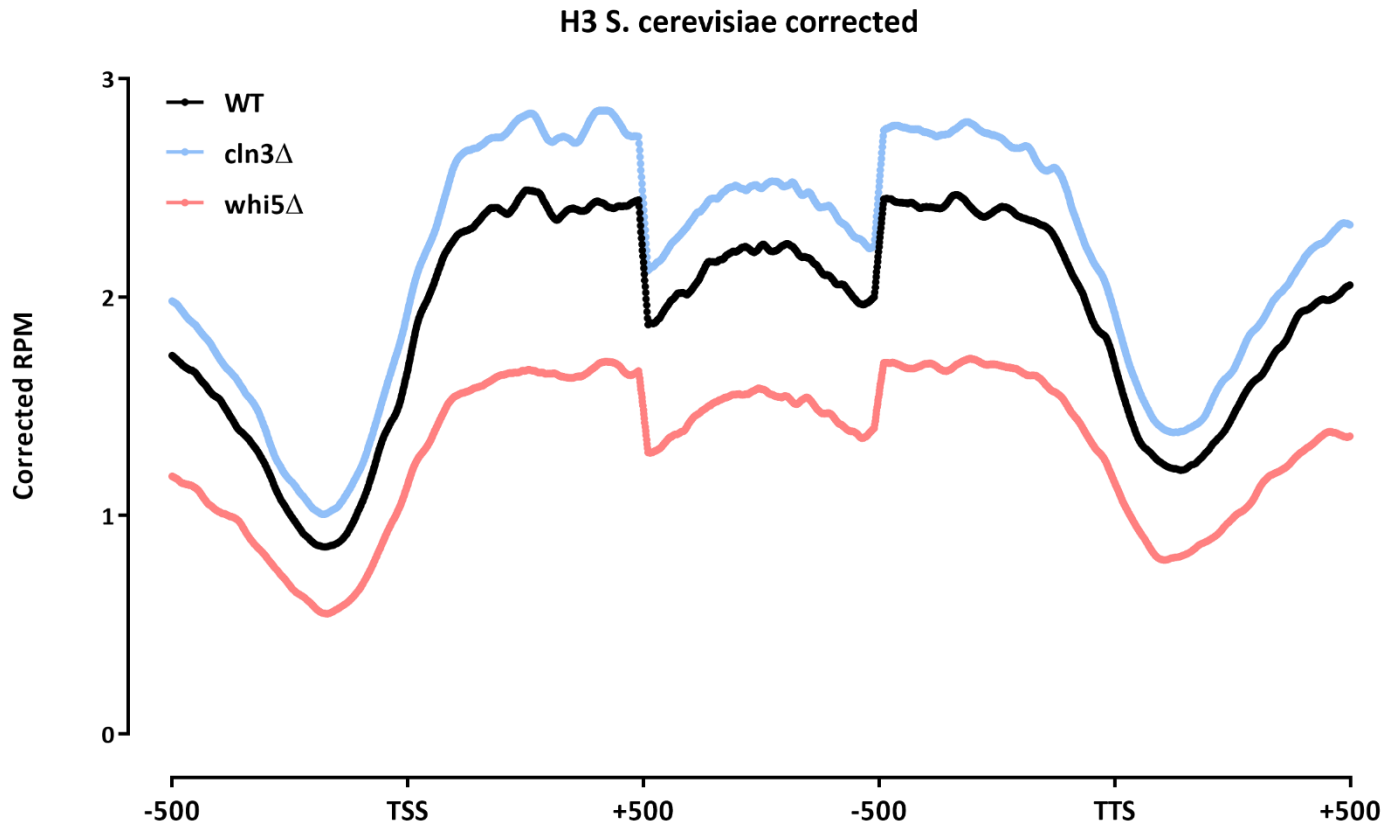
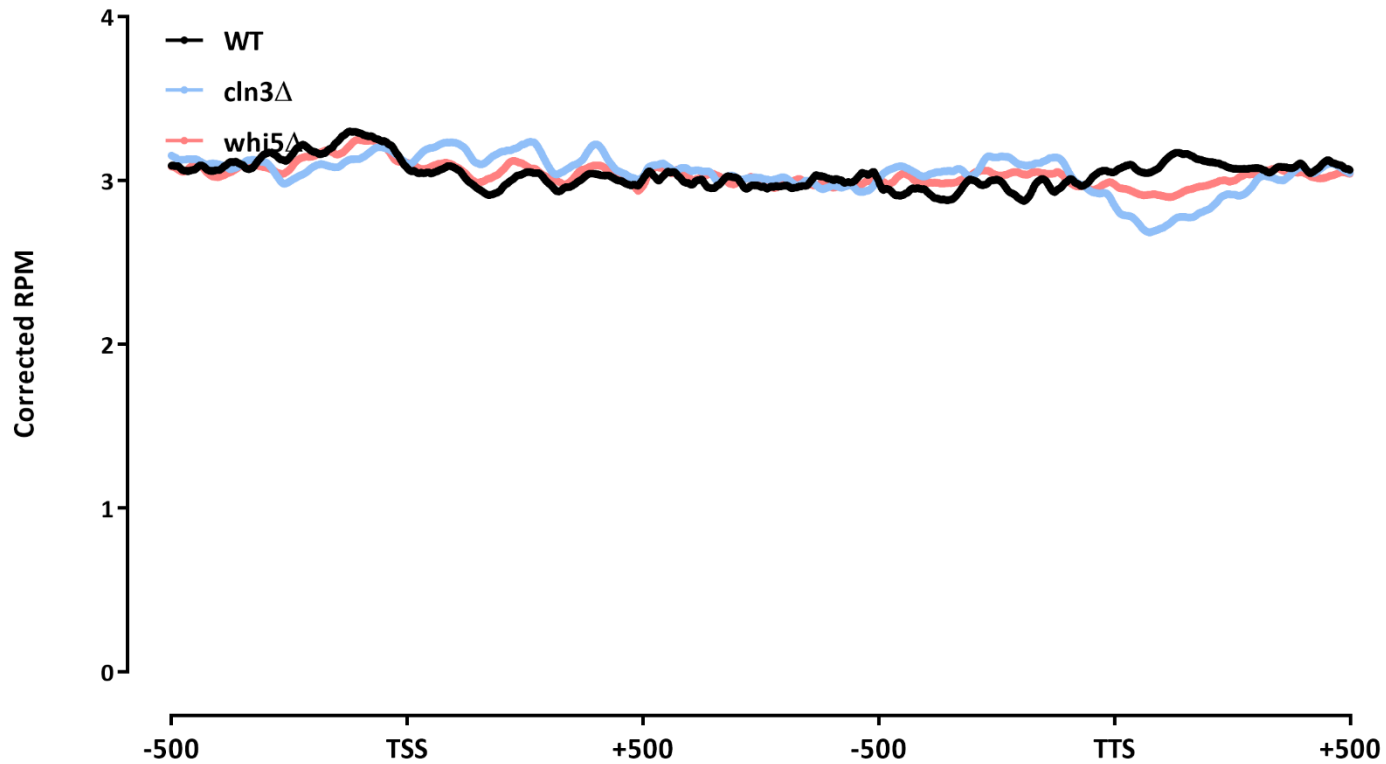


# Protein coding bioType

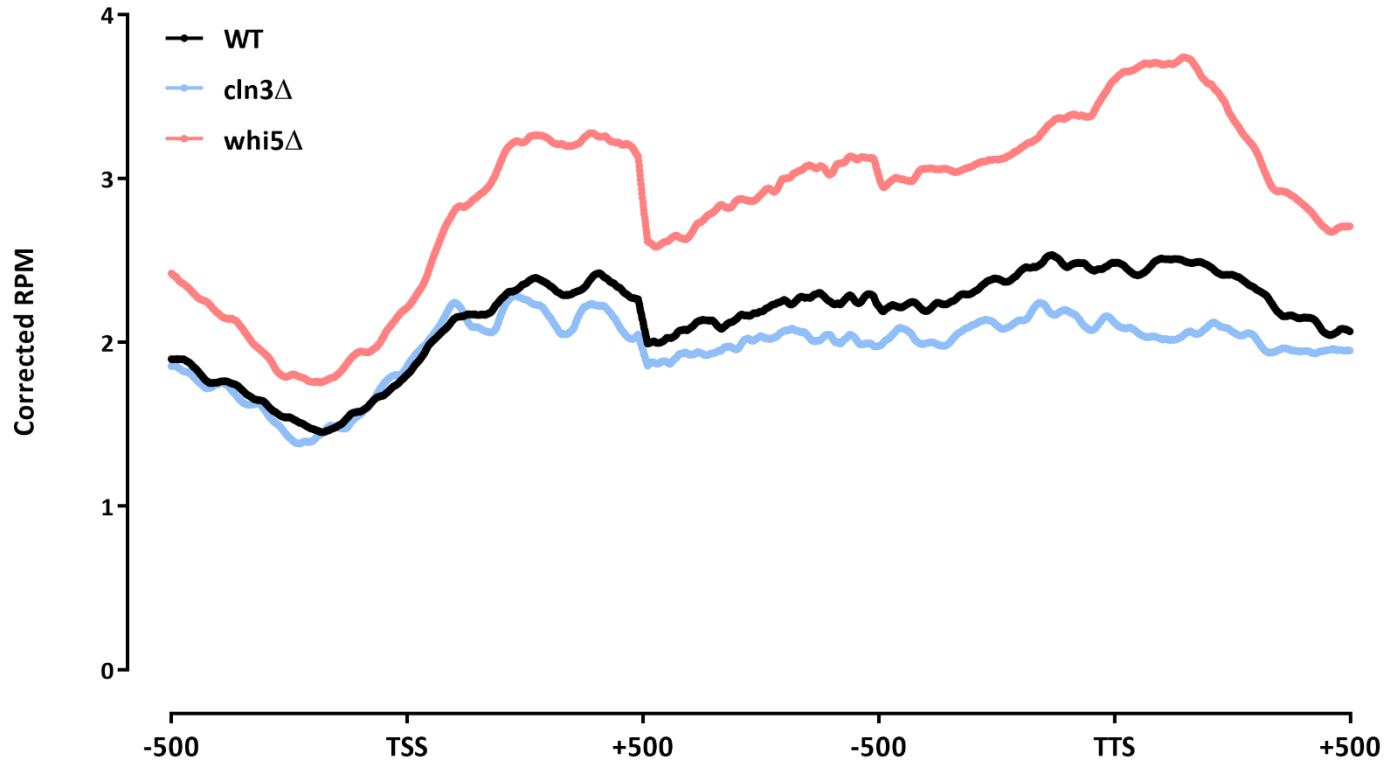
Note:  
Cln3 and Whi5 are,  
switched, but I have  
kept the old names



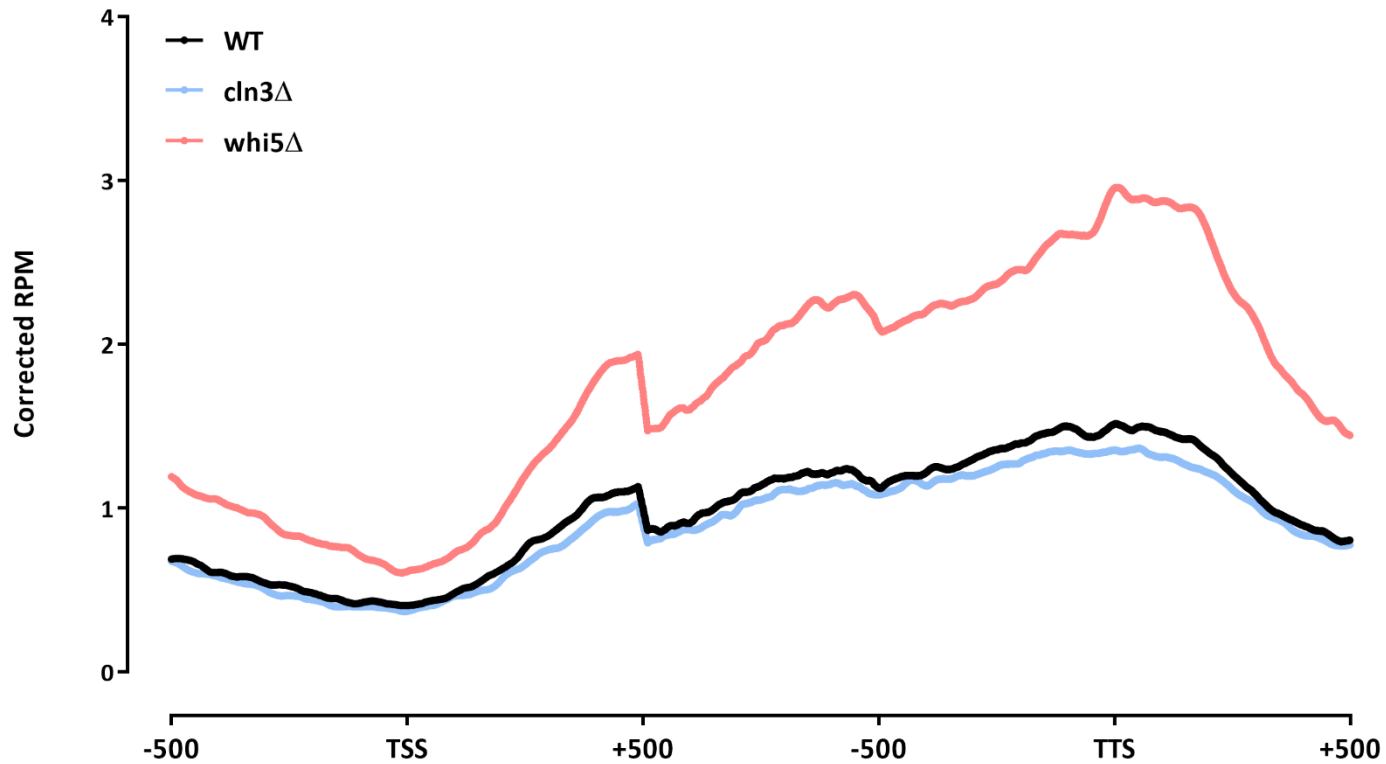
### Input *S. cerevisiae* corrected



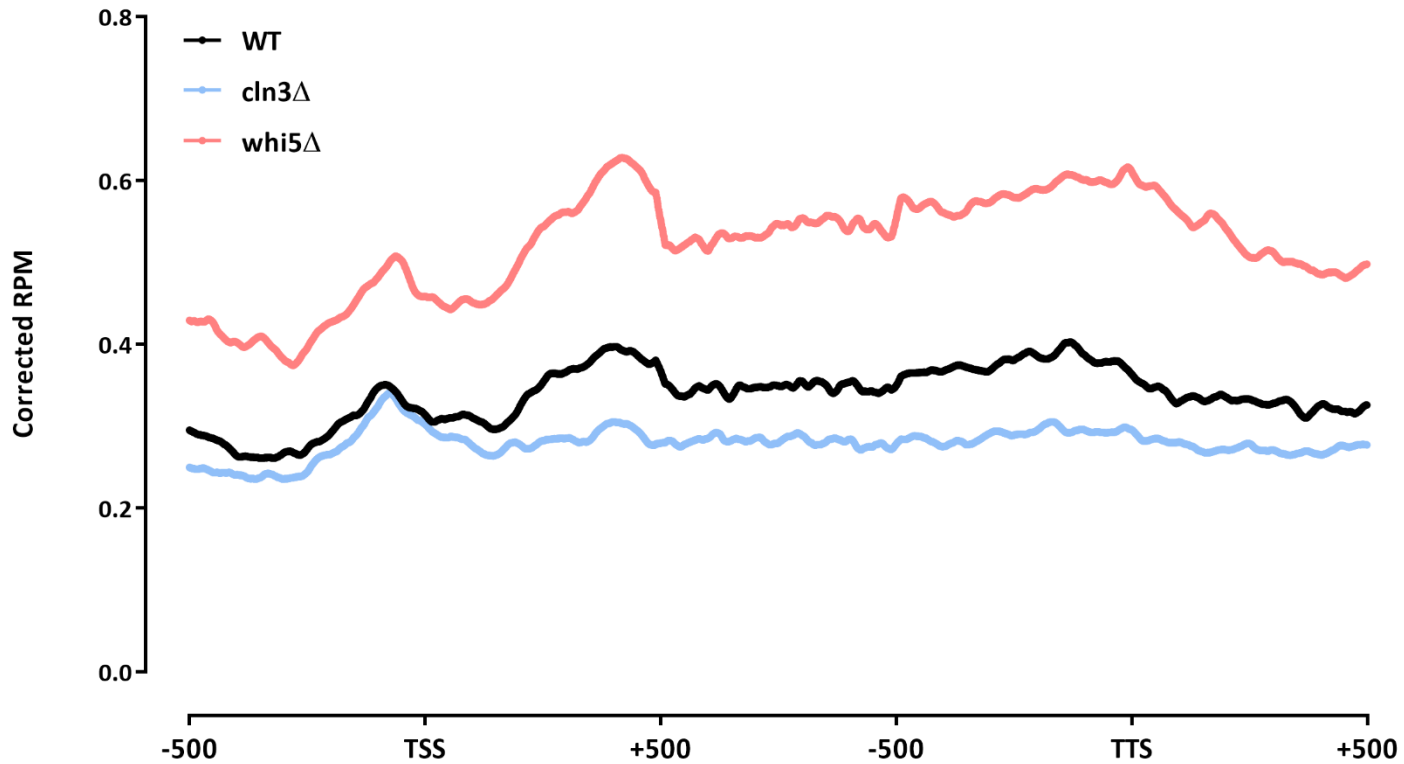
### Pol2 *S. cerevisiae* corrected



### Pol2pS2 *S. cerevisiae* corrected



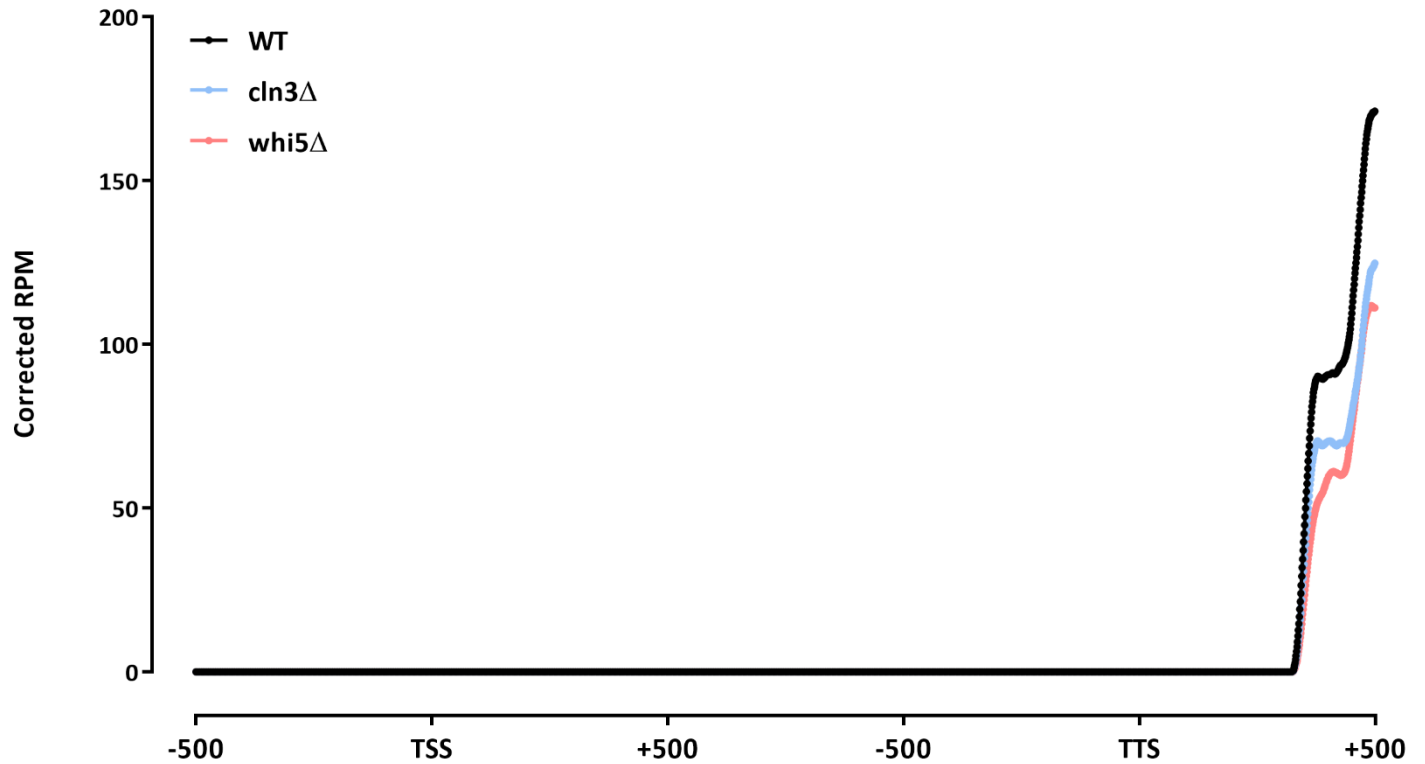
### Pol2pS5 S. cerevisiae corrected



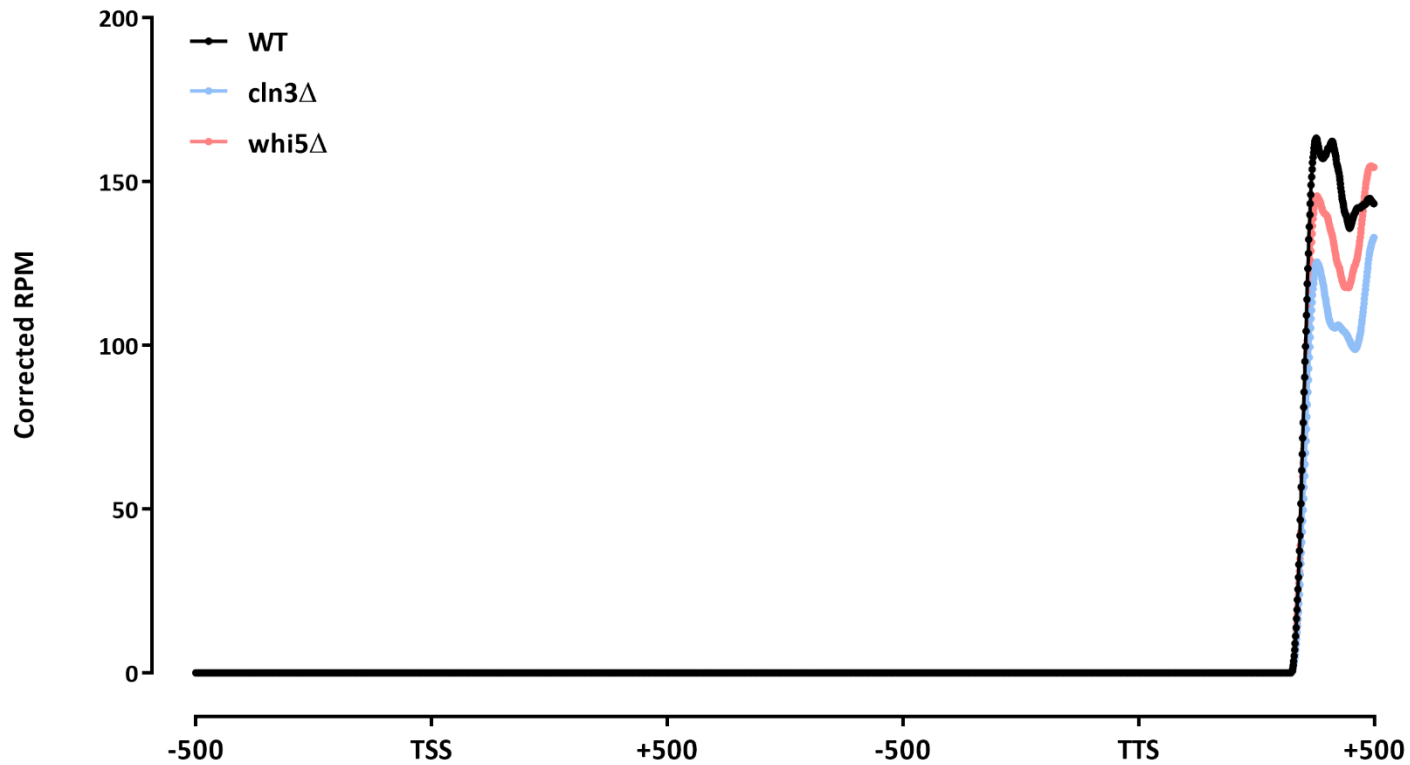
# rRNA bioType

Note:  
There is a  
mappability issue  
with the rDNA loci

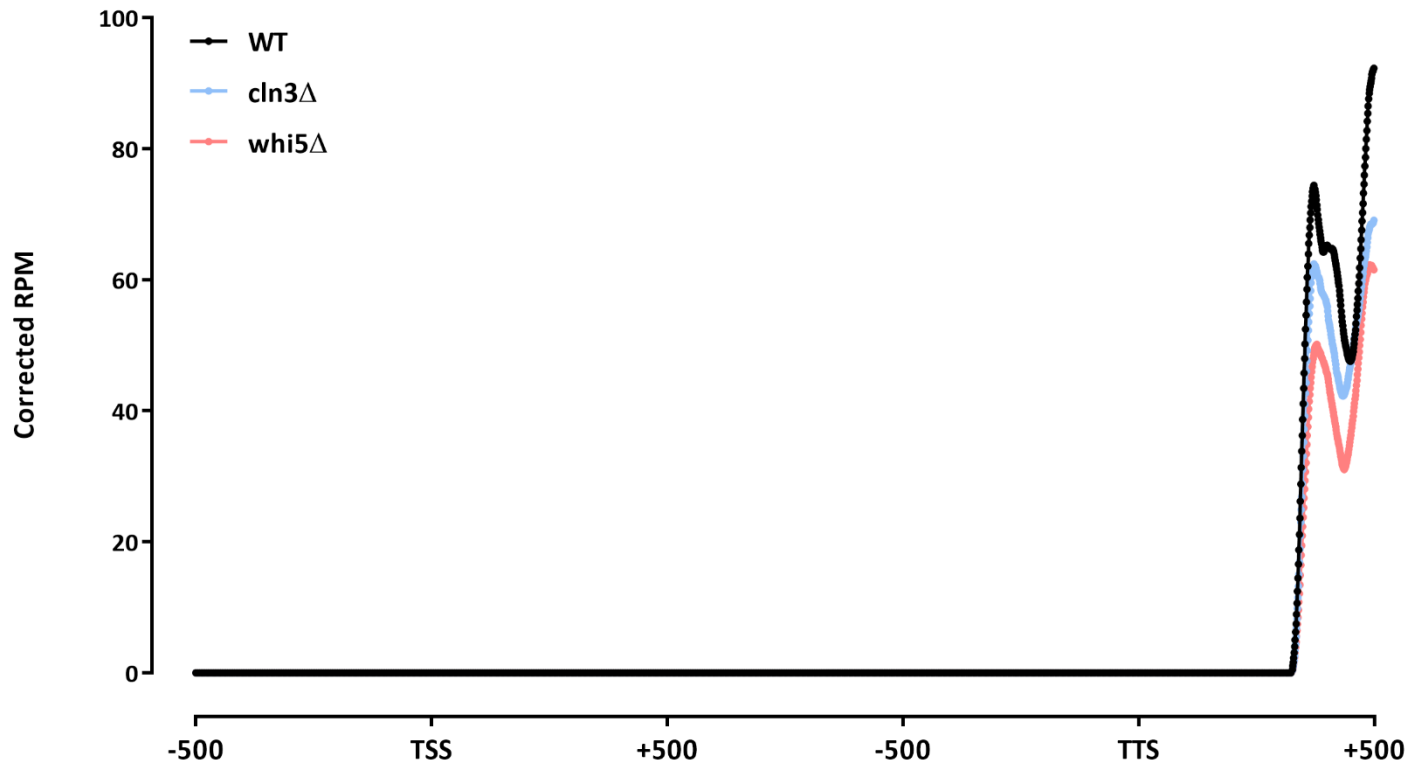
H3 *S. cerevisiae* corrected



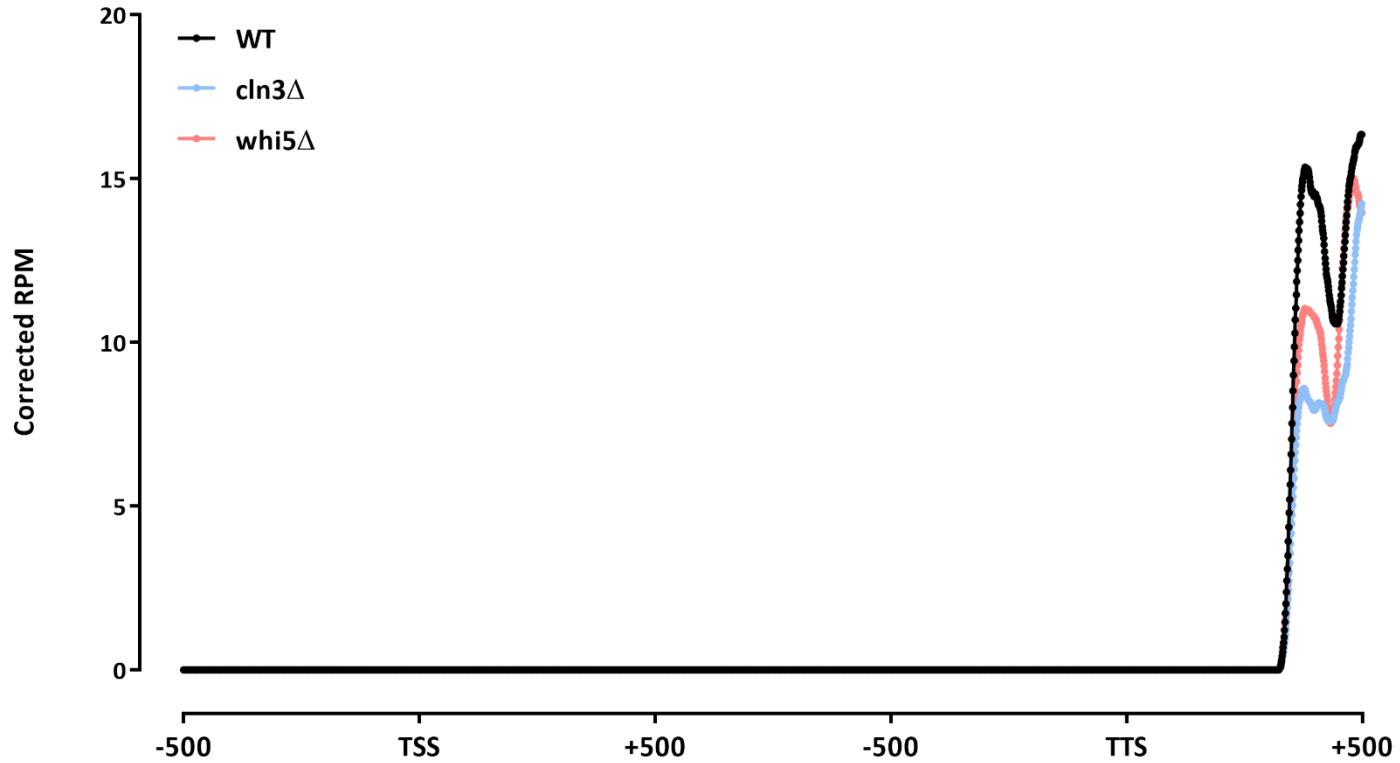
### Input *S. cerevisiae* corrected



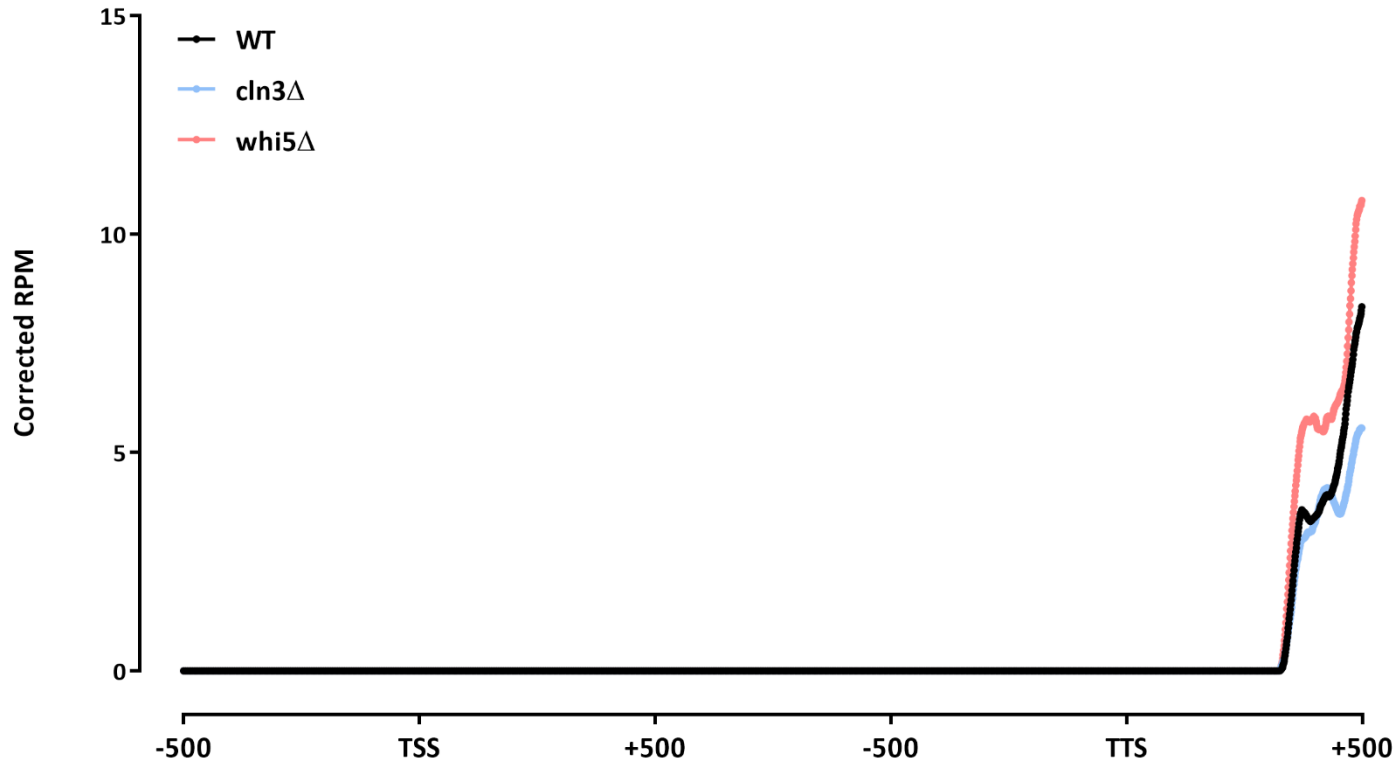
### Pol2 *S. cerevisiae* corrected



### Pol2pS2 *S. cerevisiae* corrected

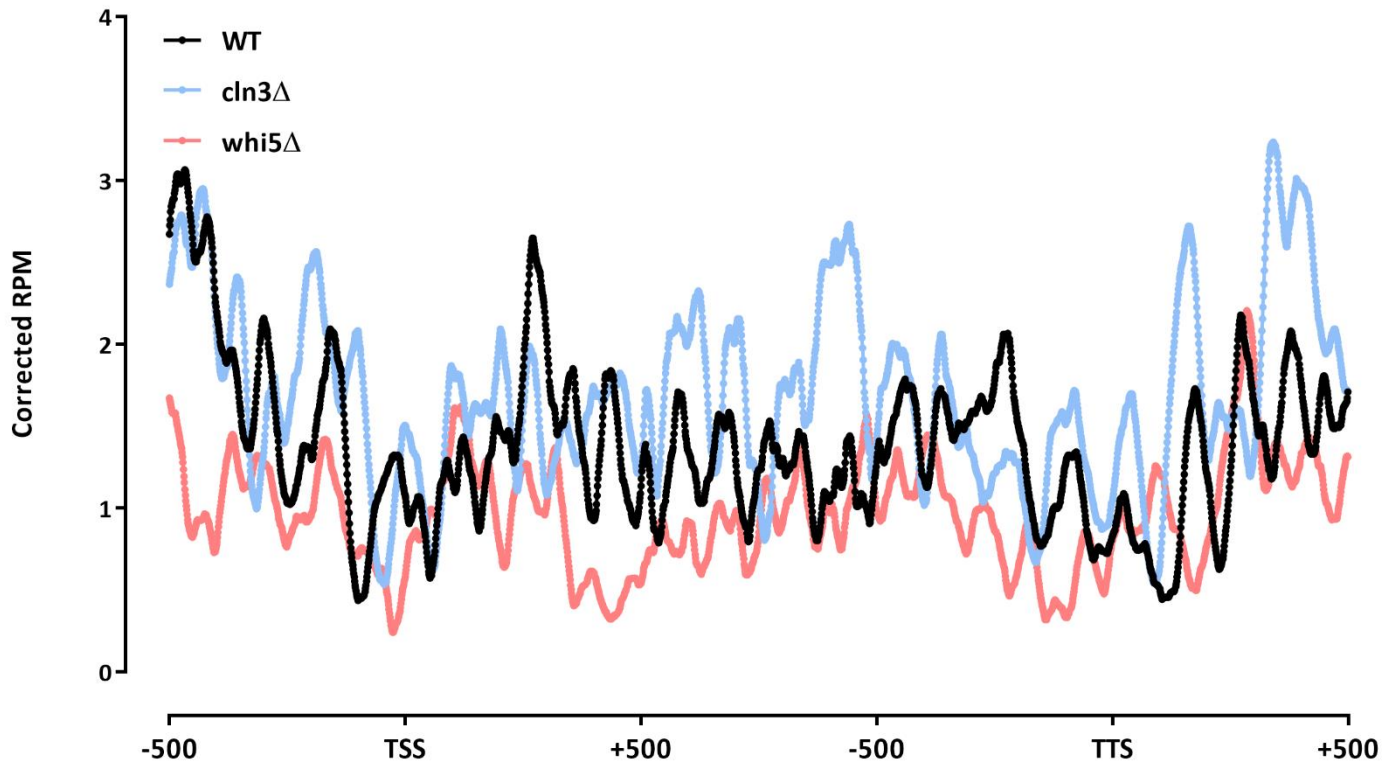


### Pol2pS5 S. cerevisiae corrected

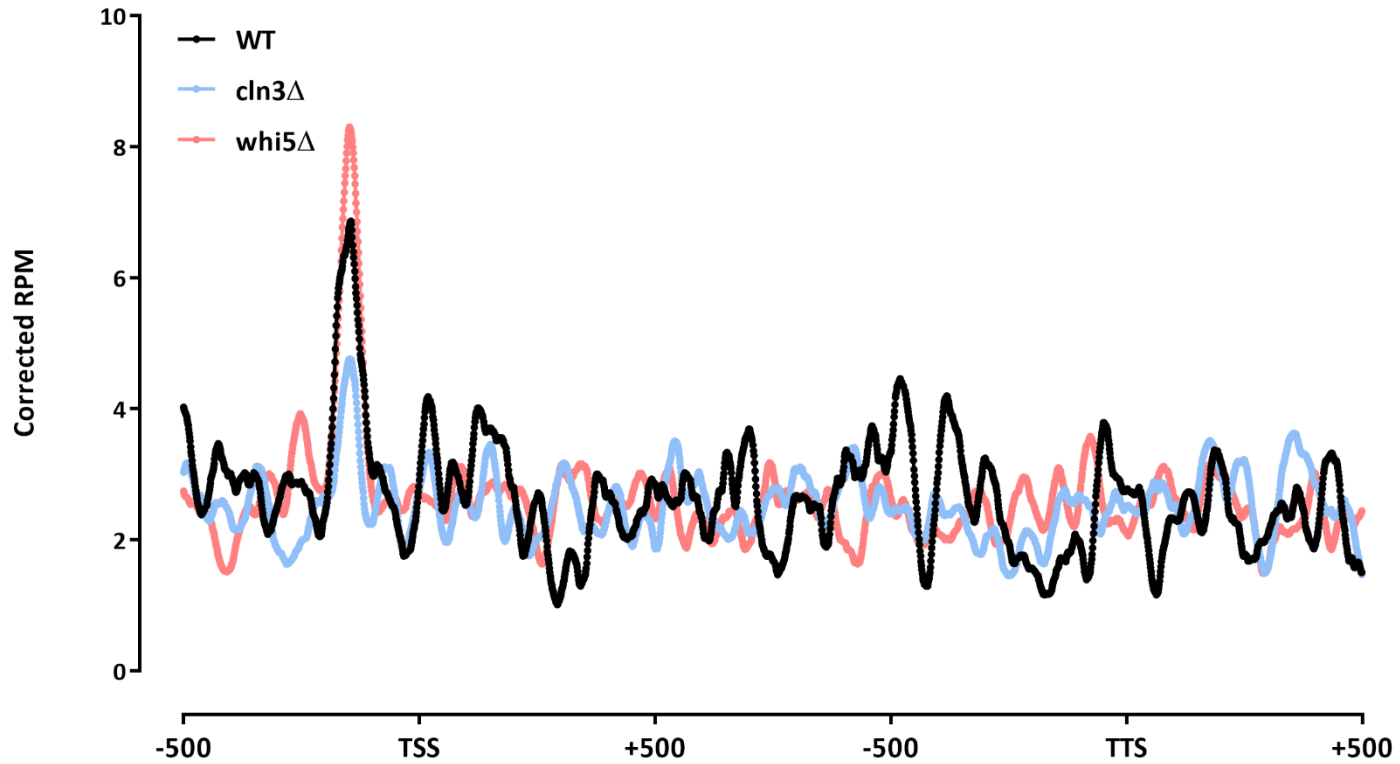


# ncRNA bioType

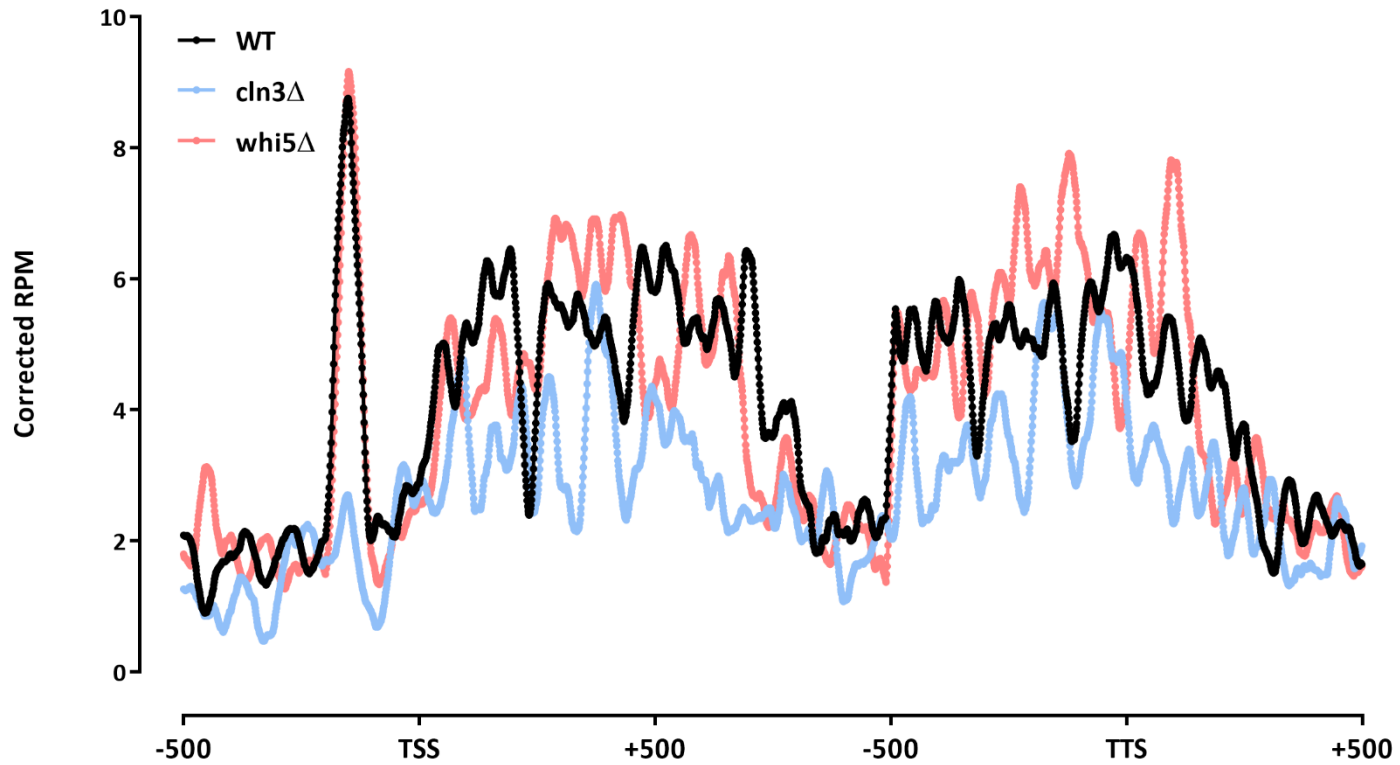
H3 *S. cerevisiae* corrected



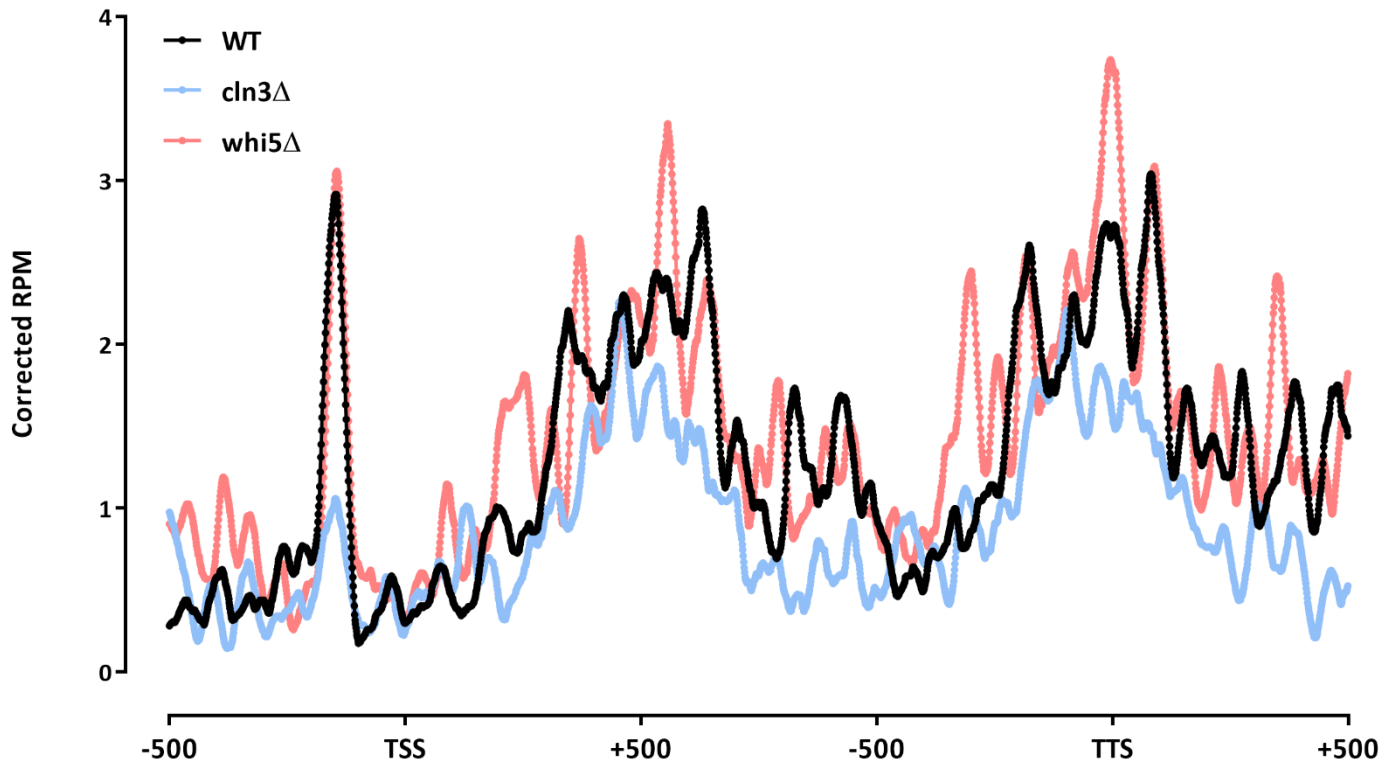
### Input *S. cerevisiae* corrected



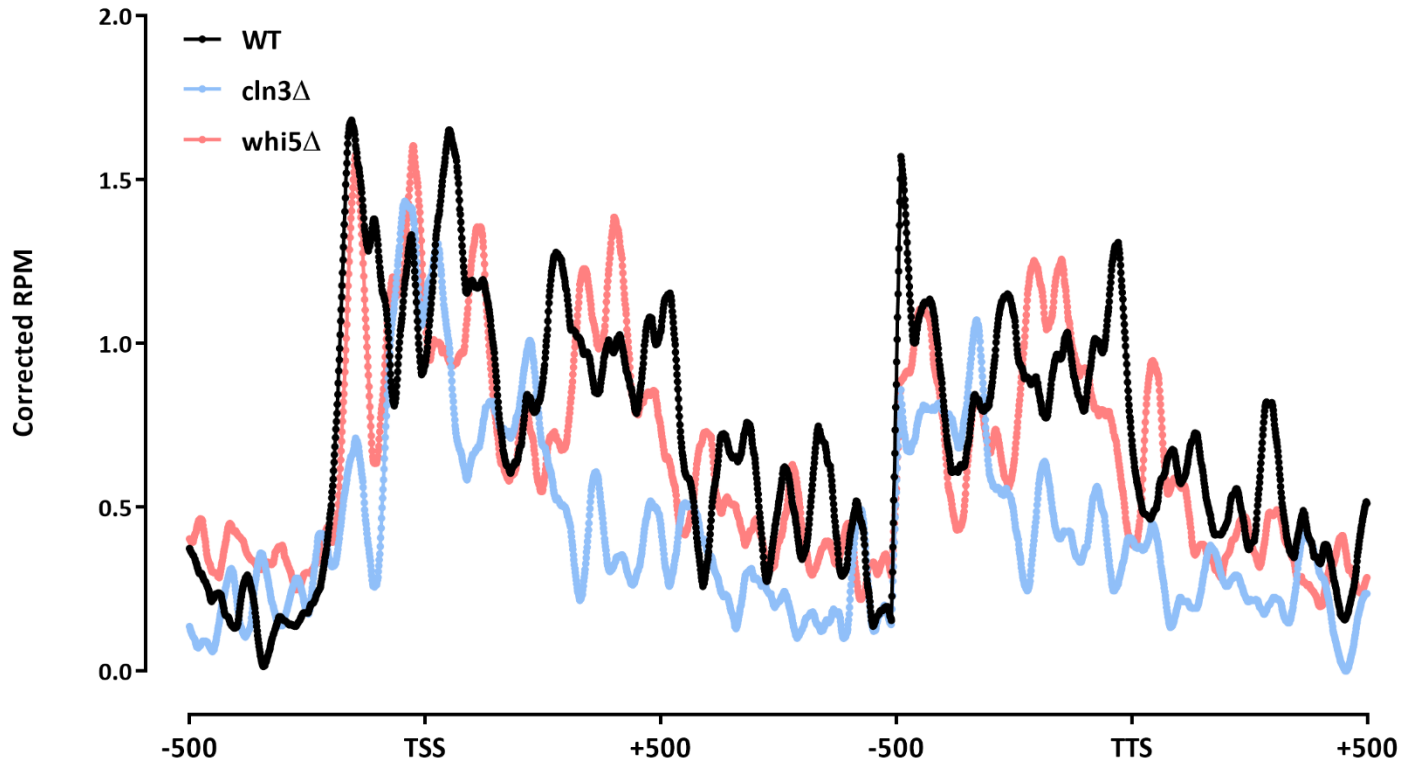
### Pol2 *S. cerevisiae* corrected



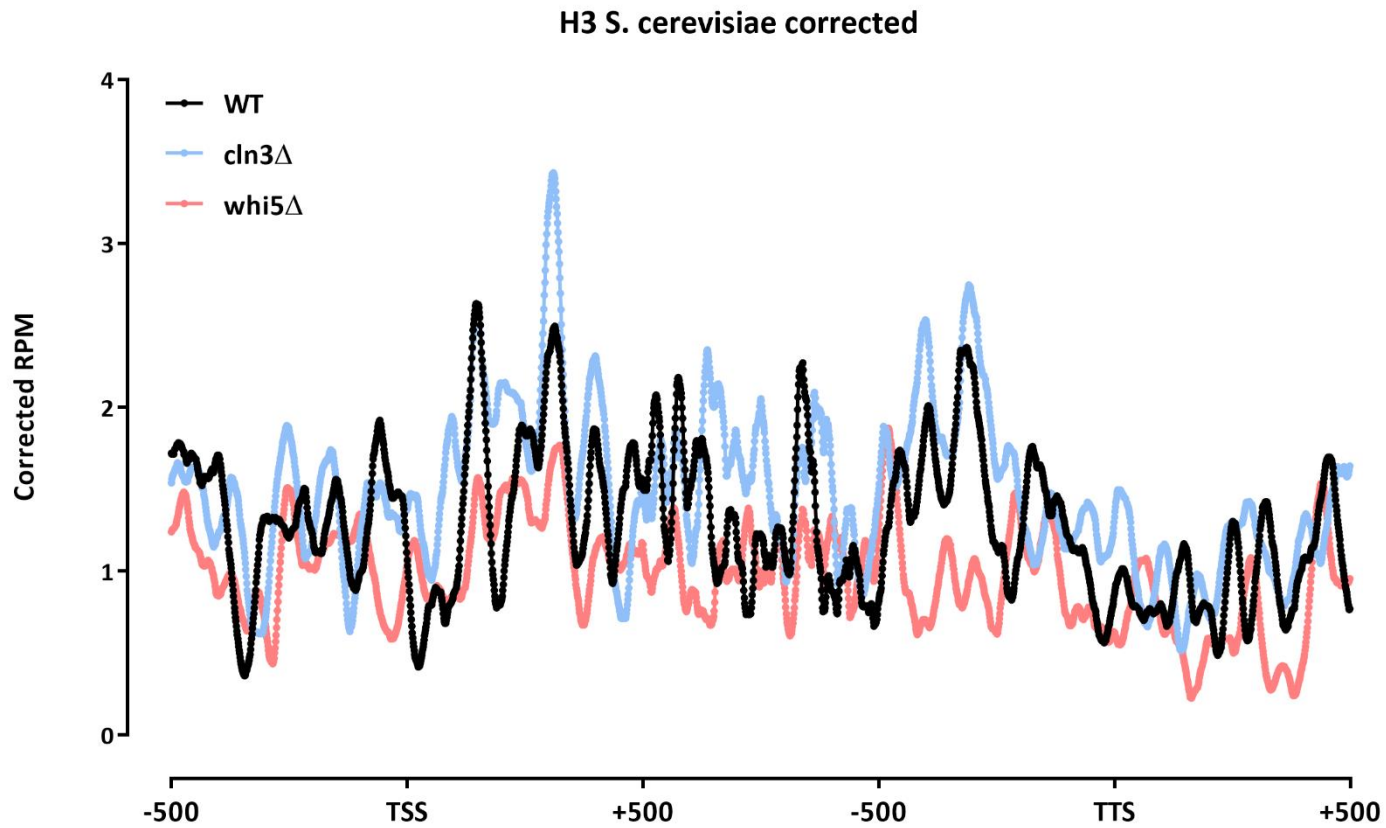
### Pol2pS2 *S. cerevisiae* corrected



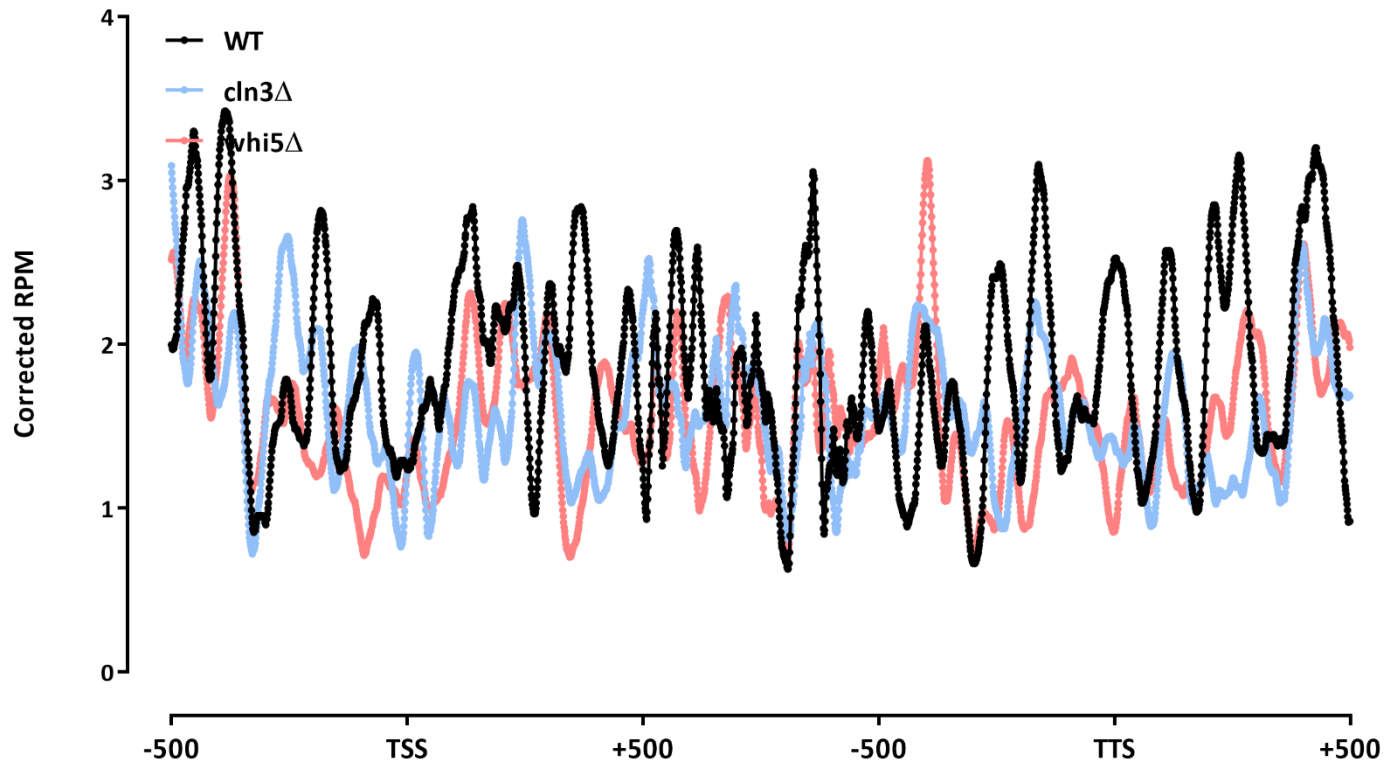
Pol2pS5 S. cerevisiae corrected



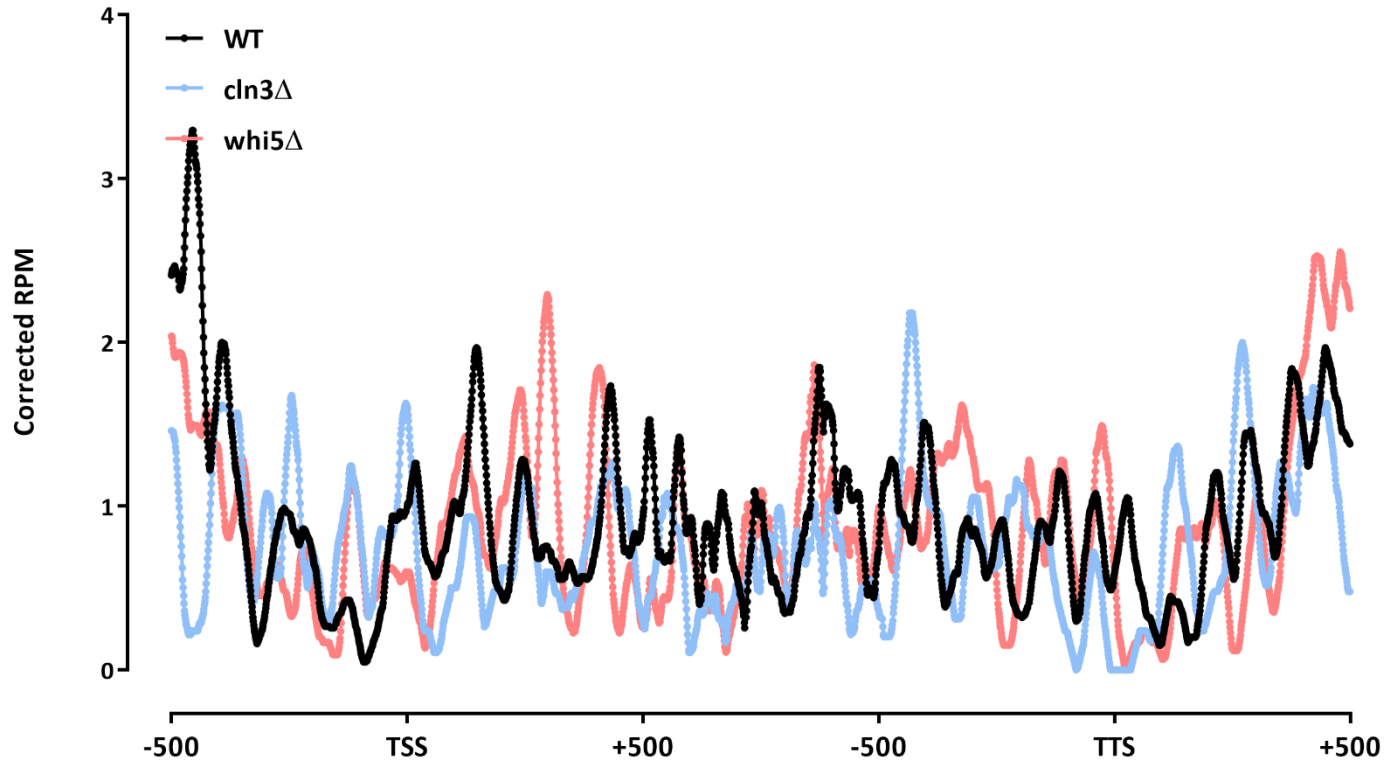
# pseudogene bioType



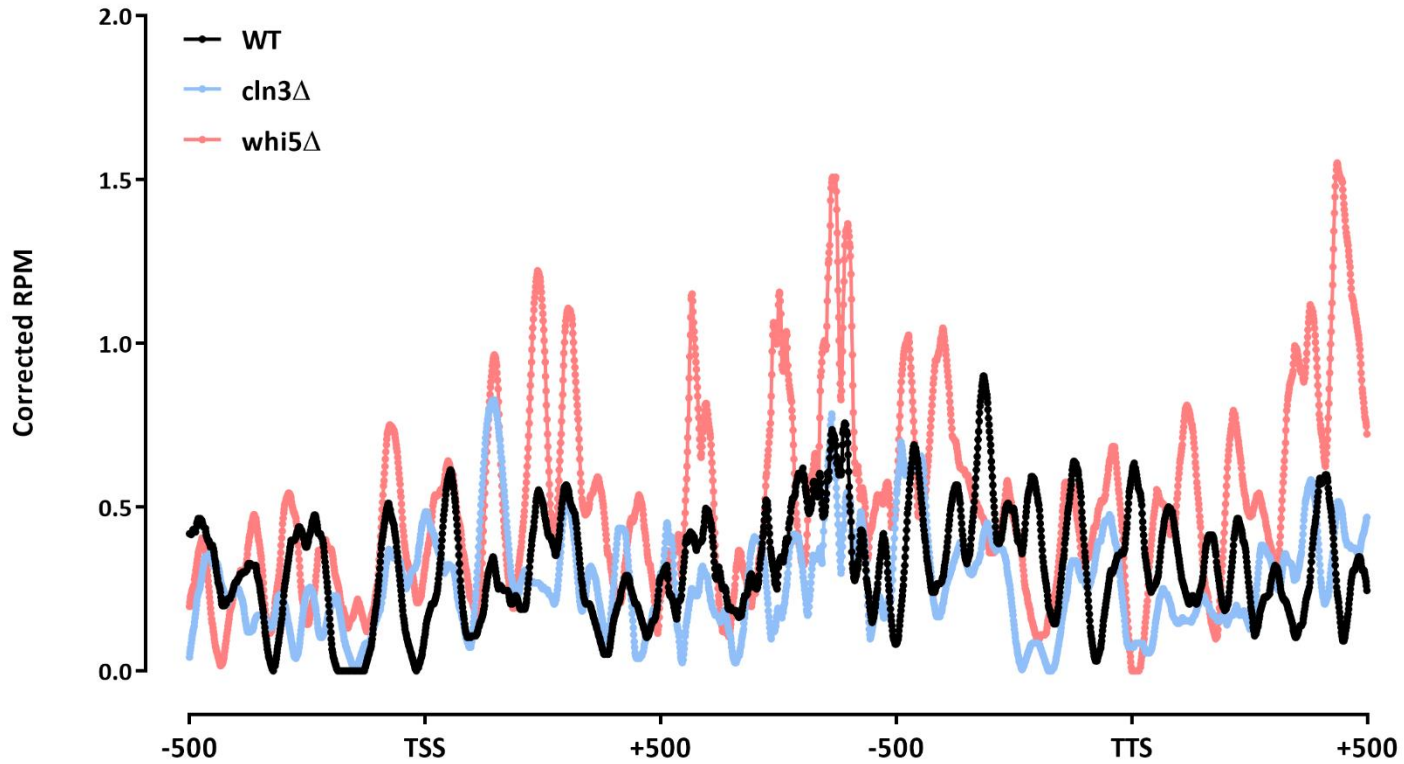
### Input *S. cerevisiae* corrected



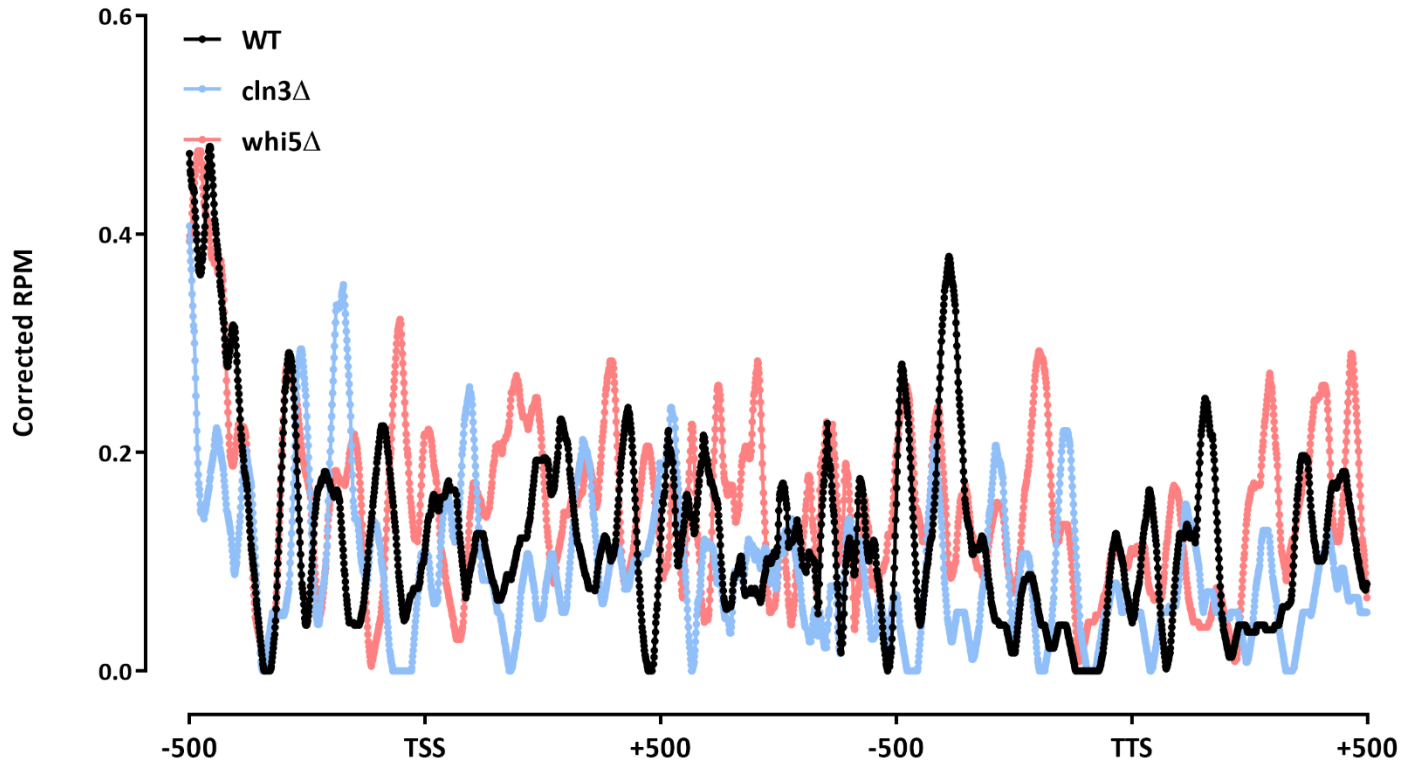
### Pol2 *S. cerevisiae* corrected



### Pol2pS2 *S. cerevisiae* corrected

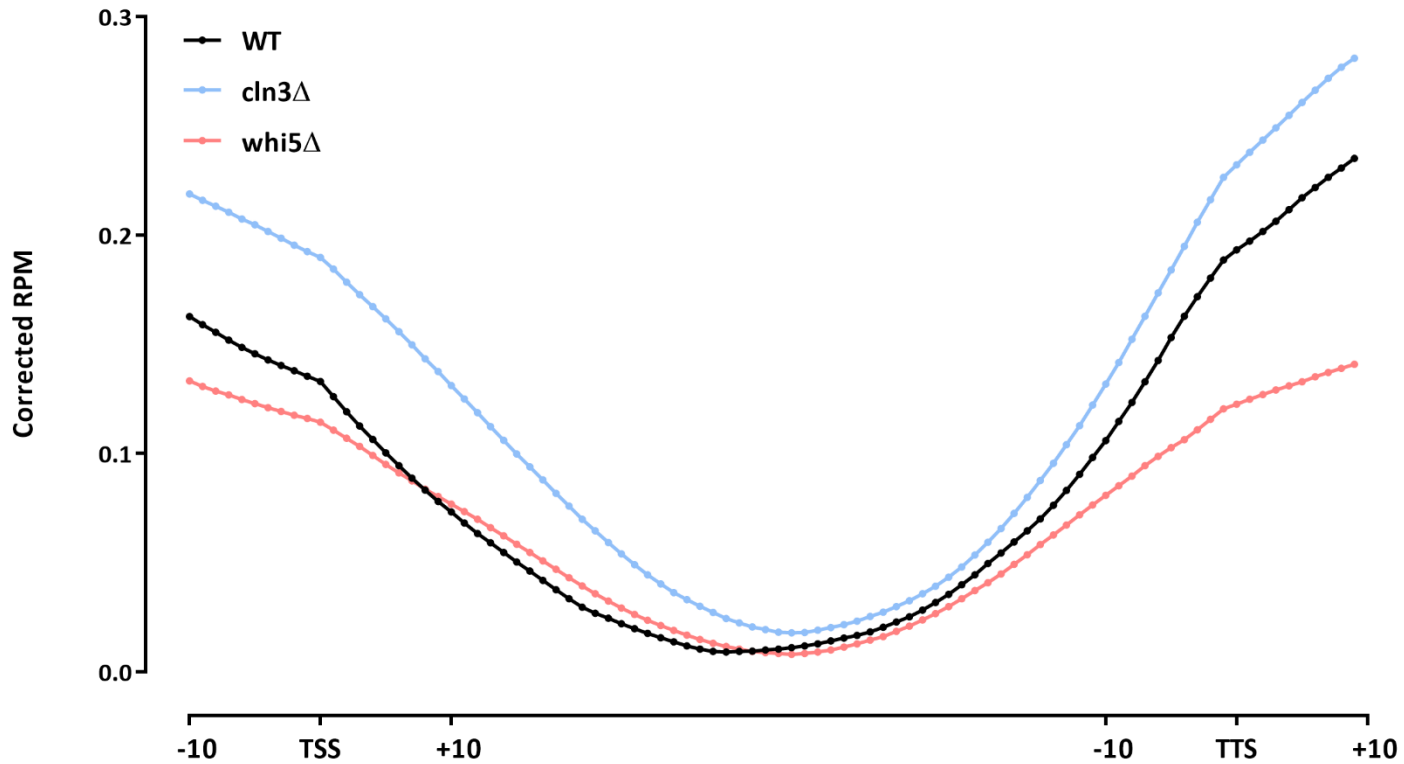


### Pol2pS5 S. cerevisiae corrected

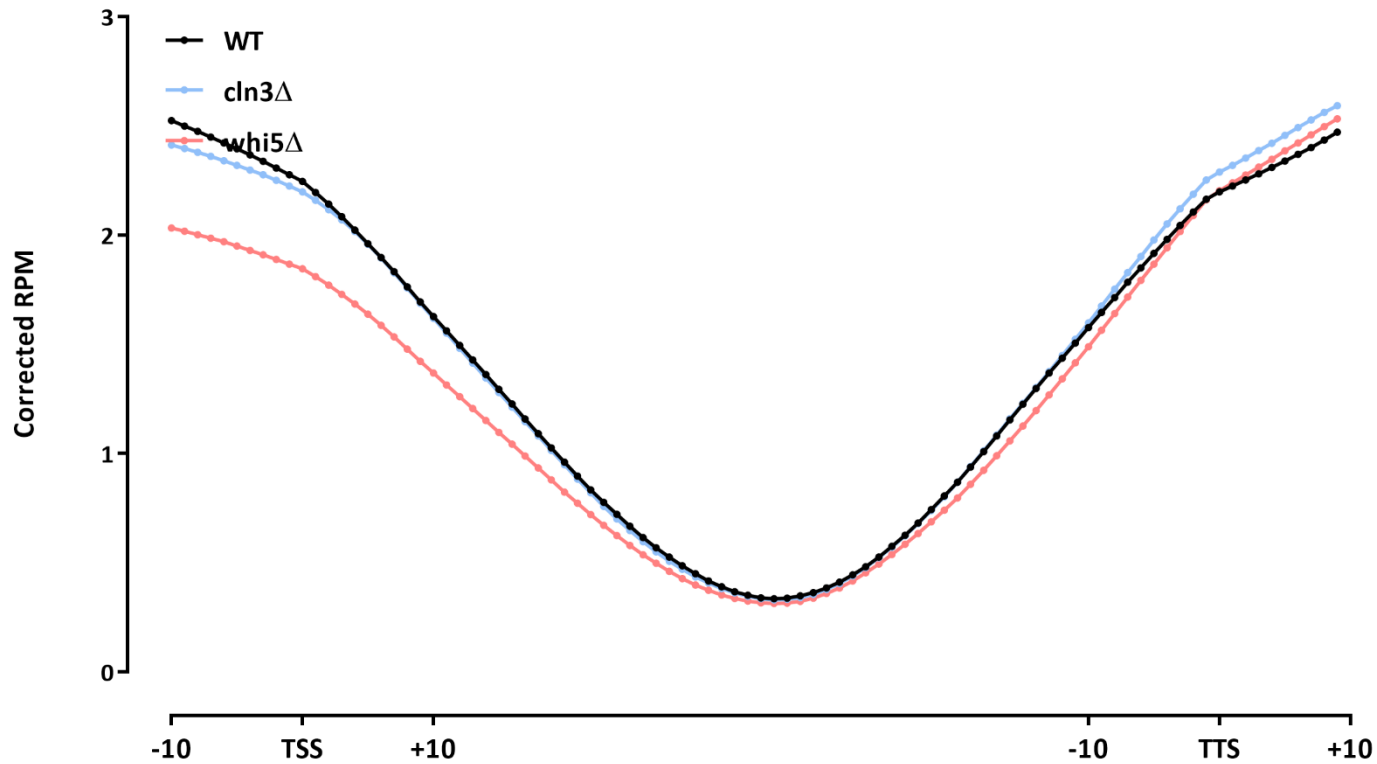


# tRNA bioType

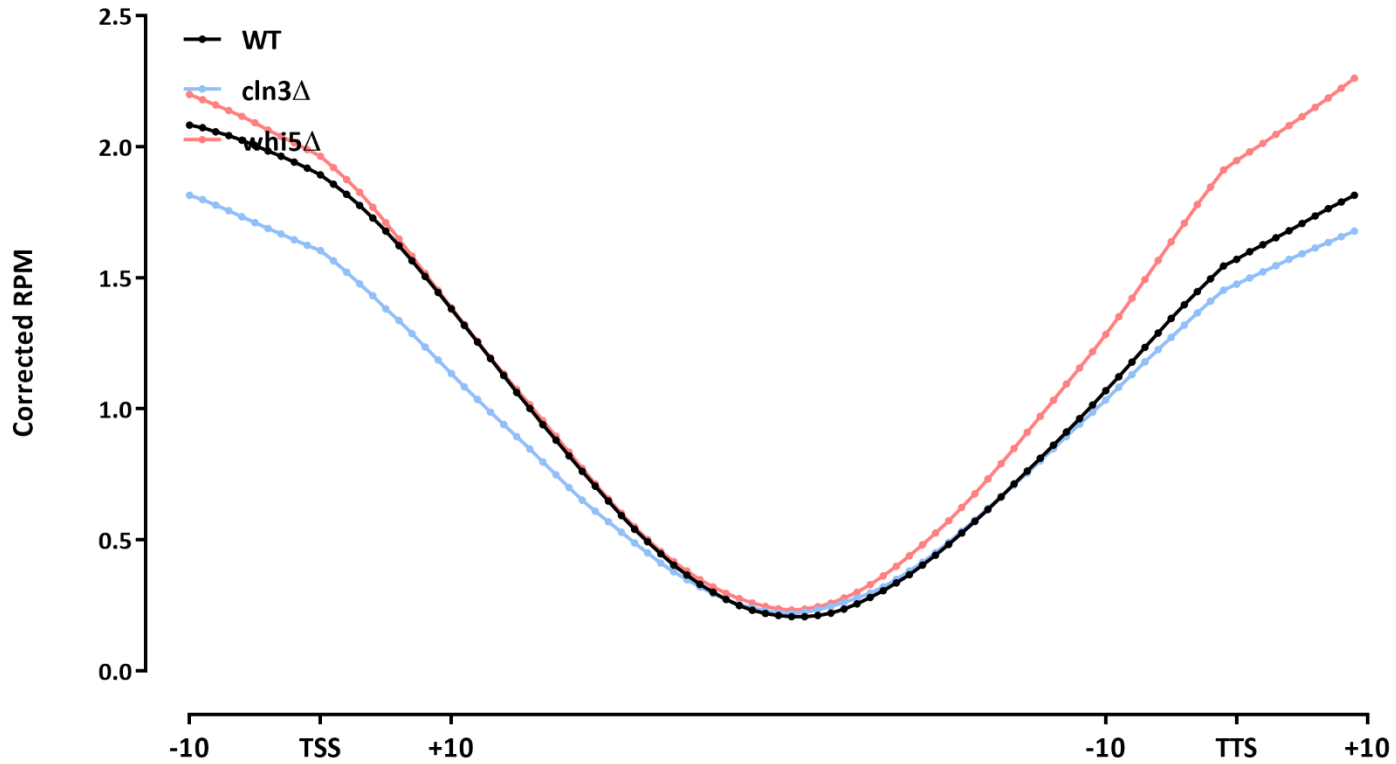
H3 *S. cerevisiae* corrected



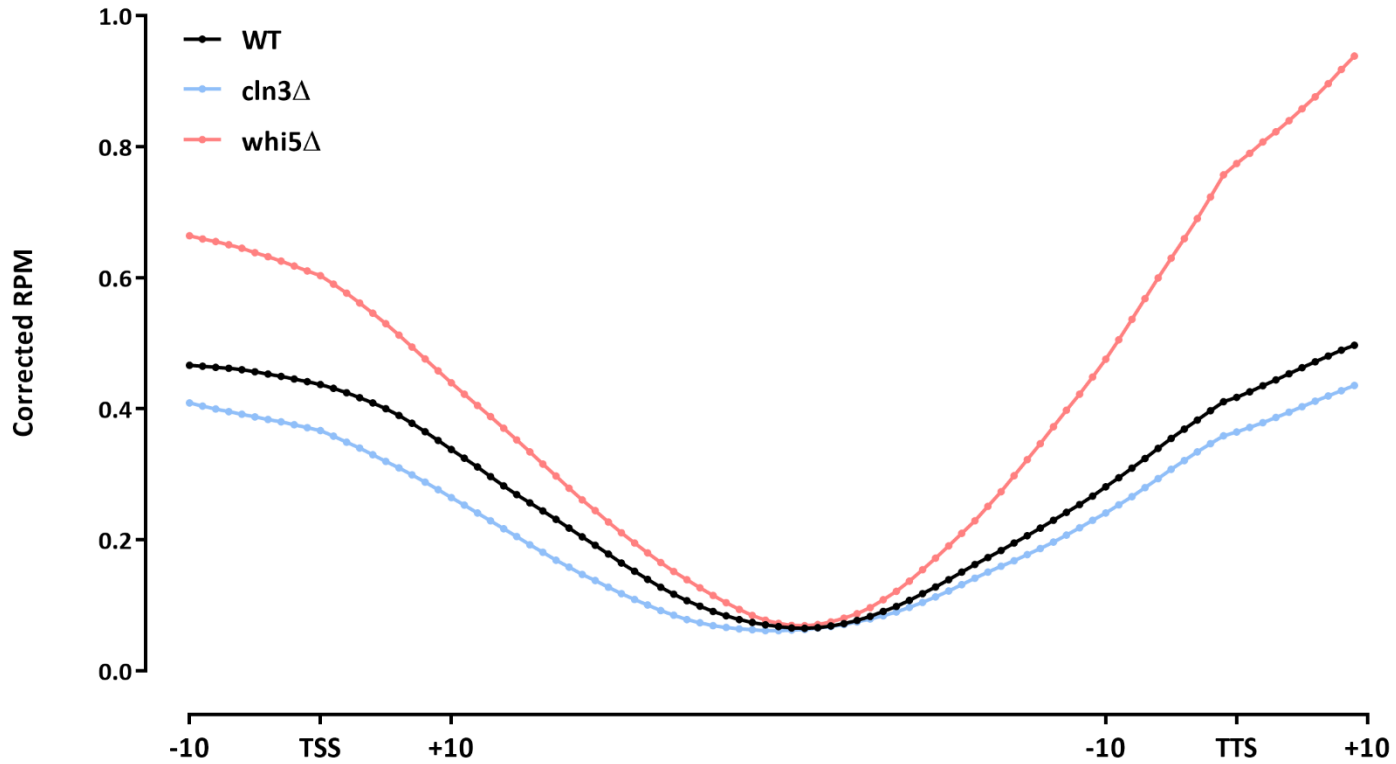
### Input *S. cerevisiae* corrected



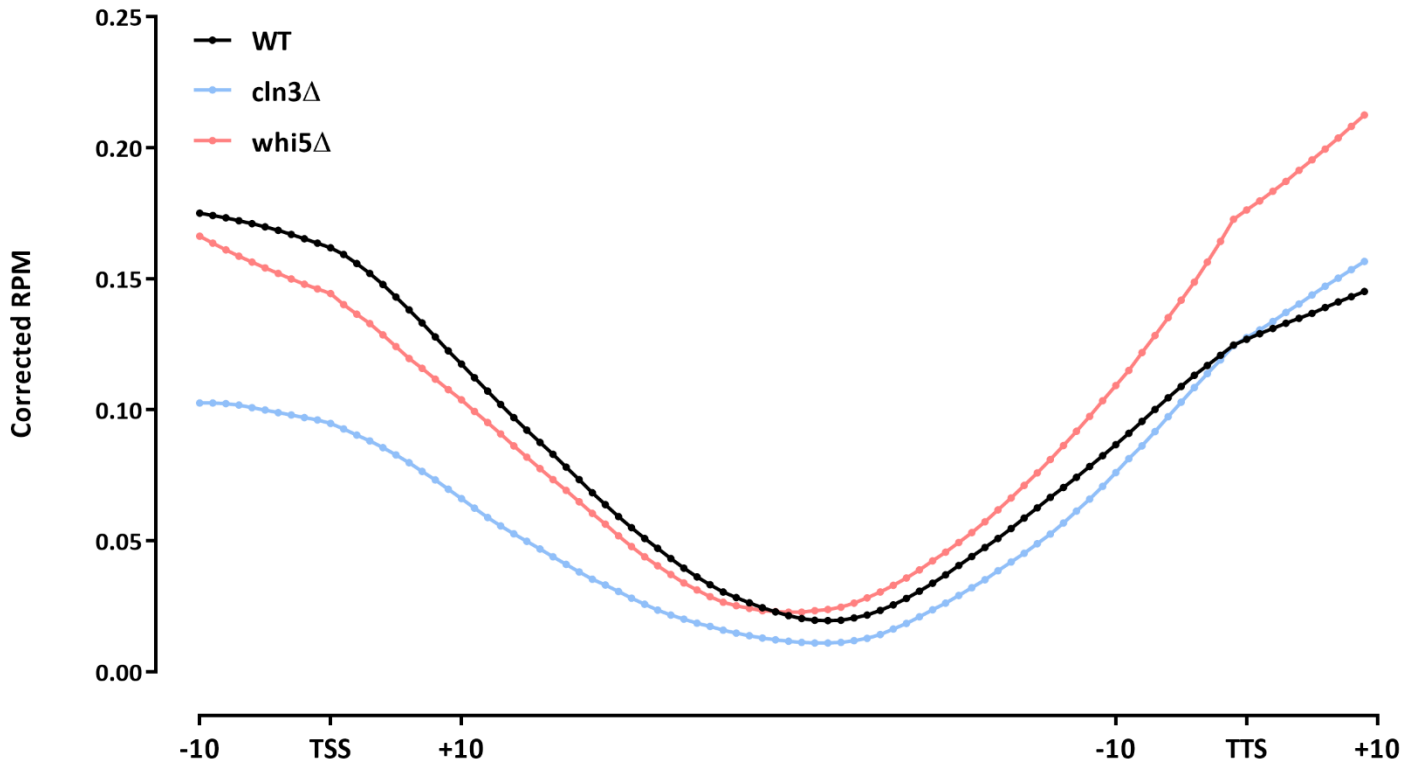
### Pol2 *S. cerevisiae* corrected



### Pol2pS2 *S. cerevisiae* corrected

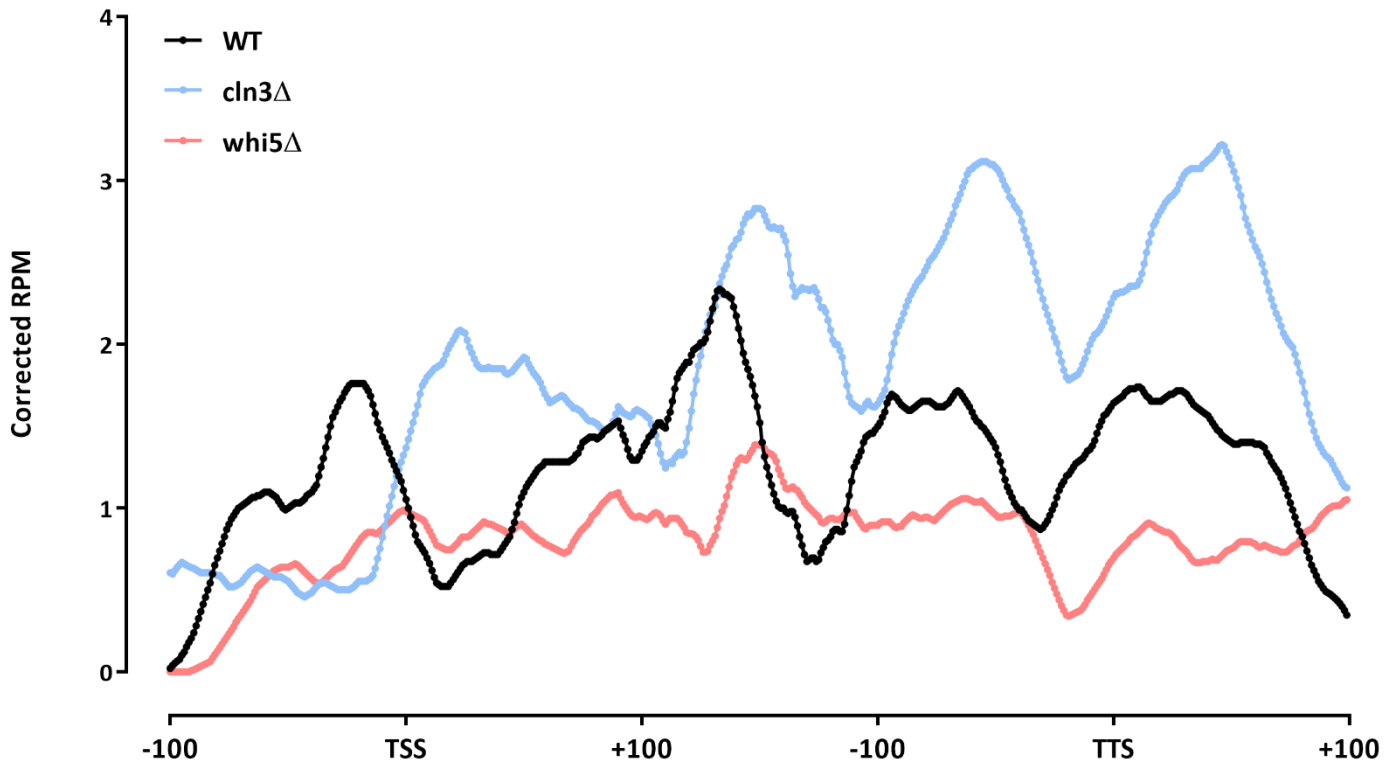


### Pol2pS5 S. cerevisiae corrected

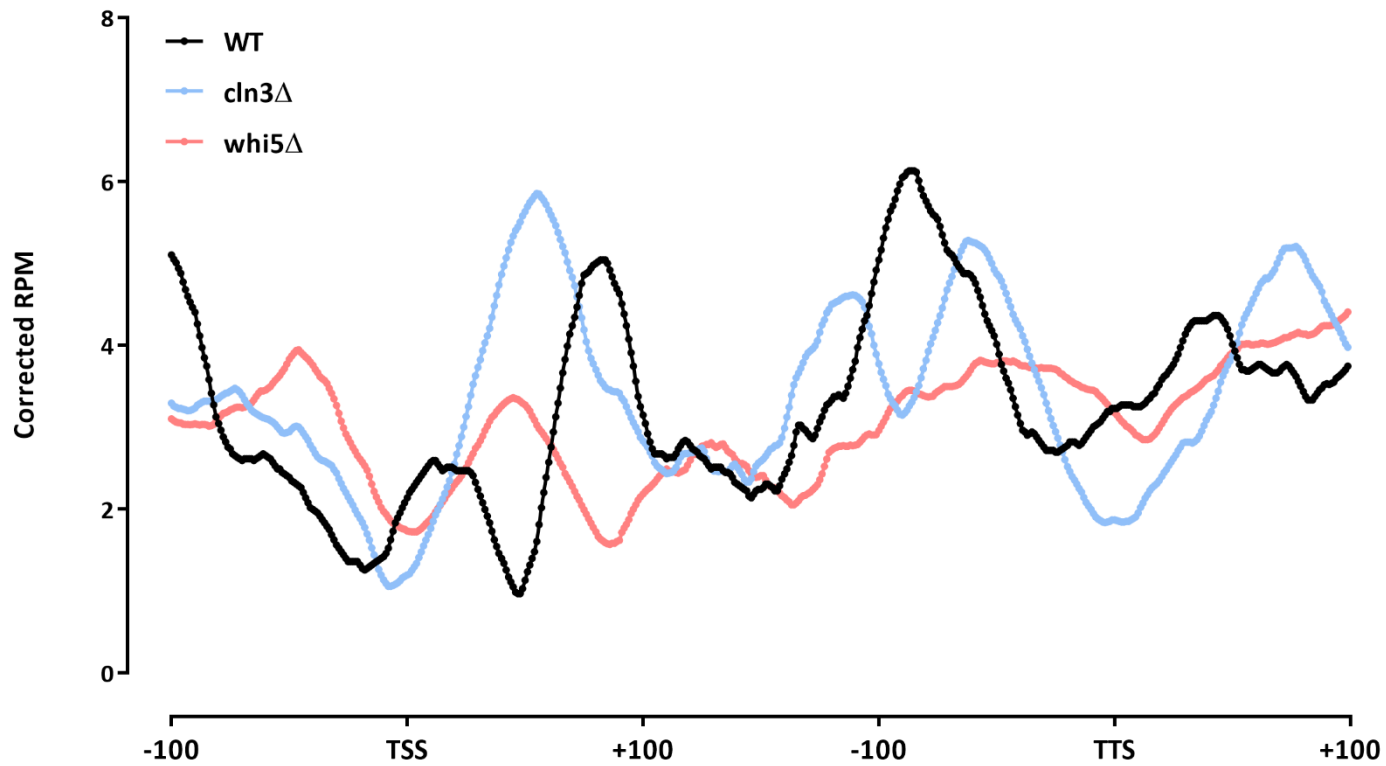


# snRNA bioType

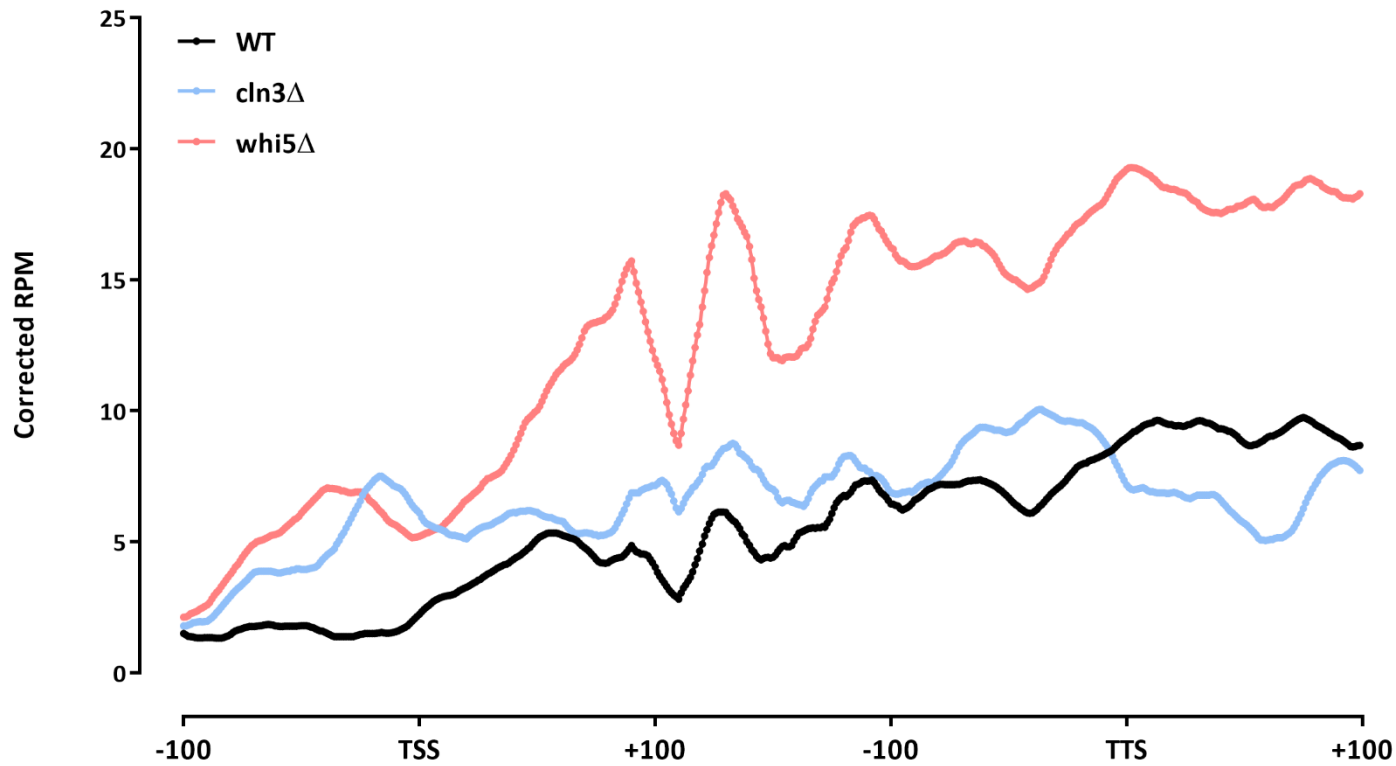
H3 *S. cerevisiae* corrected



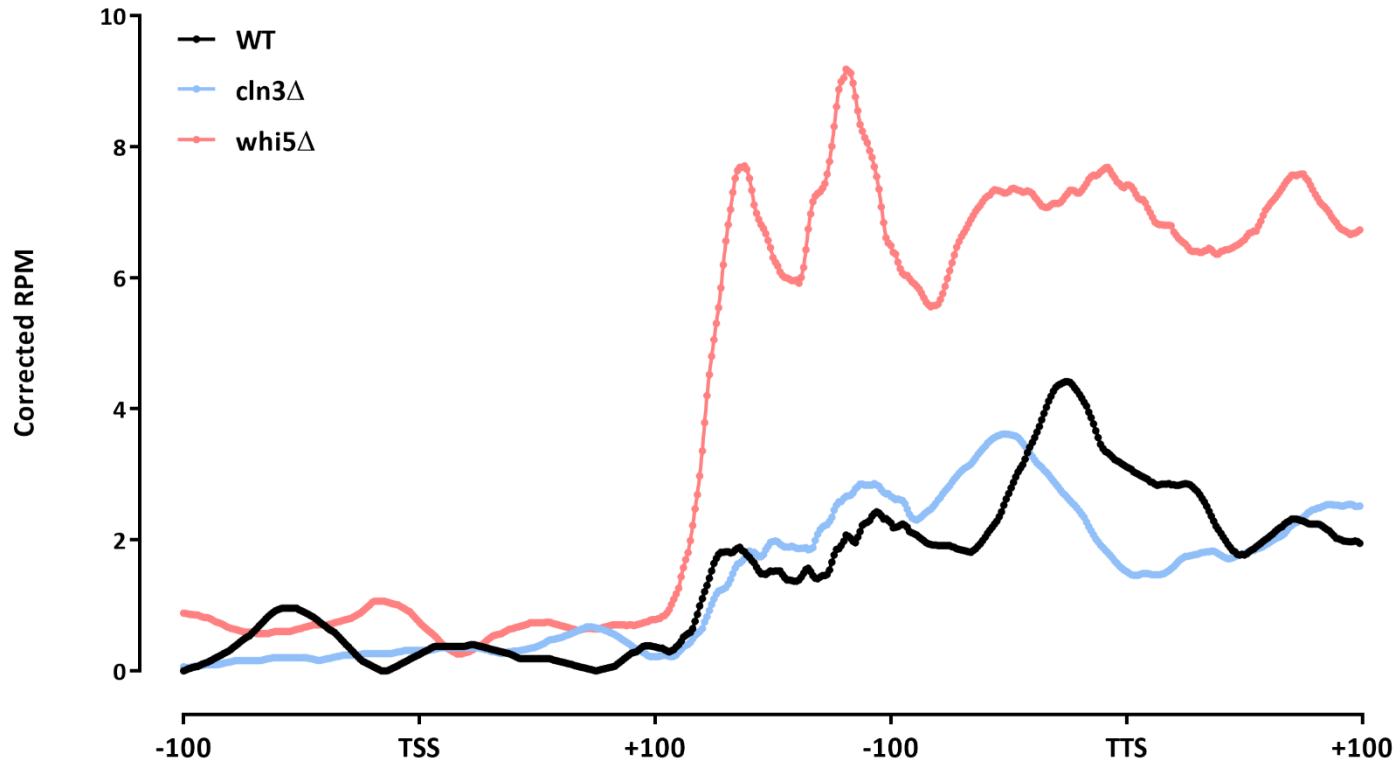
### Input *S. cerevisiae* corrected



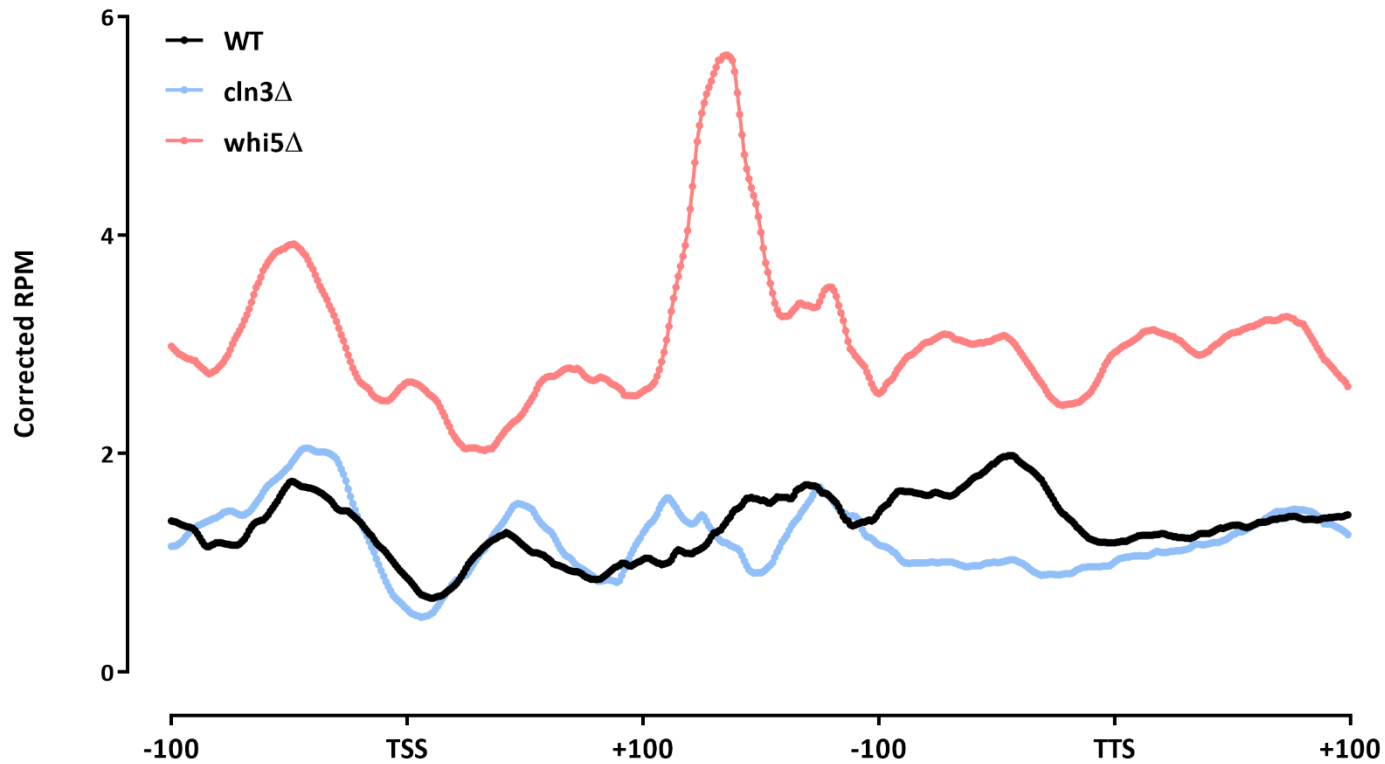
### Pol2 *S. cerevisiae* corrected



### Pol2pS2 *S. cerevisiae* corrected

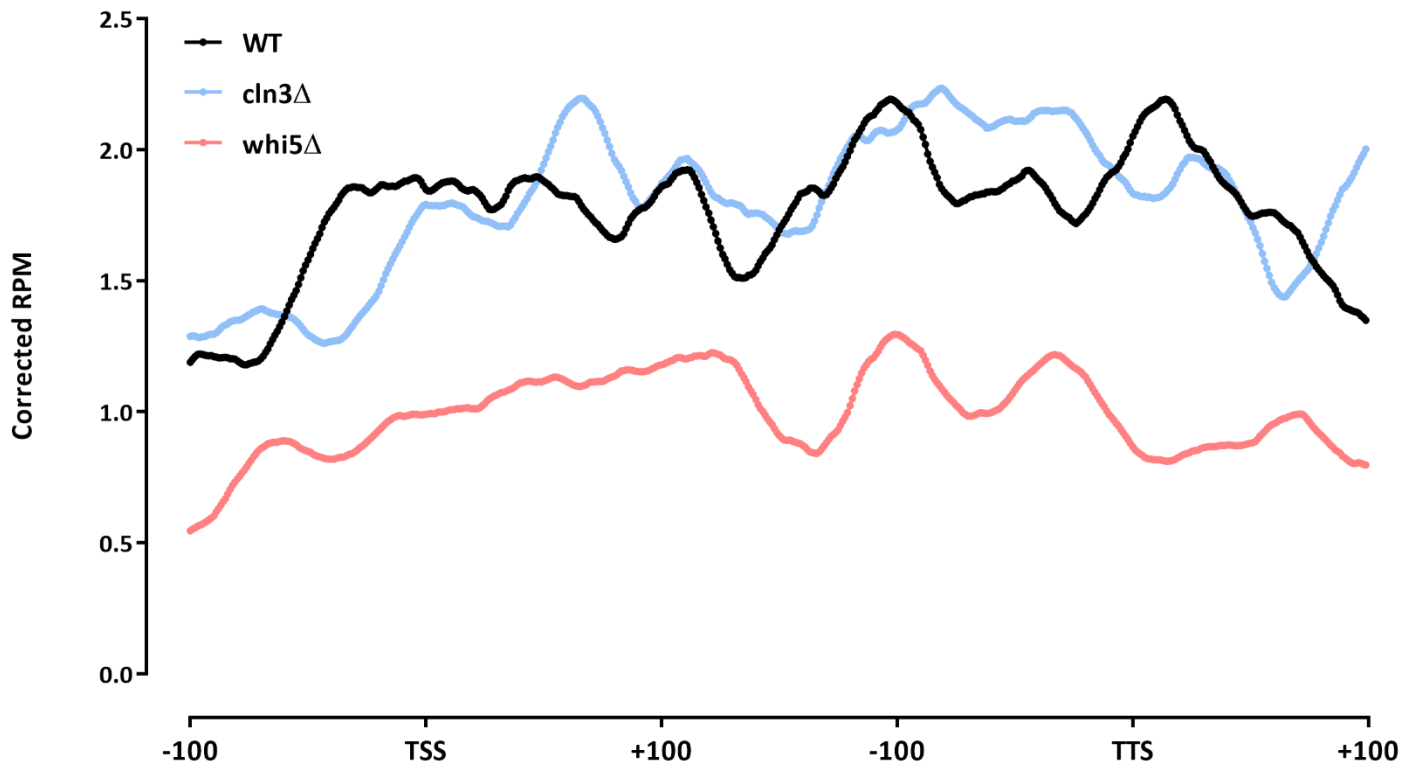


### Pol2pS5 *S. cerevisiae* corrected

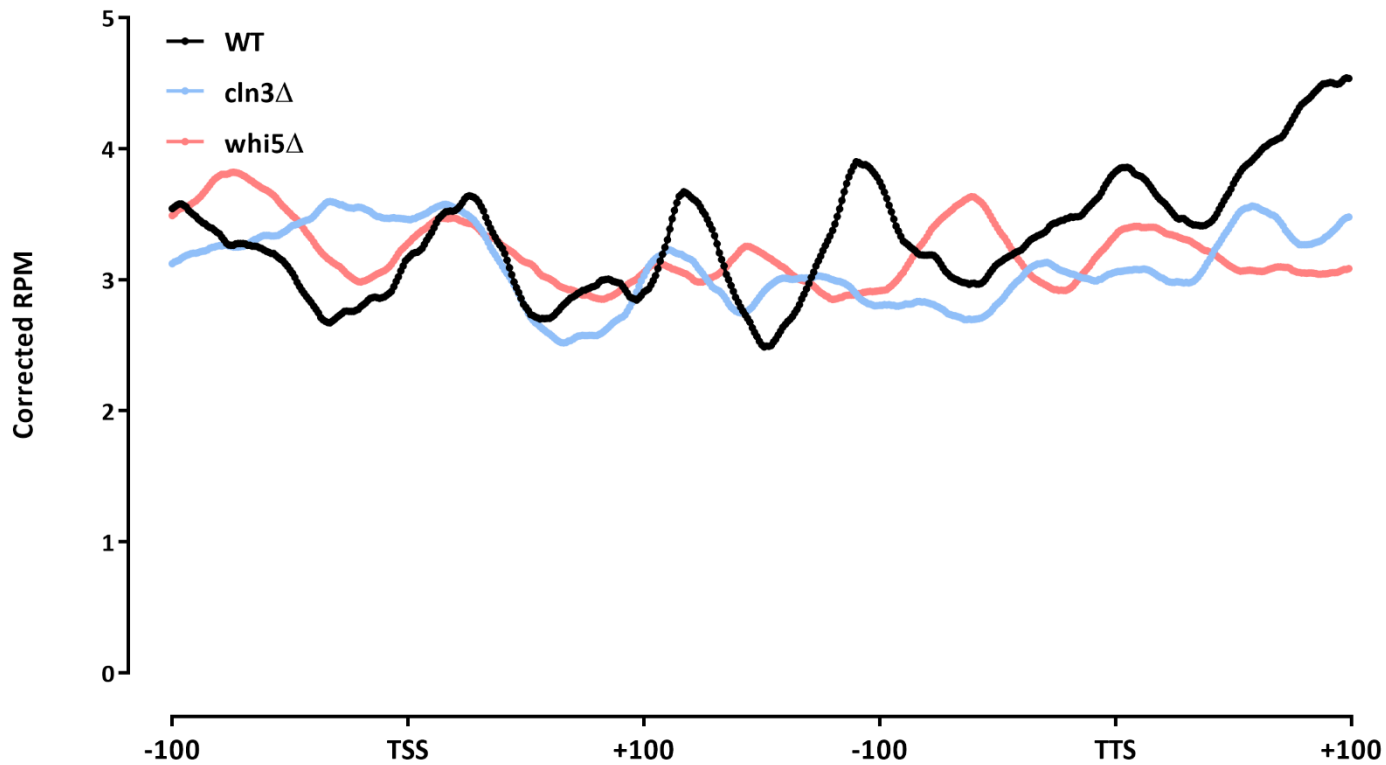


# snoRNA bioType

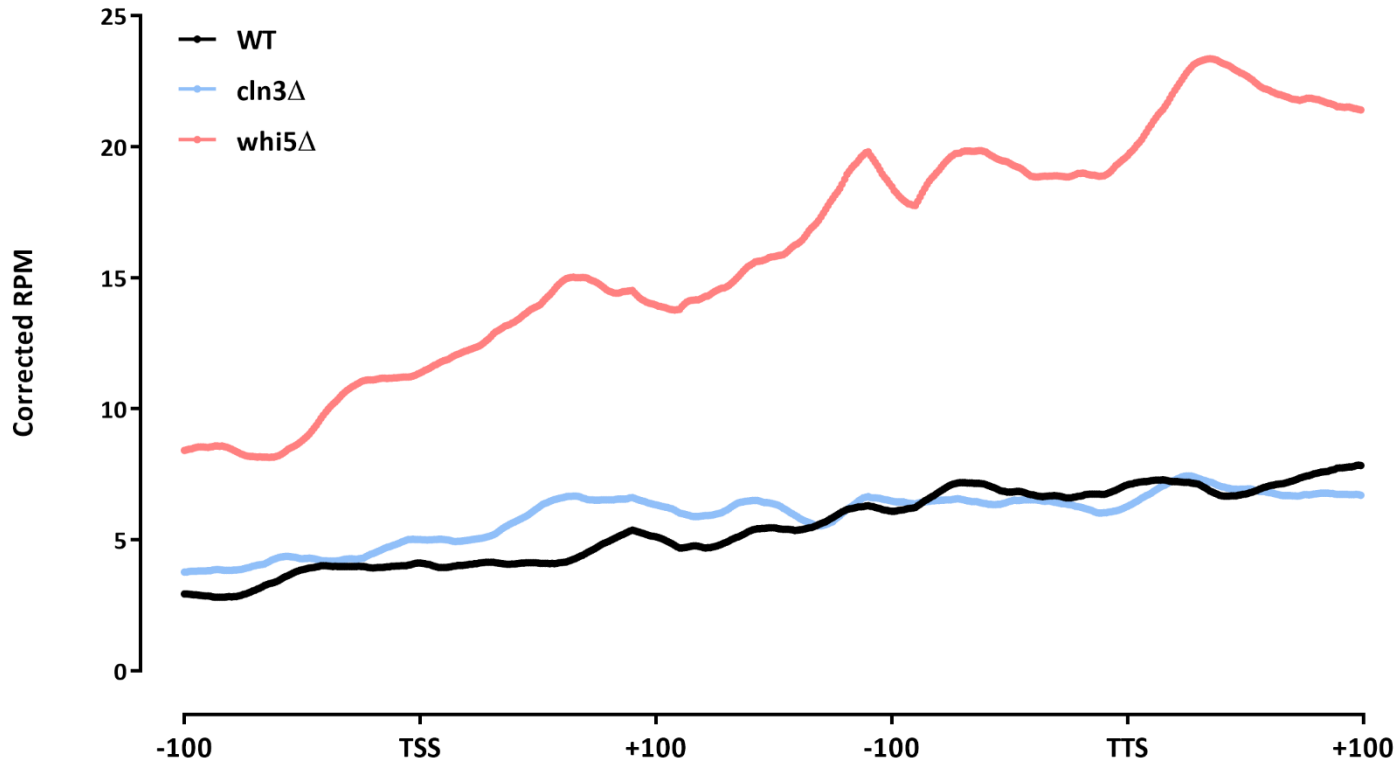
H3 *S. cerevisiae* corrected



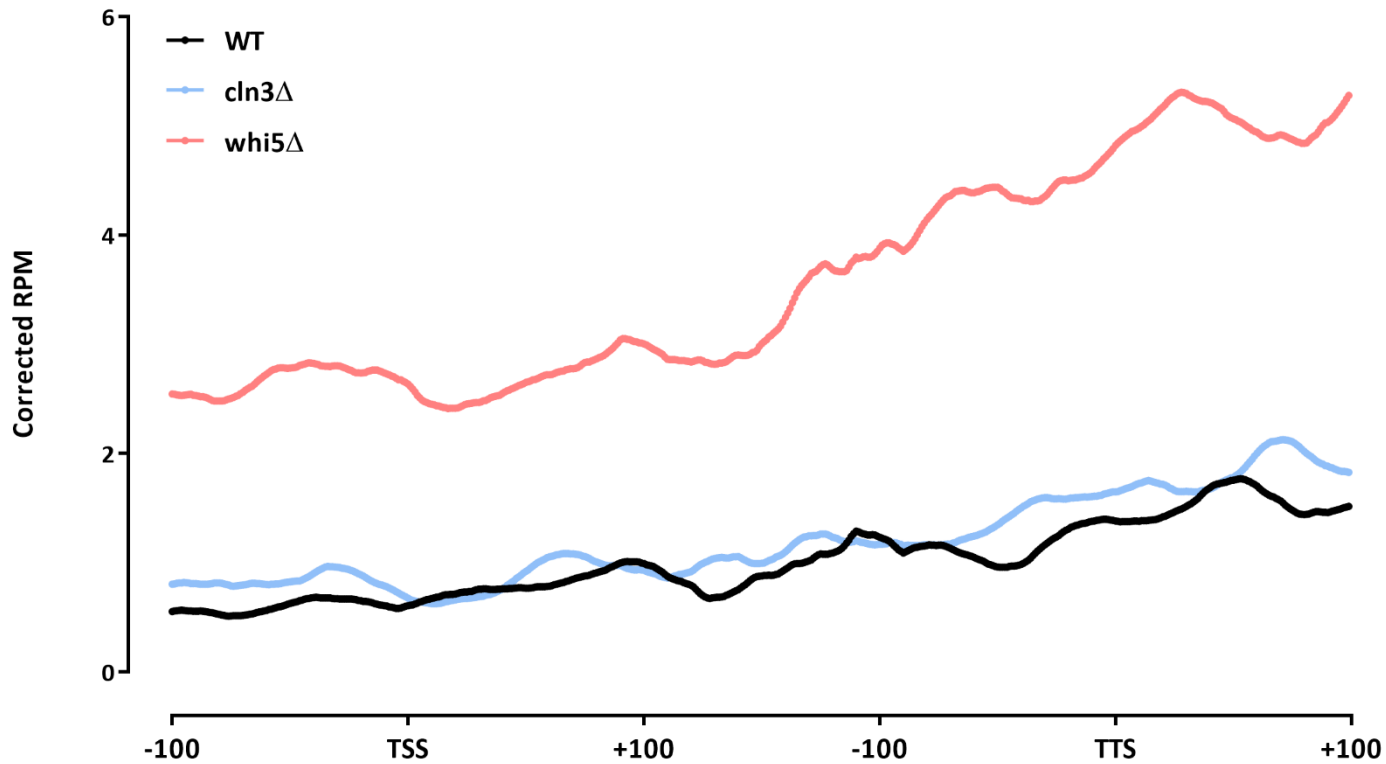
### Input *S. cerevisiae* corrected



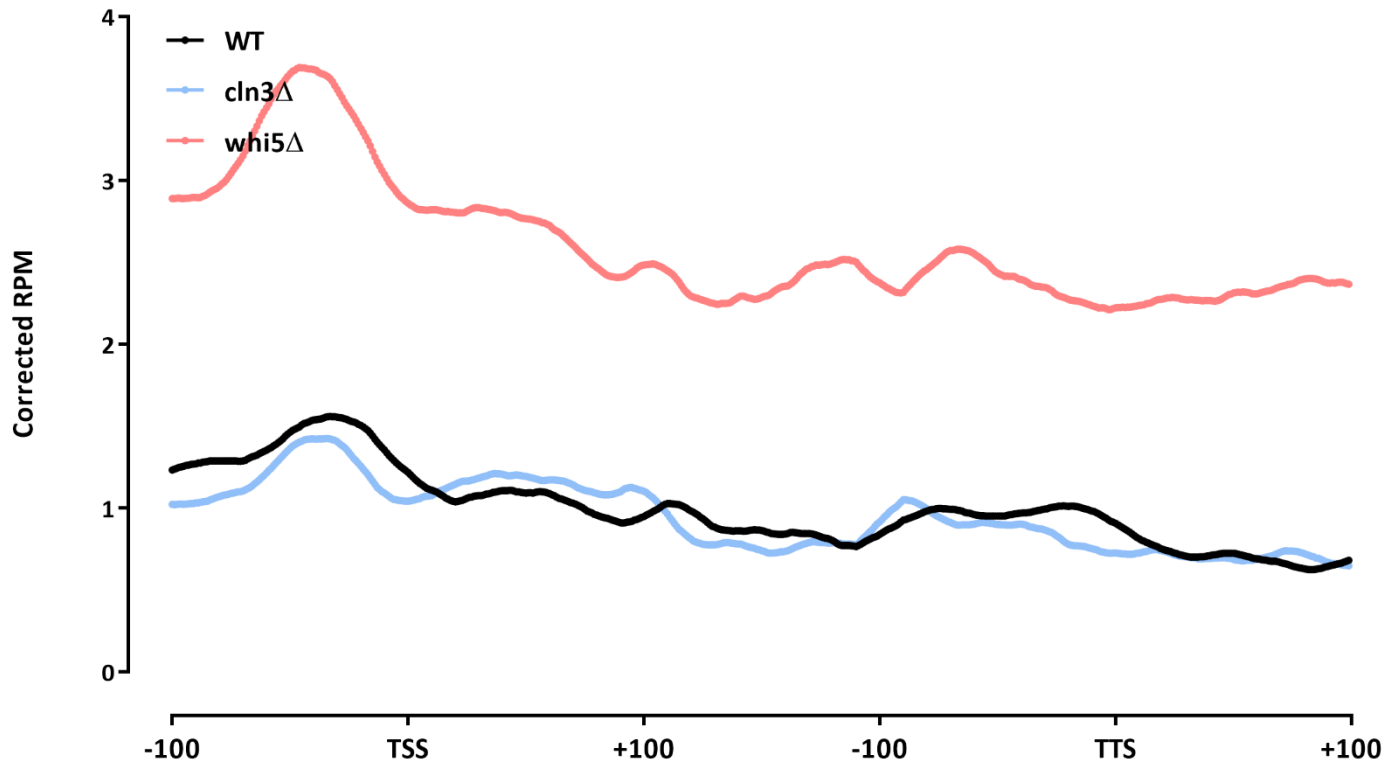
### Pol2 *S. cerevisiae* corrected



### Pol2pS2 *S. cerevisiae* corrected

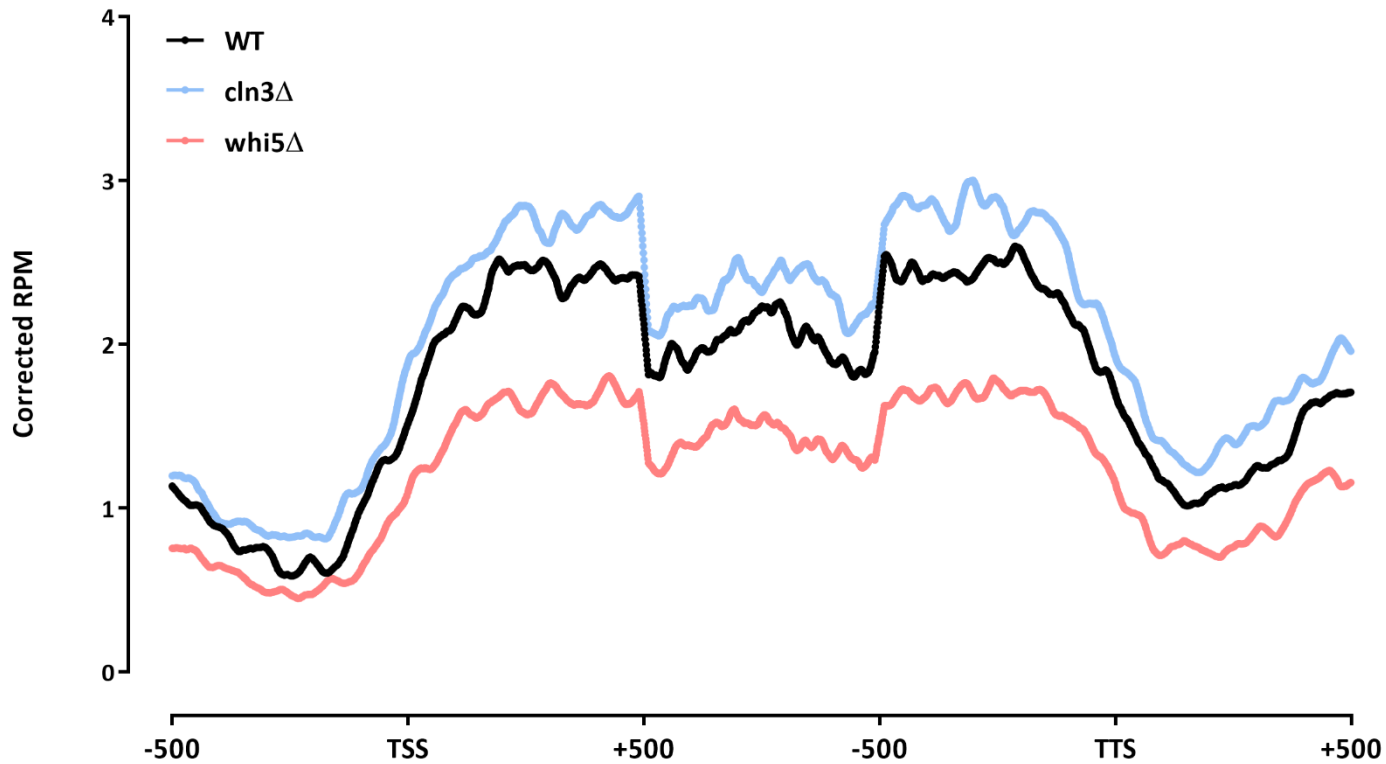


### Pol2pS5 *S. cerevisiae* corrected

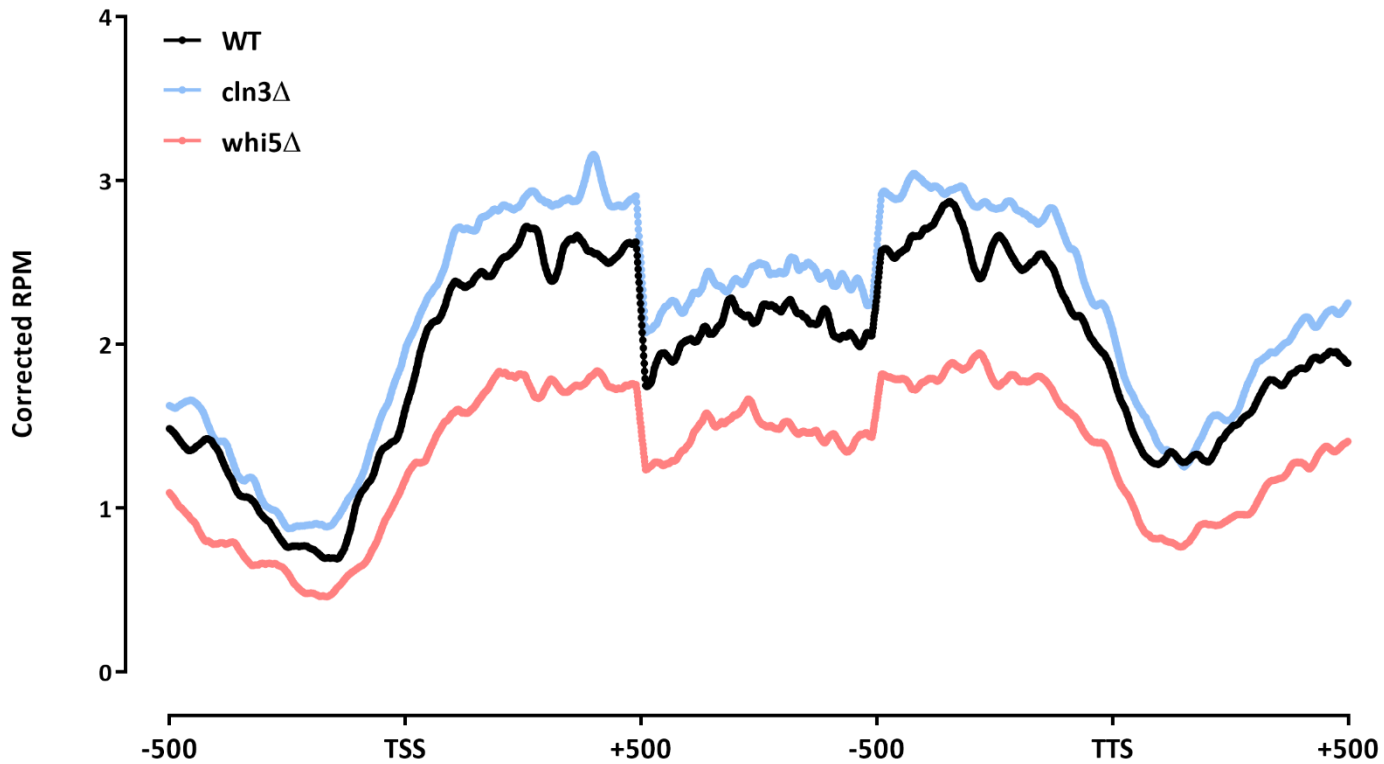


# Pol2 deciles

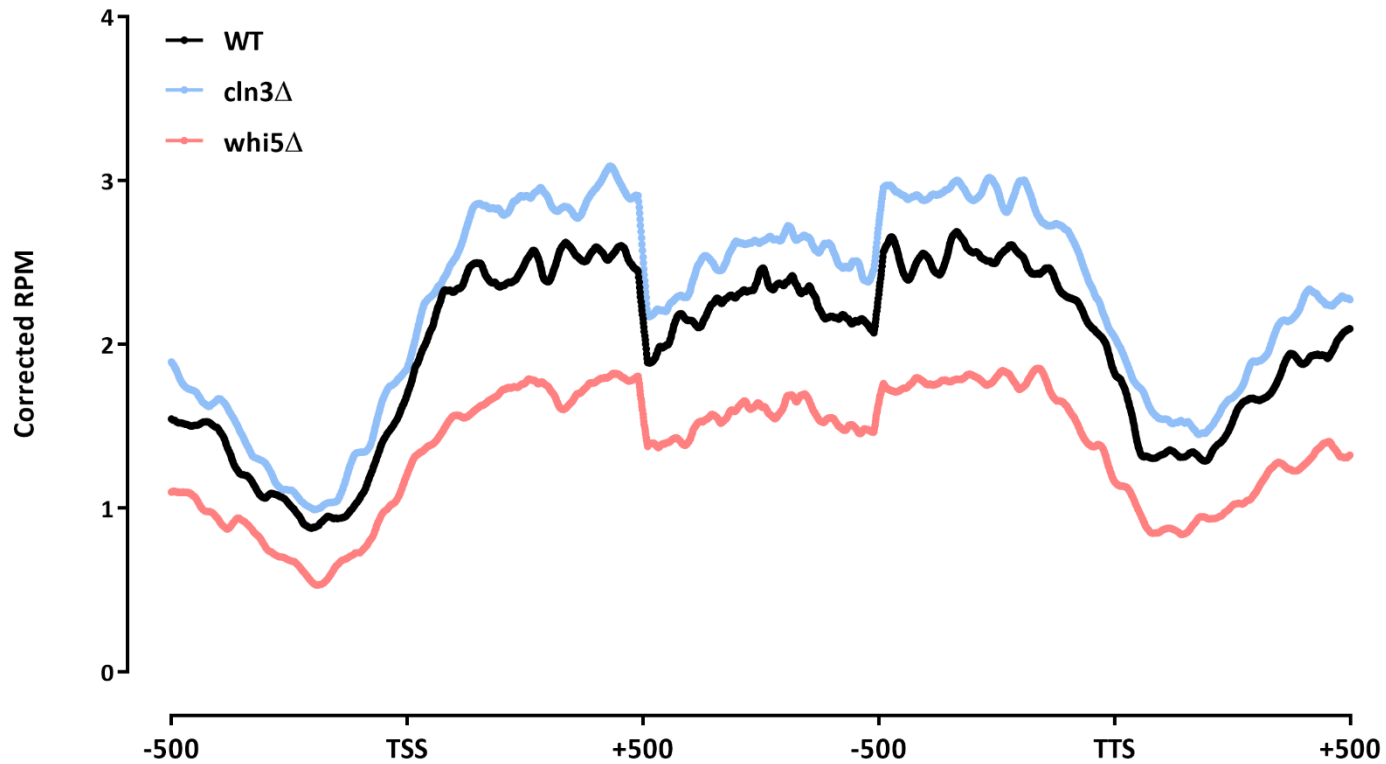
H3 *S. cerevisiae* corrected 10%th



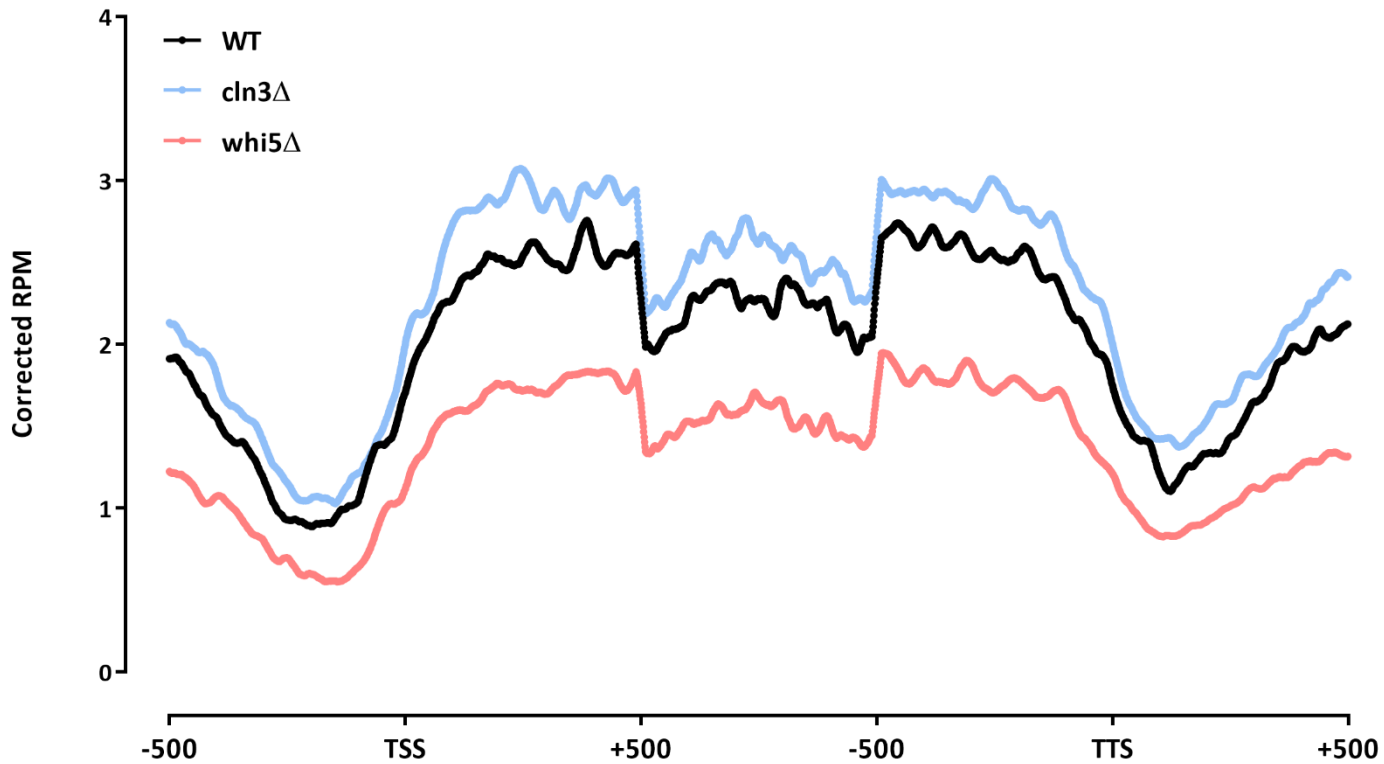
### H3 *S. cerevisiae* corrected 20%th



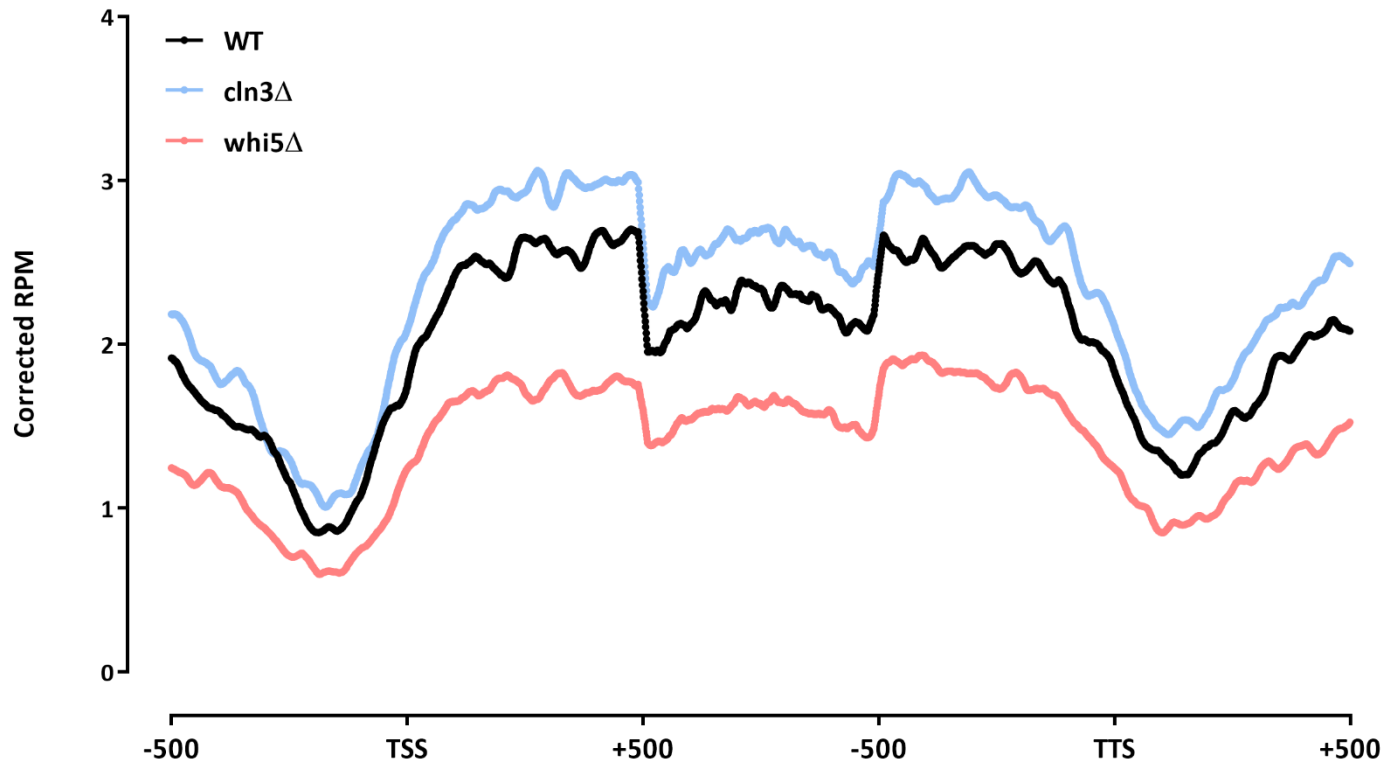
### H3 *S. cerevisiae* corrected 30%th



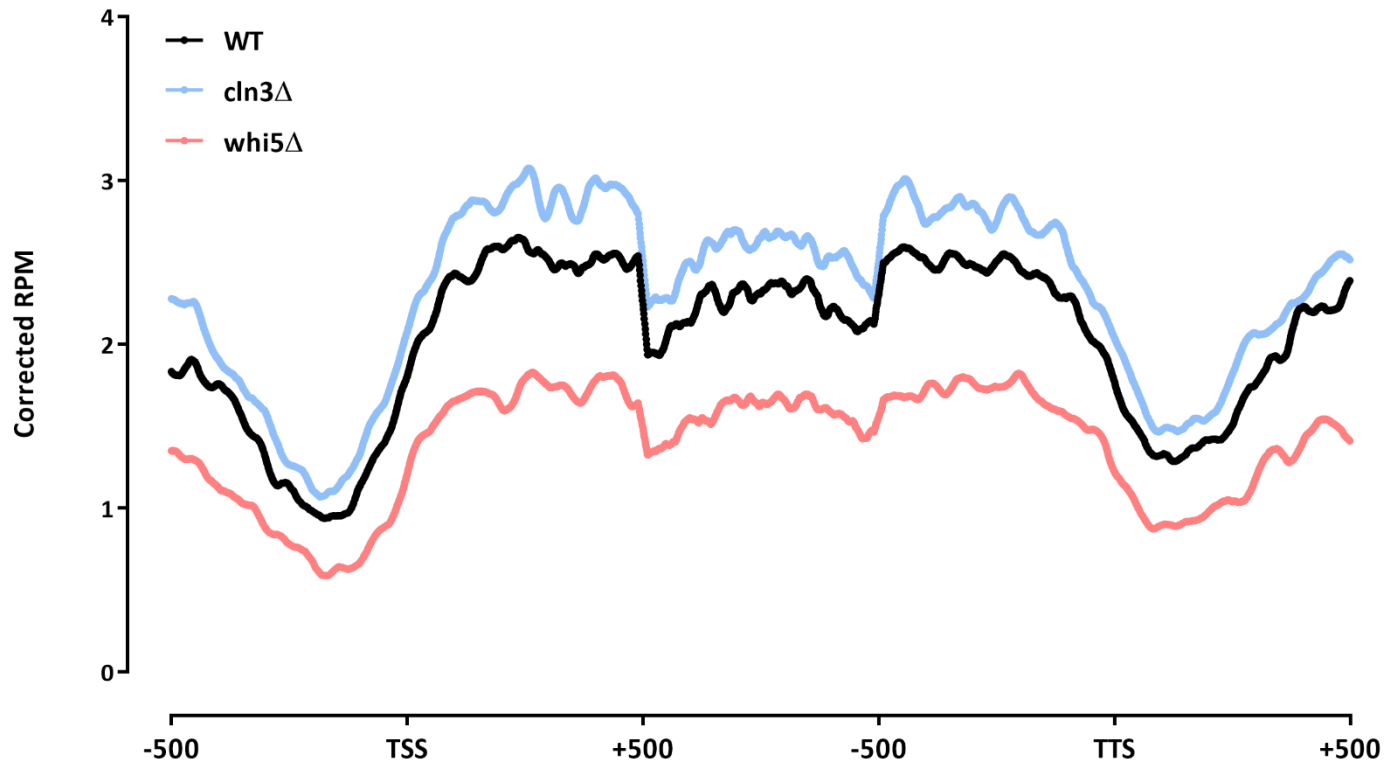
### H3 *S. cerevisiae* corrected 40%th



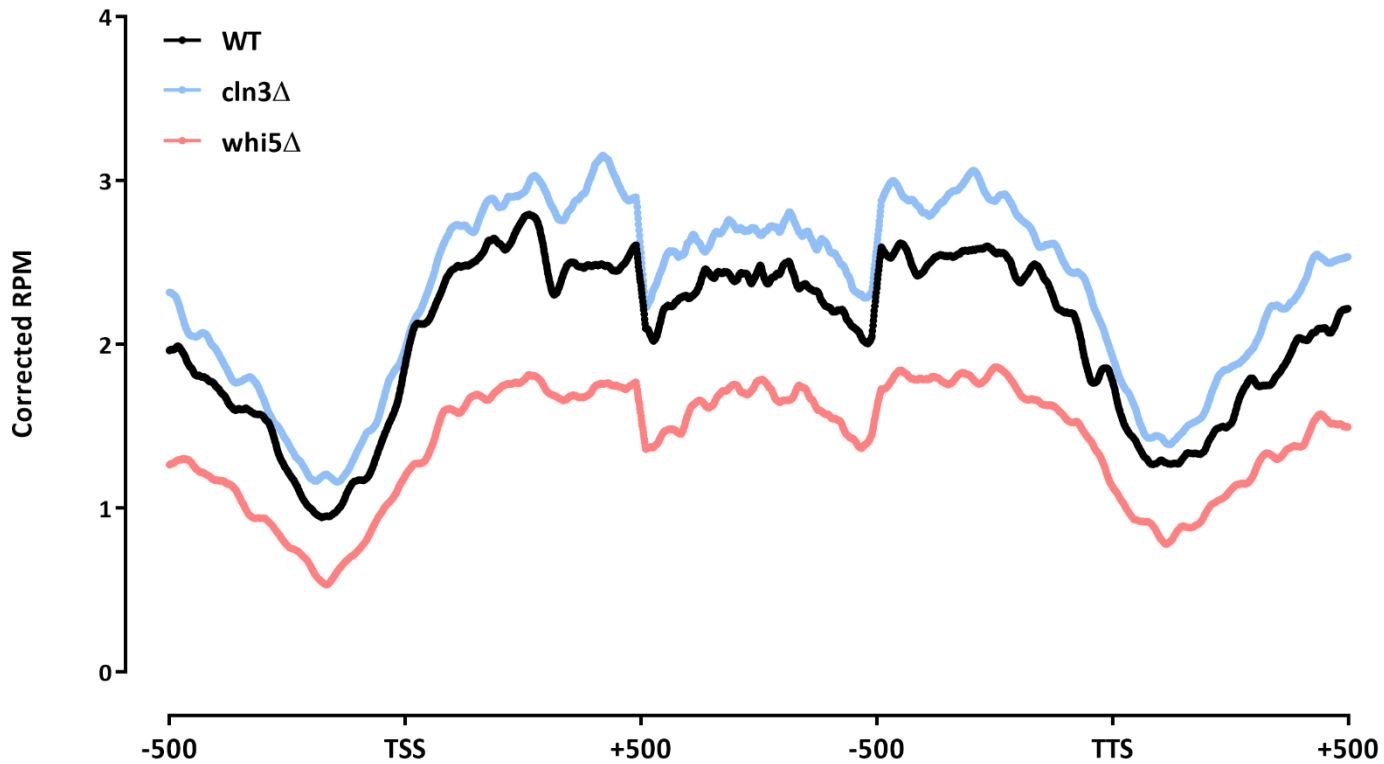
### H3 *S. cerevisiae* corrected 50%th



### H3 S. cerevisiae corrected 60%th

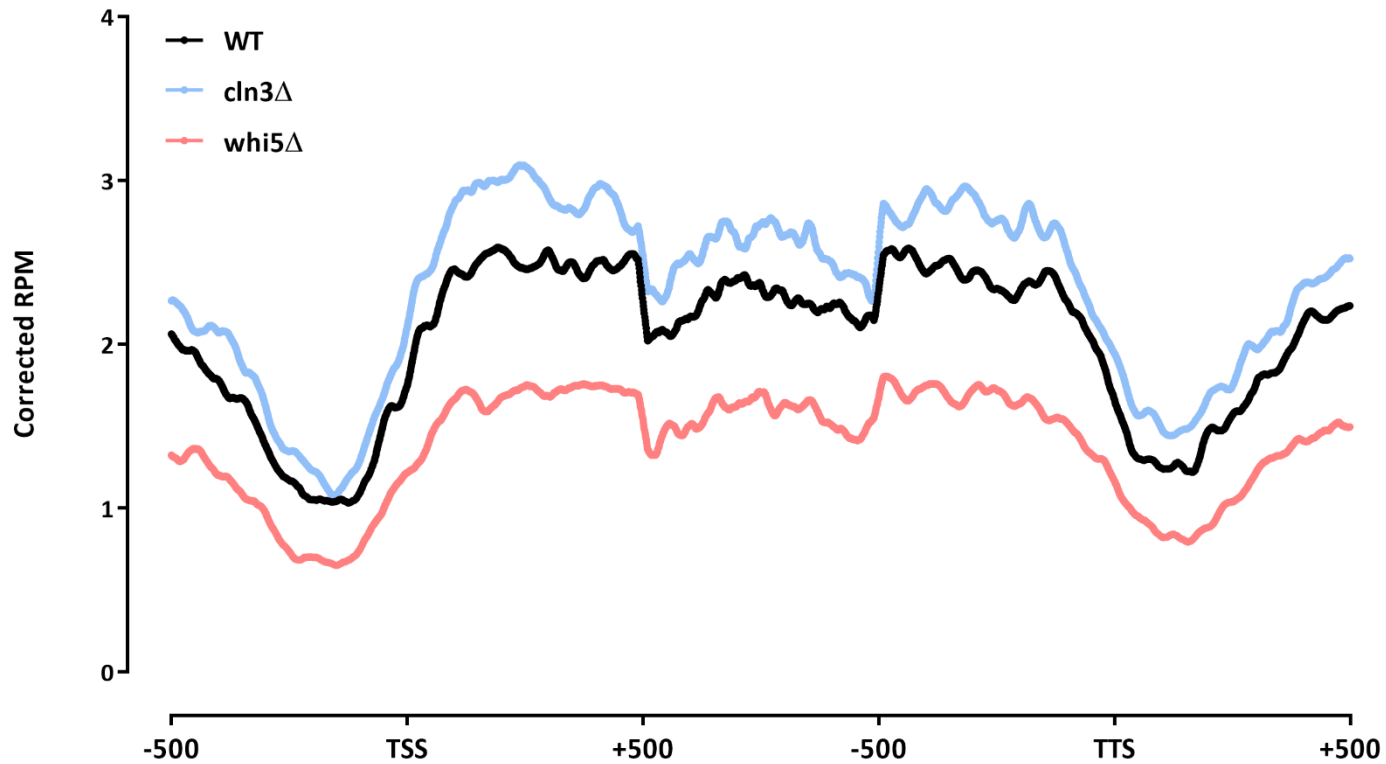


### H3 *S. cerevisiae* corrected 70%th

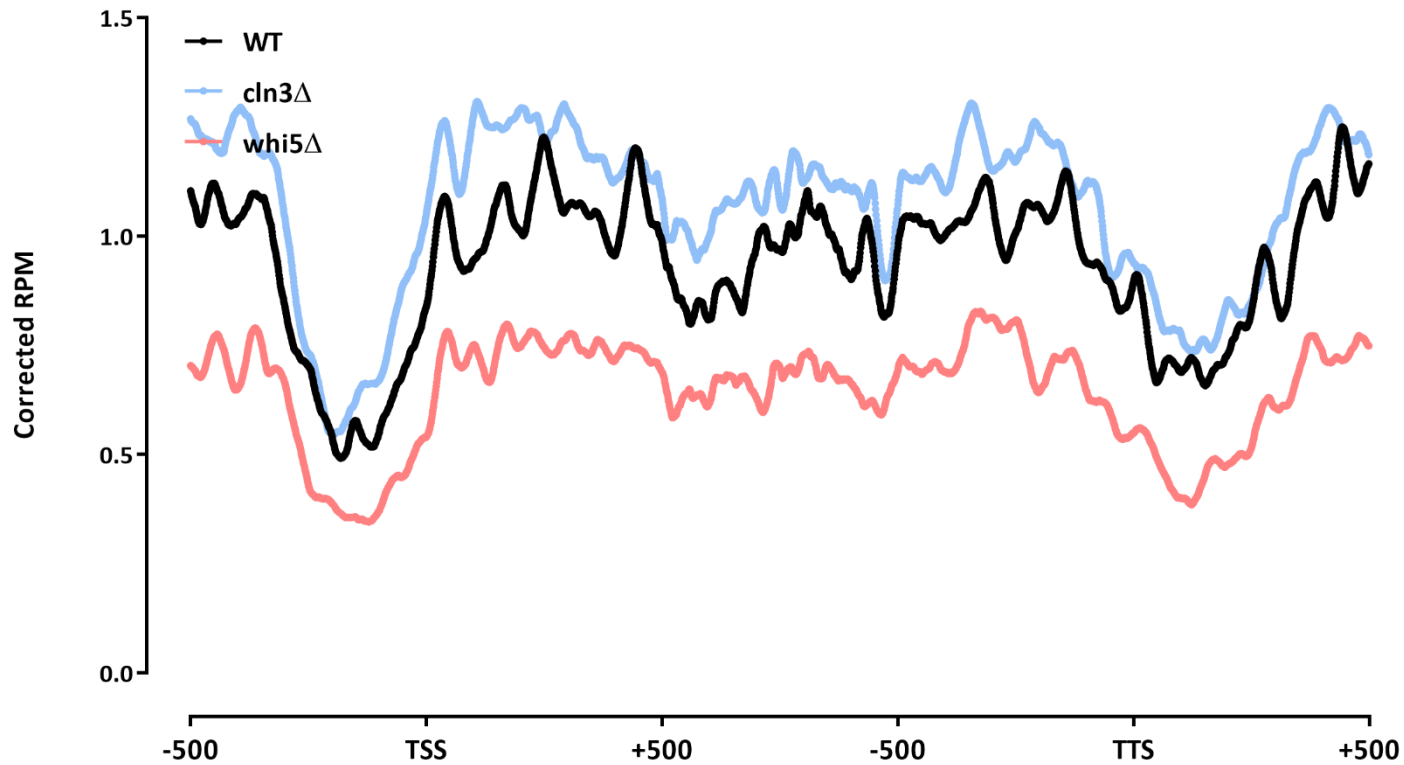




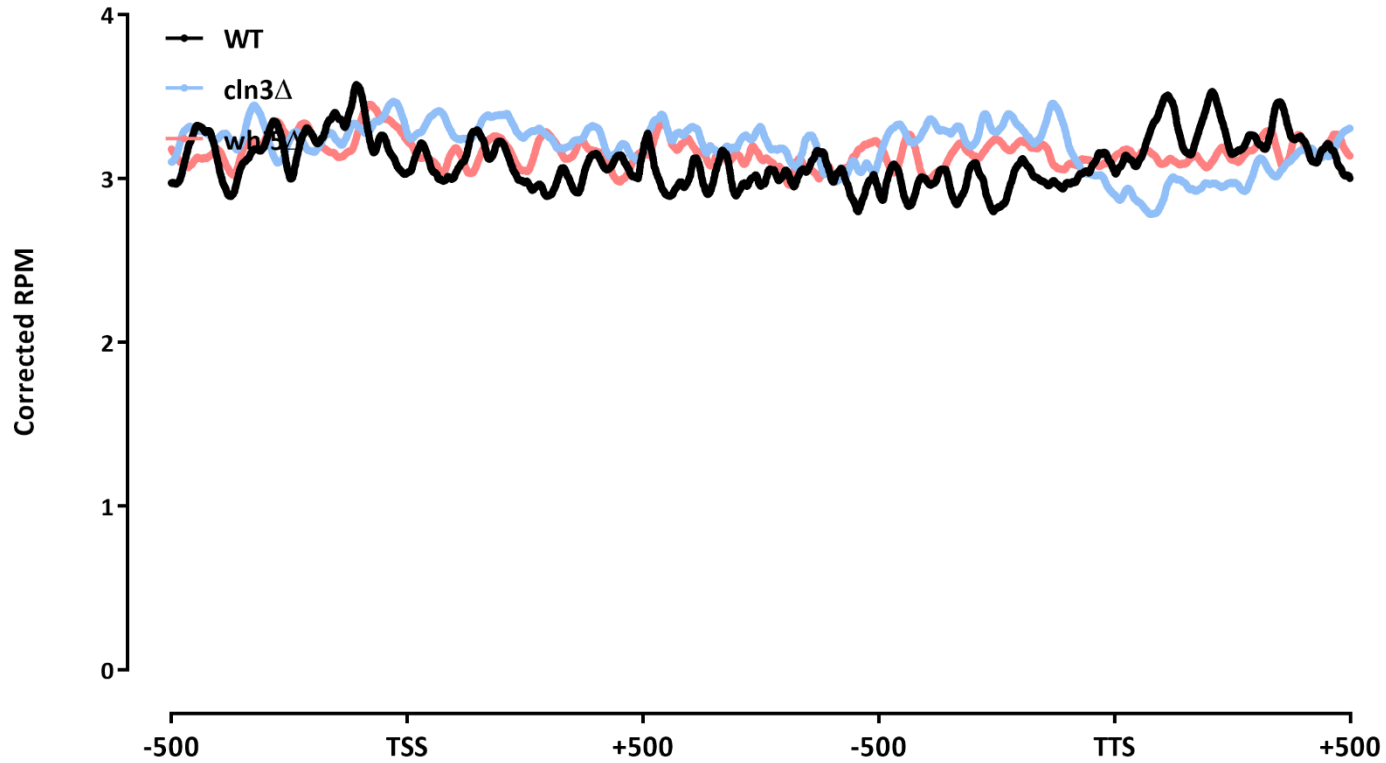
### H3 S. cerevisiae corrected 90%th



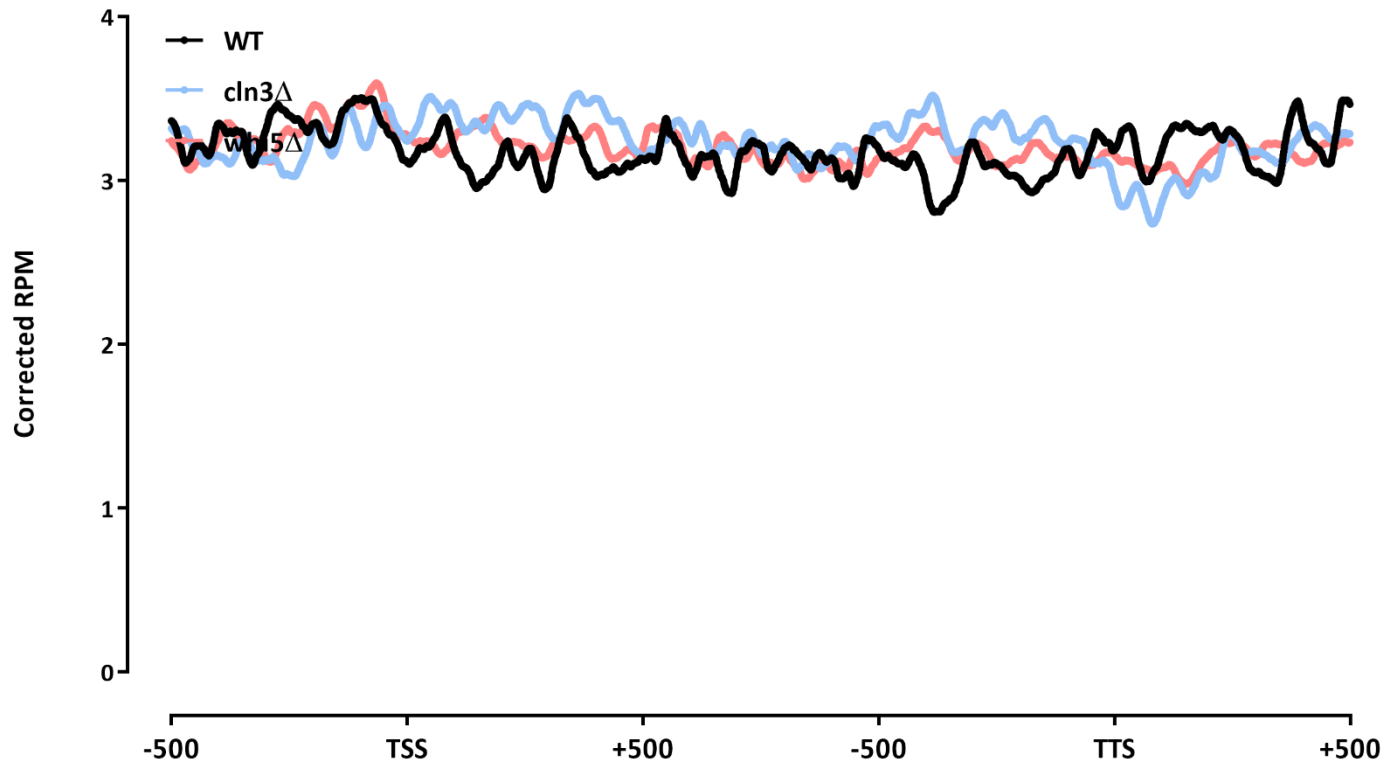
### H3 *S. cerevisiae* corrected 100%th



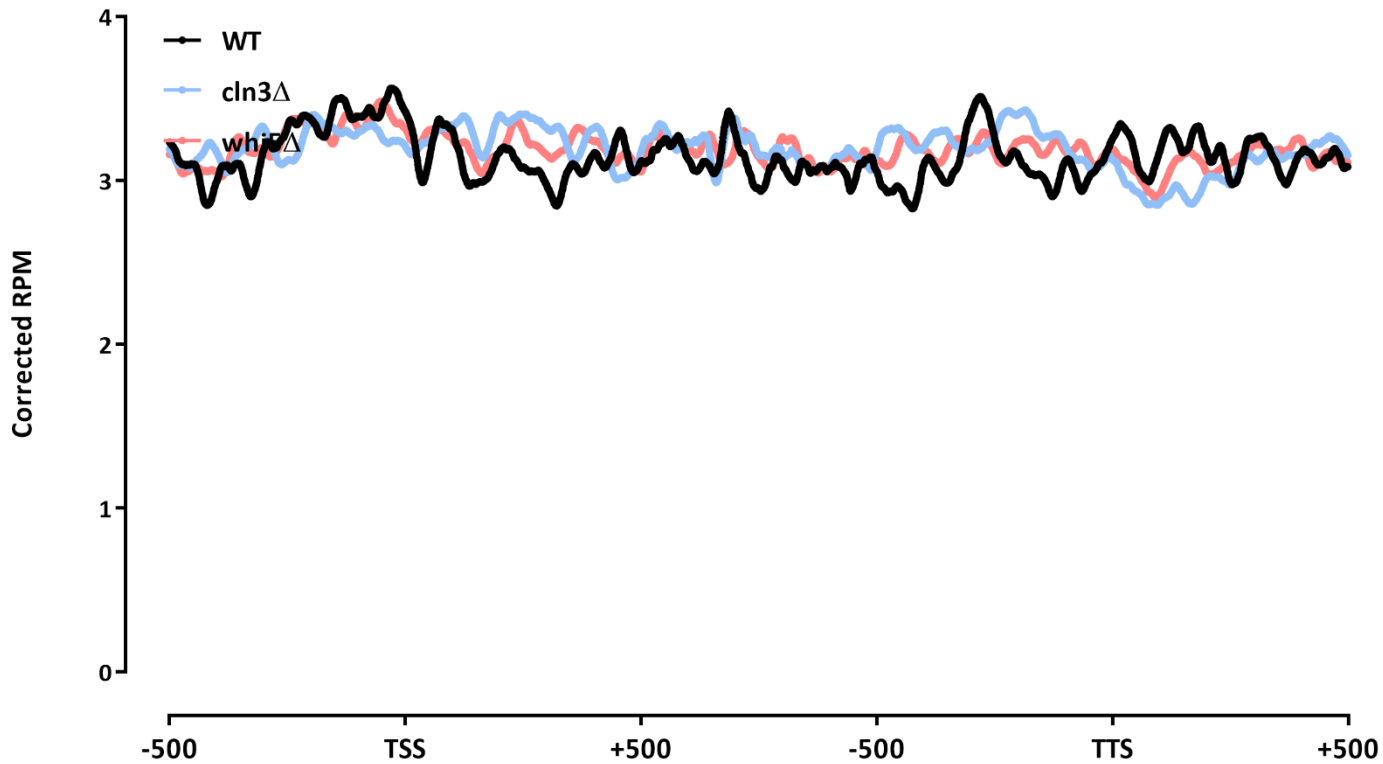
Input *S. cerevisiae* corrected 10%th



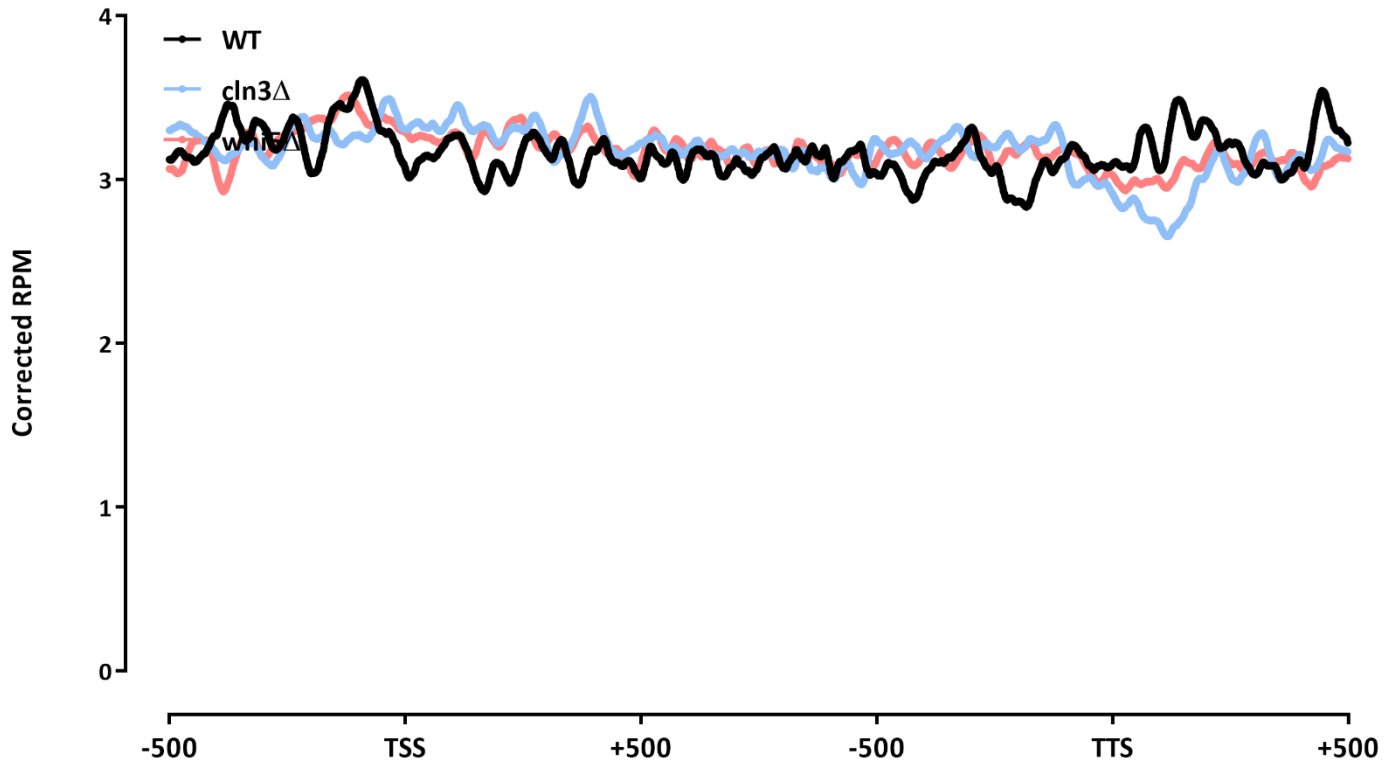
### Input *S. cerevisiae* corrected 20%th



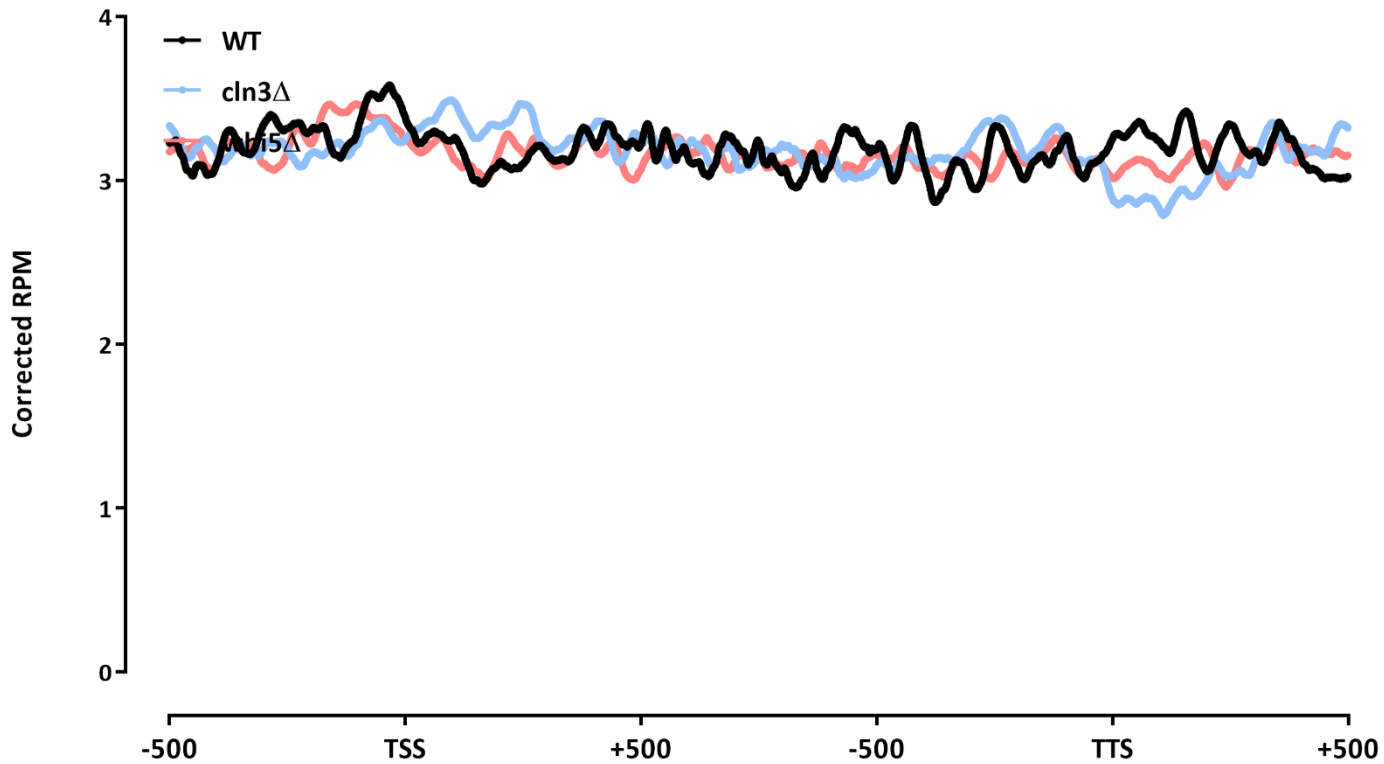
Input *S. cerevisiae* corrected 30%th



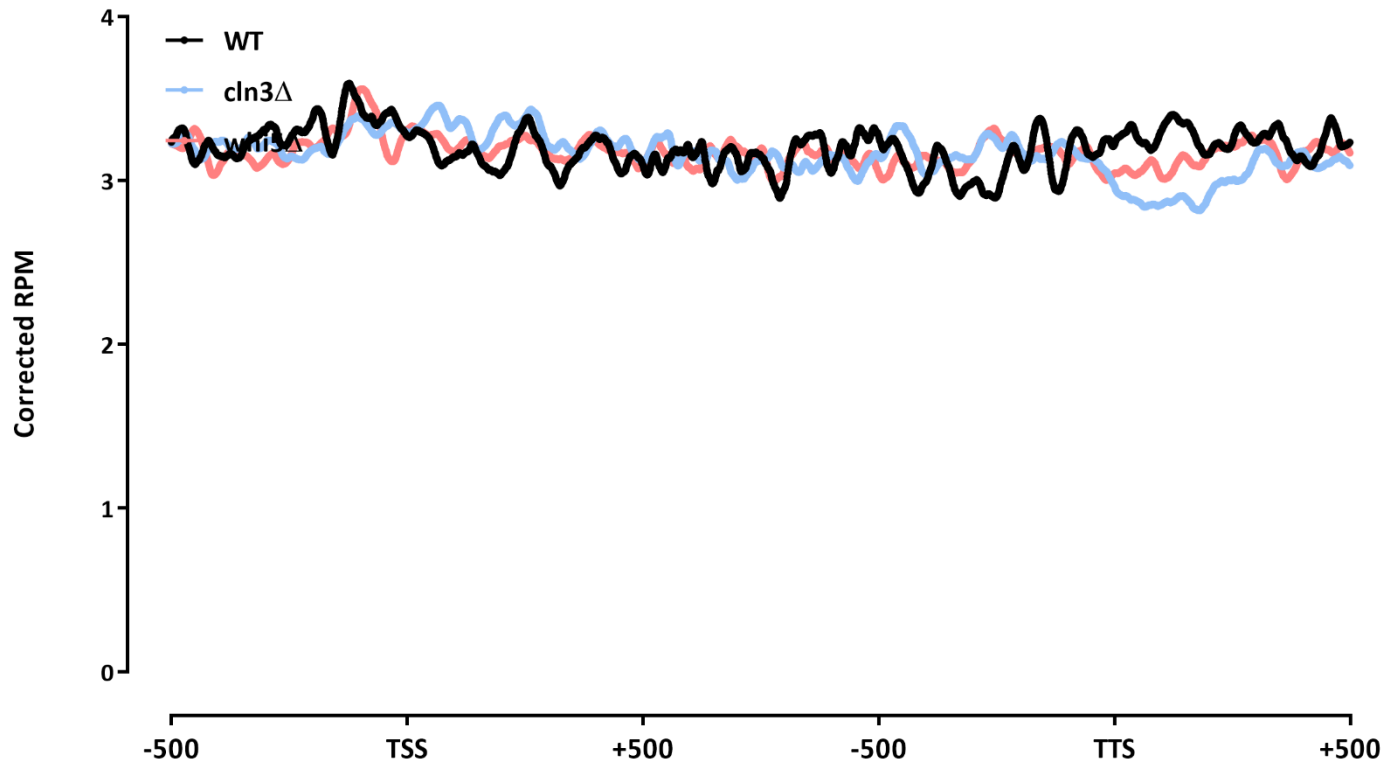
### Input *S. cerevisiae* corrected 40%th



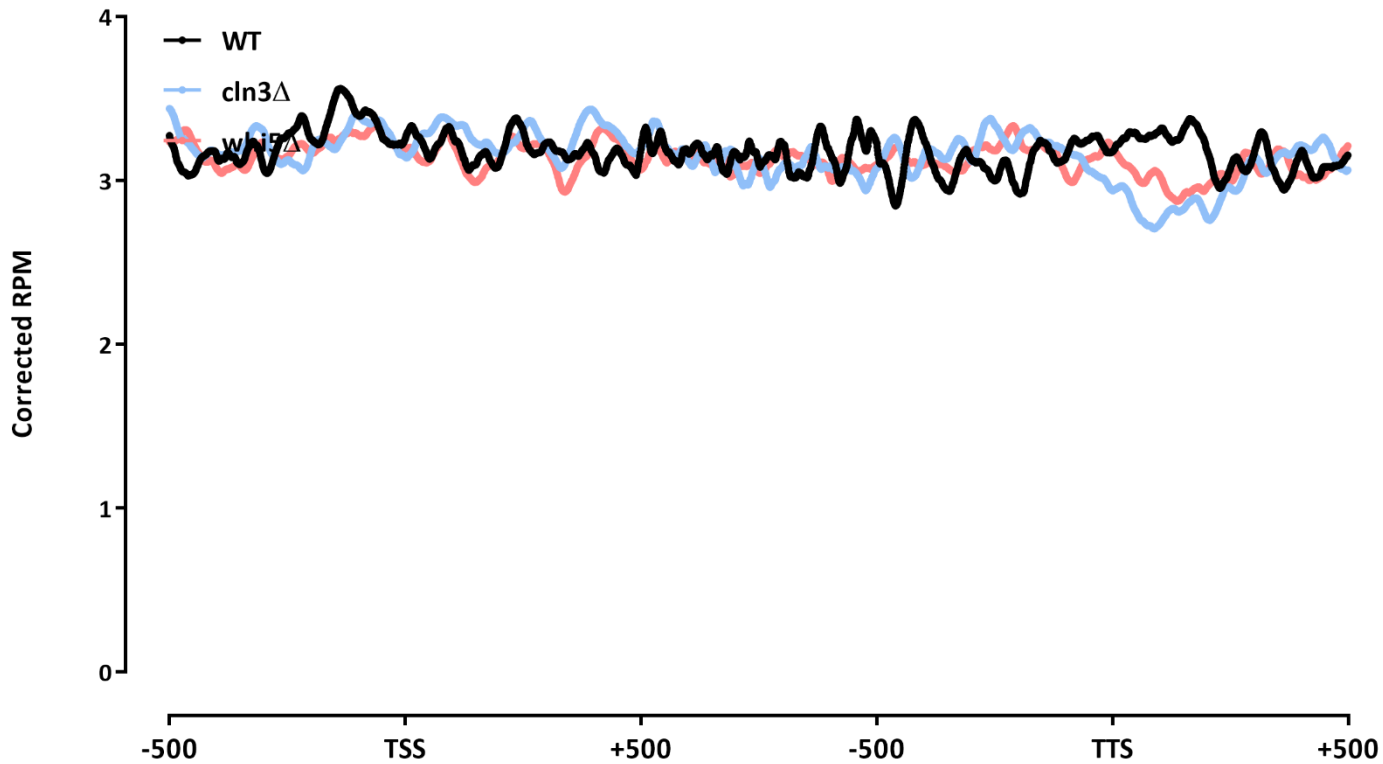
Input *S. cerevisiae* corrected 50%th



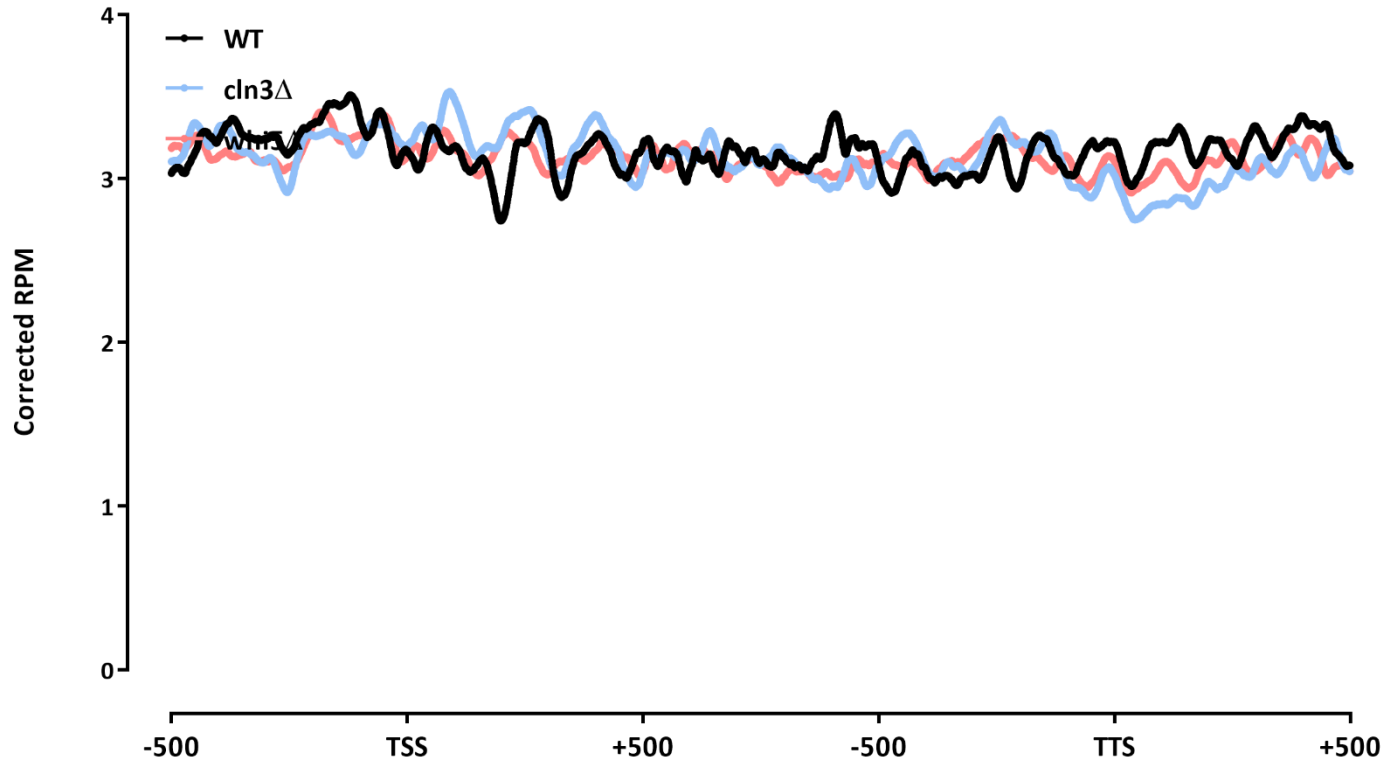
### Input *S. cerevisiae* corrected 60%th



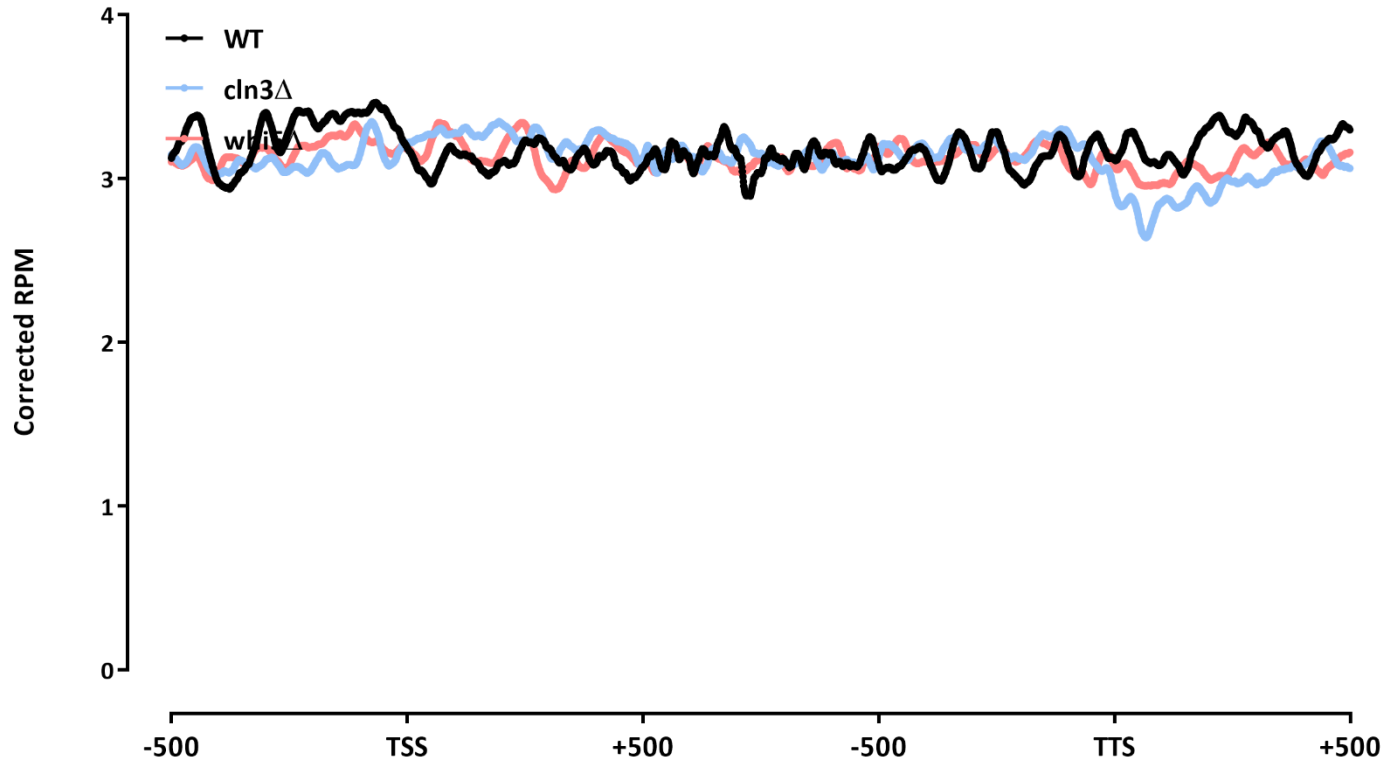
### Input *S. cerevisiae* corrected 70%th



### Input *S. cerevisiae* corrected 80%th



### Input *S. cerevisiae* corrected 90%th



### Input *S. cerevisiae* corrected 100%th

