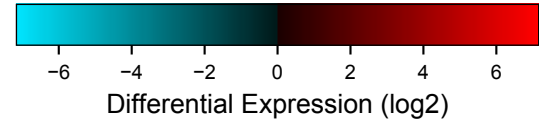
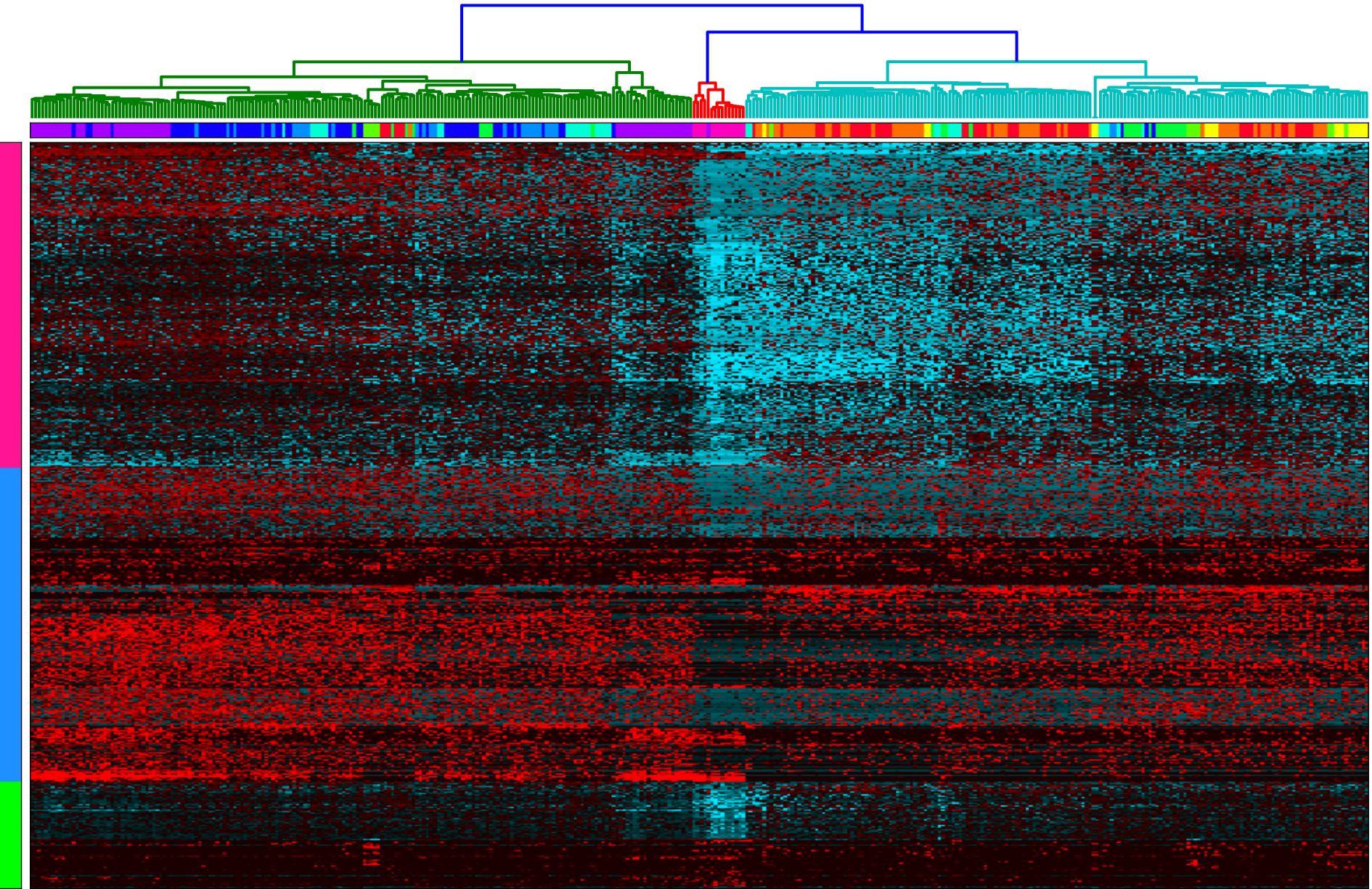


SampleLogFolds-BoneMarrow-scRNASeq-ICGS



- membrane-enclosed lumen p=5e-13 (c4)
- protein transport p=1e-13 (c4)
- intracellular transport p=4e-14 (c4)
- pigment granule p=4e-14 (c4)
- threonine-type peptidase activity p=2e-14 (c4)
- nucleotide binding p=2e-16 (c4)
- cytoplasm p=1e-17 (c4)
- nucleus p=1e-25 (c4)
- protein binding p=5e-33 (c4)
- primary metabolic process p=2e-44 (c4)
- RNA binding p=7e-51 (c4)
- cellular metabolic process p=3e-68 (c4)
- mitochondrion p=4e-76 (c4)
- macromolecular complex p=1e-84 (c4)
- intracellular organelle part p=1e-98 (c4)
- iron ion transmembrane transporter activity p=5e-06 (c2)
- immune system process p=3e-07 (c2)
- response to stress p=2e-07 (c2)
- catabolic process p=1e-07 (c2)
- organelle membrane p=5e-10 (c2)
- cytoplasm p=1e-10 (c2)
- protein binding p=9e-16 (c2)
- nitrogen compound metabolic process p=4e-18 (c2)
- intracellular organelle part p=1e-18 (c2)
- cytoplasmic part p=2e-19 (c2)
- catalytic activity p=1e-19 (c2)
- primary metabolic process p=9e-24 (c2)
- cellular metabolic process p=1e-25 (c2)
- organelle p=4e-34 (c2)
- ribosome p=2e-23 (c1)
- cytosolic large ribosomal subunit p=3e-37 (c1)
- cytosolic small ribosomal subunit p=1e-41 (c1)
- translation p=2e-71 (c1)
- structural constituent of ribosome p=4e-82 (c1)



- Med21
- Sigf1
- Slc12a3
- Wdr21
- Ctss
- Ifi302a1
- Scp2
- Ubr1p3
- Ubr1p3
- Slc15a3
- Mapk13
- Tanc1

