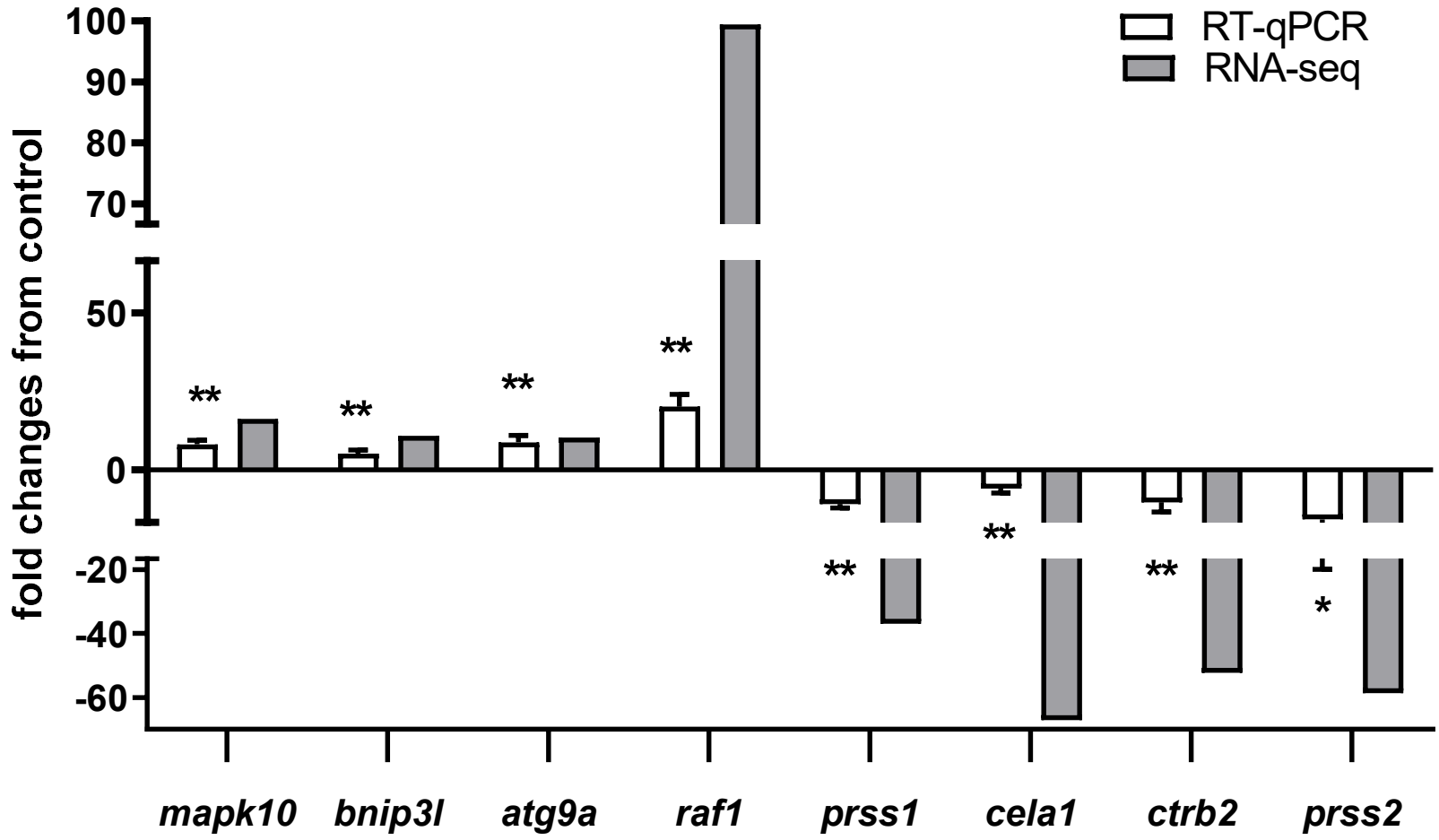
RT-qPCR
RNA-seq
fold changes from control
100
90
80
70
50
0
-20
-40
-60
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*
mapk10
bnip3l
atg9a
raf1
prss1
cela1
ctrb2
prss2