

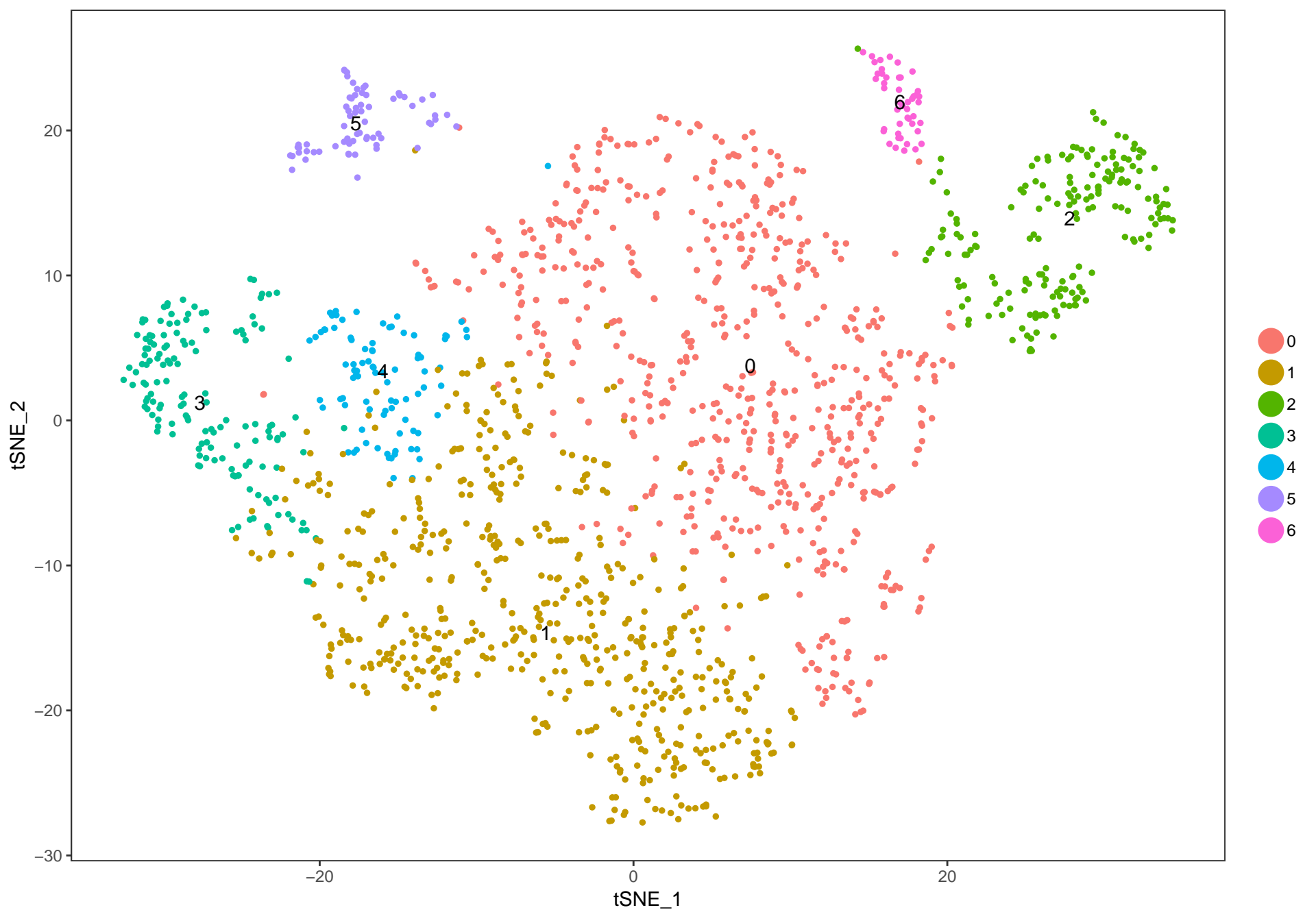
T-cell subcluster vs Response group

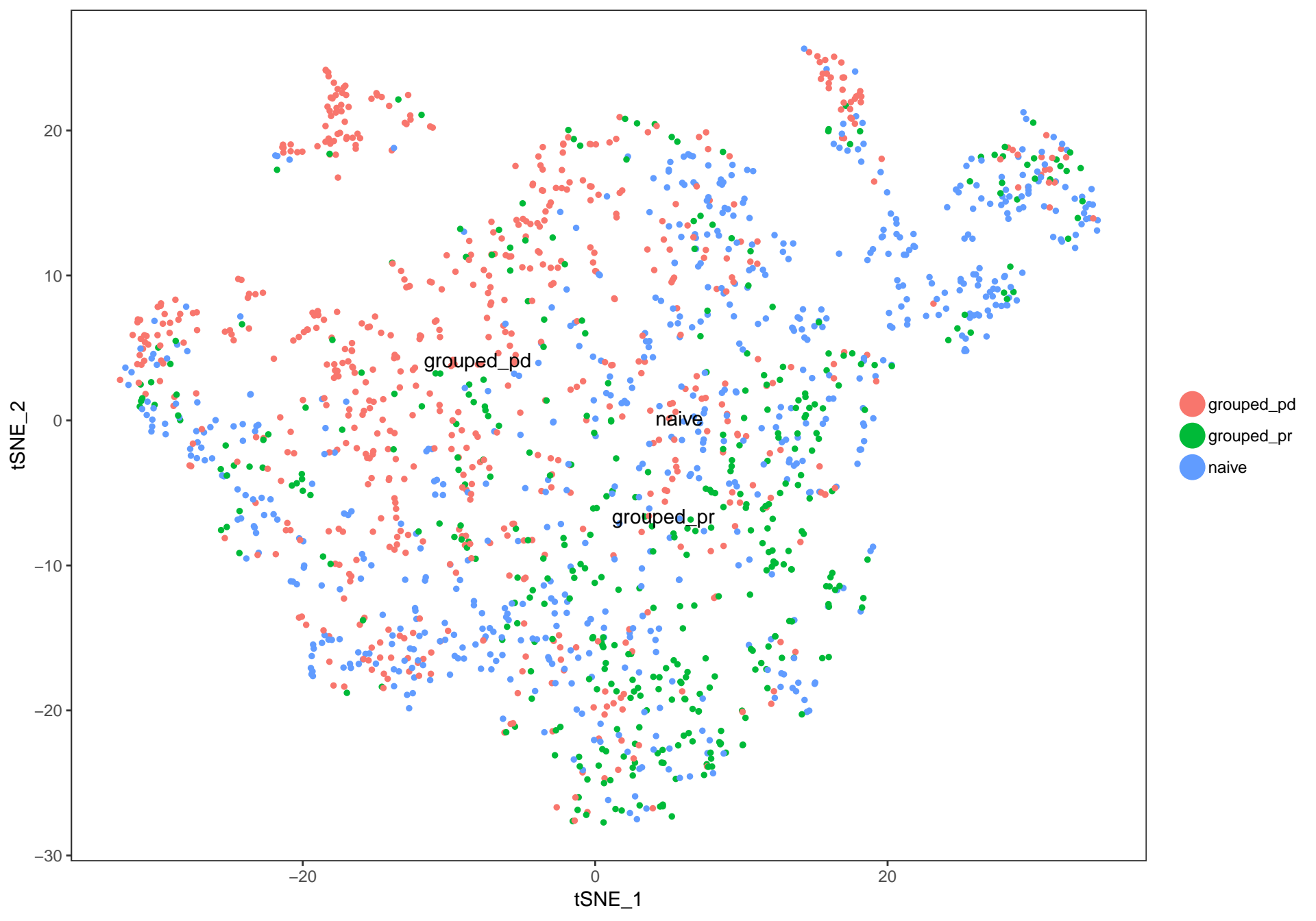


grouped_pd	213	185	21	64	79	61	30
grouped_pr	165	169	29	23	6	4	5
naive	250	208	135	60	3	4	11

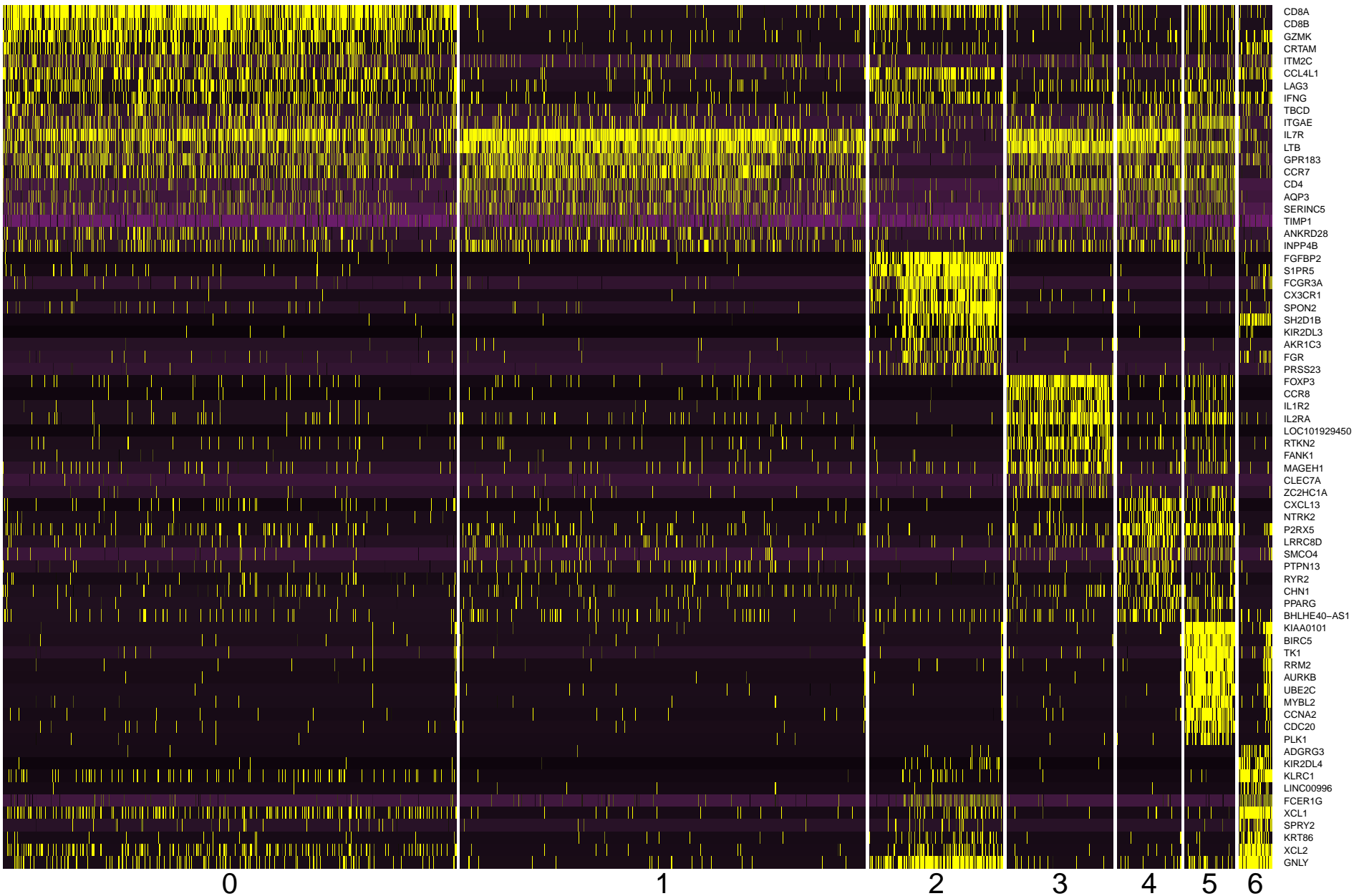
Patient ID vs T-cell cluster

	0	1	2	3	4	5	6
TH158	38	29	4	1			2
TH169	10	6	1	3	1		
TH179	147	128	4	21	66	49	17
TH179_NAT	11	18	16		5	3	1
TH226	110	99	20	48	8	11	15
TH231	1	3					1
TH236	62	74	5	14	5	2	
TH238	132	112	24	39	1	3	1
TH238_NAT	55	31	74	1		1	7
TH248	62	62	37	20	2		2

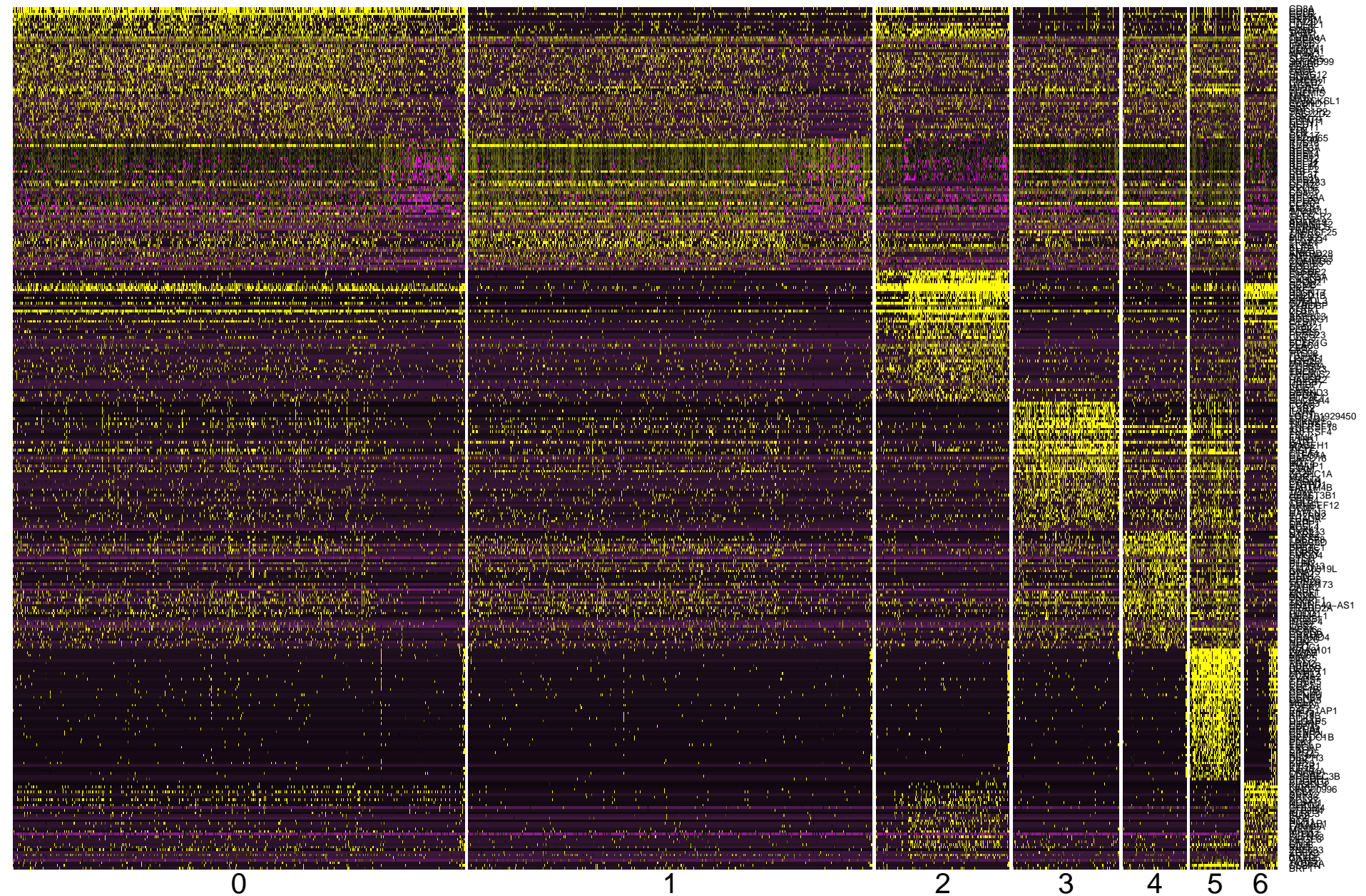


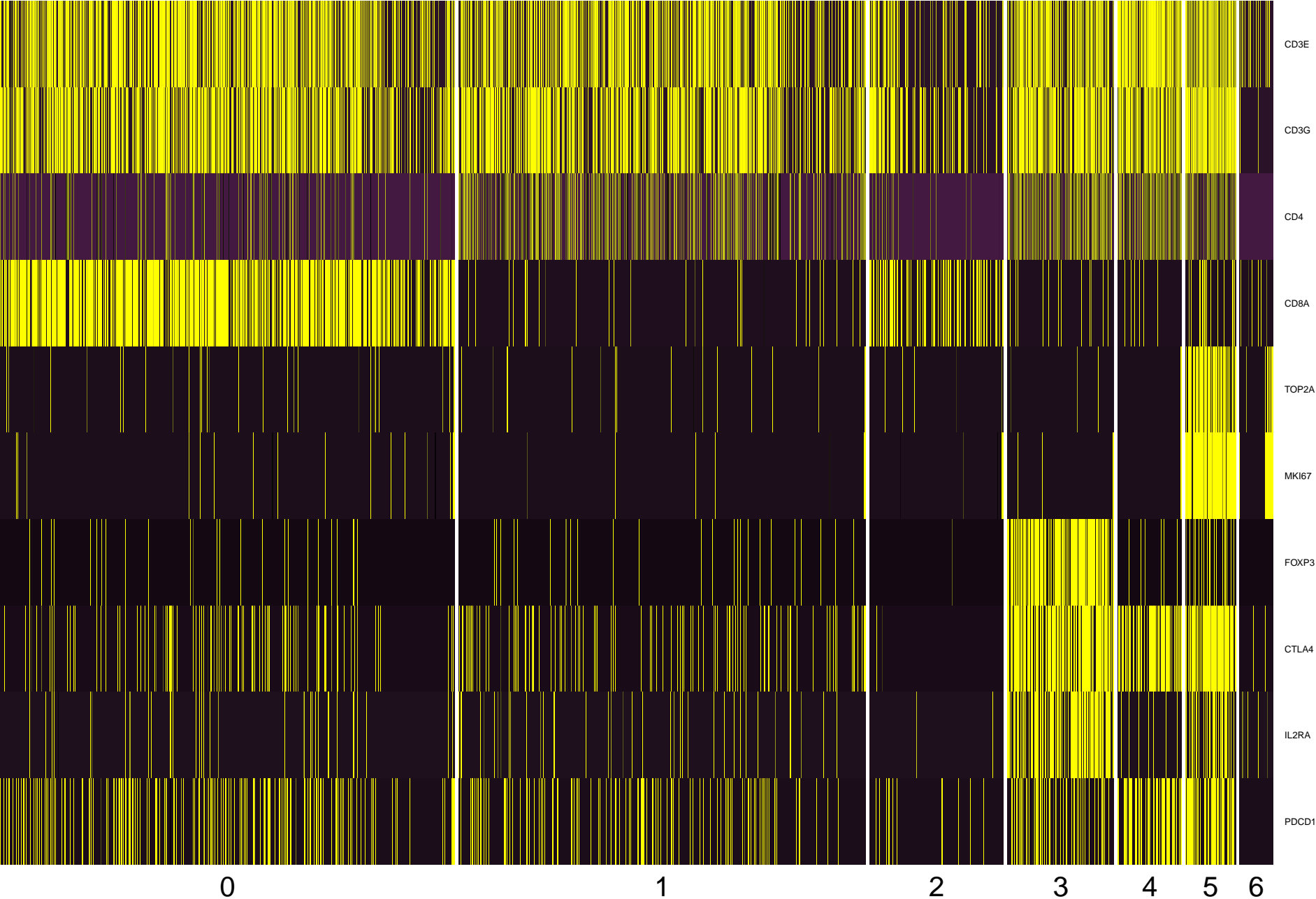


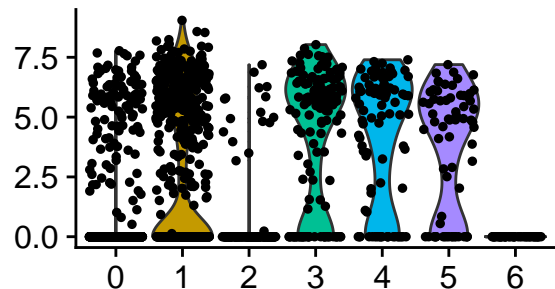
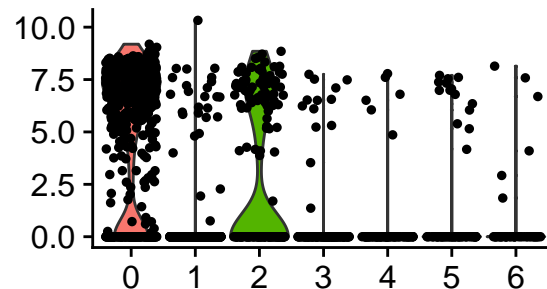
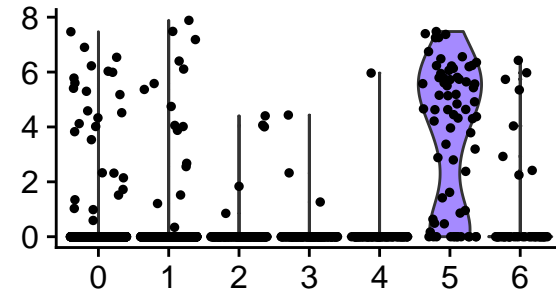
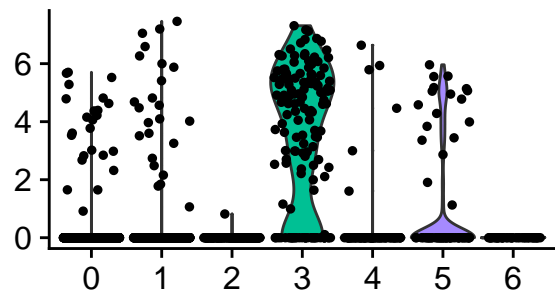
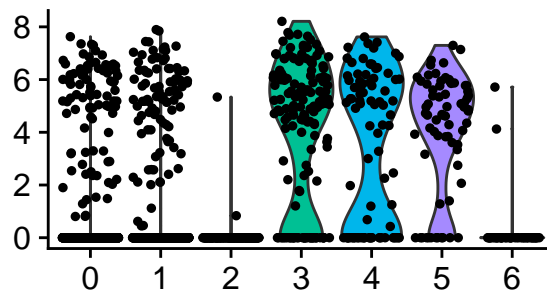
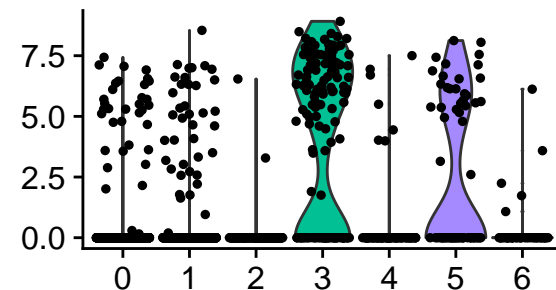
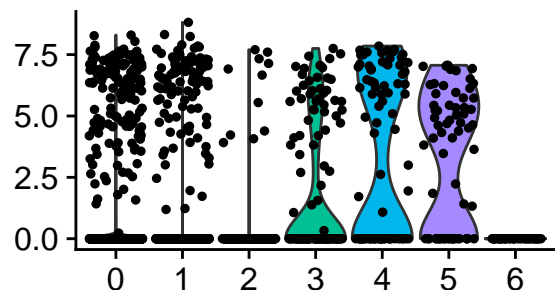
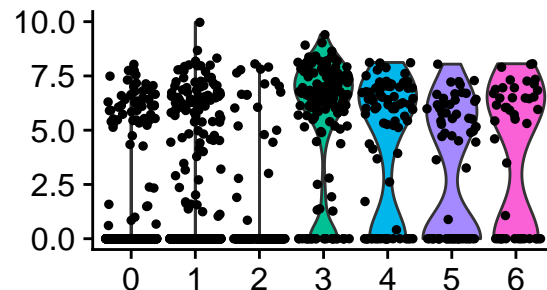
Top 10 DE genes



Top 50 DE genes





CD4**Identity****CD8A****Identity****TOP2A****Identity****FOXP3****Identity****CTLA4****Identity****IL2RA****Identity****PDCD1****Identity****TNFRSF18****Identity**

PC 1



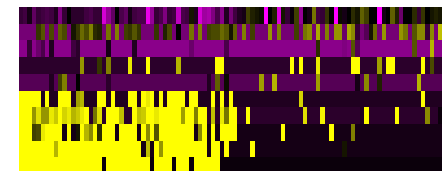
CST3
GPR5
GZMB
SH2D1B
TNFRSF4
TNFRSF25
CCR6

PC 2



EGFBBP2
S1PR5
GZMB
SH2D1B
TNFRSF4
TNFRSF25
CCR6

PC 3



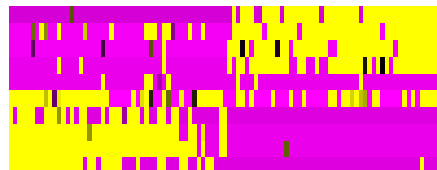
FTL
ANXA1
CST3
ZNF165
DHP53
TNFRSF1
TNFRSF4
CCR6
CCR6

PC 4



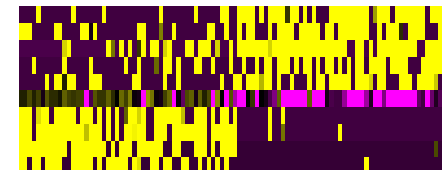
EGFBBP2
S1PR5
CCR8
SH2D1B
TNFRSF4
GZMK
ZNF683
CRTAM
CD8A
CD8B

PC 5



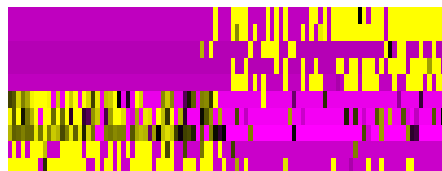
CCR8
TIGIT
TNFRSF9
LAYN
CRTAM
TNFRSF25
HIST1H2AJ
RPM2
GPR25

PC 6



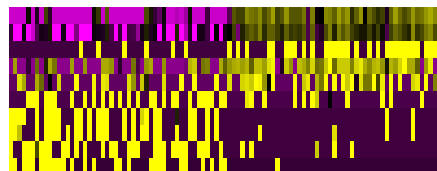
GZMK
CRTAM
CD8A
GZMH
EOMES
HSPA1A
XCL2
XCL1
KLRC1
ZNF683

PC 7



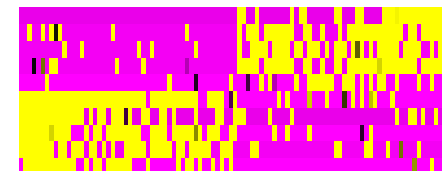
RPM2
MK167
KLRC1
HIST1H2AJ
CDC20
SLC7A5P2
TNFRSF8
CCR6

PC 8



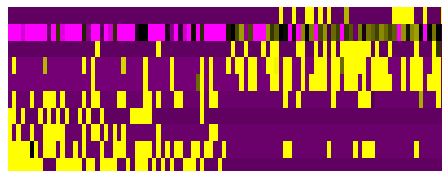
HSPA1B
HSPA1A
IFNG
DEDD2
DOK2
CCR6
XCL2
XCL1
CXCR6
CXCL13

PC 9



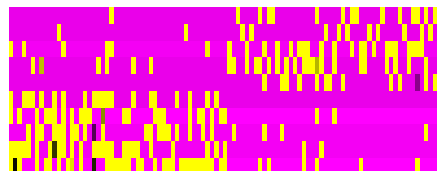
SH2D1B
XCL2
EOMES
XCL1
GZMK
CD8A
ZNF683
CD8B
GZMH
CXCR6

PC 10



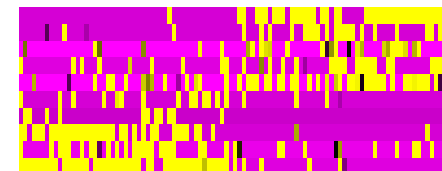
CCR8
NMDA
KLRC1
CD8A
GZMH
GPR25
FCRL6
IFNG
CXCL13

PC 11



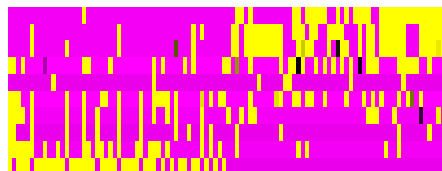
CXCL13
CCR8
IFNG
ZNF683
ZBED2
KLRC1
GZMK
EOMES
FASLG
CXCR6

PC 12



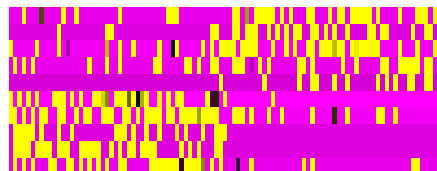
ZNF683
FCRL6
CCR8
CRTAM
OAS1
IL23R
FASLG
CXCR6
IFNG

PC 13



FCRL6
XCL2
CXCL13
CCR8
GZMB
FASLG
KLRC1
GZMH
CXCL13

PC 14



CXCR6
IFNG
CCR8
ZBED2
TNFRSF25
CD8A
NCR3
TARP

PC 15



FASLG
KLRC1
NCR3
NCR3
NCR3
XCL2
GZMH
KLRC1

