

RNAPII ChIP-seq amplicon coverage

10

1

0.1

0 0.1

1

5

average ChIP qPCR fold enrichment over input

+ Maximum amplicon coverage depth
 $r(s) = 0.683$
 $p < 0.01$

● Average amplicon coverage depth
 $r(s) = 668$
 $p < 0.01$