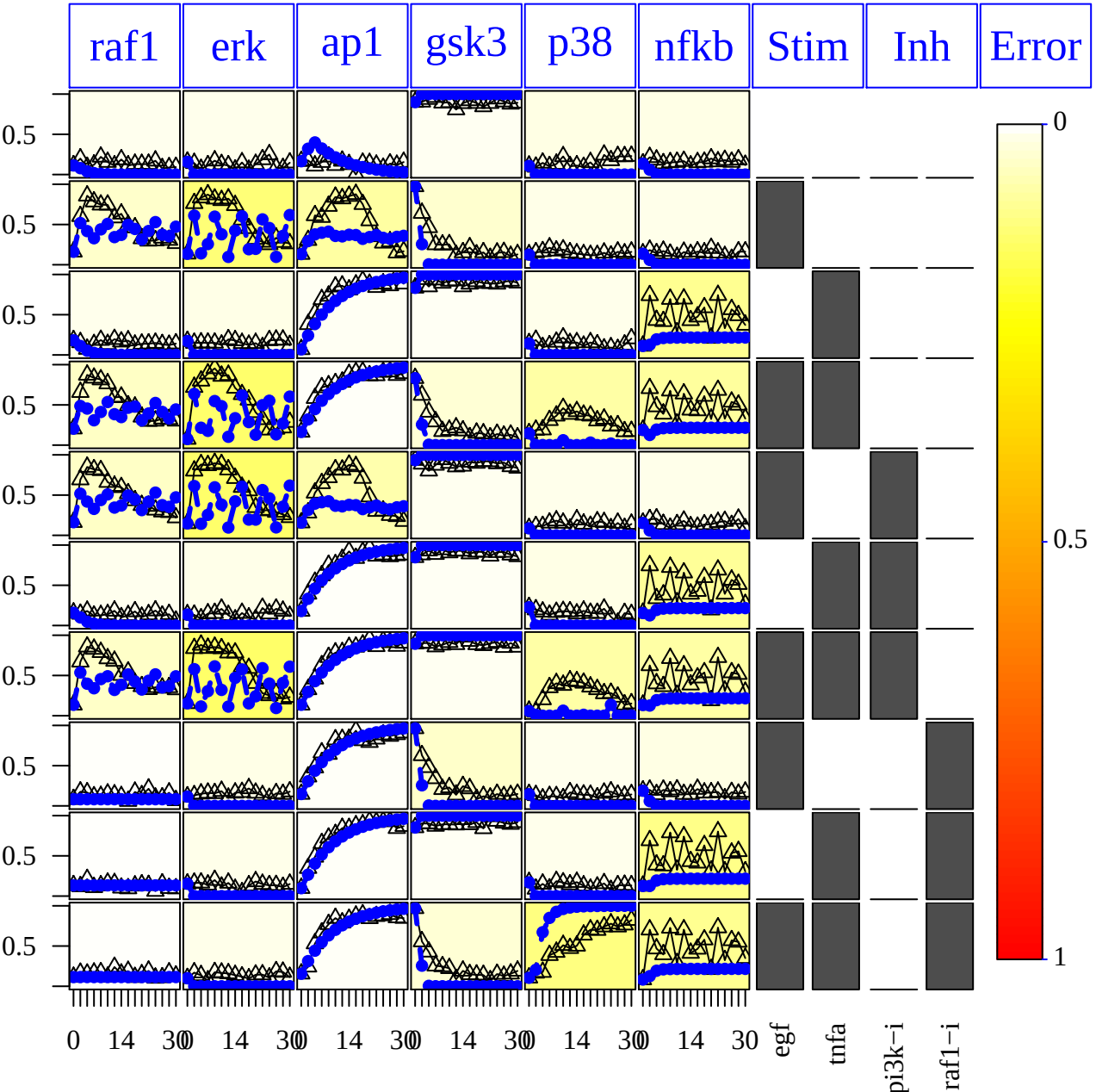
raf1
erk
ap1
gsk3
p38
nfkb
Stim
Inh
Error
0.5
0
14
30
egf
tnfa
pi3k−i
raf1−i
0
0.5
1