Clouaire et al. Additional File 7. Figure S5.

A

H3K4me3

wt  Cfp1−/−

±3 kb ±3 kb

H3K9,K14ac

wt  Cfp1−/−

±3 kb ±3 kb

B

Most active genes

H3K4me3

wt Doxo  Cfp1−/− Doxo

Distance from TSS (bp)

H3K9,K14ac

wt Doxo  Cfp1−/− Doxo

Distance from TSS (bp)

C

H3K4me3

wt  Cfp1−/−

±1.5 kb ±1.5 kb

H3K9,K14ac

wt  Cfp1−/−

±1.5 kb ±1.5 kb

D

Ectopic sites

H3K4me3

wt Input  Cfp1−/− Input  Cfp1−/− Doxo

Distance from peak centre (bp)

H3K9,K14ac

wt Input  Cfp1−/− Input  Cfp1−/− Doxo

Distance from peak centre (bp)

E

Most active genes

H3K4me3

Normalised Read Count (x10⁴)

H3K9,K14ac

Normalised Read Count (x10⁴)

F

Ectopic sites

H3K4me3

Normalised Read Count (x10⁴)

H3K9,K14ac

Normalised Read Count (x10⁴)