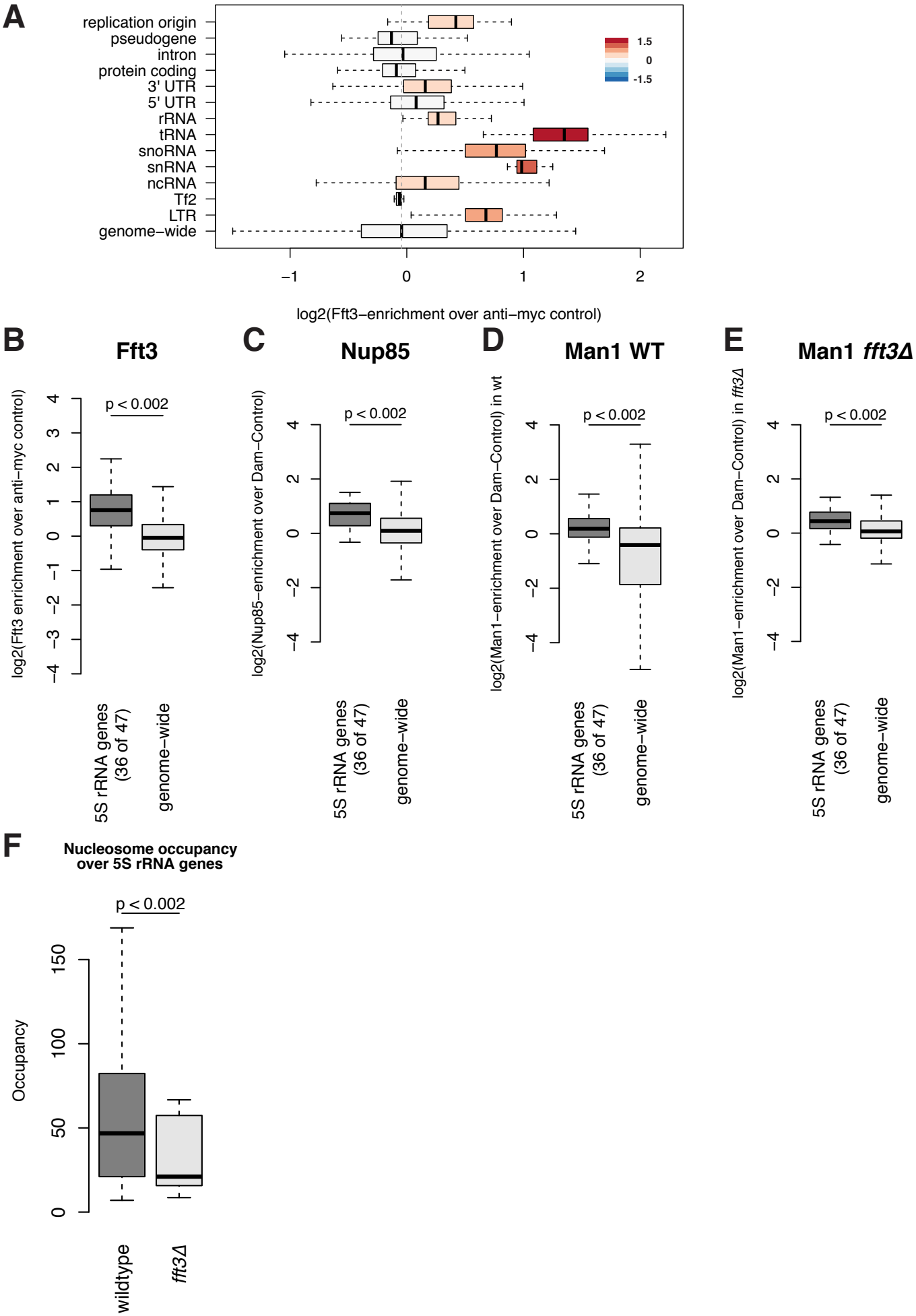
Figure S8


A
replication origin
pseudogene
intron
protein coding
3' UTR
5' UTR
rRNA
tRNA
snoRNA
snRNA
ncRNA
Tf2
LTR
genome–wide
1.5
0
-1.5
−1
0
1
2
log2(Fft3–enrichment over anti–myc control)
B
Fft3
log2(Fft3 enrichment over anti–myc control)
p < 0.002
5S rRNA genes (36 of 47)
genome–wide
C
Nup85
log2(Nup85–enrichment over Dam–Control)
p < 0.002
5S rRNA genes (36 of 47)
genome–wide
D
Man1 WT
log2(Man1–enrichment over Dam–Control) in wt
p < 0.002
5S rRNA genes (36 of 47)
genome–wide
E
Man1 fft3Δ
log2(Man1–enrichment over Dam–Control) in fft3Δ
p < 0.002
5S rRNA genes (36 of 47)
genome–wide
F
Nucleosome occupancy over 5S rRNA genes
Occupancy
p < 0.002
wildtype
fft3Δ