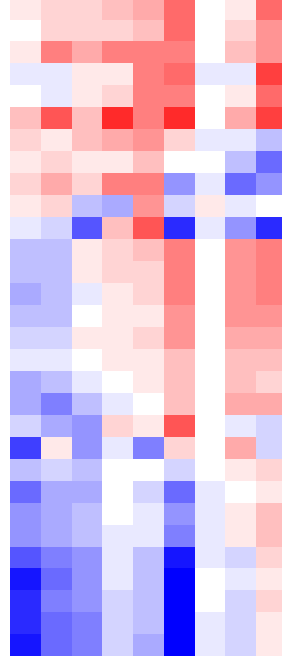
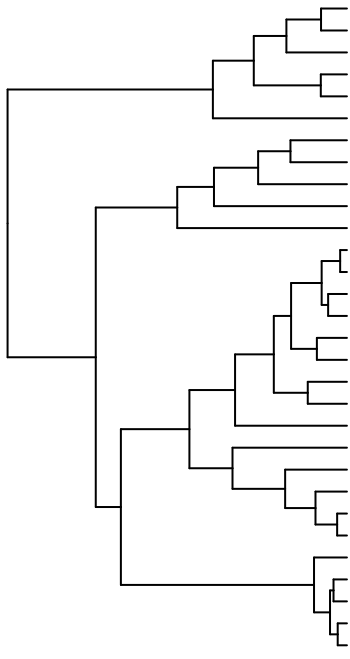
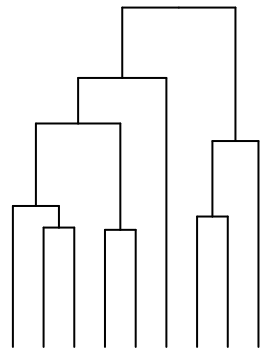
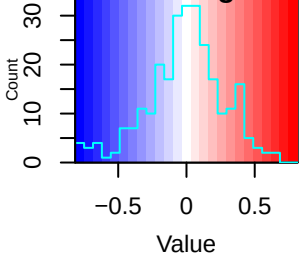


Color Key
and Histogram



s.CD14_._Mono_ CC
s.FCGR3A_._Mono_ s.DC
s.Macrophage
s.CD14_._Mono_
s.NK.2._IL7R_ s.T
s.B
s.NK.1._FCGRA3A_

- GO:0042605 peptide antigen binding
- GO:0098553 luminal side of endoplasmic reticulum membrane
- GO:0001916 positive regulation of T cell mediated cytotoxicity
- GO:0042613 MHC class II protein complex
- GO:0023026 MHC class II protein complex binding
- GO:0042612 MHC class I protein complex
- GO:0001975 response to amphetamine
- GO:1904628 cellular response to phorbol 13-acetate 12-myristate
- GO:0051412 response to corticosterone
- GO:0043330 response to exogenous dsRNA
- GO:0097398 cellular response to interleukin-17
- GO:0045271 respiratory chain complex I
- GO:0008137 NADH dehydrogenase (ubiquinone) activity
- GO:0042776 proton motive force-driven mitochondrial ATP synthesis
- GO:0009060 aerobic respiration
- GO:0032981 mitochondrial respiratory chain complex I assembly
- GO:0005743 mitochondrial inner membrane
- GO:1902600 proton transmembrane transport
- GO:0045333 cellular respiration
- GO:0140507 granzyme-mediated programmed cell death signaling pathway
- GO:0034975 protein folding in endoplasmic reticulum
- GO:0006413 translational initiation
- GO:0005840 ribosome
- GO:0003735 structural constituent of ribosome
- GO:0006412 translation
- GO:0022626 cytosolic ribosome
- GO:0015935 small ribosomal subunit
- GO:0022627 cytosolic small ribosomal subunit
- GO:0022625 cytosolic large ribosomal subunit
- GO:0002181 cytoplasmic translation