

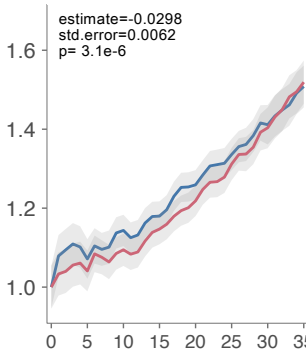


Cell line     mock     IncTAM34a

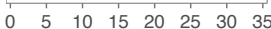
RPMI

HBSS

Fold confluency



estimate=-0.2153  
std.error=0.0061  
p= 4.3e-85



Time (hours)