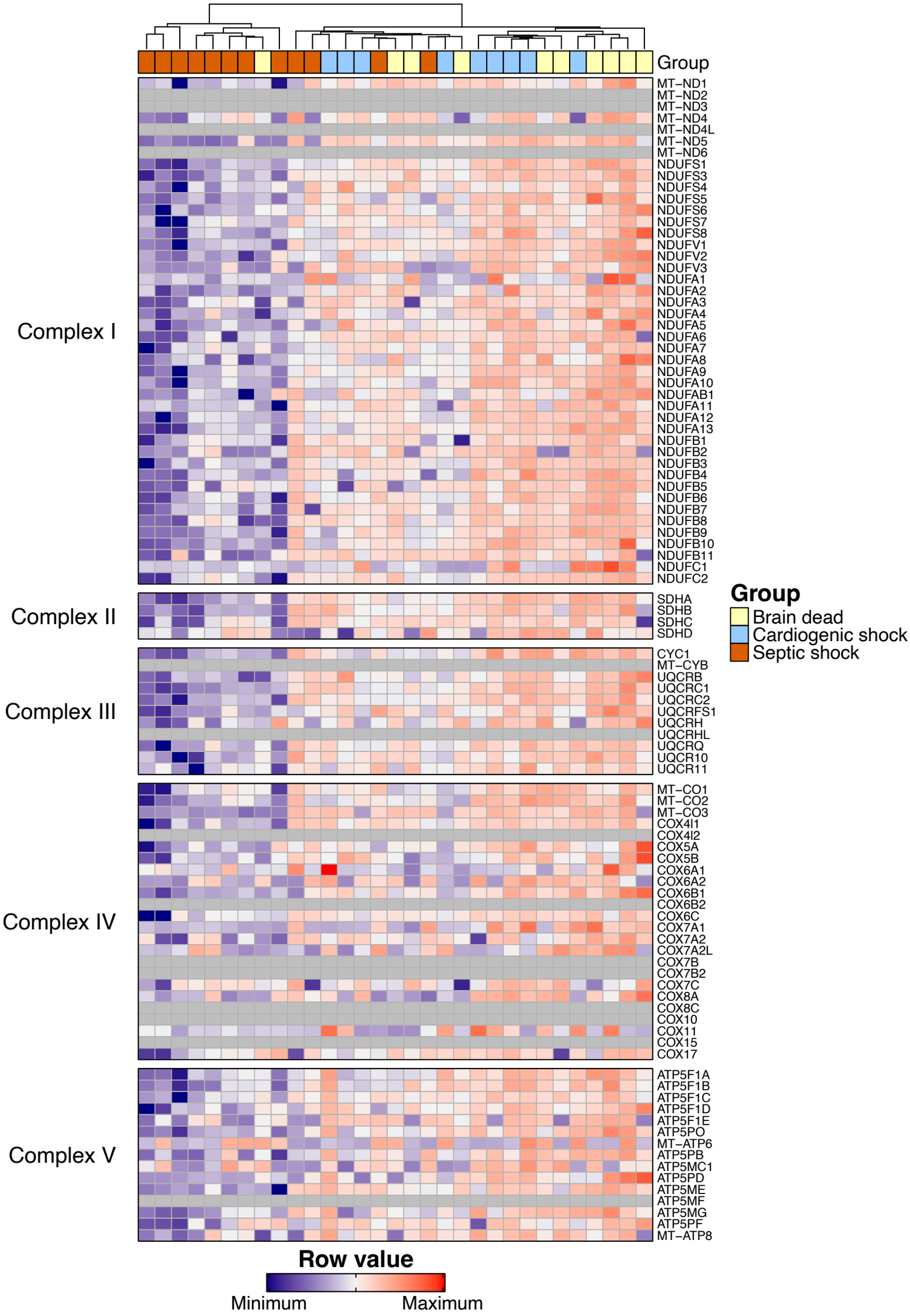
Group
Complex I
MT−ND1
MT−ND2
MT−ND3
MT−ND4
MT−ND4L
MT−ND5
MT−ND6
NDUFS1
NDUFS3
NDUFS4
NDUFS5
NDUFS6
NDUFS7
NDUFS8
NDUFV1
NDUFV2
NDUFV3
NDUFA1
NDUFA2
NDUFA3
NDUFA4
NDUFA5
NDUFA6
NDUFA7
NDUFA8
NDUFA9
NDUFA10
NDUFAB1
NDUFA11
NDUFA12
NDUFA13
NDUFB1
NDUFB2
NDUFB3
NDUFB4
NDUFB5
NDUFB6
NDUFB7
NDUFB8
NDUFB9
NDUFB10
NDUFB11
NDUFC1
NDUFC2
Complex II
SDHA
SDHB
SDHC
SDHD
Complex III
CYC1
MT−CYB
UQCRB
UQCRC1
UQCRC2
UQCRFS1
UQCRH
UQCRHL
UQCRQ
UQCR10
UQCR11
Complex IV
MT−CO1
MT−CO2
MT−CO3
COX4I1
COX4I2
COX5A
COX5B
COX6A1
COX6A2
COX6B1
COX6B2
COX6C
COX7A1
COX7A2
COX7A2L
COX7B
COX7B2
COX7C
COX8A
COX8C
COX10
COX11
COX15
COX17
Complex V
ATP5F1A
ATP5F1B
ATP5F1C
ATP5F1D
ATP5F1E
ATP5PO
MT−ATP6
ATP5PB
ATP5MC1
ATP5PD
ATP5ME
ATP5MF
ATP5MG
ATP5PF
MT−ATP8
Group
Brain dead
Cardiogenic shock
Septic shock
Row value
Minimum
Maximum