NR1D2	1.186489
STX10	1.505713	SLC2A11	1.44915	ACSF3	1.38713	DTD2	1.3417	IRF5	1.28961	HK2	1.23369	TTC7A	1.186384
C4orf27	1.505542	LRP5	1.44867	TMCC1	1.3853	NCKIPSD	1.34165	LANCL1	1.28959	LCP1	1.23362	APOA	1.186073
PCYOX1	1.505212	GAL3ST4	1.44547	DOPEY2	1.38356	SH2D3C	1.34073	STAMBPL1	1.28796	KLF12	1.23282	TMEM141	1.184657
VPS13C	1.504844	FAM53C	1.44537	MC1R	1.38303	ABHD15	1.33811	PLEKHA2	1.28789	CEP131	1.23242	R3HDM1	1.184164
WDR66	1.504304	C11orf21	1.44487	TTC21A	1.38293	SLC2A8	1.33739	EIF4EBP1	1.28705	ACYP1	1.23164	SKP2	1.184103
CORO2A	1.503338	CD99L2	1.44478	CERS6	1.38238	ARMC10	1.33557	MED13L	1.28598	RTN4IP1	1.23144	HIBCH	1.183917
CCDC86	1.503094	C15orf52	1.44424	RFXAP	1.38022	FKBP15	1.33541	NT5DC1	1.28538	NT5DC3	1.23064	FUCA1	1.182975
TRAPPC12	1.50199	UMAD1	1.44379	FANCI	1.38004	IER2	1.3337	RECK	1.28506	TPCN1	1.23059	KBTBD7	1.182947
TEP1	1.50142	UBE2L6	1.44369	NHLRC4	1.37898	TMEM109	1.33358	ATF3	1.28267	NCOA2	1.23033	RHOB	1.182849
PEX7	1.500362	SCRN3	1.44338	KDM1B	1.37804	ATP10D	1.33264	POC1A	1.28256	UBE2S	1.22977	SNAPC3	1.18245
SLC2A6	1.499813	BBS9	1.44321	GGH	1.37799	DTX3	1.33227	EIF4G3	1.28051	TRIM8	1.22924	CEBPA	1.1822
TMEM209	1.499631	CDH23	1.44266	CASP8AP2	1.37701	SLC2A9	1.33211	ZNF703	1.28042	ZNRF2	1.2283	TSEN15	1.181335
SLC25A43	1.49809	STAU2	1.43939	ZDHHC21	1.37658	STXBP4	1.33156	ALKBH8	1.2793	CCDC138	1.22811	ZNF362	1.180281
PIP4K2A	1.497586	ZNF276	1.43881	CCND3	1.3762	PTPRJ	1.33133	GABRR2	1.27821	USP6NL	1.22708	MAST3	1.179821
RP11-164J13.1	1.497582	RBCK1	1.43778	MAP4K1	1.3747	SRPX	1.33113	TRIM44	1.27736	NCOA1	1.22556	TMTC4	1.178246
DEF8	1.497466	CNTRL	1.43607	NUP210	1.37412	MTMR12	1.33066	SORD	1.2772	RABL2B	1.22491	MAPRE2	1.177751
BLNK	1.496863	FBXW5	1.4351	RP11-298J23.10	1.37371	PON2	1.3305	FLNB	1.27664	GHDC	1.22439	SETDB2	1.177463
CLEC10A	1.496382	PPP3CA	1.43464	SLC47A1	1.37351	FGD3	1.33006	BTBD2	1.27578	HHAT	1.22406	KCNK1	1.176484
NADK	1.496107	TMEM251	1.43457	INTS10	1.37249	PRMT7	1.32966	PYURF	1.27549	OGG1	1.22364	PIP5K2B	1.175532
LDAH	1.49577	NSD1	1.4336	RASGRP2	1.37094	ARHGEF4	1.3294	ITPR2	1.27357	DERL3	1.22284	ARV1	1.175095
NIPSNAP1	1.493412	ZDHHC17	1.43348	MPPE1	1.37058	NSL1	1.32626	DACT1	1.27254	GCSAM	1.22265	PLCG2	1.174994
POP5	1.49192	CYSLTR1	1.43306	ALPK1	1.37057	STARD10	1.32512	CHD9	1.27196	TMEM128	1.22265	IFT140	1.173759
GJC2	1.491771	MYO1F	1.43281	MSH3	1.37051	CRTAP	1.32492	YPEL1	1.27095	SH3BP2	1.22128	ACBD6	1.173456
OSM	1.48953	KRCC1	1.43191	XKR6	1.36983	ZNF296	1.32474	PACS2	1.27079	GGCT	1.22029	FBXO44	1.172072
ADGRE2	1.488556	STRA13	1.43048	FBXO4	1.36943	ABI2	1.32467	RNF123	1.26771	C11orf80	1.21999	ZRANB3	1.171514
BANK1	1.484983	GLT25D1	1.42992	CNNM3	1.36918	SUMF1	1.32452	MLLT1	1.26655	CORO1B	1.21989	GZF1	1.171435
SLC7A6	1.484981	MBNL3	1.42966	PARN	1.36816	MTAP	1.32316	ATP6V0A1	1.26628	ADD1	1.21896	CUL4B	1.171112
TNF	1.483749	FTCDNL1	1.42892	KCNK17	1.36651	LRRC16A	1.32209	CREB3L2	1.26579	SPATA5	1.21787	PCYT2	1.170932
EHBP1	1.482924	MLH3	1.42873	RAP1GDS1	1.36577	FIG4	1.32071	NLRC5	1.2646	ATPIF1	1.21717	SCFD2	1.170774
KIAA0226L	1.481743	CAT	1.42804	SKAP2	1.36361	RNASEH2B	1.32037	DERA	1.26397	FHL1	1.21629	C4orf3	1.169135
CSRP2BP	1.480028	PRKCB	1.42731	LILRA2	1.36344	SNUPN	1.31963	SERPINI1	1.26364	GIMAP2	1.21599	RASGRP4	1.167615
IRF8	1.47941	VCL	1.42666	CECR1	1.36319	SOCS3	1.31816	CCDC183	1.26361	NAA38	1.21581	SNRNP25	1.167608
PLCL2	1.479323	KHDRBS2	1.42619	CCDC152	1.36038	PAM16	1.3165	GLIPR1	1.26102	RHBDD1	1.21514	DTX2	1.167594
BBS10	1.478157	SLC25A42	1.42553	FAM210B	1.36036	CPPED1	1.31636	ACAP1	1.26067	TMEM156	1.21448	ZNF540	1.167283
PIGH	1.477987	PPM1K	1.42522	TMEM91	1.36026	SPIB	1.31525	KIAA1109	1.26065	RFX3	1.21425	ATAD5	1.166984

CLUSTER 4													
DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)
IER5L	1.888718	DAB2	1.82526	VWA8	1.76562	IFFO2	1.7073	EPHB1	1.66958	TTC39C	1.62444	EMILIN1	1.583532
LYRM7	1.886612	DCPS	1.82297	POU4F1	1.76279	SLC18A2	1.70706	PTGS2	1.66908	LPGAT1	1.62373	IGSF22	1.580745
BIN2	1.883158	RSPH1	1.81947	FCER1A	1.76214	APC2	1.70675	QDPR	1.66878	USP46	1.62358	CDPF1	1.580136
TBC1D14	1.880879	SMIM5	1.81901	REER	1.75862	RAD51AP1	1.7037	TM7SF3	1.66867	PACSN1	1.6227	SPATA2L	1.579178
ZBTB18	1.877709	LIMD1	1.81851	SASH3	1.75674	MEF2C	1.7033	PNPLA4	1.66642	CCR5	1.62034	TCTA	1.579128
EPS8L2	1.877525	RPS6KL1	1.81843	CALHM2	1.75661	MYCBP2	1.70245	PRKCD	1.6658	MGAT4A	1.62031	TMEM182	1.578299
AFF3	1.875791	RNF213	1.81838	VWA2	1.75566	PIGK	1.70144	SORL1	1.66383	MYO9A	1.61778	C12orf66	1.577605
C2CD2	1.873632	LRRC26	1.81807	KLF13	1.7527	CYB561A3	1.70113	POLG2	1.66326	LDLRAD2	1.61685	DSN1	1.577044
PTGDS	1.87226	LDOC1	1.81616	FOXRED2	1.75108	EMB	1.7006	PAQR8	1.66255	GLO1	1.61532	CHEK2	1.573243
ARL11	1.868935	NGFR	1.81538	IKBKE	1.74228	SLC38A1	1.70025	C7orf31	1.66096	SLC37A1	1.61459	TYSND1	1.5732
ANXA2R	1.866682	CBFA2T3	1.81484	PFKFB2	1.74188	HPS3	1.6986	ARHGAP17	1.65998	FCER1G	1.61418	ZNF594	1.57316
EME1	1.86617	NOTCH3	1.80938	RASSF7	1.74131	PSD4	1.69803	CD79B	1.65971	KRI1	1.61302	CSGALNACT1	1.571502
ANKRD44	1.863976	METRN	1.80861	CTSO	1.74054	MAPKAPK3	1.697	DOCK10	1.6593	ATXN7L1	1.6104	DGKZ	1.57048
JAK2	1.86384	DHFR	1.80676	ST6GAL1	1.74023	BCL11A	1.69342	FBXO41	1.65913	NBEA	1.60724	THEM4	1.570199
AEBP1	1.860769	PTER	1.80444	VPS36	1.73997	CD164	1.69284	OSGEPL1	1.65882	THTPA	1.60451	MPEG1	1.568859
ELMSAN1	1.855011	PYCARD	1.79871	CROT	1.73877	MPIG6B	1.6914	PLCB2	1.65786	WDR81	1.60369	RUNX2	1.560398
PTPN6	1.854454	43160	1.79791	PLK2	1.73598	ZDHH23	1.69101	C14orf159	1.65237	KLHL13	1.60041	STK11IP	1.559116
NAT8L	1.854275	PLXNA4	1.79616	ALAD	1.73312	NIN	1.68967	SNTA1	1.65172	RFLNB	1.60029	MSL2	1.558554
FAM213A	1.854115	EHMT2	1.79449	SMAD3	1.73078	PLPP1	1.68916	APOBEC3F	1.65166	POMGNT1	1.59956	TBC1D5	1.556834
SLITRK5	1.850623	SCARB1	1.7935	ZNF589	1.72949	DEF6	1.68562	EPDR1	1.6496	MOSPD2	1.5993	S100PBP	1.556288
CRIP1	1.849202	SLC2A1	1.7933	NAALAD2	1.72779	BTLA	1.68527	CPT1A	1.64752	CMTM3	1.59803	FAM160A1	1.55619
ADAMTS4	1.849086	FCRLA	1.791	CLN8	1.72475	CHURC1	1.68428	SCAI	1.64711	TECPR1	1.59765	HHLA3	1.555042
CARD11	1.846084	NREP	1.78928	SEMA4D	1.72415	PDSS2	1.68403	THRA	1.64486	UBE2E2	1.59668	ST3GAL4	1.554627
IMPACT	1.844453	VMAC	1.7846	CA8	1.72237	MARVELD1	1.6823	SLC27A3	1.64139	GATB	1.59303	WDFY4	1.554404
DHODH	1.844429	C16orf54	1.78368	TXNDC5	1.71947	TMIGD2	1.67871	TTL12	1.64035	CETP	1.59272	DBP	1.553732
HDAC9	1.843091	GNF7	1.78234	APOBR	1.71827	OAS1	1.67797	UBE2J1	1.6401	STK17B	1.59126	GUCY1A3	1.553269
MCC	1.840186	MYO1E	1.78201	FLOT2	1.71454	TMEM144	1.67659	SIRPB1	1.63585	SIGLEC6	1.59124	TMEM107	1.551555
ZNF607	1.839793	PPM1M	1.7774	FAM221B	1.71318	SORBS3	1.6764	ARHGEF6	1.63206	TMEM242	1.5902	HACD4	1.550519
KIF17	1.838859	SYNJ2BP	1.77432	ZNF438	1.71051	FAM117A	1.67529	MMP25	1.63001	ALDH3B1	1.58987	FCMR	1.55042
SCRN1	1.836454	TMEM53	1.77337	ERMP1	1.71025	STIM1	1.67422	SLC20A1	1.62694	RCCD1	1.58941	MED12L	1.549428
PPP1R13L	1.834954	BBC3	1.77144	LYL1	1.71021	C3orf58	1.67368	PREX1	1.62592	LRRC1	1.58891		
PARP10	1.833772	P3H2	1.771	ARHGDB	1.70923	SLC44A2	1.67222	FAN1	1.62464	RNASE6	1.5886		
SCML2	1.828339	PPM1F	1.76911	RNASEL	1.70734	COL24A1	1.67026	POLA1	1.62451	DUSP1	1.5875		
TMEM210	1.826608	ZNF469	1.76711										
CLUSTER 5													
DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)	DEG	FC (log ₂)
ITGB8	-7.35802	TNFRSF12A	-4.9512	NKD2	-4.2715	KHDRBS3	-3.7598	ST8SIA4	-3.52093	RGS9	-3.3157	C17orf99	-3.13554
ECCL1	-6.51225	CXCL8	-4.8359	RAMP1	-4.0276	TOGARAM2	-3.7274	RAB7B	-3.5052	ANKRD33B	-3.2748	NINJ1	-3.1196
AVP	-6.31372	C15orf48	-4.6744	IL4I1	-4.0051	KSR1	-3.7216	CLECL1	-3.48558	MACC1	-3.2643	SMARCA1	-3.09589
TNFRSF9	-5.81803	SPSB1	-4.5451	PEL1	-3.9999	LMNA	-3.6825	TNFSF4	-3.48496	NUAK2	-3.2569	AIM1	-3.08527
TRH	-5.74118	CTNNAL1	-4.4758	BCAN	-3.9921	DEPDC7	-3.6805	CACNA1G	-3.4564	SIPA1L1	-3.2176	ADTRP	-3.07417
CCR7	-5.69939	INHBA	-4.4576	LGALS3	-3.8467	WNT9A	-3.6771	DKK4	-3.44069	CD9	-3.2117	CD69	-3.04501
GPR157	-5.6783	BMP6	-4.4531	MARCKS	-3.819	B3GNT5	-3.6393	TGIF1	-3.37978	FAM222A	-3.2036	IKZF2	-3.03664
CCL22	-5.28399	NRP2	-4.41	NCMAP	-3.8152	KCNE5	-3.6232	AHR	-3.37825	NRIP3	-3.196	BIRC3	-3.02975
PALLD	-5.07311	DUSP4	-4.3881	SLC6A12	-3.7936	TNFRSF18	-3.5903	SERPINB9	-3.34639	TFPI	-3.1828	CTSL	-3.02006
BCL2A1	-5.06808												

*Relative fold-change (log₂ values) of DEG using non-TME samples (dLNneg) as the reference. DEG = differentially expressed genes. FC = fold-change.

Supplemental Table 5: Flow cytometry antibodies

FACS-Sorting Cocktails (Human)					
Antigen	Fluorochrome	Clone	Company	Catalog No.	Dilution
Live/Dead	Aqua (405 nm excitation)	n/a	ThermoFisher	L34965	1:100
CD45	BUV395	HI30	BD Biosciences	563792	1:40
HLA-A2	V450	BB7.2	BD Biosciences	561344	1:30
HLA-A2	PE	BB7.2	BD Biosciences	558570	1:30
HLA-DR	APC-Cy7	L243	BioLegend	307618	1:25
CD11c	BUV395	B-ly6	BD Biosciences	563787	1:20
CD11c	APC	3.9	BioLegend	301614	1:20
CD123	PE	7G3	BD Biosciences	554529	1:30
CD303	PerCP-Cy5.5	BDCA-2	BioLegend	354209	1:20
CD134/OX40	BV421	ACT35	BioLegend	350014	1:30
CD134/OX40	APC-Cy7	ACT35	BioLegend	350022	1:30
FACS-Sorting Cocktails (Mouse)					
Antigen	Fluorochrome	Clone	Company	Catalog No.	Dilution
Live/Dead	Aqua (405 nm excitation)	n/a	ThermoFisher	L34965	1:100
CD45	Pacific Blue	30-F11	BioLegend	103126	1:40
CD45R/B220	APC-Cy7	RA3-6B2	BioLegend	103236	1:50
CD317/BST2/ PDCA-1	PE	129c	ThermoFisher,	12-3171-82	1:30
CD11c	PE-Cy7	N418	BioLegend	117318	1:40
CD11b	FITC	M1/70	BioLegend	101206	1:50
Flow Cytometry Phenotyping Antibodies (Human)					
Antigen	Fluorochrome	Clone	Company	Catalog No.	Dilution
Live/Dead	Aqua (405 nm excitation)	n/a	ThermoFisher	L34965	1:100
CD45	BUV395	HI30	BD Biosciences	563792	1:40
CD3	APC-R700	UCHT1	BD Biosciences	565119	1:40
CD3	PE-CF594	SP34-2	BD Biosciences	562406	1:40
CD8	BB515	RPA-T8	BD Biosciences	564526	1:40
CD8	AlexaFluor647	RPA-T8	BD Biosciences	557708	1:40
CD4	PerCP-Cy5.5	OKT4	BioLegend	317428	1:40
HLA-DR	APC-Cy7	L243	BioLegend	307618	1:25
HLA-DR	BV421	G46-6	BD Biosciences	562805	1:30
Lineage (CD3/CD14/ CD19/CD20/ CD56)	FITC	UCHT1/ HCD14/HIB1 9/ 2H7/ HCD56	BioLegend	348701	1:40
CD11c	BUV395	B-ly6	BD Biosciences	563787	1:20
CD11c	APC	3.9	BioLegend	301614	1:20
CD123	PE	7G3	BD Biosciences	554529	1:30
CD303	PerCP-Cy5.5	BDCA-2	BioLegend	354209	1:20
CD134/OX40	BV421	ACT35	BioLegend	350014	1:30
CD134/OX40	APC-Cy7	ACT35	BioLegend	350022	1:30
CD252/OX40(L)	BV421	ik-1	BD Biosciences	563766	1:20
CD275/ICOS(L)	PE-CF594	2D3/ B7- H2	BD Biosciences	564277	1:25
CD40	PE	HB14	BioLegend	313006	1:30
CD80	APC	2D10	BioLegend	305220	1:30
CD86	AlexaFluor700	2331/ FUN-1	BD Biosciences	561124	1:30
CD25	AlexaFluor488	M-A251	BioLegend	356116	1:30
CD137/4-1BB	PE-Cy7	4B4-1	BioLegend	309818	1:30
CD253/TRAIL	BV421	RIK-2	BD Biosciences	564243	1:30
CD327/Siglec-6	APC	REA852	Miltenyi Biotec	130112711	1:20
Axl	AlexaFluor700	FAB154N	R&D	108724	1:20
CD69	APC/Fire750	FN50	BioLegend	310946	1:30
CD11b	BB515	ICRF44	BD Biosciences	564517	1:30
CD14	PerCP-Cy5.5	HCD14	BioLegend	325622	1:30
CD15	AlexaFluor700	W6D3	BioLegend	323026	1:30
CD68	PE-Cy7	Y1/82A	BioLegend	333816	1:30
Tbet	PerCP-Cy5.5	4B10	BioLegend	644806	1:25
IFN-γ	APC-Cy7	4S.B3	BioLegend	502530	1:25
FoxP3	PE	259D/C7	BD Biosciences	560046	1:30
IFN-γ	BV711	B27	BD Biosciences	564039	1:25
IL-12p70	PE	20C2	BD Biosciences	559325	1:25
Granzyme B (GzB)	AlexaFluor700	GB11	BD Biosciences	560213	1:30
GzB	FITC	GB11	BioLegend	515403	1:30

Flow Cytometry Phenotyping Antibodies (Mouse)					
Antigen	Fluorochrome	Clone	Company	Catalog No.	Dilution
CD45	BUV395	30 F11	BD Biosciences	564279	1:40
CD3	AlexaFluor700	17A2	BioLegend	100216	1:40
CD11b	FITC	M1/70	BioLegend	101206	1:50
CD11c	PE-Cy7	N418	BioLegend	117318	1:40
CD45R/B220	PerCP-Cy5.5	RA3-6B2	BioLegend	103236	1:50
CD317/BST2 PDCA-1	PE	129c	ThermoFisher	12-3171-82	1:30
CD8a	BV421	53-6.7	BioLegend	100737	1:40
CD4	APC-Cy7	GK1.5	BioLegend	100414	1:50
CD134/OX40	APC	OX86	BioLegend	119414	1:30
CD252/OX40(L)	eFluor710	RM134L	ThermoFisher	46590580	1:25
CD252/OX40(L)	PE-Cy7	RM134L	BioLegend	108813	1:20
CD103	PE-CF594	M290	BD Biosciences	565849	1:20
Ly-6G	APC	1A8	BioLegend	127614	1:25
Ly6c	FITC	HK1.4	BioLegend	128006	1:40
Gr1	APC-Cy7	RB6-8C5	BioLegend	108424	1:40
FoxP3	PE	259D/C7	BD Biosciences	560046	1:30
TNF- α	APC	MP6-XT22	BioLegend	506308	1:40
IL-12p40	PE	C15.6	BioLegend	505204	1:30
IFN- γ	PE-Cy7	XMG1.2	BioLegend	505826	1:30