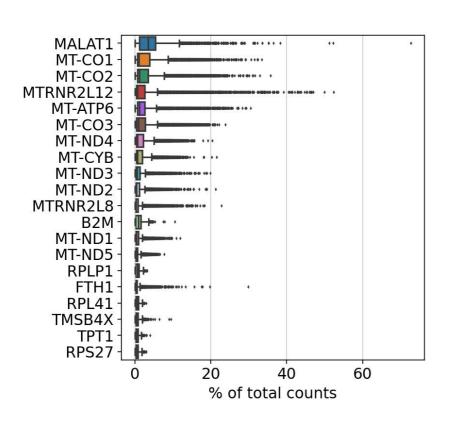
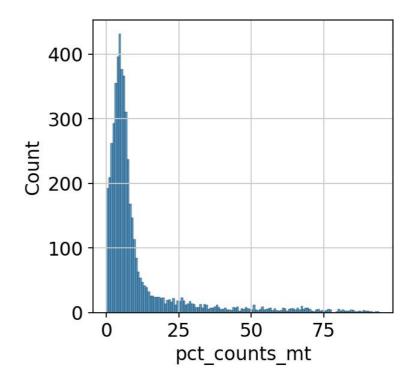
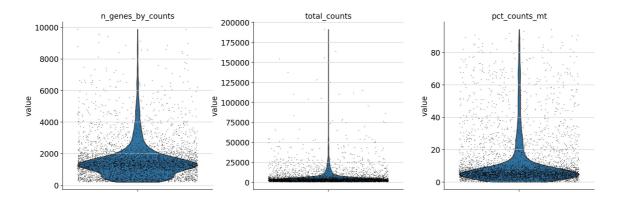
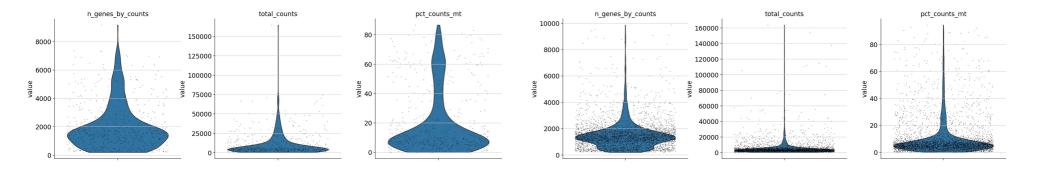
Preprocessing



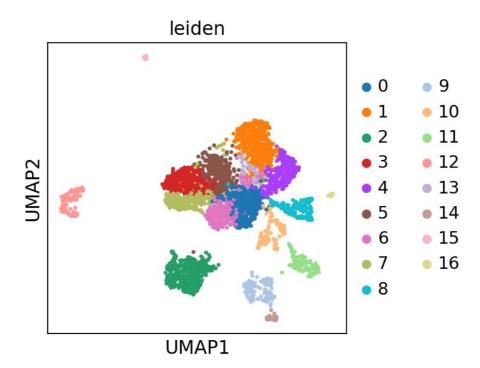


Violin plots of Quality Measures

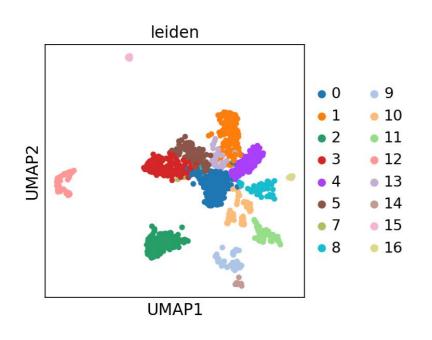


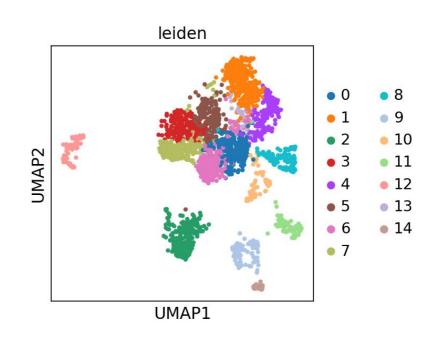


Clustering

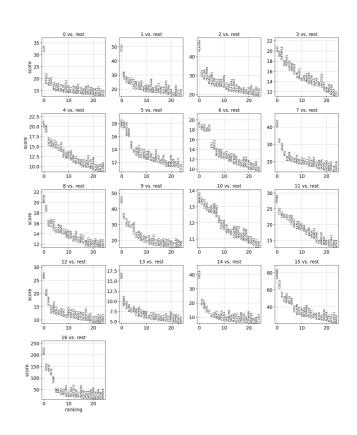


76M cells vs 76V cells

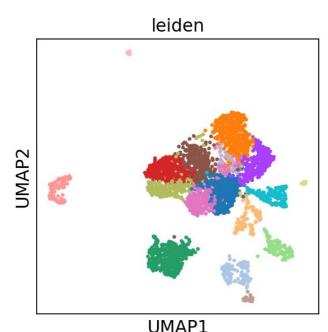




Ranking gene Groups



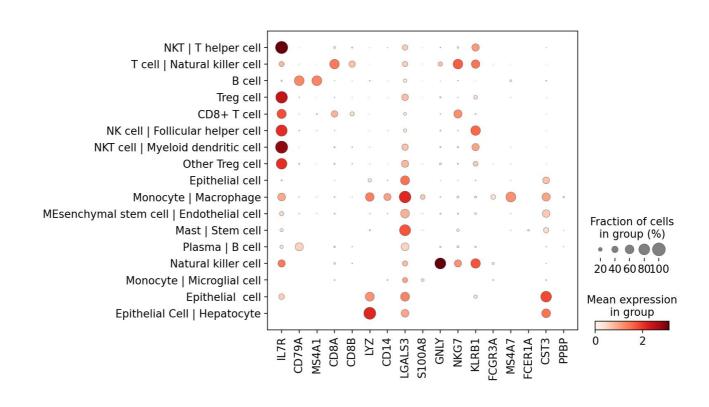
Labelling Clusters through SCSA



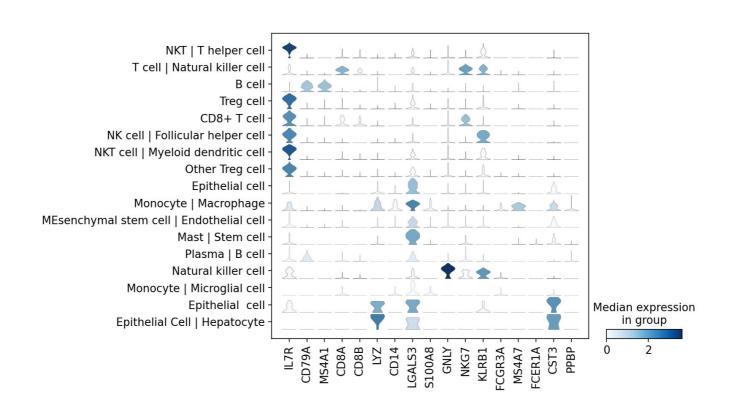
- NKT | T helper cell
- T cell | Natural killer cell
- B cell
- Treg cell
- CD8+ T cell
- NK cell | Follicular helper cell
- NKT cell | Myeloid dendritic cell
- Other Treg cell
- Epithelial cell

- Monocyte | Macrophage
- MEsenchymal stem cell | Endothelial cell
- Mast | Stem cell
- Plasma | B cell
- Natural killer cell
- Monocyte | Microglial cell
- Epithelial cell
- Epithelial Cell | Hepatocyte

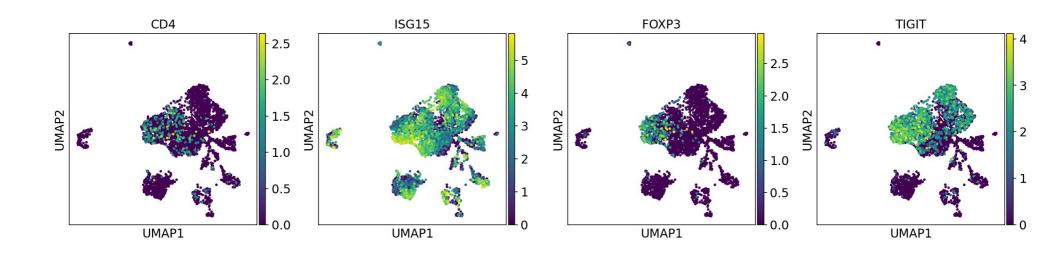
Gene Markers of labelled clusters



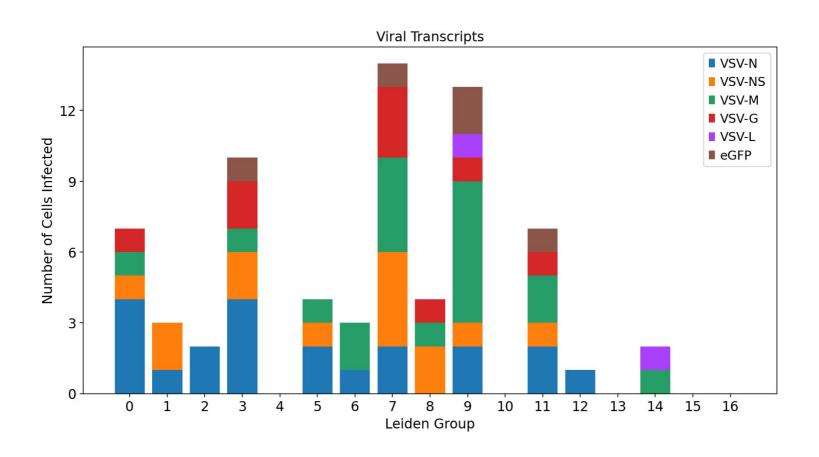
Cont.



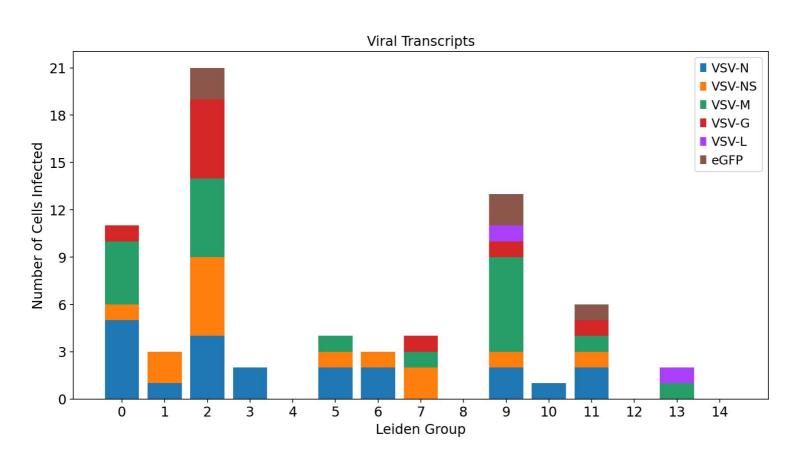
Viral Infection with VSVd51



Counts of Viral Transcripts



76V Transcripts



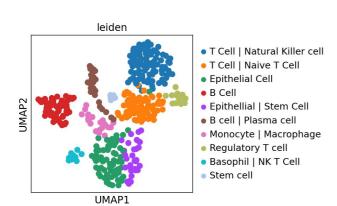
76 Aggregate Upregulated Genes by Cluster

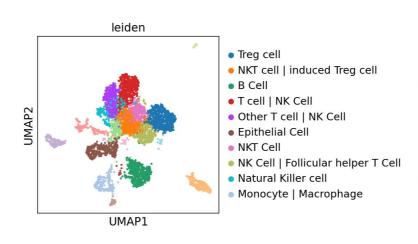
NKT T helper cell	T cell Natural killer cell	B cell	Treg cell	CD8+ T cell	NK cell Follicular helper cell	NKT cell Myeloid dendritic cell	Other Treg cell	Epithelial cell	Monocyte Macrophage	MEsenchymal stem cell Endothelial cell	Mast Stem cell	Plasma B cell	Natural killer cell	Monocyte Microglial cell	Epithelial cell	Epithelial Cell Hepatocyte
IL7R		HLA-DRA			FKBP5					MARCKS						MT1G
ZFP36L2																MT2A
RPL13																MT1X
CXCR4																MT1E
RPS12																TXNIP
RPS3A																UBC
ANXA1										HSP90B1						JUN
RPL32																EGR1
RPL34																HES1
LEPROTL1																SLC30A1
RPL39		HLA-DPB1														IER2
TPT1																PHLDA2
RPS18			PMAIP1													HIST1H1C
RPL30																KRT18
TNFAIP3						EIF2AK2										ATF3
RPL21																HSP90AB1
RPS6											GCSAML					H3F3B
SNHG6																ZFAS1
RPS27A				LEPROTL1												SAP18
RPS16																\$100A6

76 Downregulated Genes By Cluster

NKT T helper cell	T cell Natural killer cell	B cell	Treg cell	CD8+ T cell	NK cell Follicular helper cell	NKT cell Myeloid dendritic cell	Other Treg cell	Epithelial cell	Monocyte Macrophage	MEsenchymal stem cell Endothelial cell	Mast Stem cell	Plasma B cell	Natural killer cell	Monocyte Microglial cell	Epithelial cell	Epithelial Cell Hepatocyte
HLA-DRA																
										PCED1B-AS1						
					B MARCKS	S HLA-DQB1										TSC22D3
HLA-DRB1						1 HLA-DRB1										
HLA-DPA1					HLA-DQA1	1 HLA-DPB1										
HLA-DQA2					1 HLA-DRA	A HLA-DQA1										

Leiden Clustering of 56 and Total Aggregate





- B cell | plasma cell
- ?
- Mesenchymal stem cell | Neutrophil
- Mast cell | stem cell
- Monocyte | Epithelial Cell
- Hepatocyte | Epithilial Cell
- Other Epithelial cell
- NKT cell
- B Cell | Myeloid Dendritic cell

Aggregated Upregulated Genes by Cluster

Treg cell	NKT cell induced Treg cell	B Cell	T cell NK Cell	Other T cell NK Cell	Epithelial Cell	NKT Cell	NK Cell Follicular helper T Cell	Natural Killer cell	Monocyte Macrophage	B cell plasma cell	Mesenchymal stem cell Neutrophil	Mast cell stem cell	Monocyte Epithelial Cell	Hepatocyte Epithilial Cell	Other Epithelial cell	NKT cell	B Cell Myeloid Dendritic cell
							KIAA0319L										
								TMSB4X			MARCKS						
		HLA- DQB1															
														DNAJB1			
		HLA- DRB1															
12 MAF						TMSB4X											
		HLA- DPB1															
														DNAJB4			
		HLA- DPA1															
		HLA- DQA1															

Agreggate Downregulated Genes By Cluster

Treg cell	NKT cell induced Treg cell	B Cell	T cell NK Cell	Other T cell NK Cell	Epithelial Cell	NKT Cell	NK Cell Follicular helper T Cell	Natural Killer cell	Monocyte Macrophage	B cell plasma cell	Mesenchymal stem cell Neutrophil	Mast cell stem cell	Monocyte Epithelial Cell	Hepatocyte Epithilial Cell	Other Epithelial cell	NKT cell	B Cell Myeloid Dendritic cell
SPINT2						HLA- DQB1								PCED1B-AS1			
									PCED1B-AS1								
	HLA-DQA2				SAMSN1	RAB13				SPOCK2							
MARCKS						HLA- DQA2					PCED1B-AS1						
ANXA1																	

Sample Normalized Ratios per Cluster

	56M	56V	76M	76V
0	0.092767	0.127132	0.309367	0.470733
1	0.362067	0.068703	0.356873	0.212357
2	0.303835	0.165068	0.291444	0.239653
3	0.126079	0.172785	0.303351	0.397785
4	0.223763	0.383318	0.227456	0.165464
5	0.525381	0.309178	0.058795	0.106647
6	0.010482	0.594723	0.000000	0.394795
7	0.106918	0.075356	0.405101	0.412626
8	0.348259	0.286361	0.203666	0.161714
9	0.188605	0.436172	0.193916	0.181306
10	0.271450	0.407590	0.174341	0.146618
11	0.213259	0.242800	0.330445	0.213497
12	0.186501	0.345046	0.318284	0.150169
13:	0.151524	0.103828	0.440878	0.303770
14	0.000000	0.184275	0.263333	0.552392
15	0.422909	0.069549	0.340755	0.166787
16	0.000000	0.000000	0.968400	0.031600
17	0.286510	0.235587	0.336661	0.141242
18	0.000000	0.764607	0.052031	0.183362

56 Upregulated Genes

4.2	. II I M I W. II II	TC-011 N . TC 0	e-51-11-1 e-0	D.C. !!	F :- 11: 11: 51 - 5 W	B		D	B. Lillaux Bell	C
	ell Natural Killer cell			B Cell			Monocyte Macrophage	Regulatory T cell		Stem cell
0	CCL5	RPL32	C19orf33	HLA-DRA	S100A14	SSR4	FTH1	B2M	CD63	PFDN5
	PTPRC	RPS12	KRT19	CD74	C19orf33	MZB1	C15orf48	MAF	ANXA5	GPAT4
2	TNFAIP3	RPS3A	CEACAM5	RPS11	KRT18	TXNDC5	FTL	IL2RA	DAD1	CREM
	TMSB4X	RPL13	LCN2	TNFRSF13C	YWHAZ	DERL3	SAT1	SRGN	SEC61G	EEF1A1
	CD3D	RPS2	CLDN4	RPL18A	ACTG1	HERPUD1	SOD2	LINC01943	C11orf96	FTH1
	ZFP36L2	RPL34	LMO7	RPS23	SPINT2	LY96	CXCL8	HLA-A	FGFR1	VIM
	CD8A	PTPRC	EPCAM	CD37	EPCAM	SEC11C	TYROBP	AC017002.3	TMED5	PLIN2
	CST7	RPL27A	SPINT2	RPL13A	CYSTM1	IGHA1	TXN	TNFRSF18	PDIA3	RPL34
	PTPN22	RPL28	KRT8	MS4A1	S100A6	PRDX4	CTSD	ITM2B	GEM	PSMB1
	CD3G	RPS6	LAMB3	CD83	GPRC5A	GADD45A	CREG1	UQCRB	PHLDA1	RPS4X
10	CD7	RPS3	GPRC5A	HLA-DRB1	S100P	HIST1H1C	HLA-DRA	S100A4	LMAN1	CD63
11	NKG7	RPS23	IFI27	RPL23A	ELF3	ANKRD28	G0S2	TIGIT	HSP90B1	ENO1
12	CXCR4	RPS15A	ELF3	RPS18	CTTN	JCHAIN	CXCL3	IL32	CALU	ALKBH5
13	TRBC2	IL7R	DSTN	RPS27	CLDN3	TENT5C	GAPDH	RPL24	TCEAL9	TAGLN2
14	SRGN	SARAF	S100A14	RPL9	KRT19	FKBP11	CD14	CD2	SQSTM1	SAMSN1
15	CD2	RPL13A	ANXA3	CD79A	KRT8	VOPP1	PLEK	CALM1	VEGFA	HNRNPC
16	SRSF7	RPL11	CLDN7	MEF2C	ANXA2	CUTA	LGALS1	C17orf49	ARF4	SRGN
17	PIK3R1	RPS8	CEACAM6	LAPTM5	S100A16	DANCR	SPP1	GBP5	GSTO1	TPSB2
18	RUNX3	RPL36	SDCBP2	RPS27A	CEACAM5	SPCS2	FCER1G	CD247	NEAT1	EMP3
19	FYN	RPS27A	IGFBP3	HLA-DPB1	RNF114	FCRL5	NPC2	TMSB10	IER3	PTP4A2

56 Downregulated Genes

T Cell Natural Killer cell	T Cell Naive T Cell	Epithelial Cell	B Cell	Epithellial Stem Cell	B cell Plasma cell	Monocyte Macrophage	Regulatory T cell	Basophil NK T Cell	Stem cell
EPCAM	FAM3C	CD52	CAMK2N1	GMFG	TNFAIP3	GSPT1	TUBA1C	SAMSN1	MRPL41
C19orf33	KRT8	LAPTM5	KRT18	LAPTM5	PIK3R1	CD7	TSC22D1	MBP	PPA1
S100A14	CD63	CD53	CD3G	TUBA1A	CD3D	CD2	NPEPPS	IL2RG	KRAS
CLDN4	GPRC5A	CYTIP	ID2	CYTIP	CD7	NSMCE3	CD9	CD74	CD47
KRT18	CD74	CREM	CD2	VIM	CD3E	CNOT6L	TES	OCIAD2	MT1X
KRT8	C19orf33	TSC22D3	CD3E	CD53	REL	PIK3IP1	suco	LAPTM5	ANXA2
VEGFA	MARCKS	PTPRC	CD7	CXCR4	CD52	CD3D	GARS	CYTIP	STK17A
KRT19	VEGFA	SRGN	CD3D	TSC22D3	TRBC2	SYNE2	SPINT2	STK17B	CHURC1
MARCKS	HLA-DRB1	CXCR4	ANXA1	SRGN	FOXP1	TUBA1A	HMGA1	CD53	IL32
SPINT2	HLA-DRA	VIM	IL32	PTPRC	ETS1	RESF1	RHOB	PTPRC	ISG20

Heatmap of Upregulated Genes

