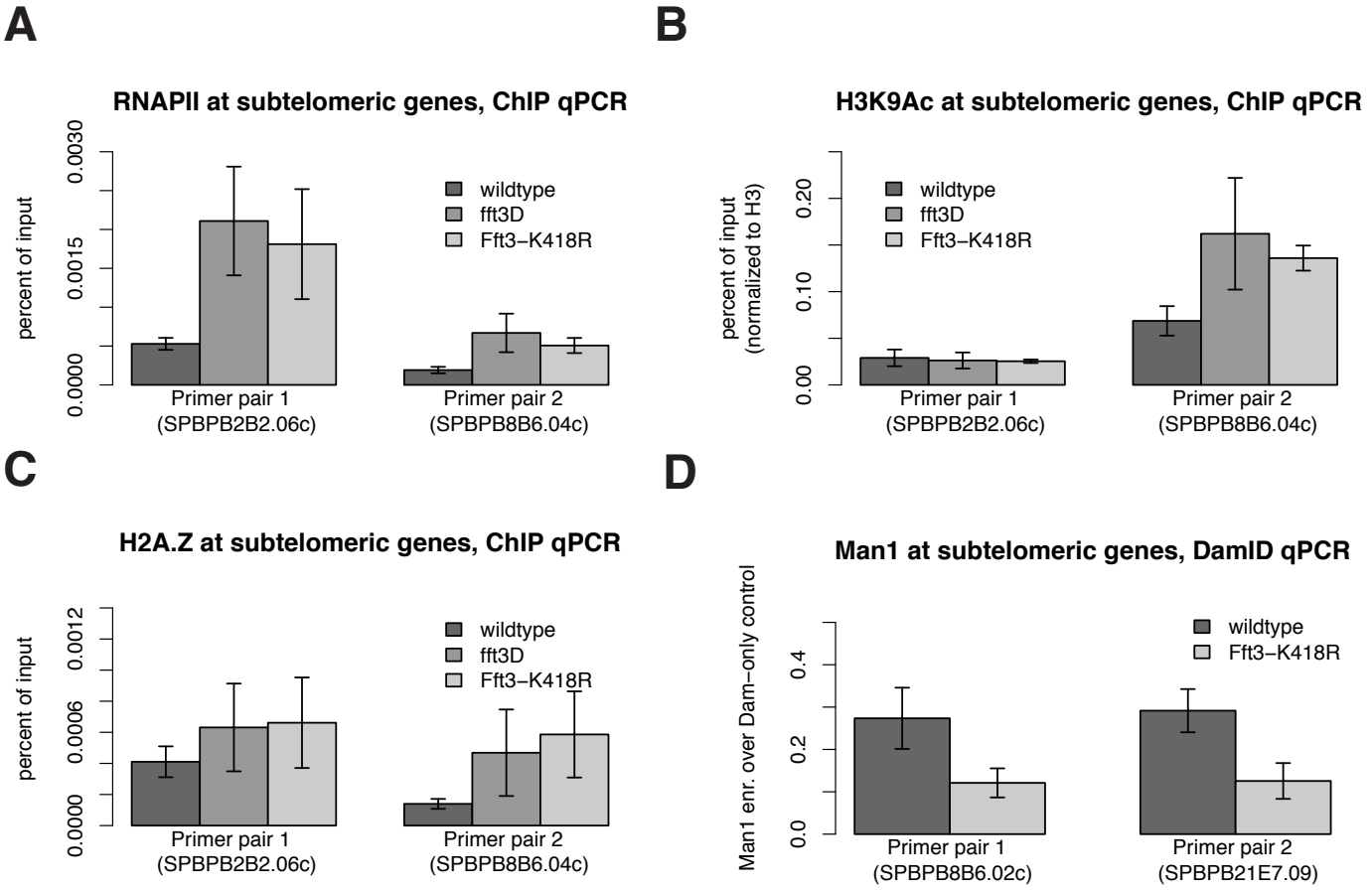
Figure S7


A
RNAPII at subtelomeric genes, ChIP qPCR
percent of input
0.0000
0.0015
0.0030
wildtype
fft3D
Fft3−K418R
Primer pair 1
(SPBPB2B2.06c)
Primer pair 2
(SPBPB8B6.04c)
B
H3K9Ac at subtelomeric genes, ChIP qPCR
percent of input
(normalized to H3)
0.00
0.10
0.20
wildtype
fft3D
Fft3−K418R
Primer pair 1
(SPBPB2B2.06c)
Primer pair 2
(SPBPB8B6.04c)
C
H2A.Z at subtelomeric genes, ChIP qPCR
percent of input
0.0000
0.0006
0.0012
wildtype
fft3D
Fft3−K418R
Primer pair 1
(SPBPB2B2.06c)
Primer pair 2
(SPBPB8B6.04c)
D
Man1 at subtelomeric genes, DamID qPCR
Man1 enr. over Dam−only control
0.0
0.2
0.4
wildtype
Fft3−K418R
Primer pair 1
(SPBPB8B6.02c)
Primer pair 2
(SPBPB21E7.09)