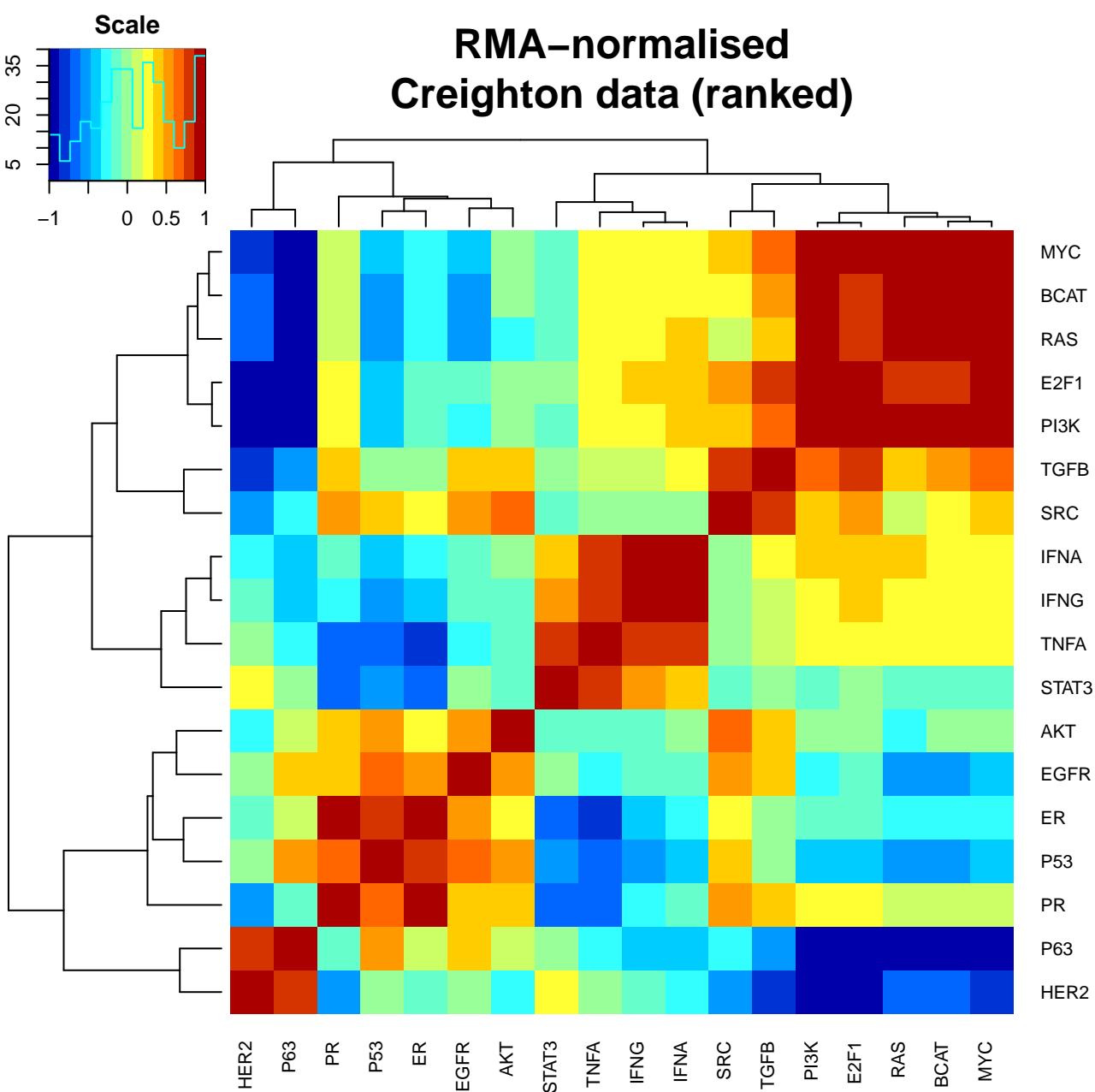
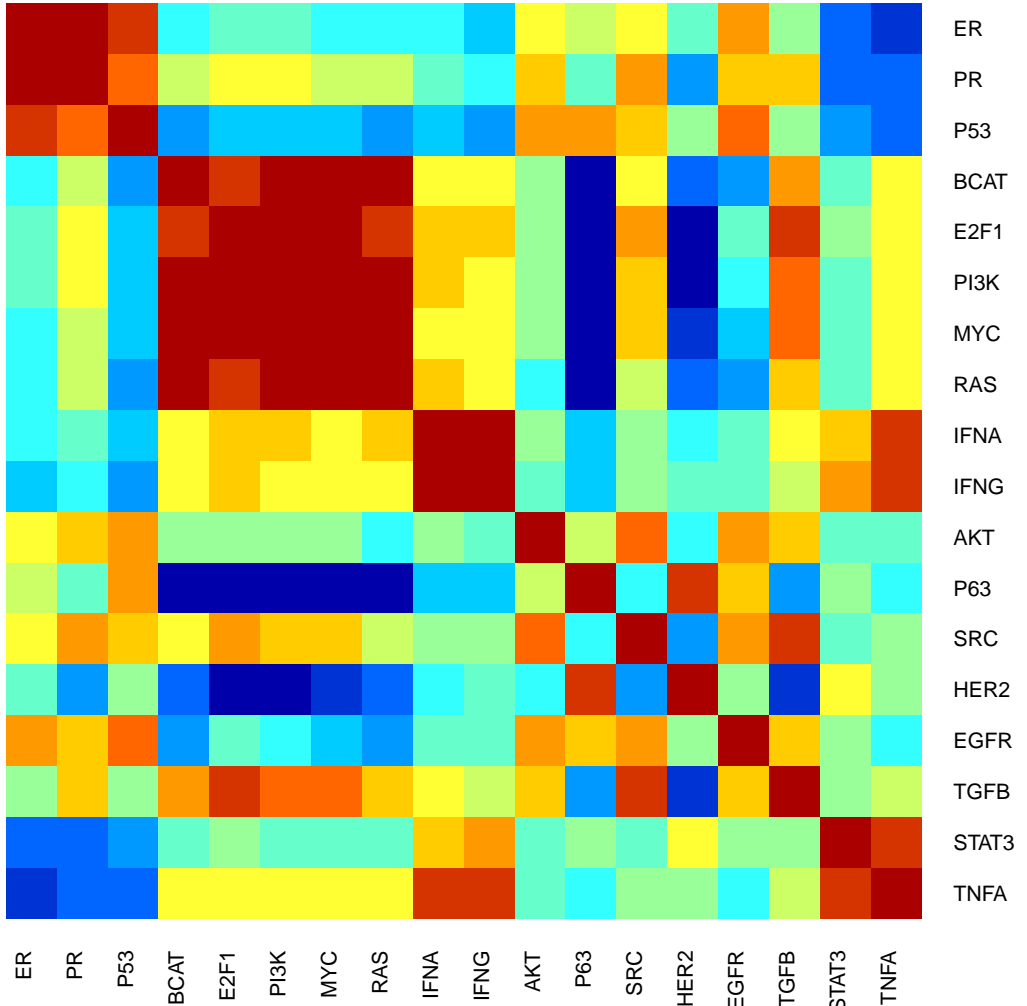
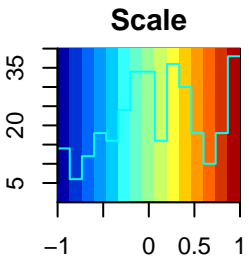
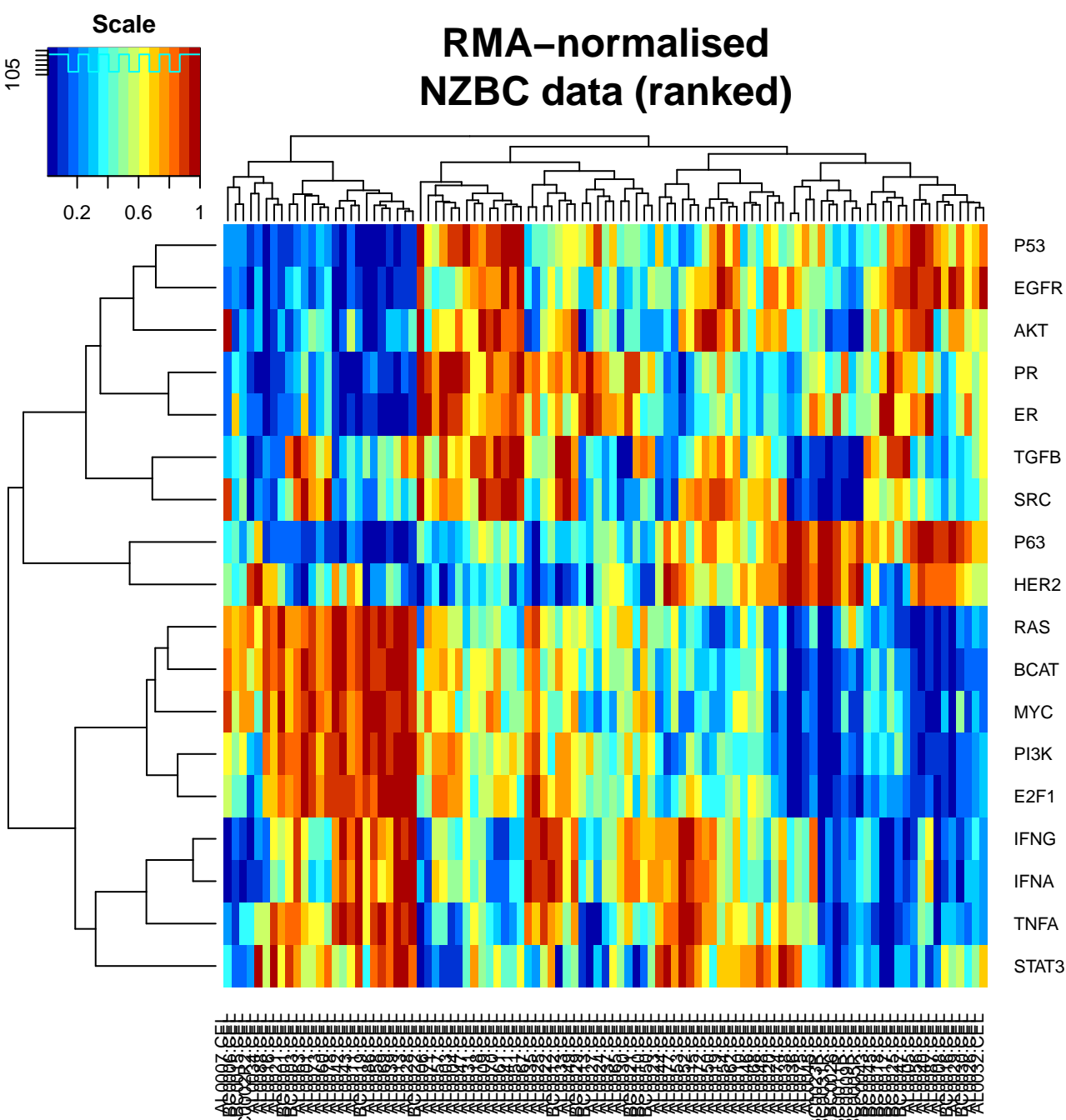
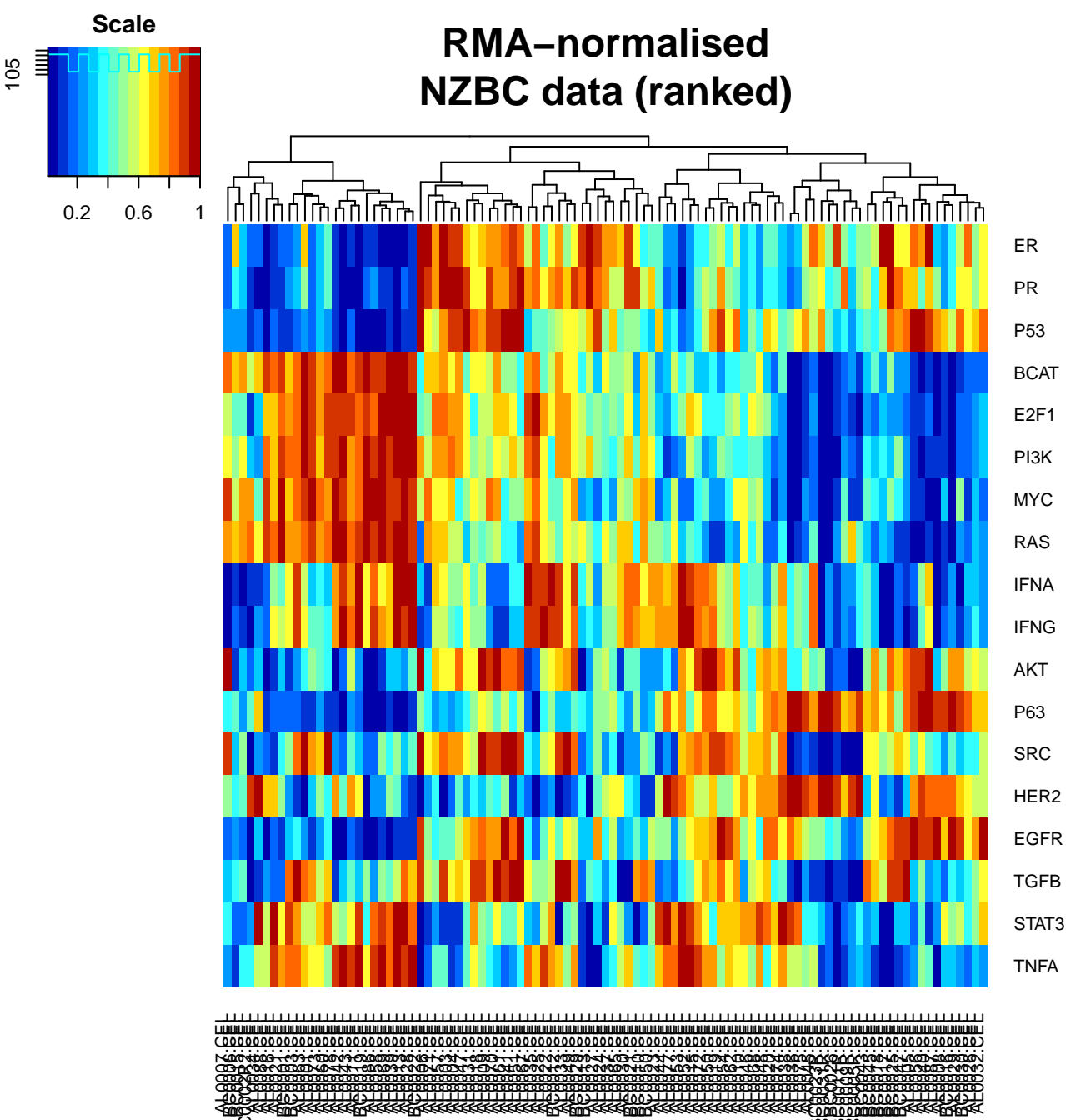
[illegible]



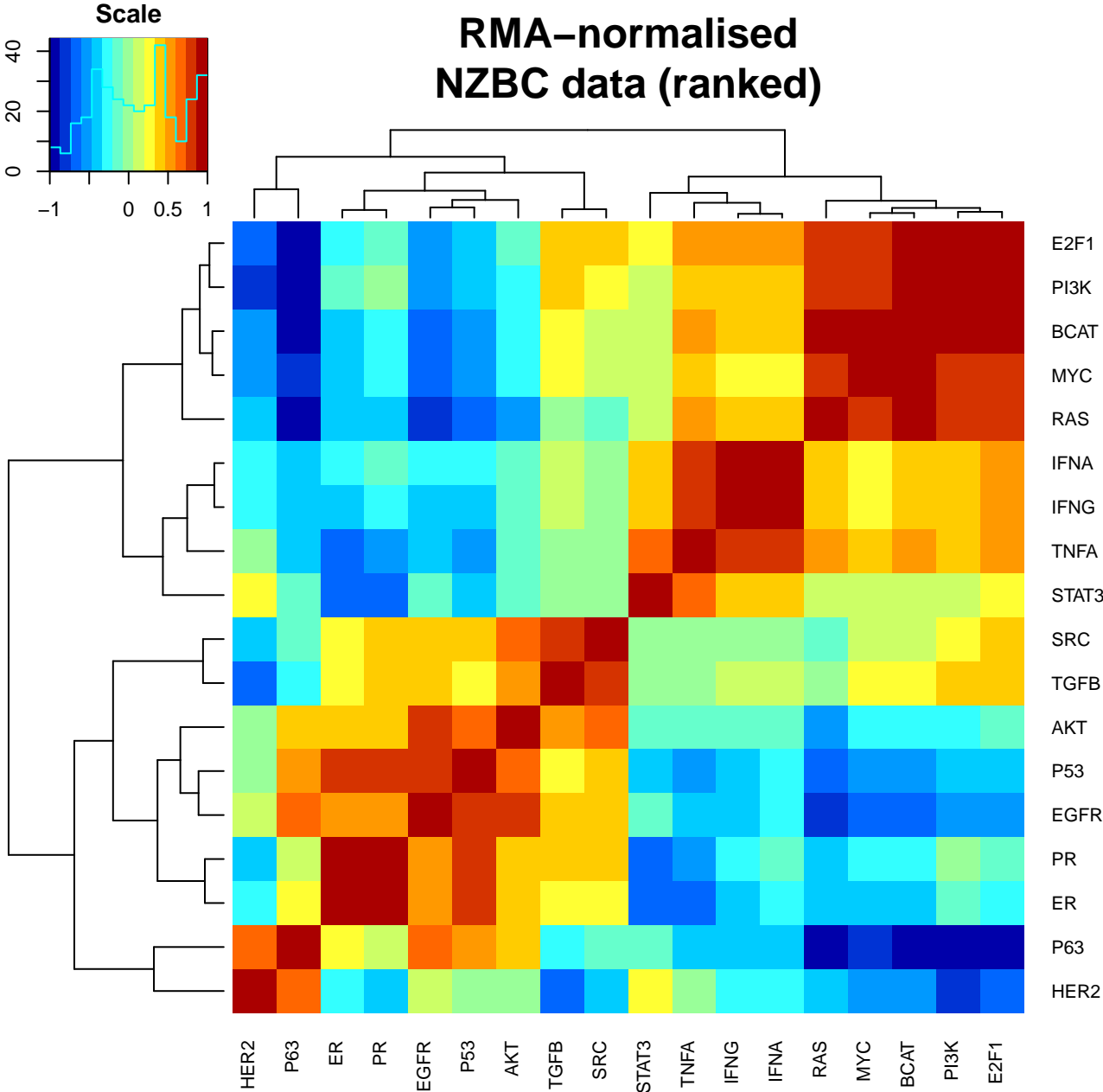
RMA-normalised Creighton data (ranked)



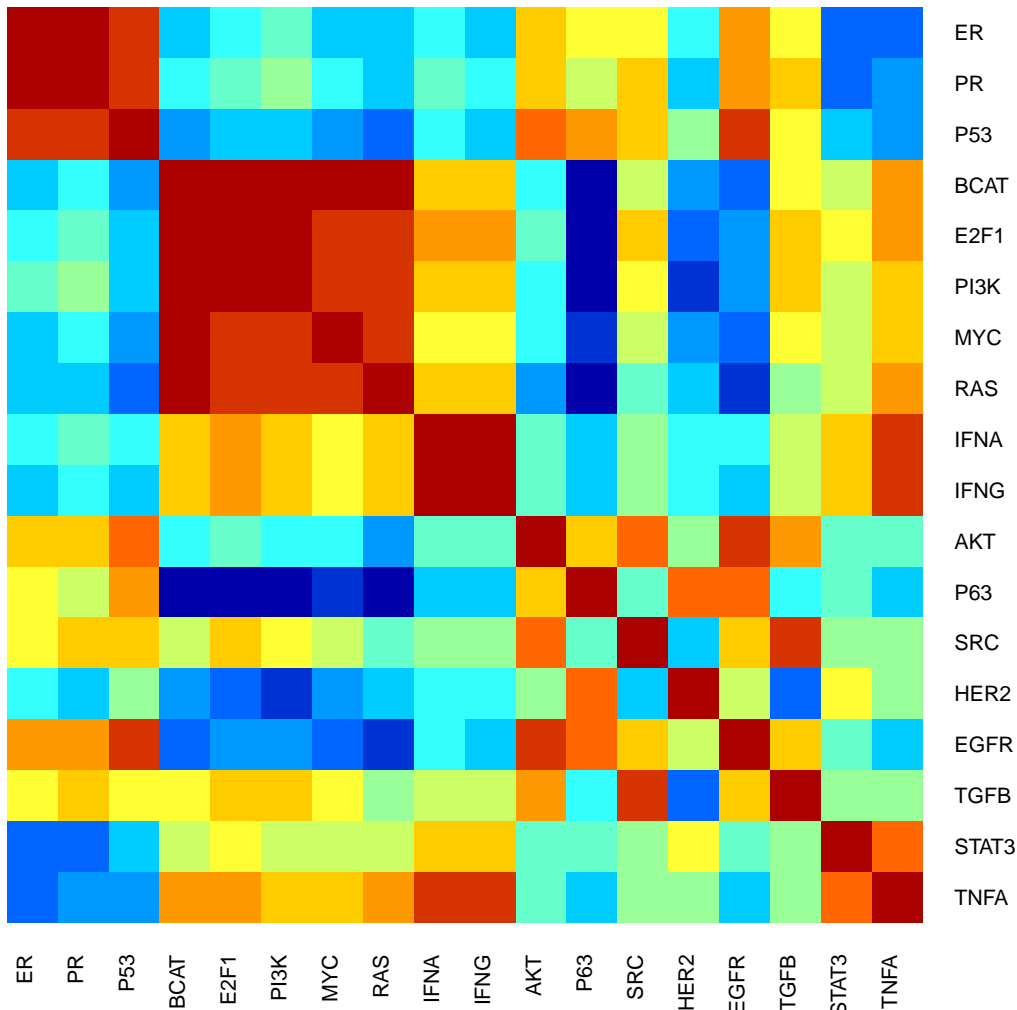
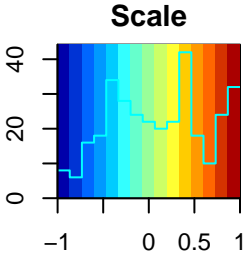




RMA-normalised NZBC data (ranked)

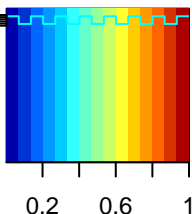


RMA-normalised NZBC data (ranked)

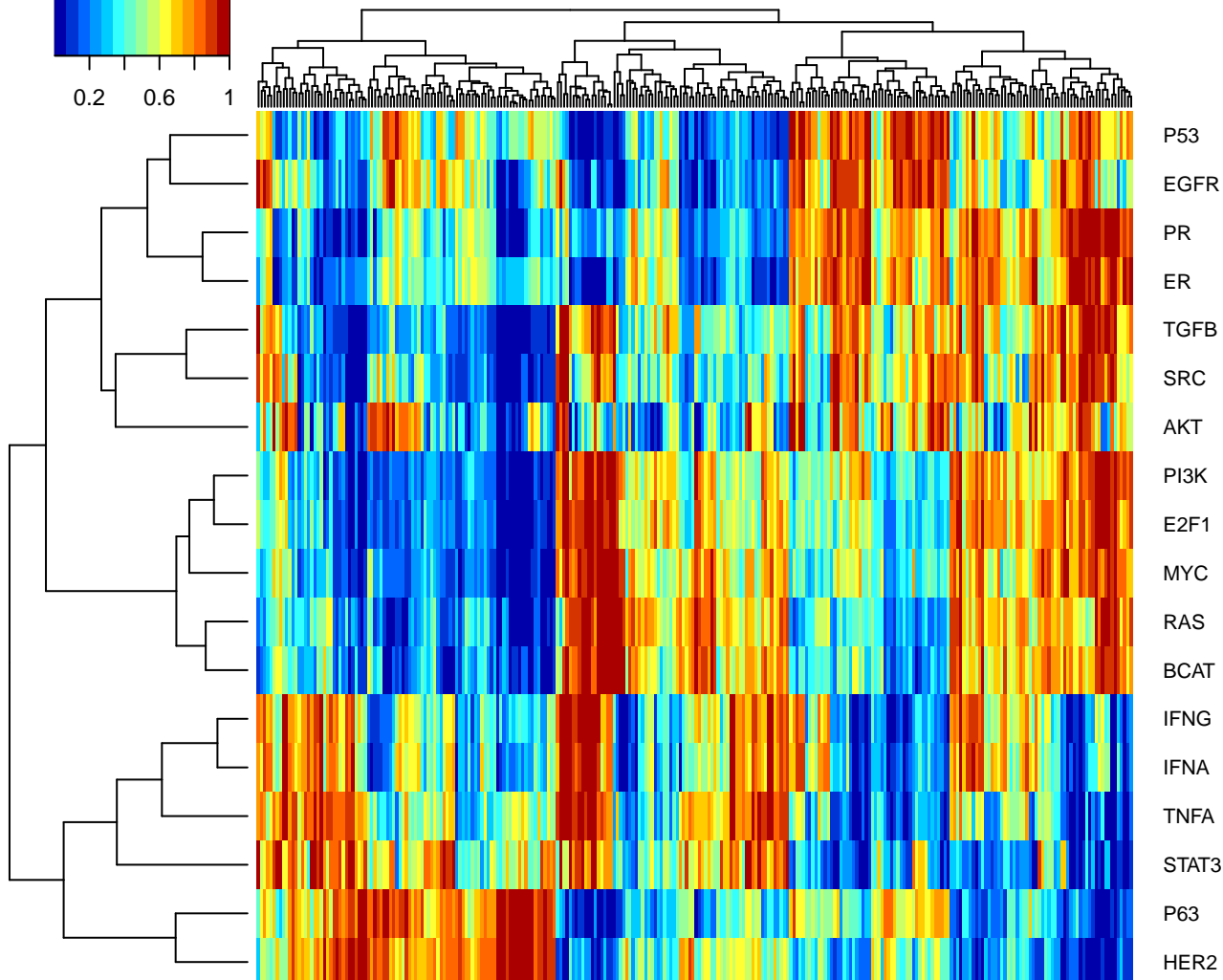


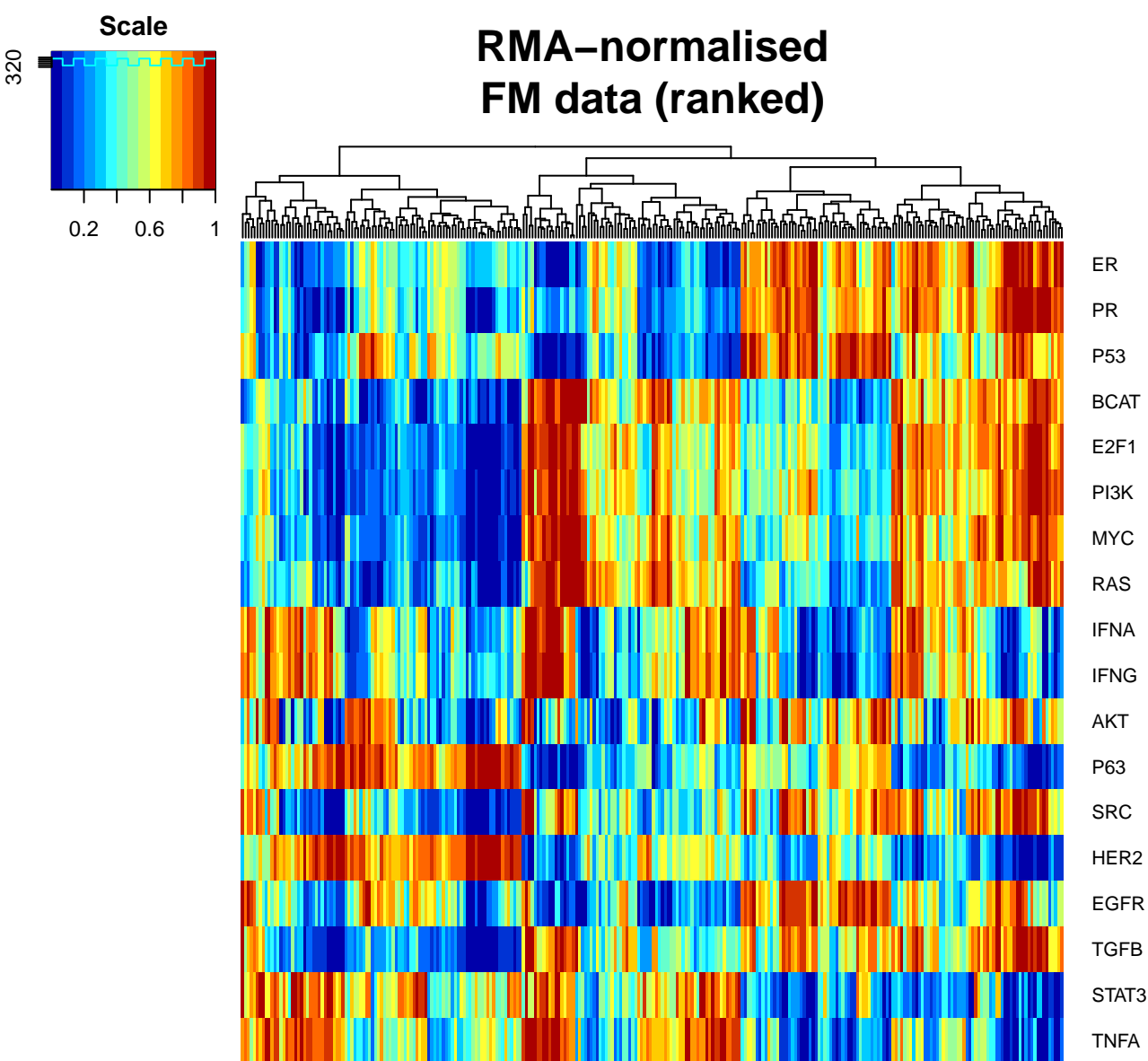
320

Scale



RMA-normalised FM data (ranked)





320

0.2 0.6 1

RMA-normalised FM data (ranked)

ER
PR
P53
BCAT
E2F1
PI3K
MYC
RAS
IFNA
IFNG
AKT
P63
SRC
HER2
EGFR
TGFB
STAT3
TNFA

Scale

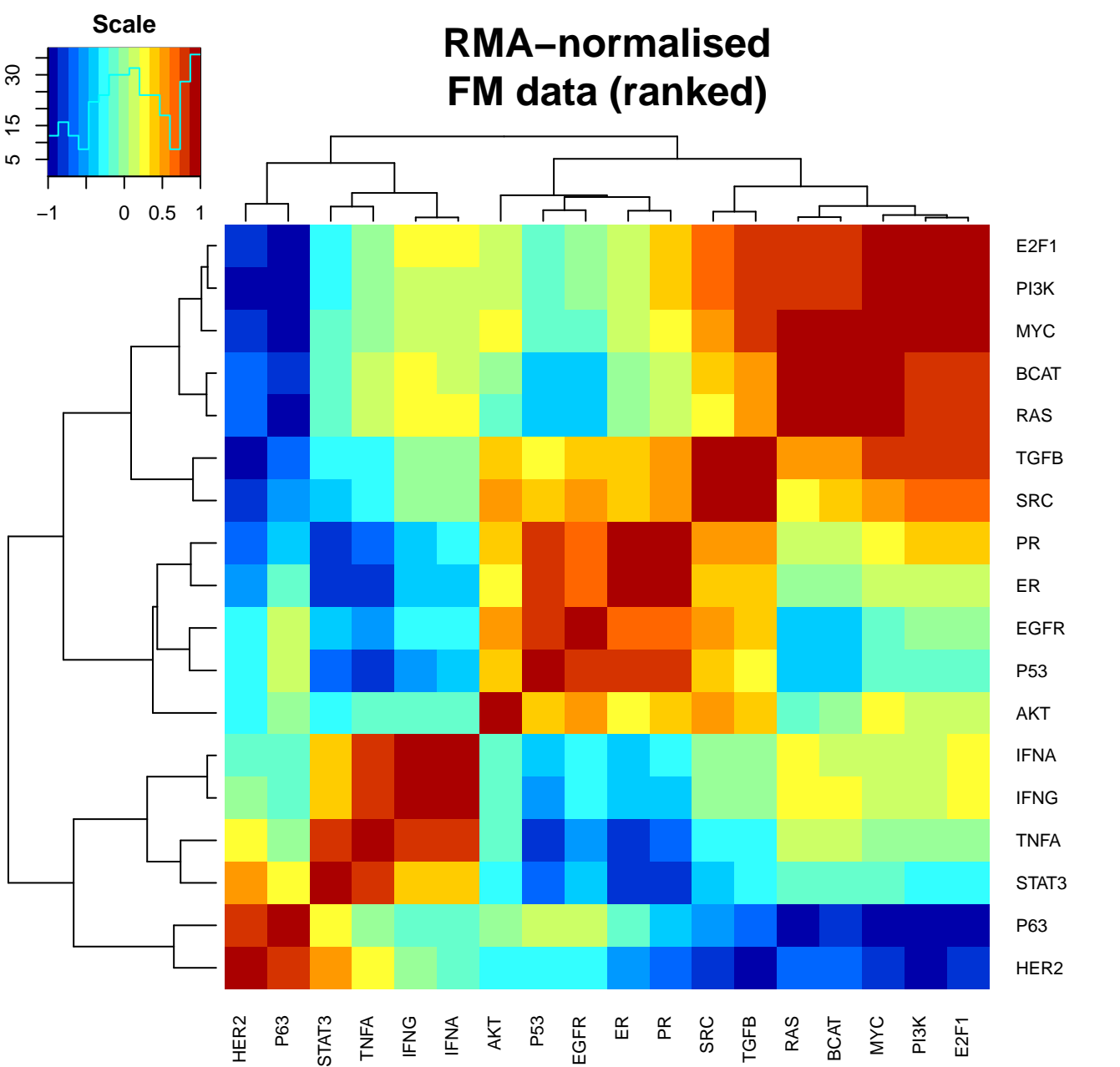
RMA-normalised FM data (ranked)

Scale plot showing values from -1 to 1, with a color gradient from blue to red. The y-axis is labeled 5, 15, 30.

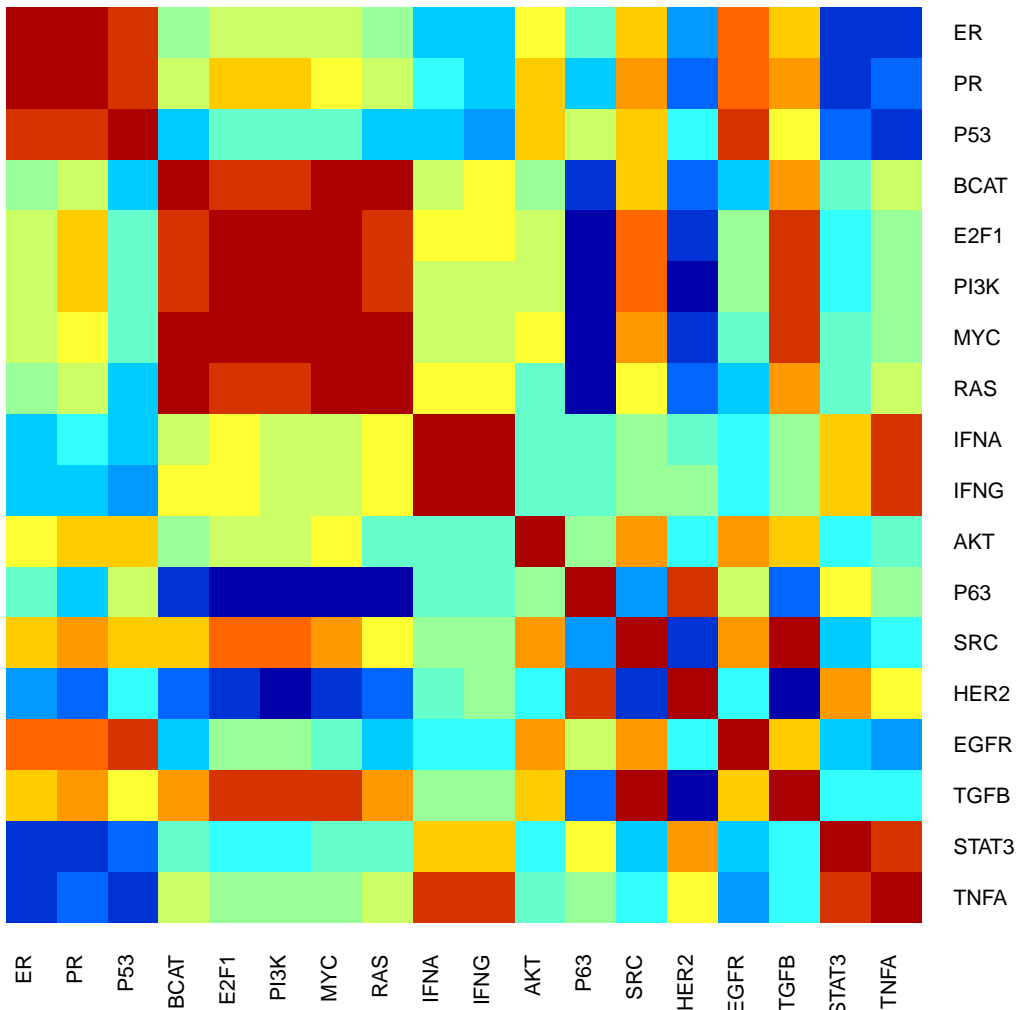
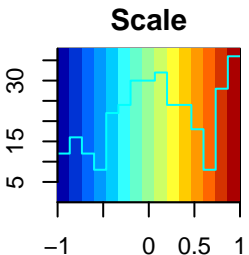
Heatmap showing RMA-normalised FM data (ranked) for 16 genes (rows) across 16 samples (columns). The color scale ranges from -1 (blue) to 1 (red). Hierarchical clustering dendrograms are shown on the top and left.

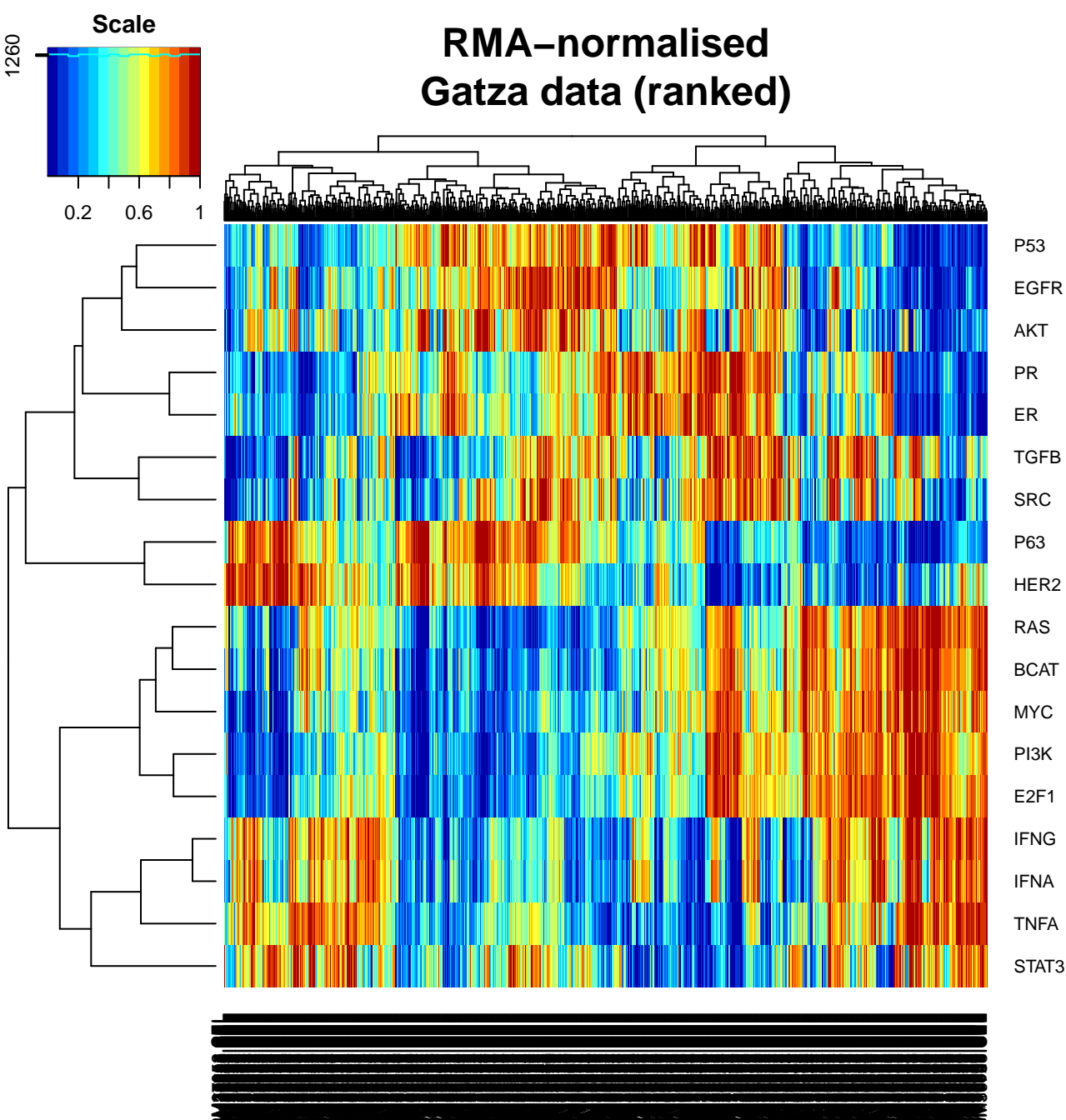
Gene labels (rows): E2F1, PI3K, MYC, BCAT, RAS, TGFB, SRC, PR, ER, EGFR, P53, AKT, IFNA, IFNG, TNFA, STAT3, P63, HER2.

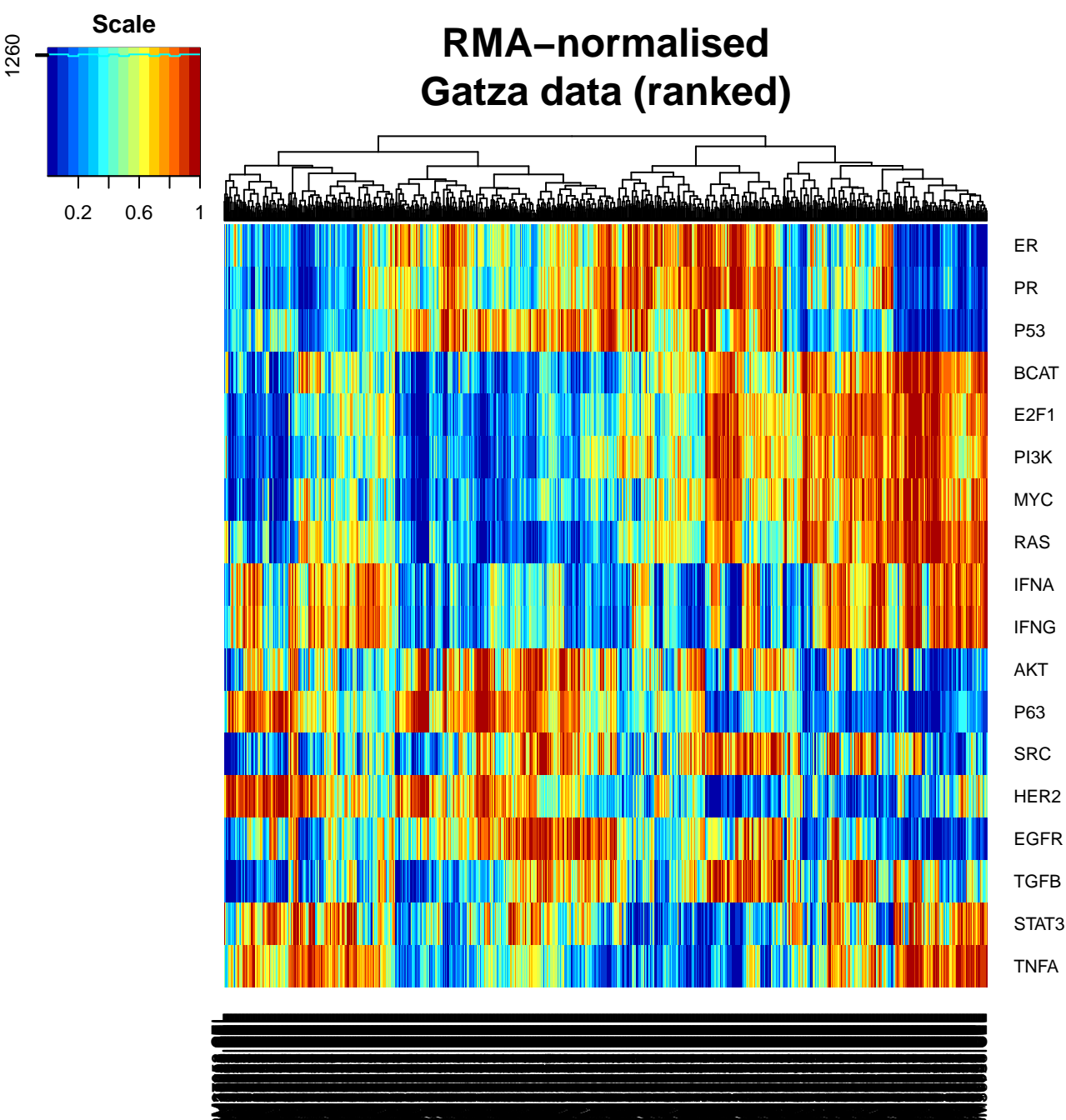
Sample labels (columns): HER2, P63, STAT3, TNFA, IFNG, IFNA, AKT, P53, EGFR, ER, PR, SRC, TGFB, RAS, BCAT, MYC, PI3K, E2F1.



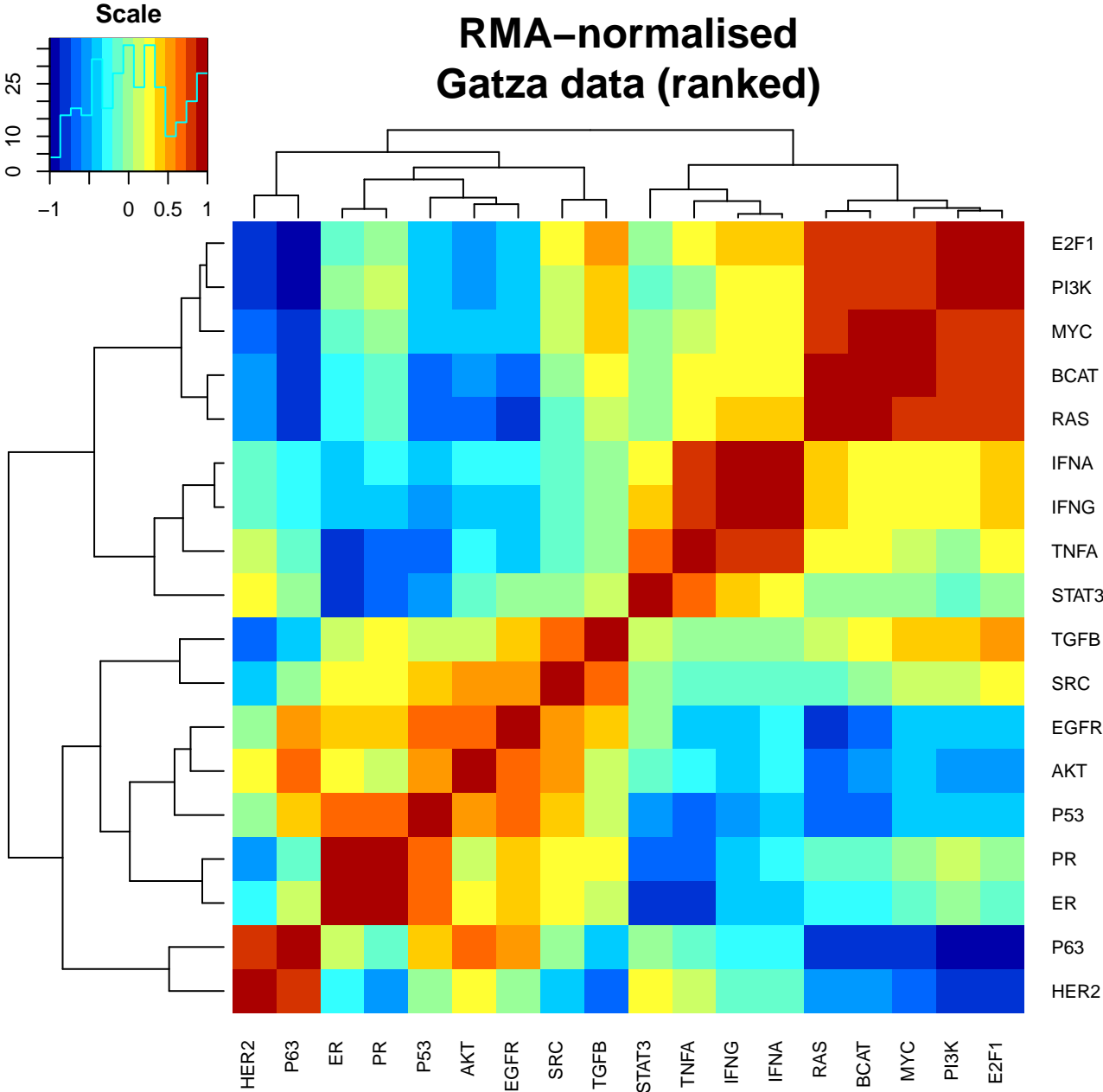
RMA-normalised FM data (ranked)

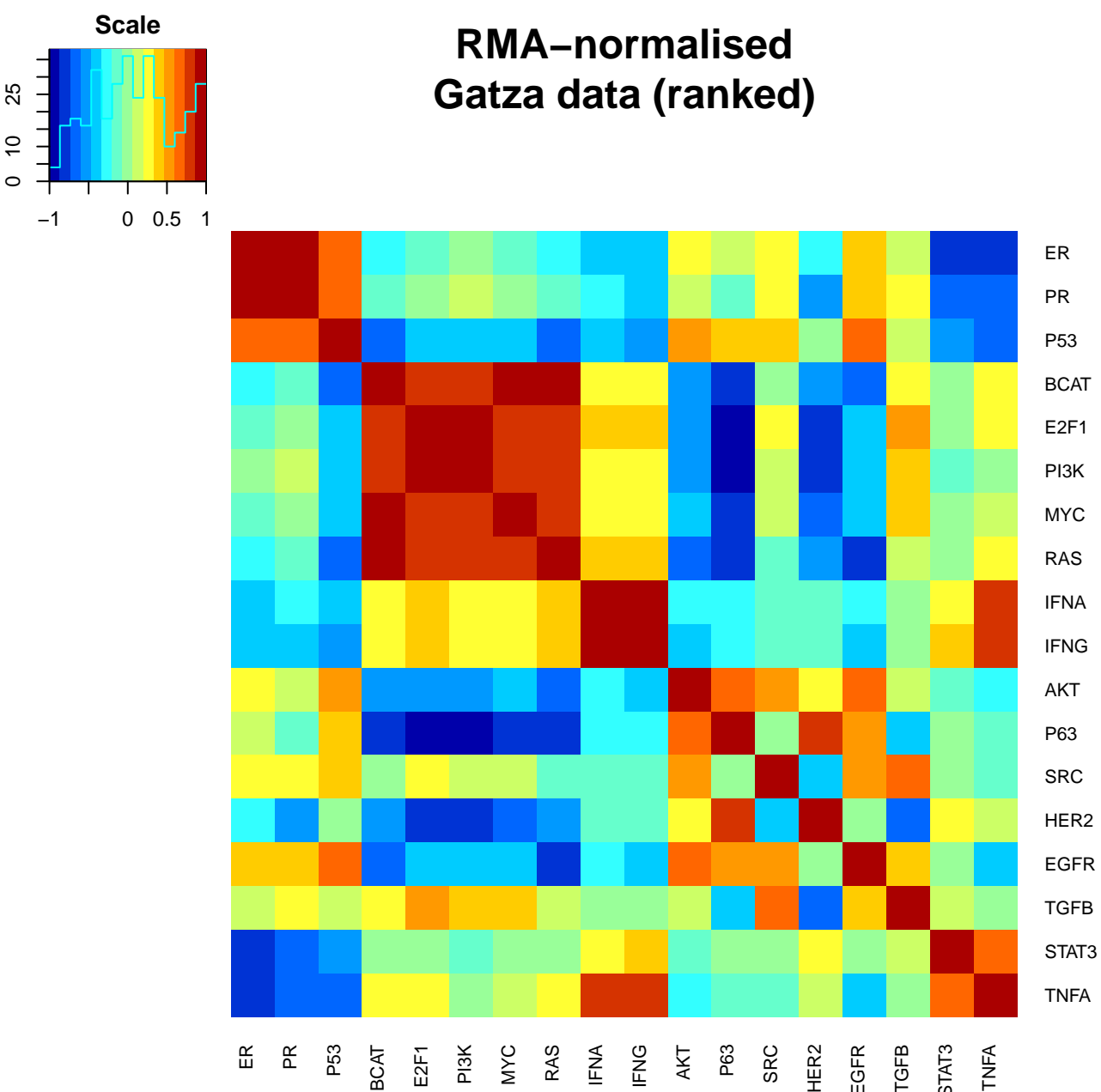


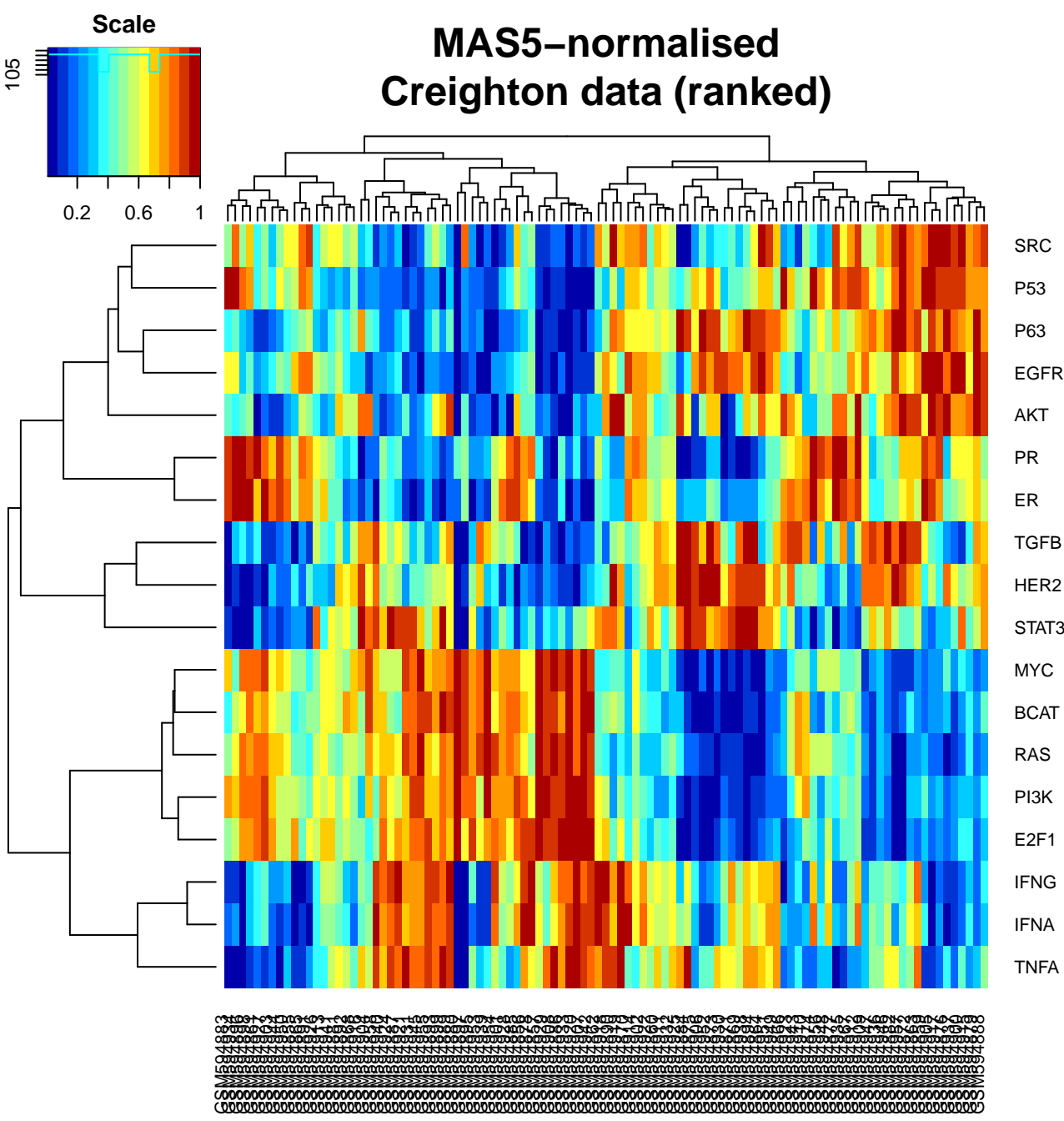


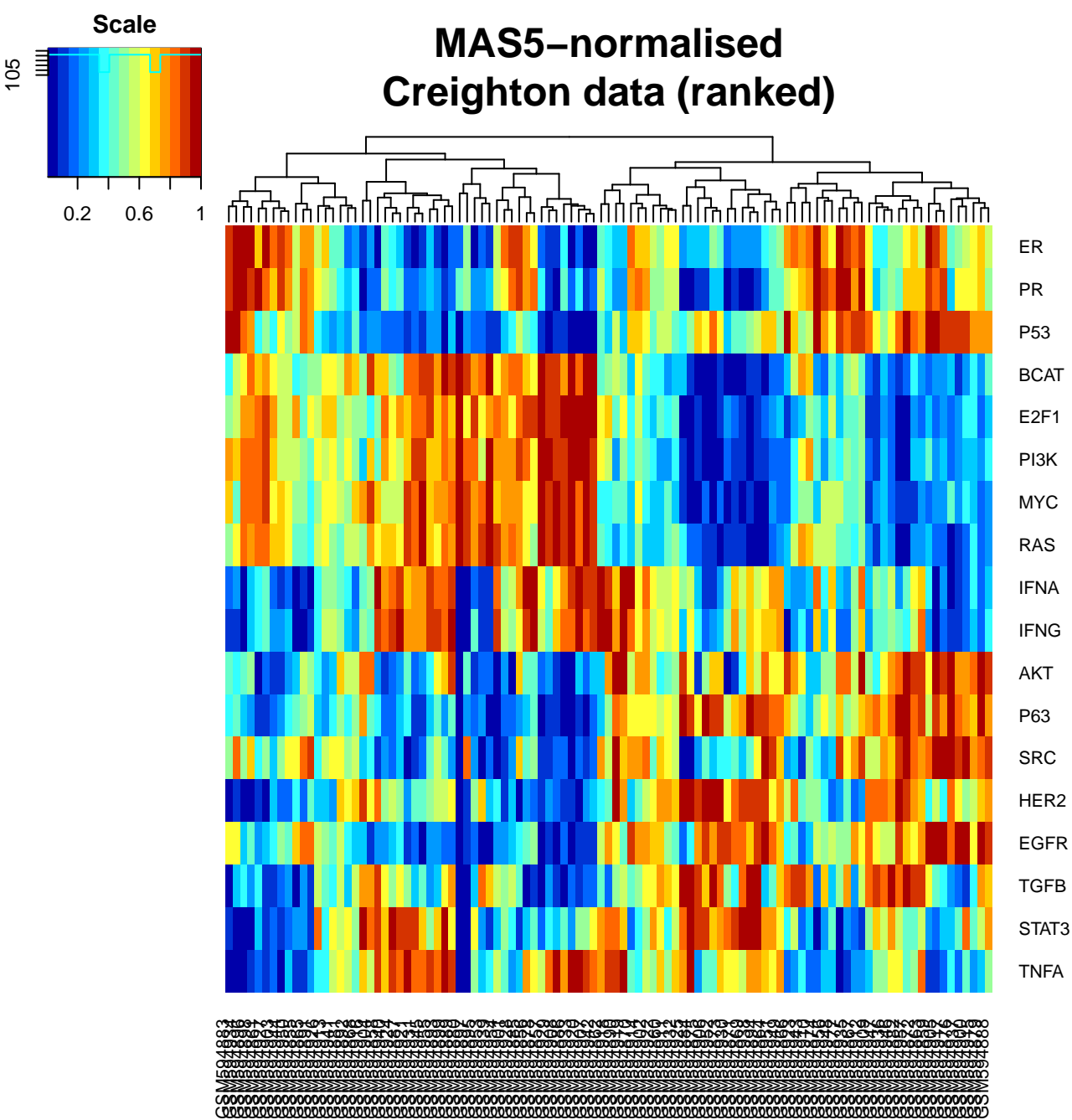


RMA-normalised Gatza data (ranked)

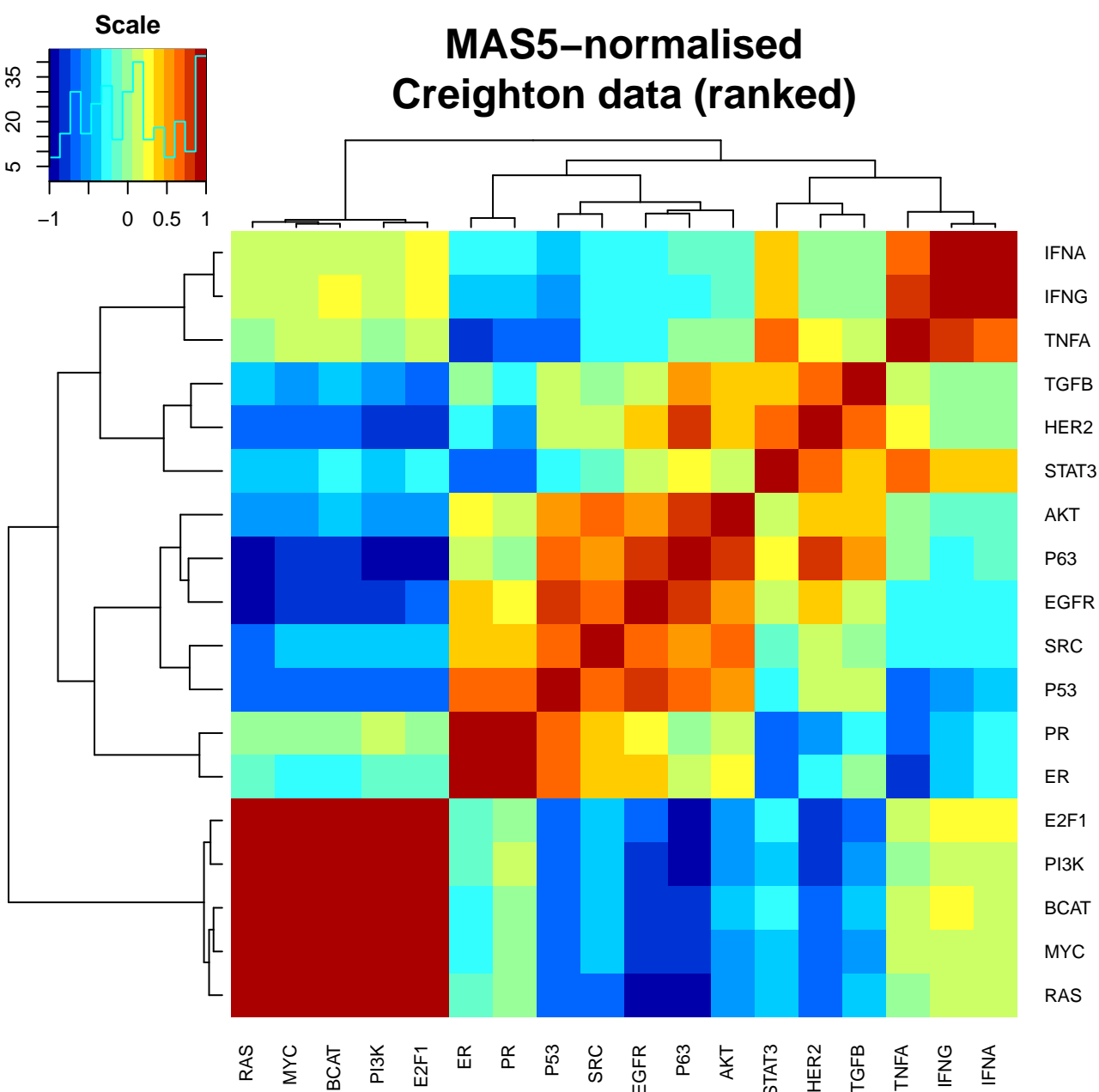




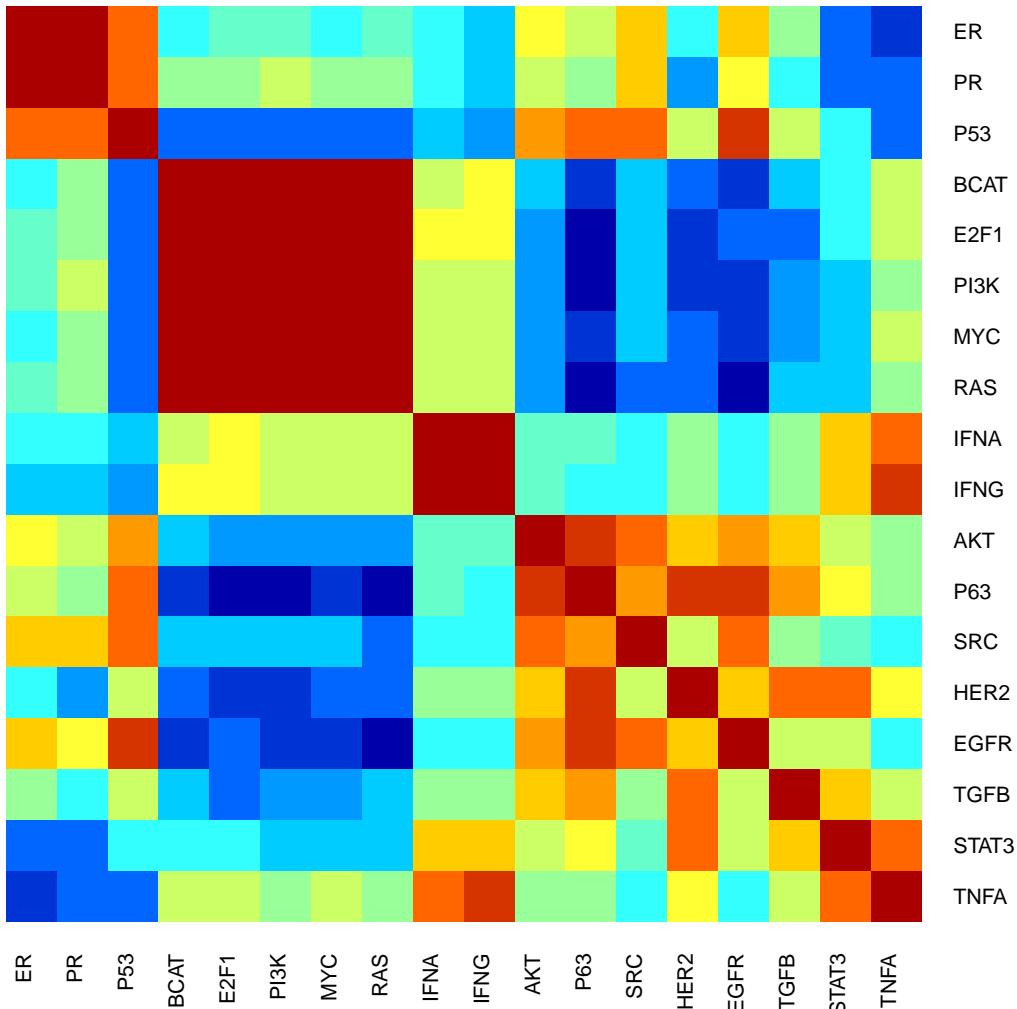
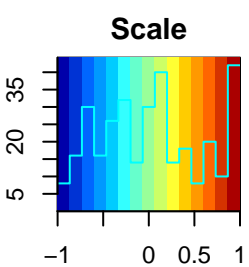




MAS5-normalised Creighton data (ranked)



MAS5-normalised Creighton data (ranked)

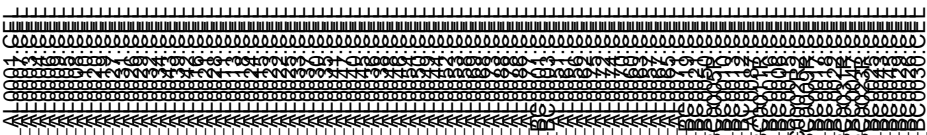


**MAS5-normalised
NZBC data (ranked)**

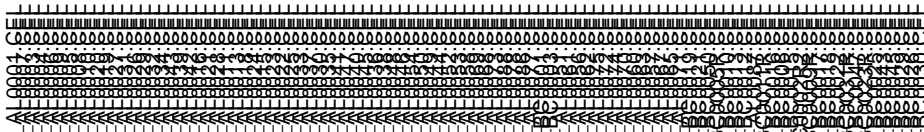
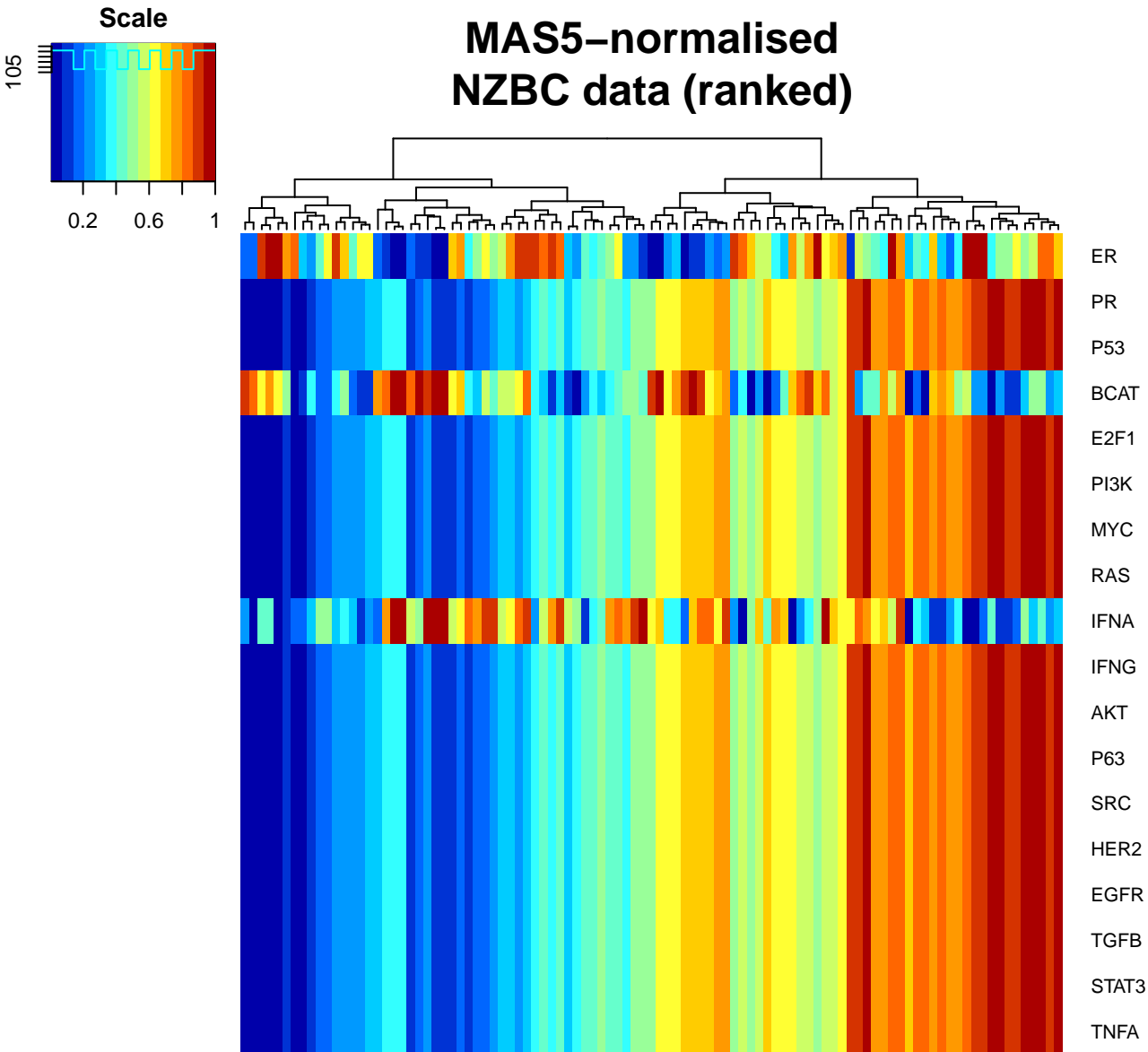
Scale: 0.2 0.6 1

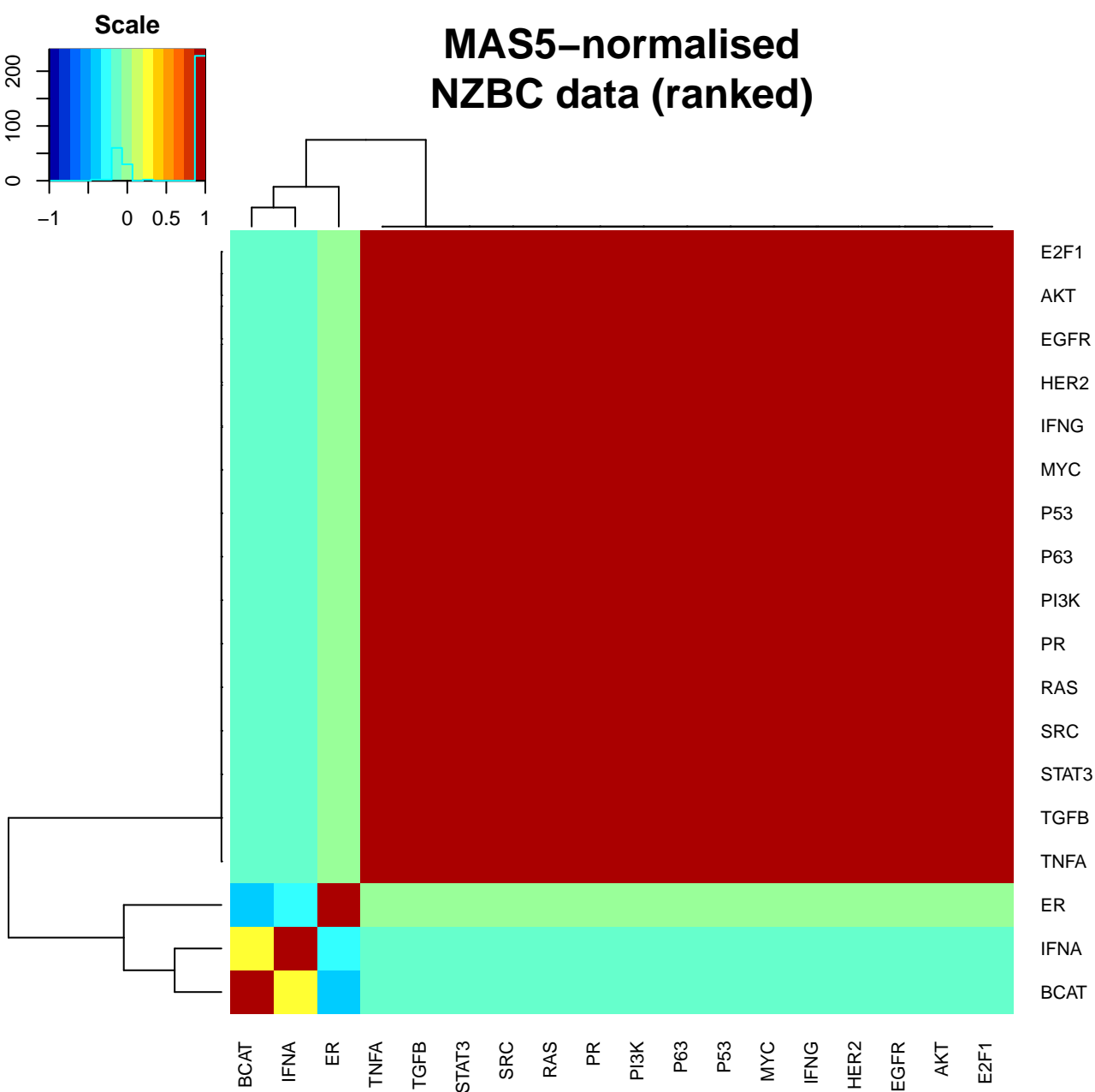
Genes (from top to bottom): E2F1, AKT, EGFR, HER2, IFNG, MYC, P53, P63, PI3K, PR, RAS, SRC, STAT3, TGFB, TNFA, ER, IFNA, BCAT.

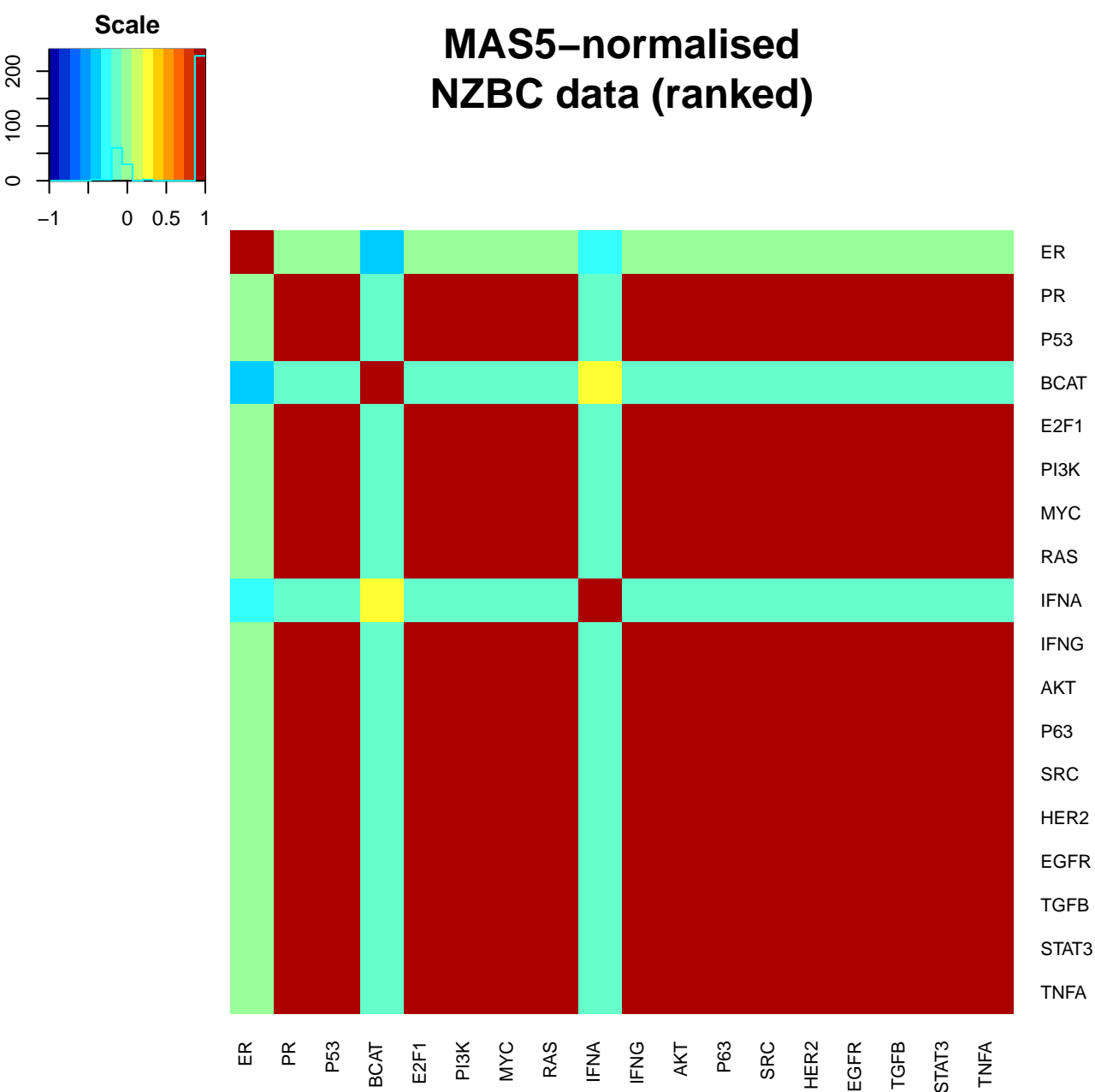
E2F1
AKT
EGFR
HER2
IFNG
MYC
P53
P63
PI3K
PR
RAS
SRC
STAT3
TGFB
TNFA
ER
IFNA
BCAT



NZBC data (ranked)









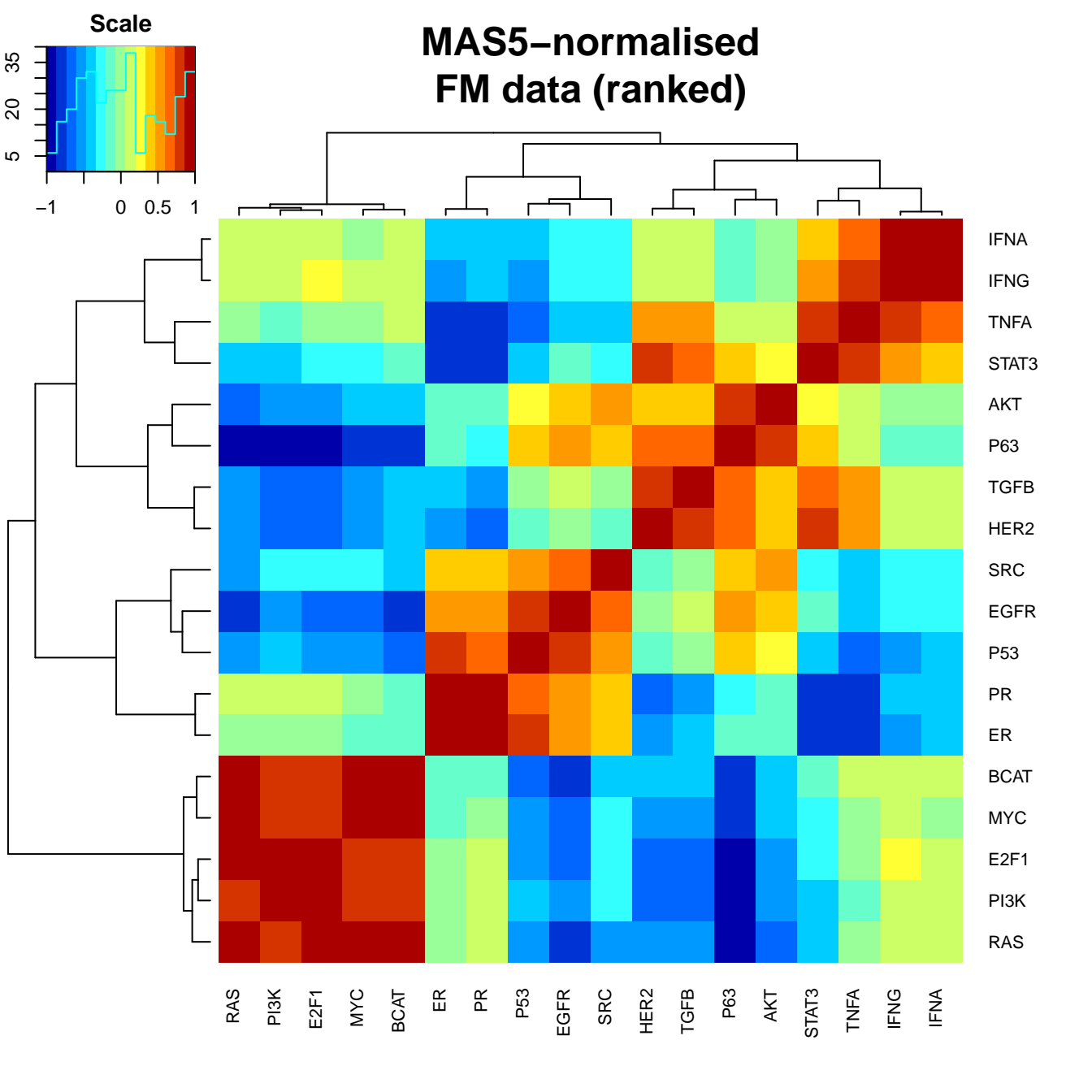
**MAS5-normalised
FM data (ranked)**

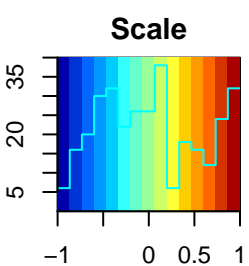
Scale

35
20
5
-1 0 0.5 1

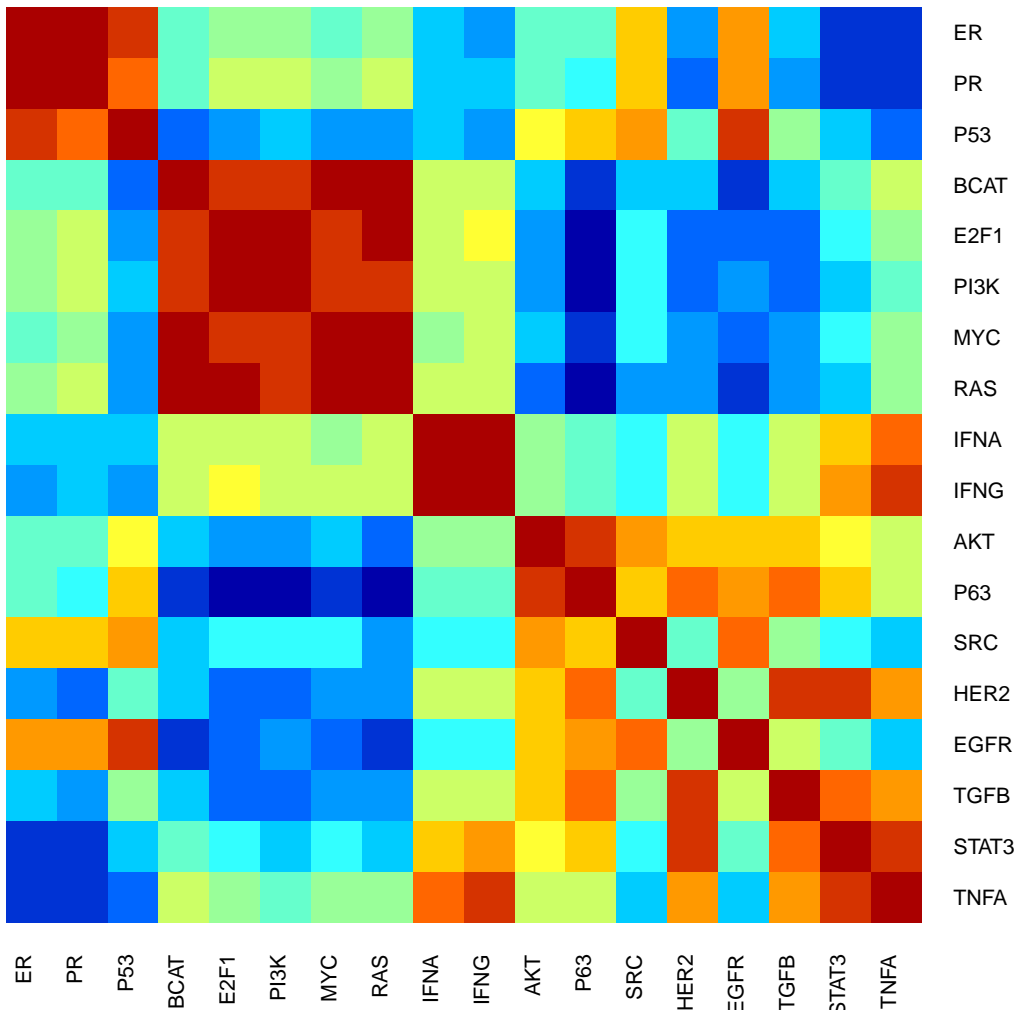
IFNA
IFNG
TNFA
STAT3
AKT
P63
TGFB
HER2
SRC
EGFR
P53
PR
ER
BCAT
MYC
E2F1
PI3K
RAS

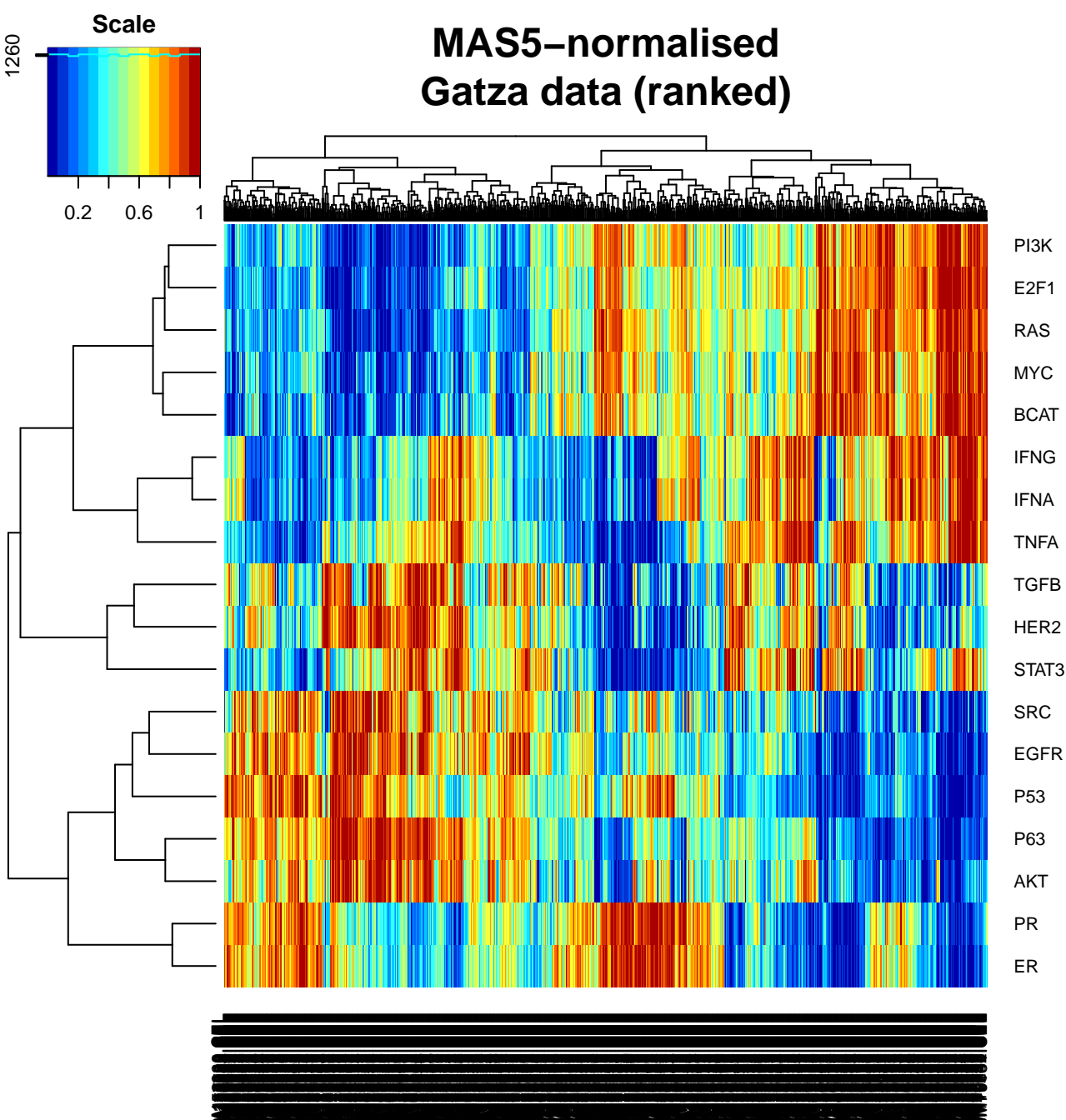
RAS
PI3K
E2F1
MYC
BCAT
ER
PR
P53
EGFR
SRC
HER2
TGFB
P63
AKT
STAT3
TNFA
IFNG
IFNA

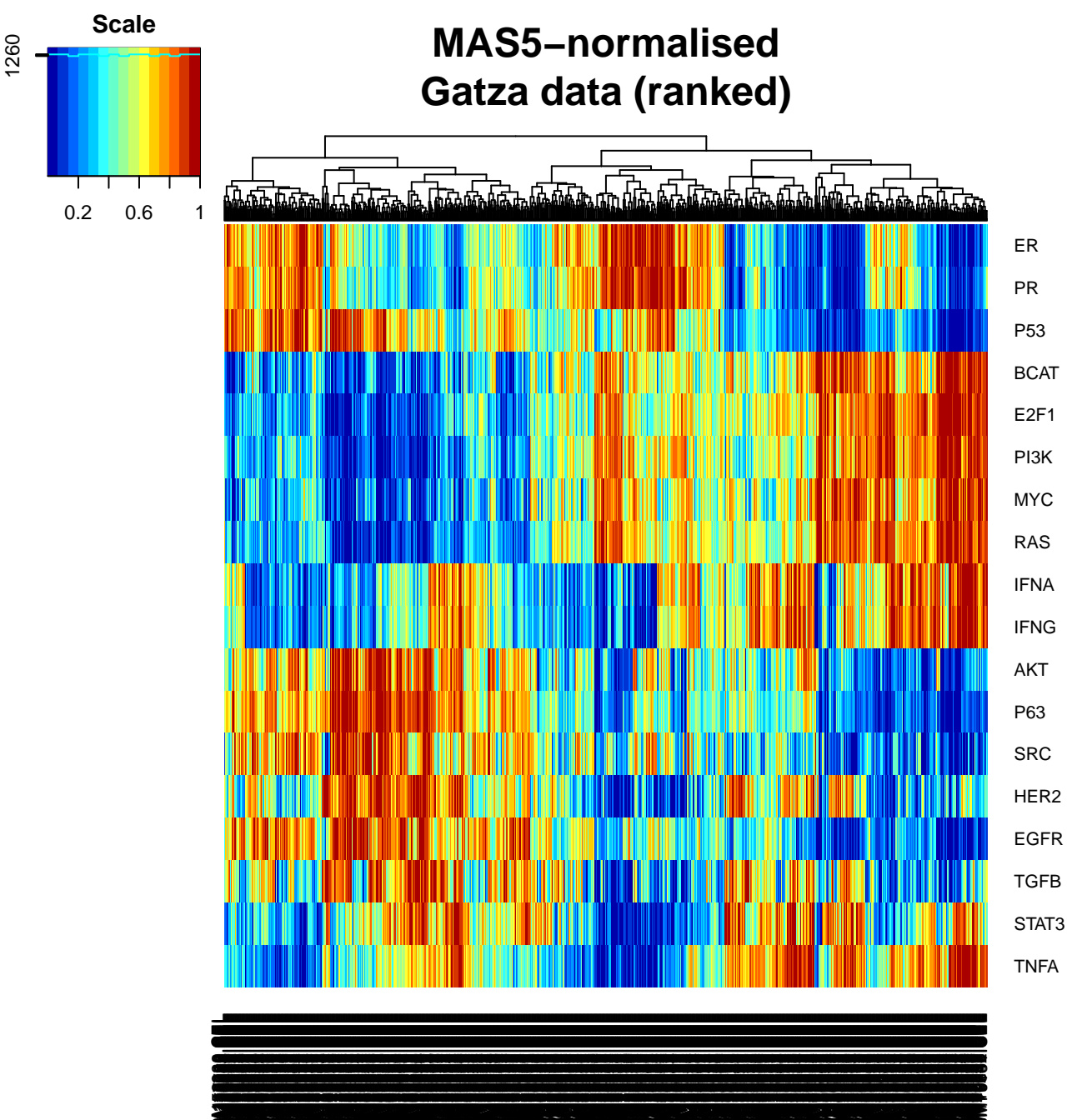




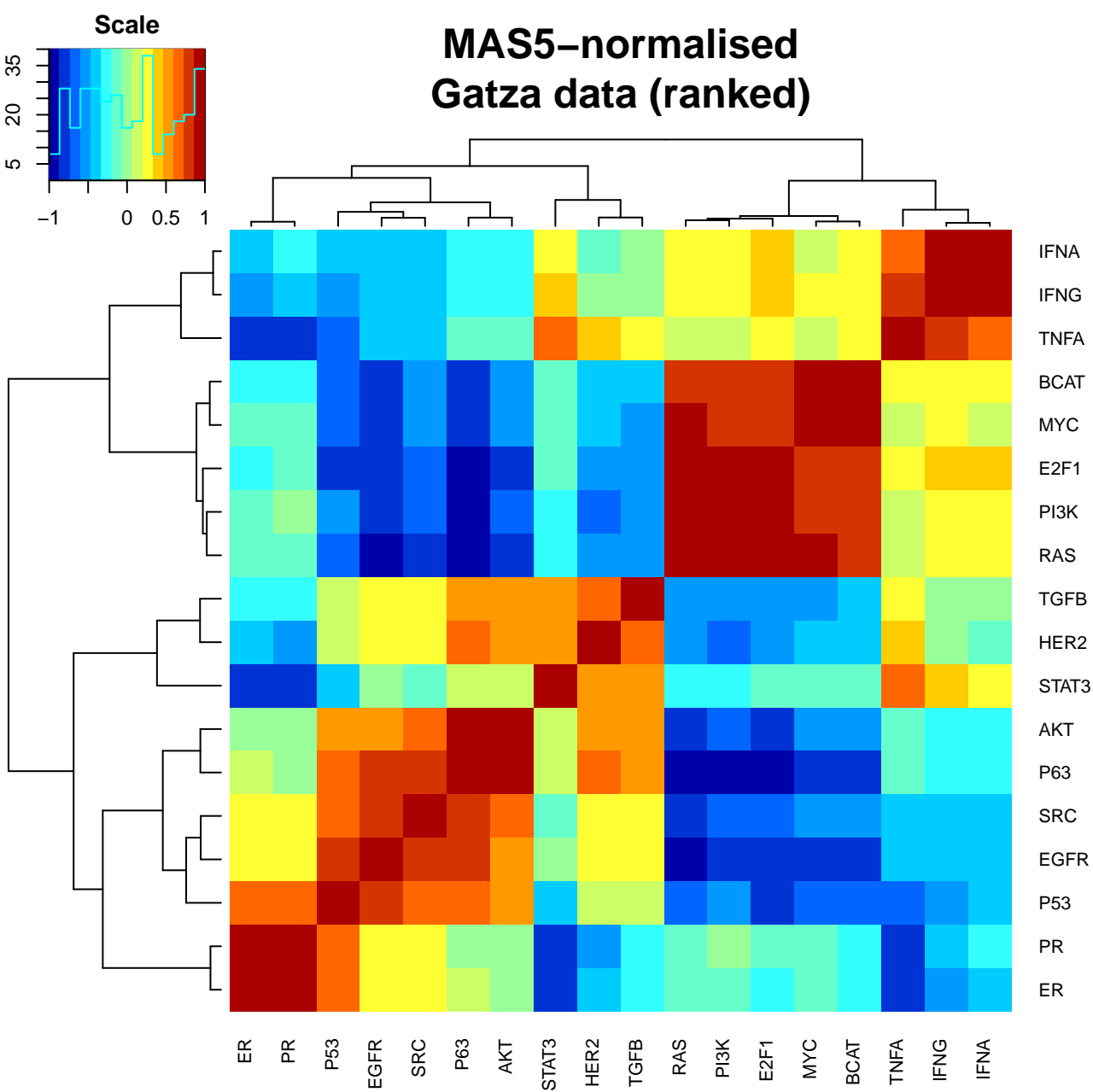
MAS5-normalised FM data (ranked)

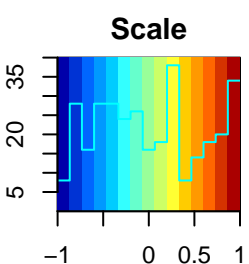






MAS5-normalised Gatza data (ranked)





MAS5-normalised Gatza data (ranked)

