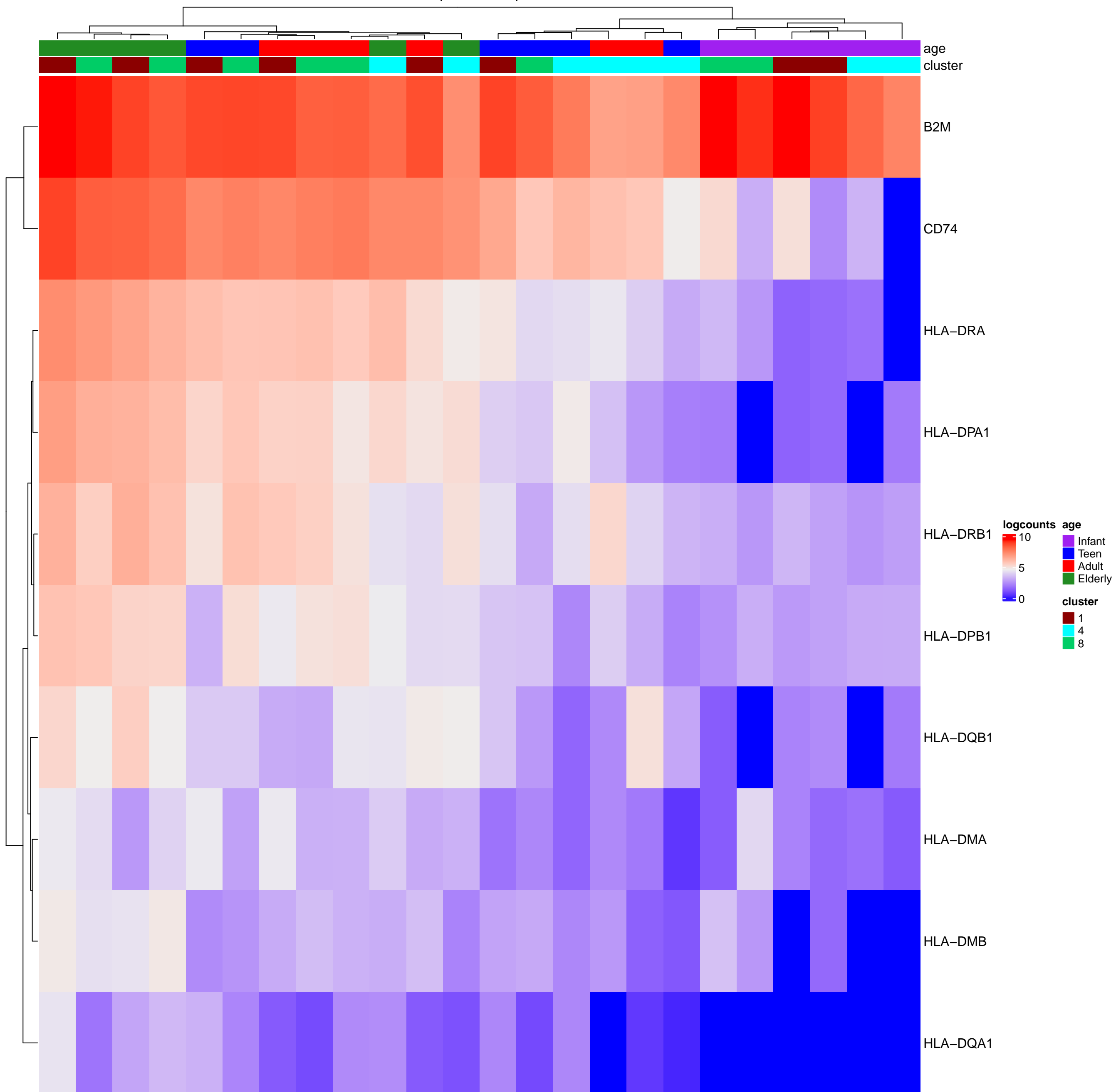
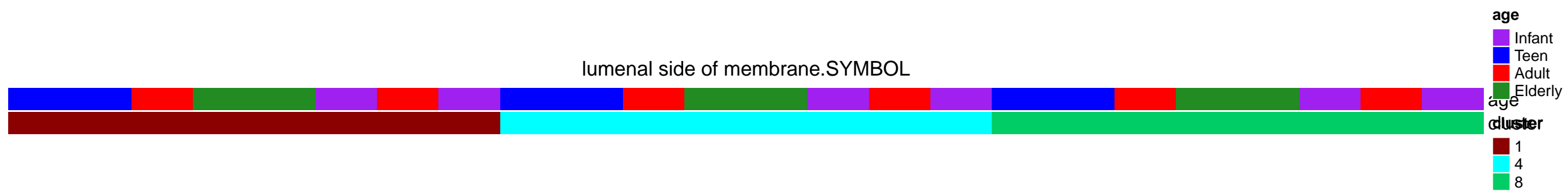
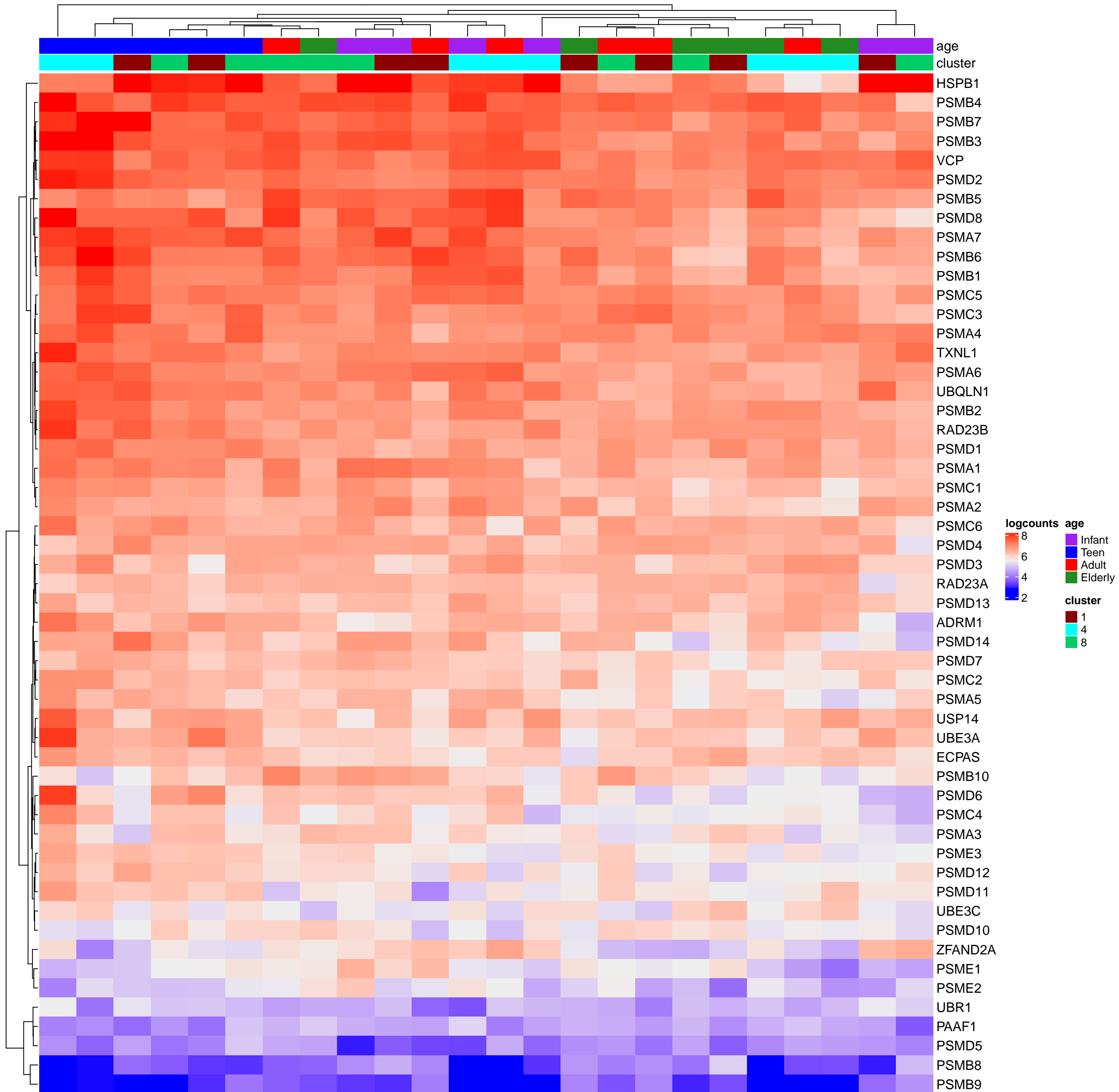


MHC class II protein complex.SYMBOL

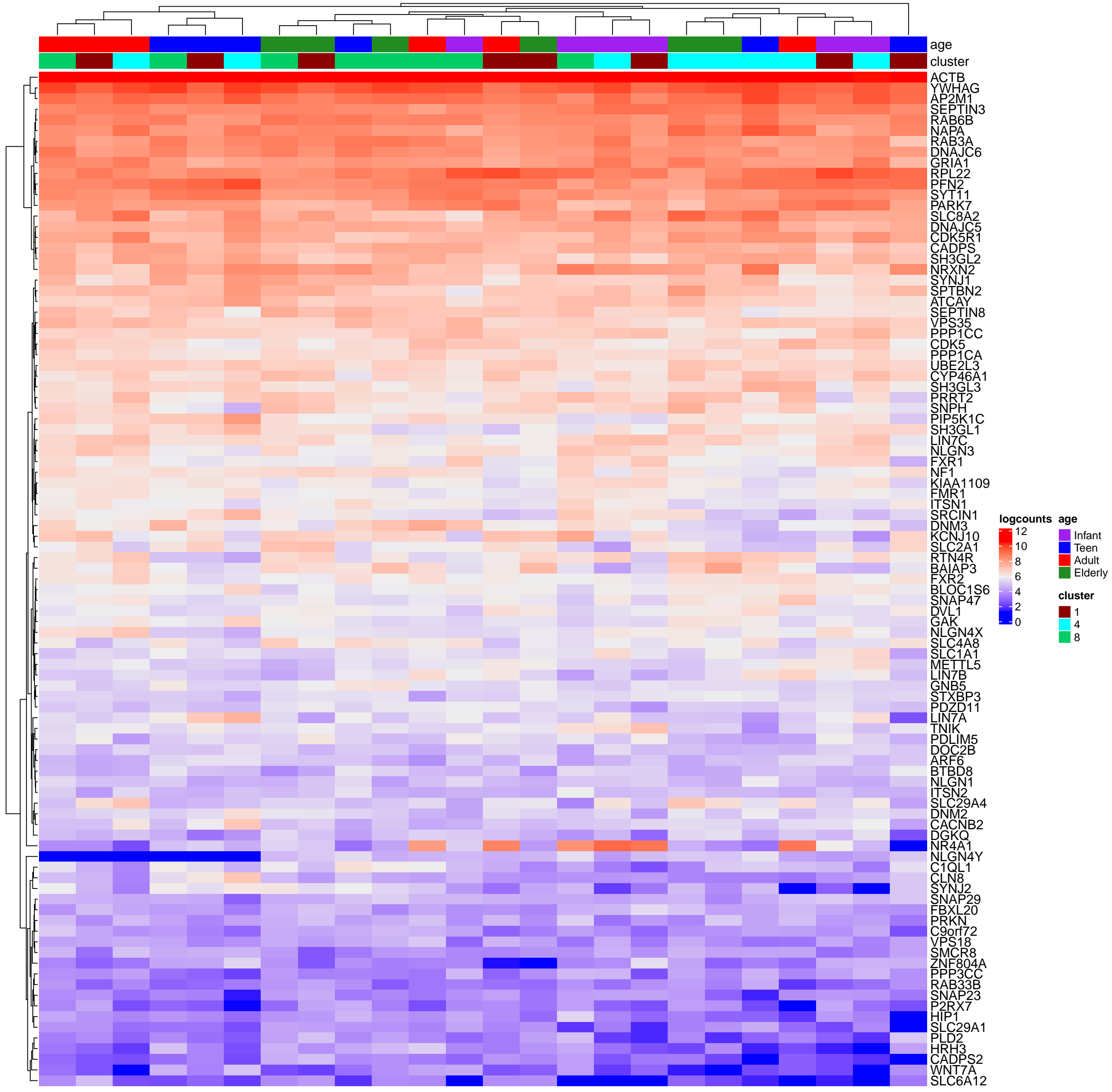




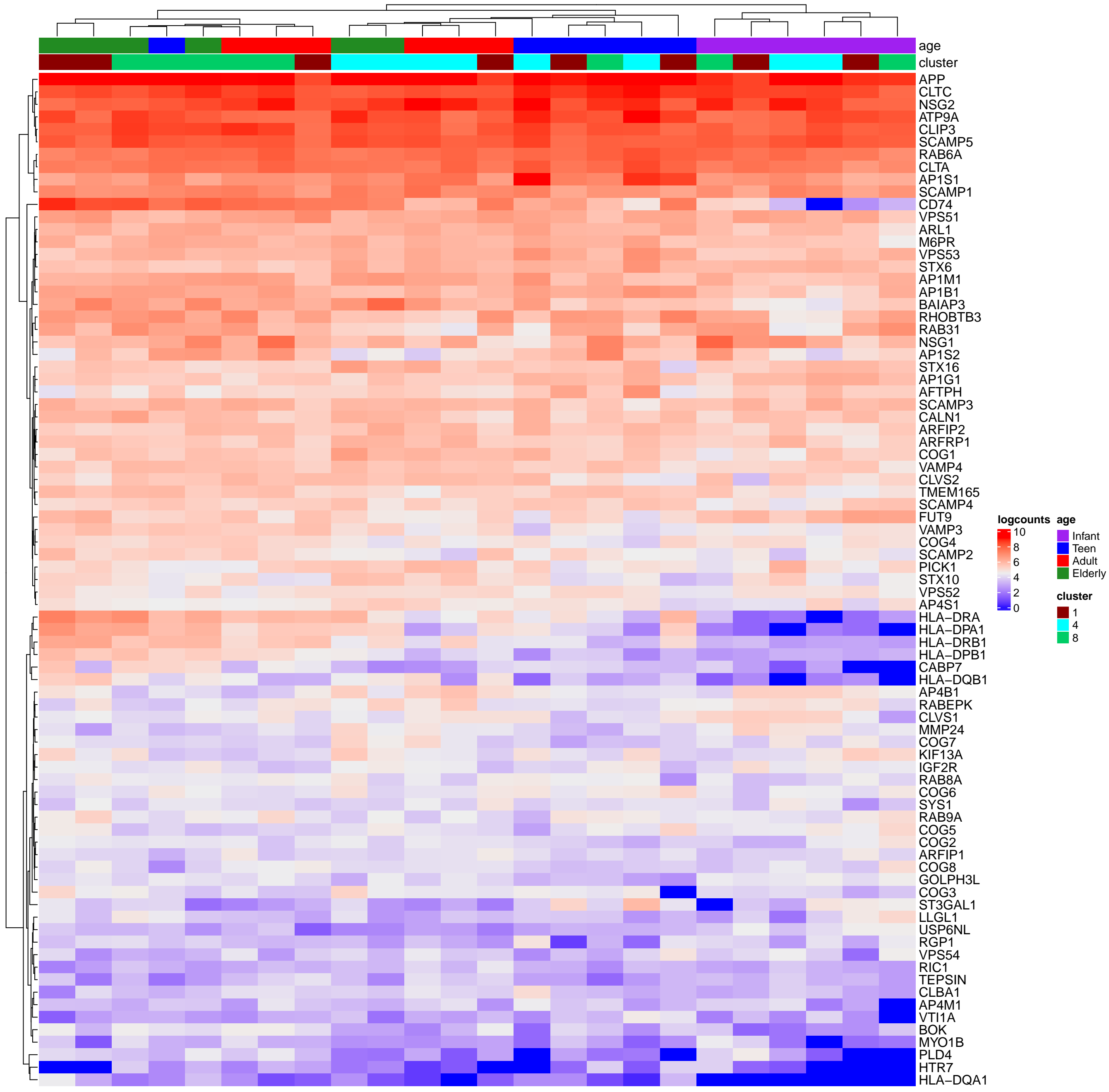
proteasome complex.SYMBOL



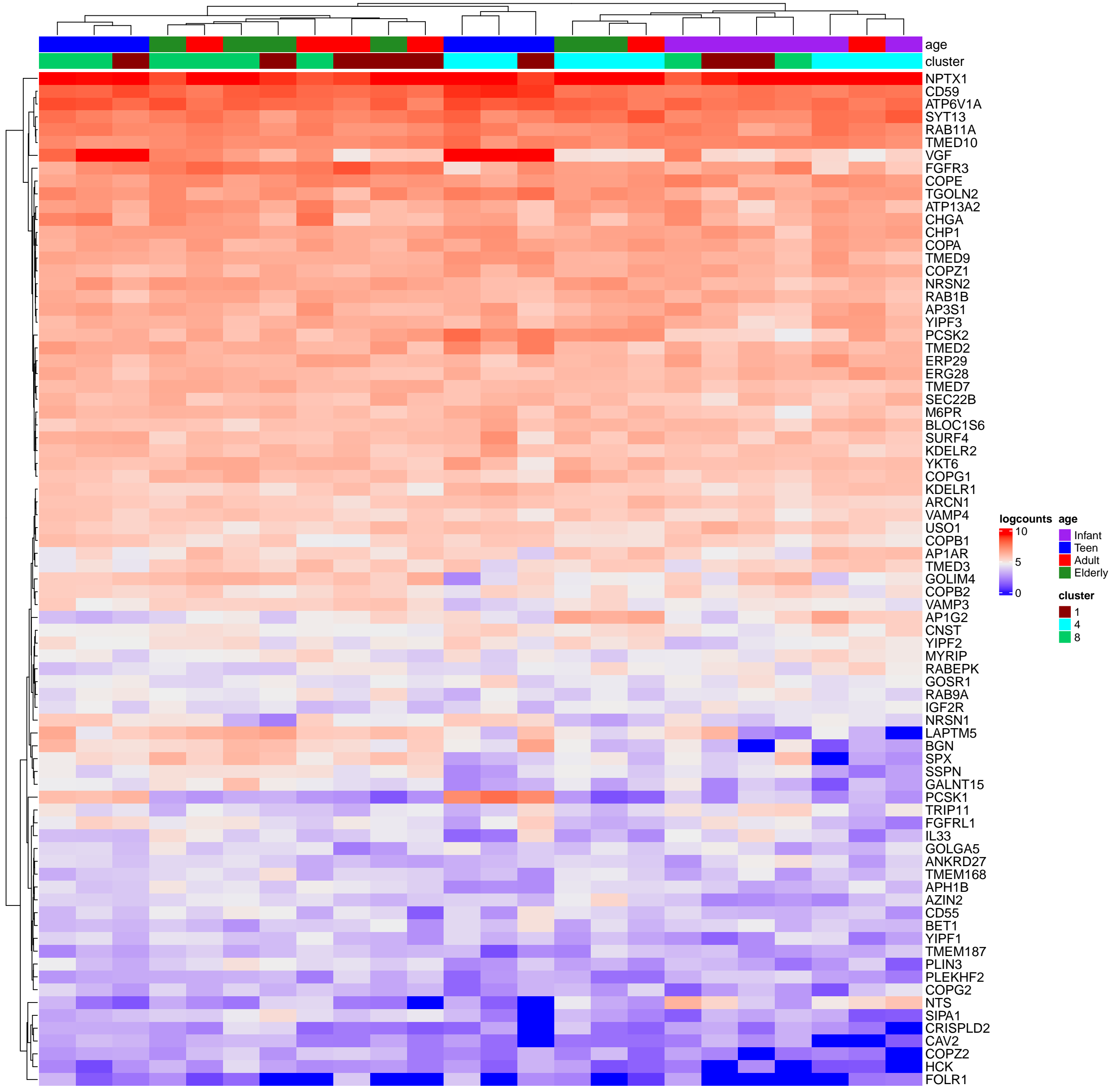
presynapse.SYMBOL



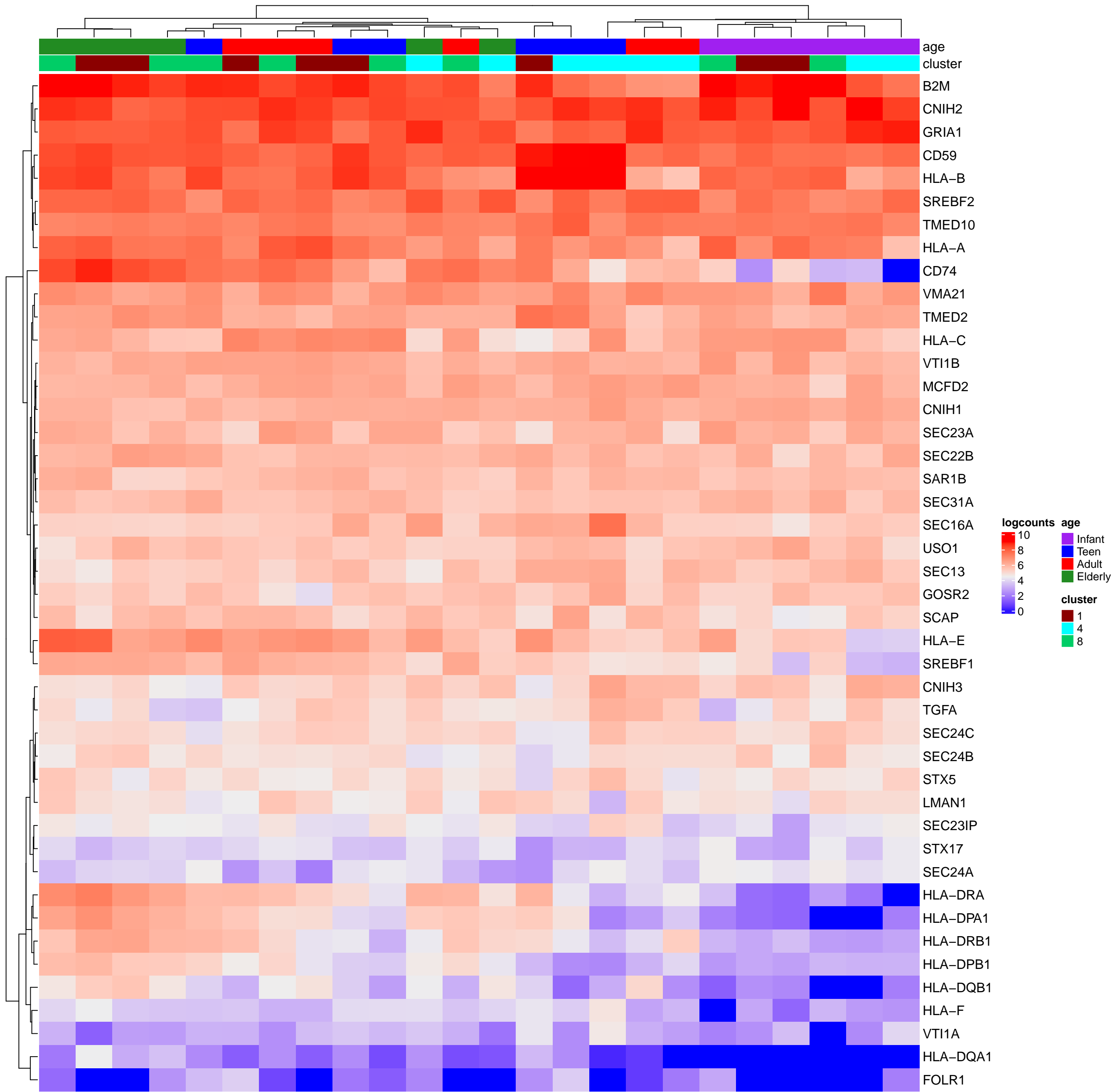
trans-Golgi network membrane.SYMBOL



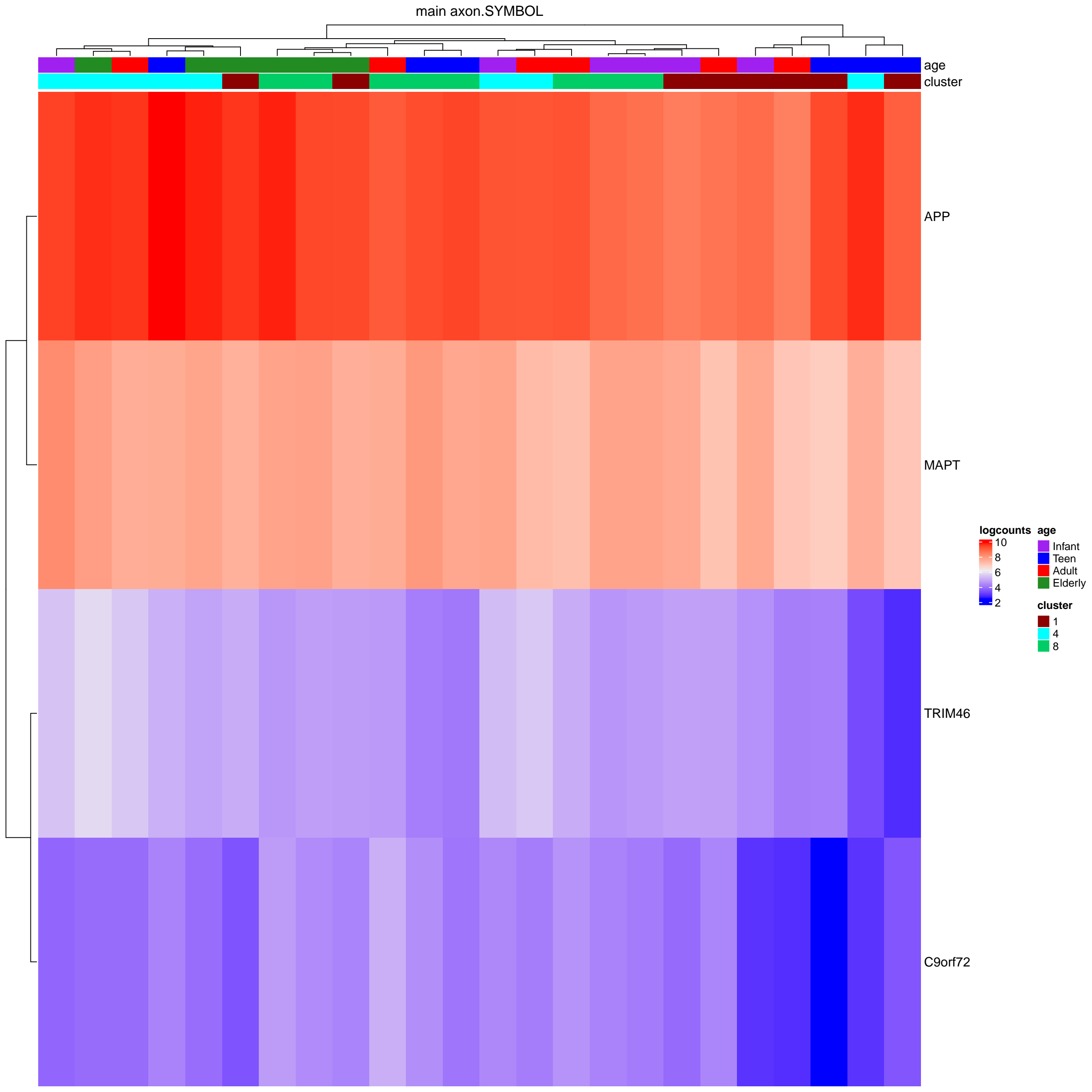
transport vesicle.SYMBOL



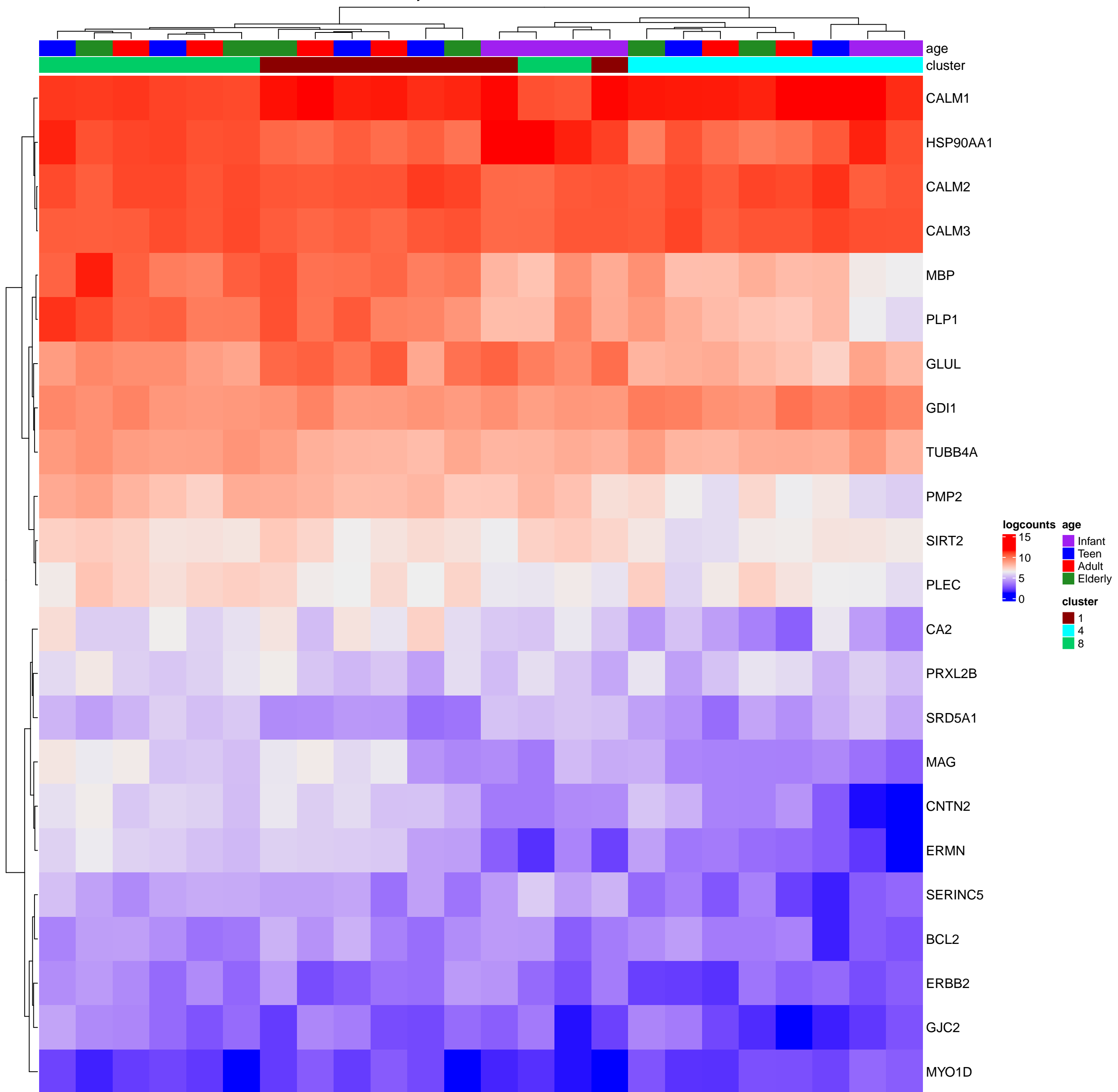
ER to Golgi transport vesicle membrane.SYMBOL



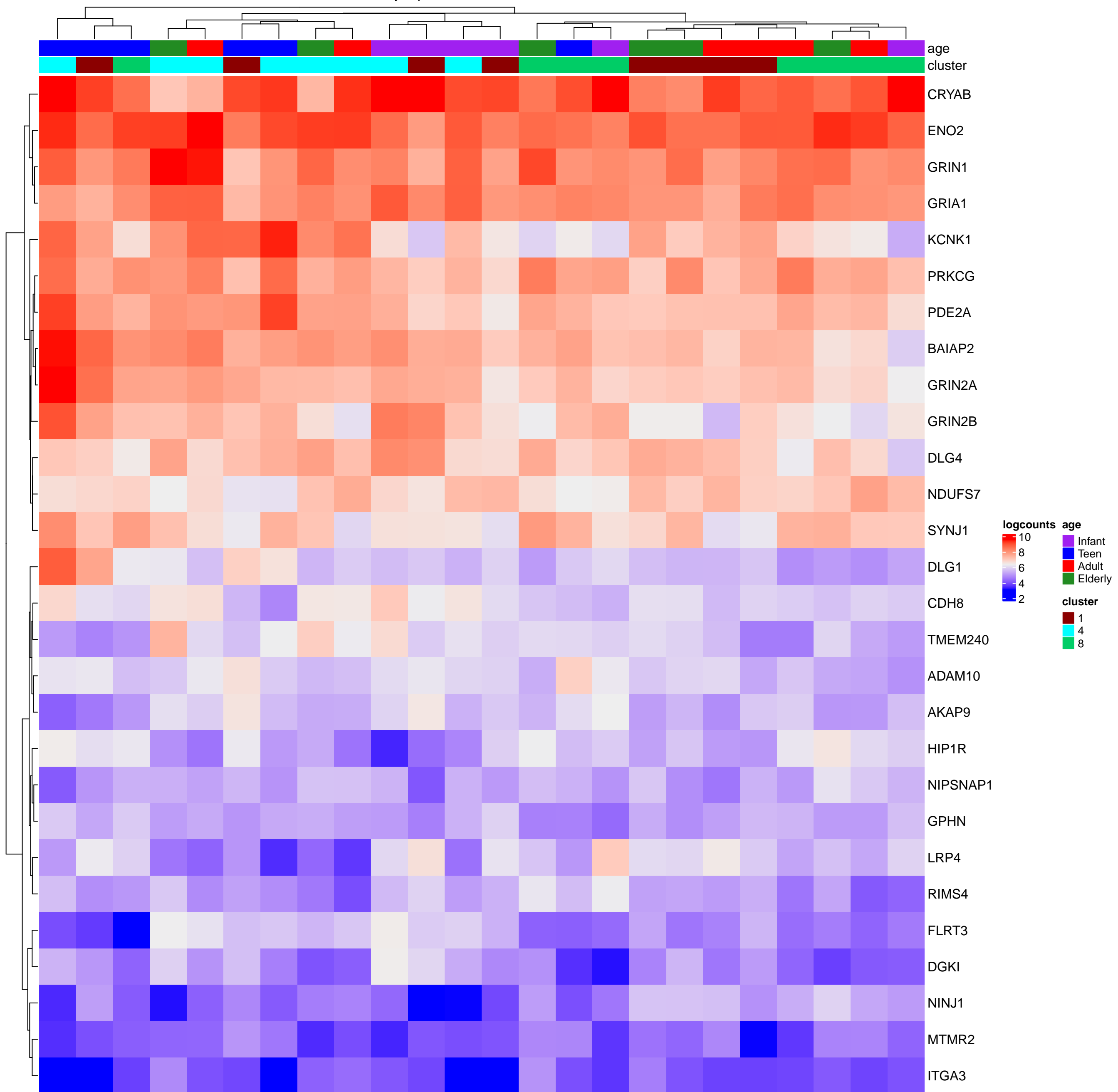
main axon.SYMBOL



myelin sheath.SYMBOL

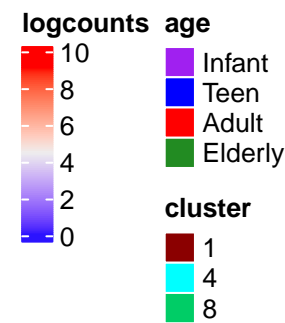


synaptic membrane.SYMBOL

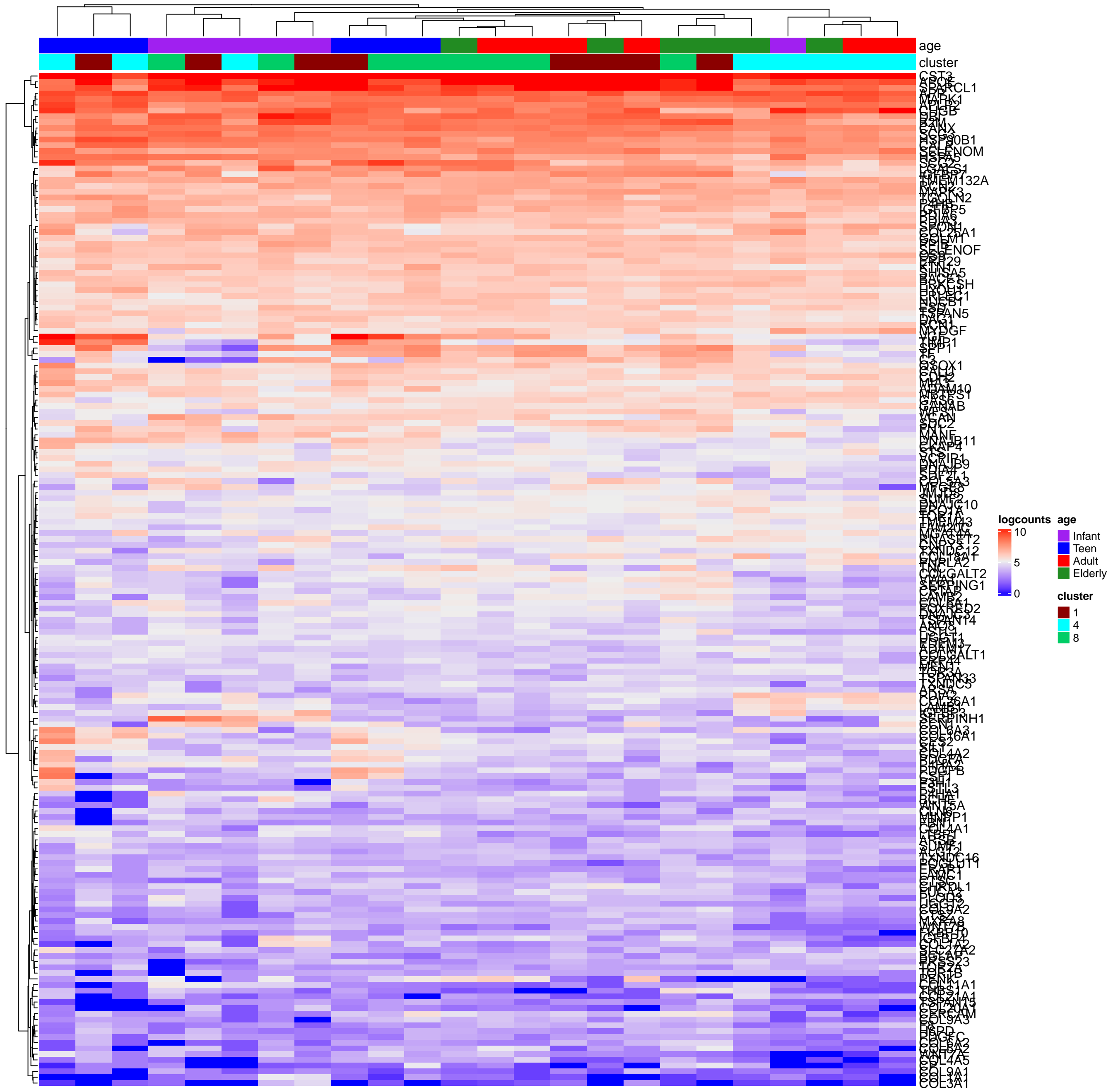


This figure displays a heatmap visualization of a dataset, likely representing gene expression data across various samples. The color scale ranges from dark blue (low values) to dark red (high values). The top dendrogram shows hierarchical clustering of columns, and the left dendrogram shows hierarchical clustering of rows. The heatmap itself is a grid of colored cells representing the data values.

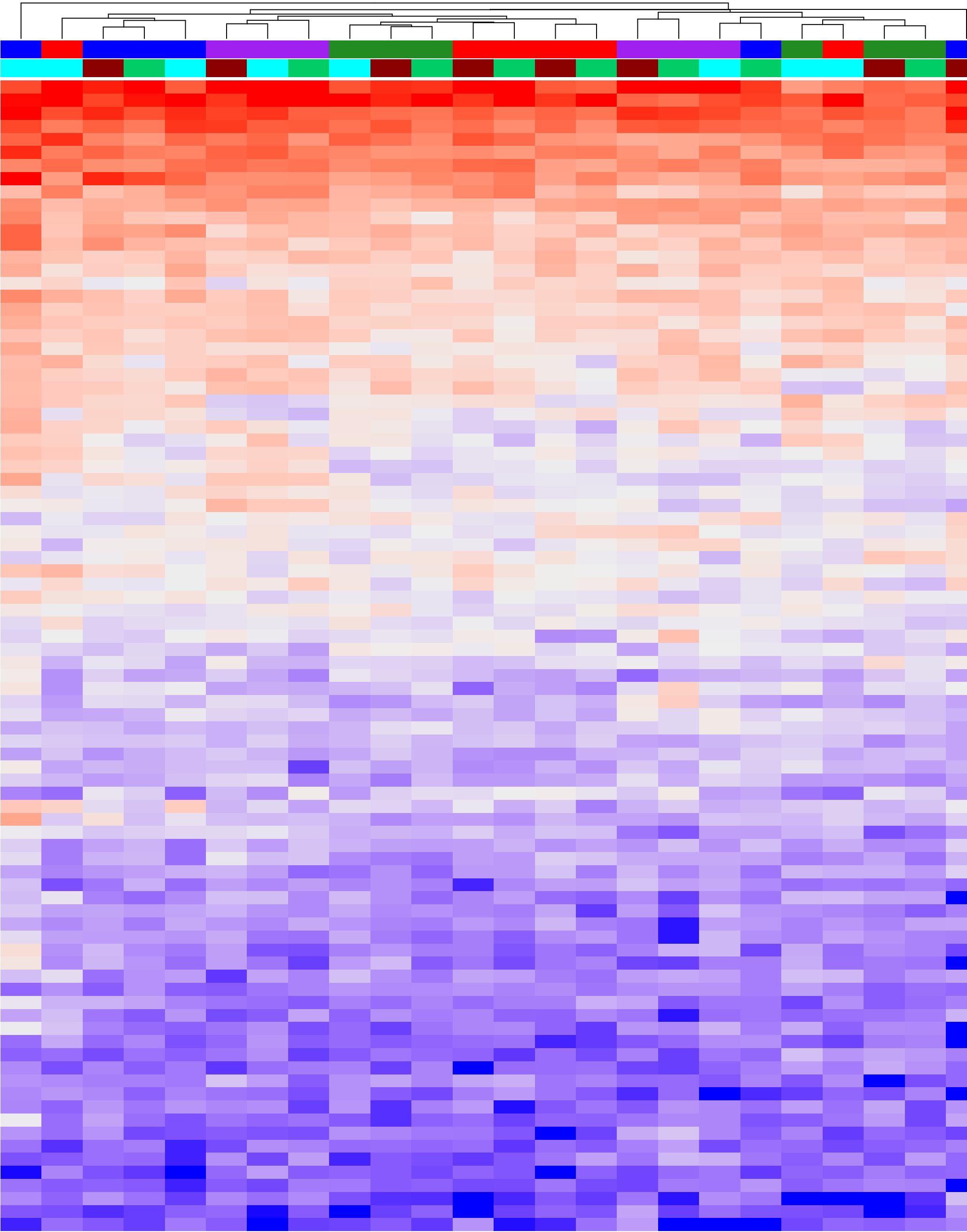
LEF1
PDLIM1



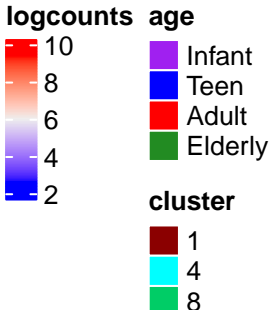
endoplasmic reticulum lumen.SYMBOL



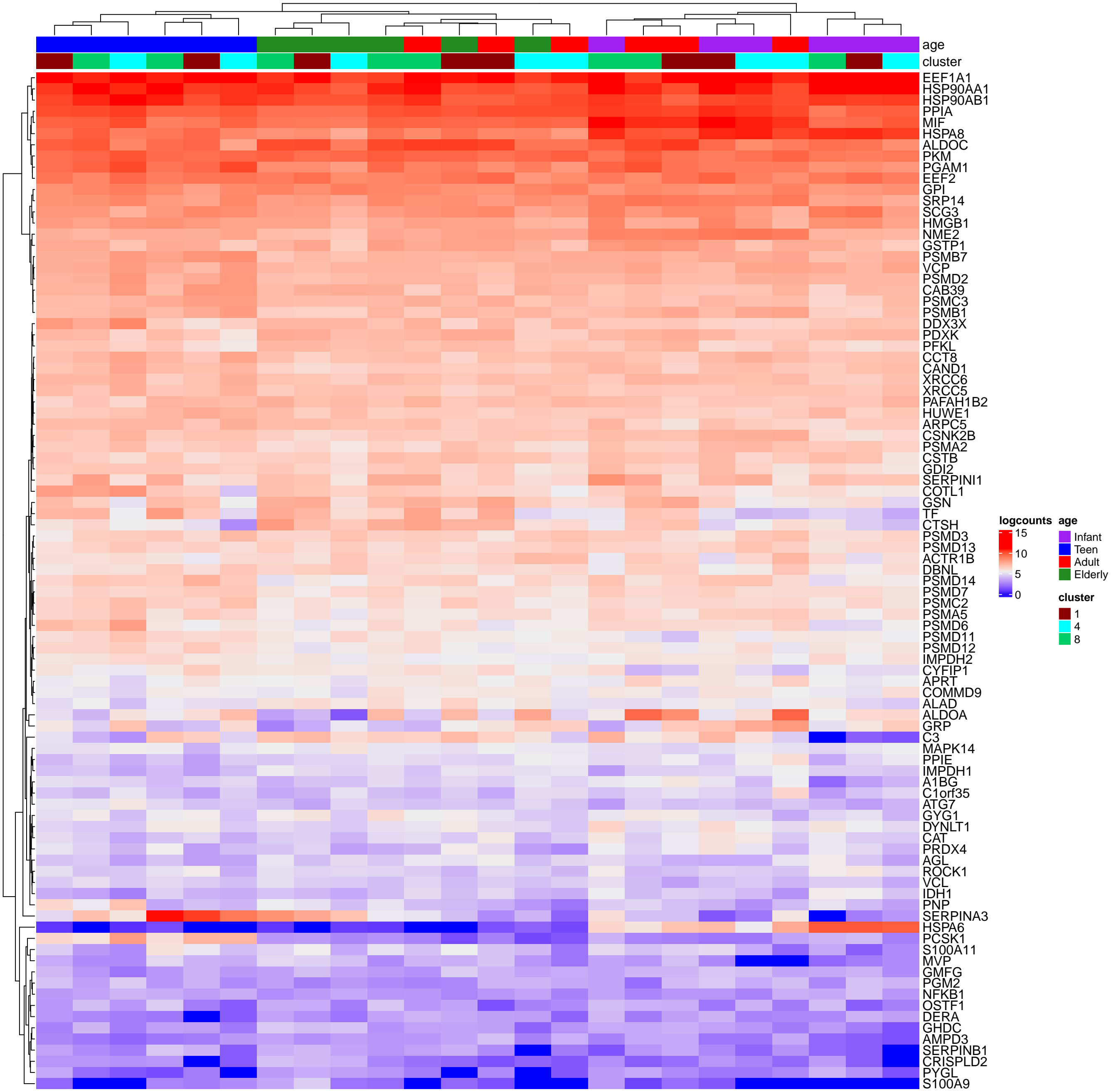
spliceosomal complex.SYMBOL



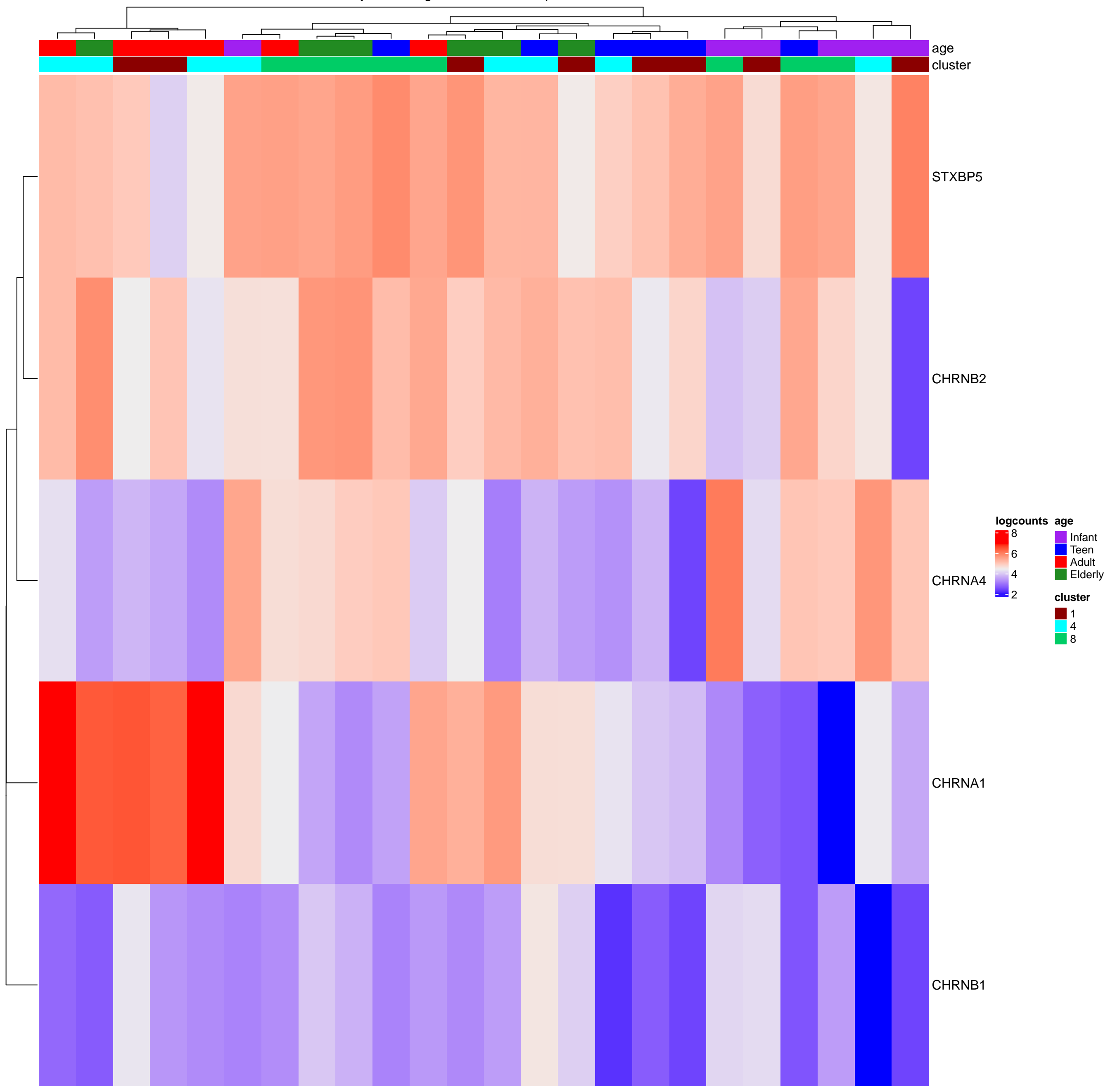
age
cluster
HSPA8
SNRPN
HNRNPA2B1
HNRNPA1
CIRBP
HNRNPC
SNRPD2
RHEB
TXNL4A
NCL
HNRNPH3
SRSF2
SF1
SF3B2
WAC
PRPF19
HNRNPM
DDX39B
HNRNPR
SF3B1
TRA2B
SNRNP70
BUD31
SNRPG
RBM3
API5
TRA2A
RBM5
RBMX
SNRPB2
DHX15
SNRPF
SNRPE
C9orf78
PTBP2
SF3A1
CLNS1A
BCAS2
CWC15
WBP11
SNRNP200
SLU7
SNRPD3
DDX41
RBM17
SF3A3
U2AF2
IVNS1ABP
SREK1
PRPF6
PLRG1
PPP1R8
CTNNBL1
SNW1
LGALS3
SNRPB
SNRPA
SNRPA1
SRRM1
CDC5L
DHX38
SMNDC1
U2AF1L4
AKAP17A
USP39
SF3B4
TXNL4B
CACTIN
PRKRIP1
SUGP1
CRNKL1
TFIP11
WDR83
PIIH
SF3A2
DHX8
GPKOW
DHX16
ZNF830
AAR2
PRPF3
RBM28
RBMXL1
GEMIN2
PRPF4
PRPF18
CWC22
DHX32



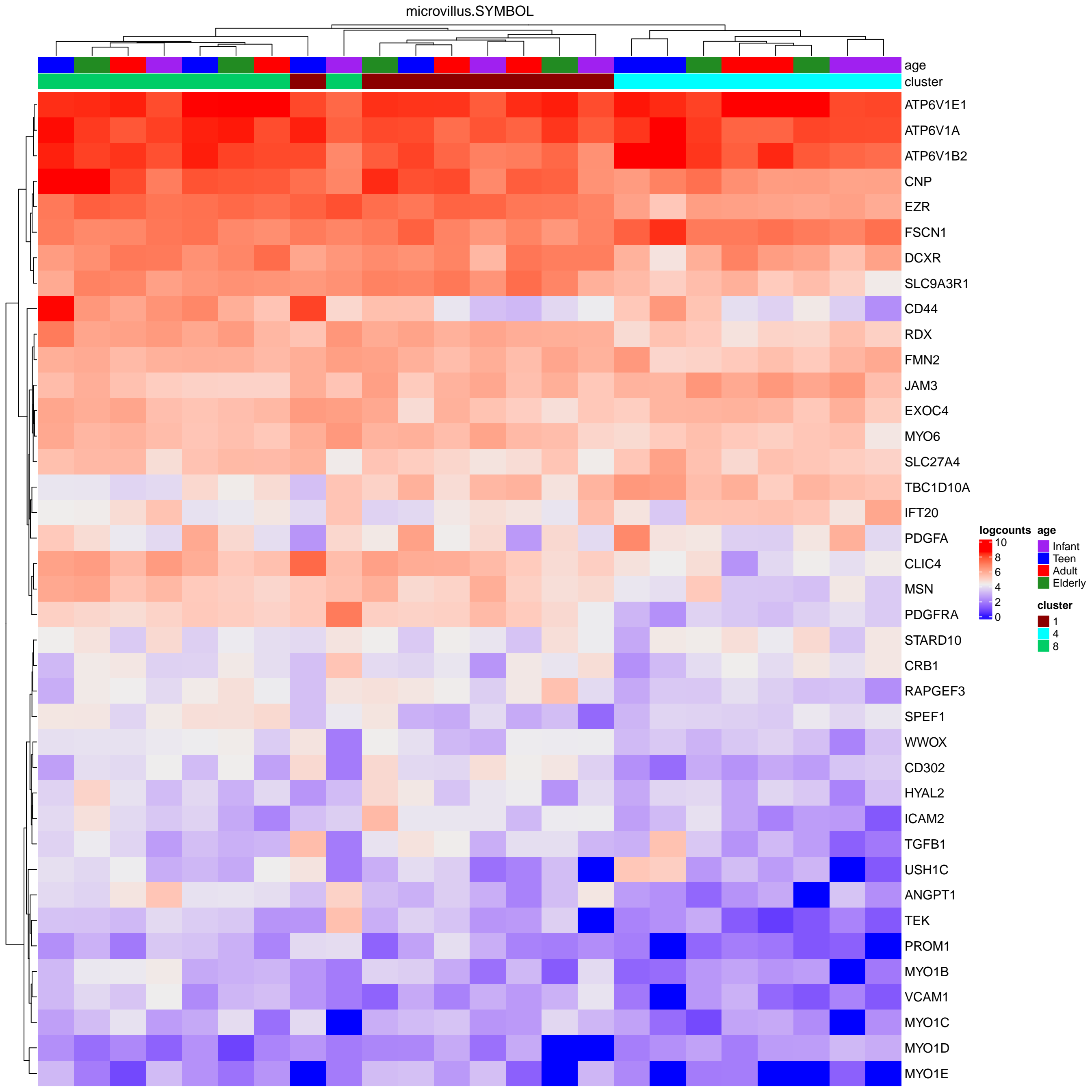
secretory granule lumen.SYMBOL



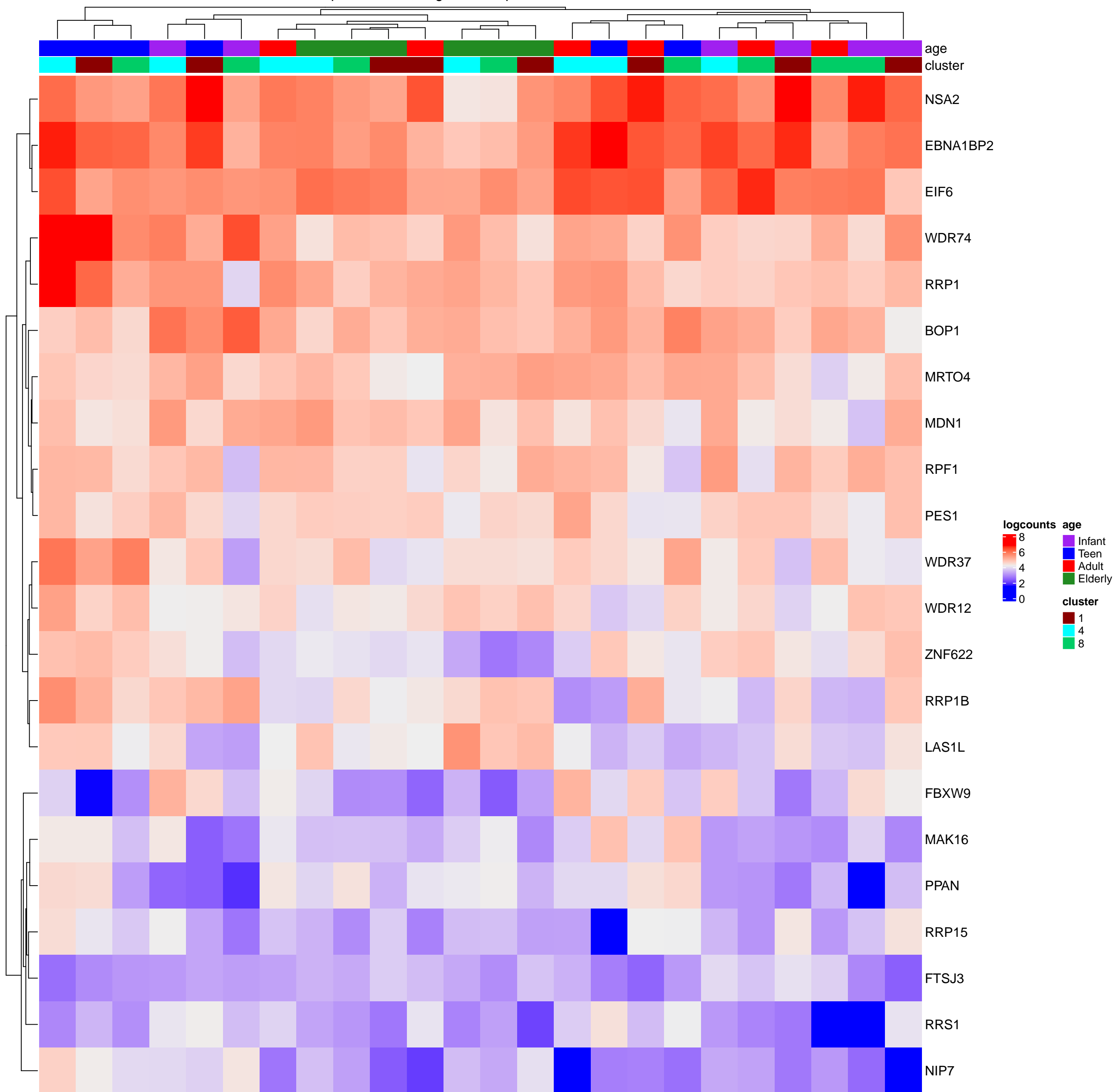
acetylcholine-gated channel complex.SYMBOL



microvillus.SYMBOL



preribosome, large subunit precursor.SYMBOL



focal adhesion.SYMBOL

