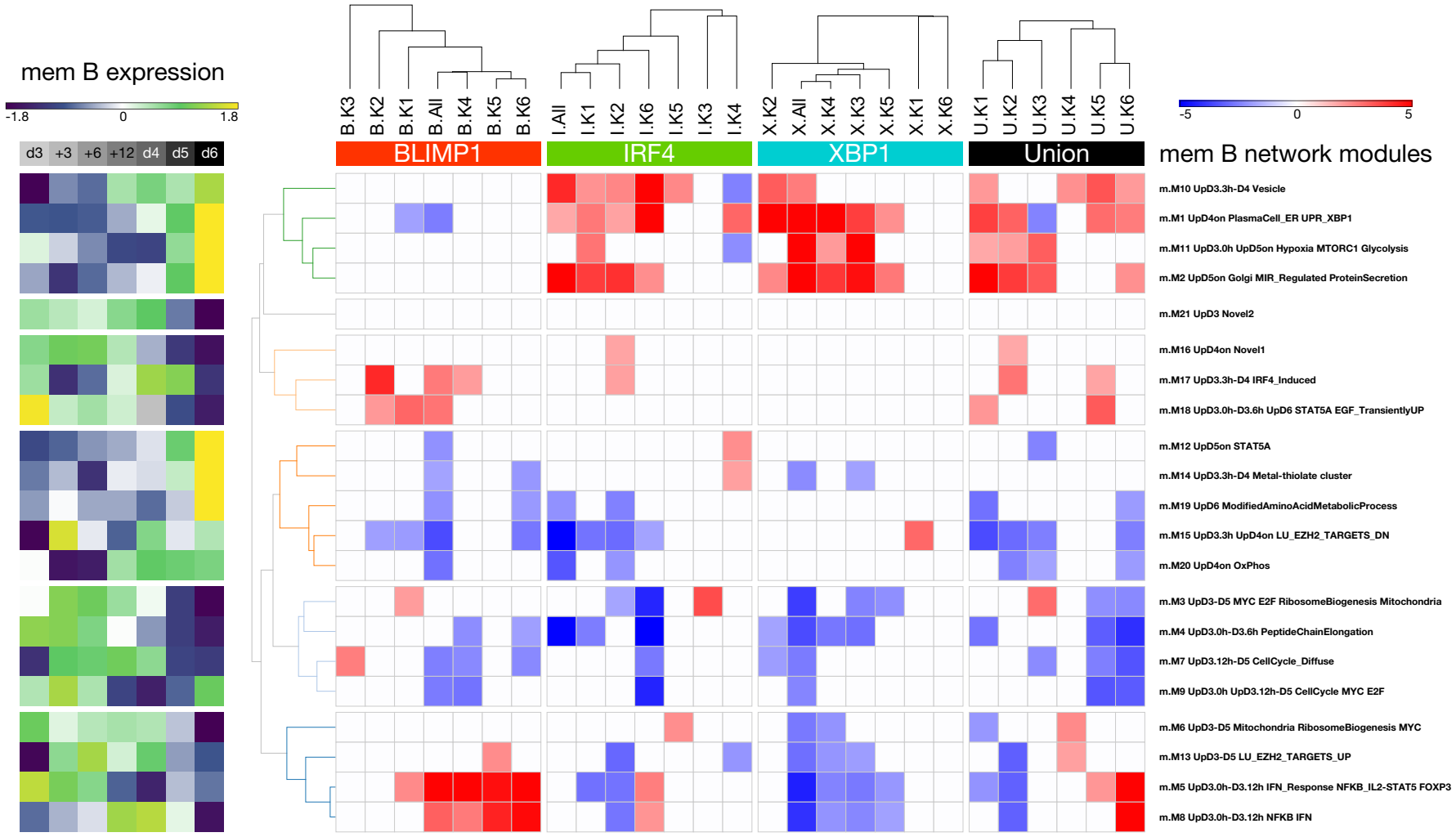
mem B expression
-1.8
0
1.8
d3
+3
+6
+12
d4
d5
d6
B.K3
B.K2
B.K1
B.All
B.K4
B.K5
B.K6
I.All
I.K1
I.K2
I.K6
I.K5
I.K3
I.K4
X.K2
X.All
X.K4
X.K3
X.K5
X.K1
X.K6
U.K1
U.K2
U.K3
U.K4
U.K5
U.K6
BLIMP1
IRF4
XBP1
Union
-5
0
5
mem B network modules
m.M10 UpD3.3h-D4 Vesicle
m.M1 UpD4on PlasmaCell_ER UPR_XBP1
m.M11 UpD3.0h UpD5on Hypoxia MTORC1 Glycolysis
m.M2 UpD5on Golgi MIR_Regulated ProteinSecretion
m.M21 UpD3 Novel2
m.M16 UpD4on Novel1
m.M17 UpD3.3h-D4 IRF4_Induced
m.M18 UpD3.0h-D3.6h UpD6 STAT5A EGF_TransientlyUP
m.M12 UpD5on STAT5A
m.M14 UpD3.3h-D4 Metal-thiolate cluster
m.M19 UpD6 ModifiedAminoAcidMetabolicProcess
m.M15 UpD3.3h UpD4on LU_EZH2_TARGETS_DN
m.M20 UpD4on OxPhos
m.M3 UpD3-D5 MYC E2F RibosomeBiogenesis Mitochondria
m.M4 UpD3.0h-D3.6h PeptideChainElongation
m.M7 UpD3.12h-D5 CellCycle_Diffuse
m.M9 UpD3.0h UpD3.12h-D5 CellCycle MYC E2F
m.M6 UpD3-D5 Mitochondria RibosomeBiogenesis MYC
m.M13 UpD3-D5 LU_EZH2_TARGETS_UP
m.M5 UpD3.0h-D3.12h IFN_Response NFKB_IL2-STAT5 FOXP3
m.M8 UpD3.0h-D3.12h NFKB IFN