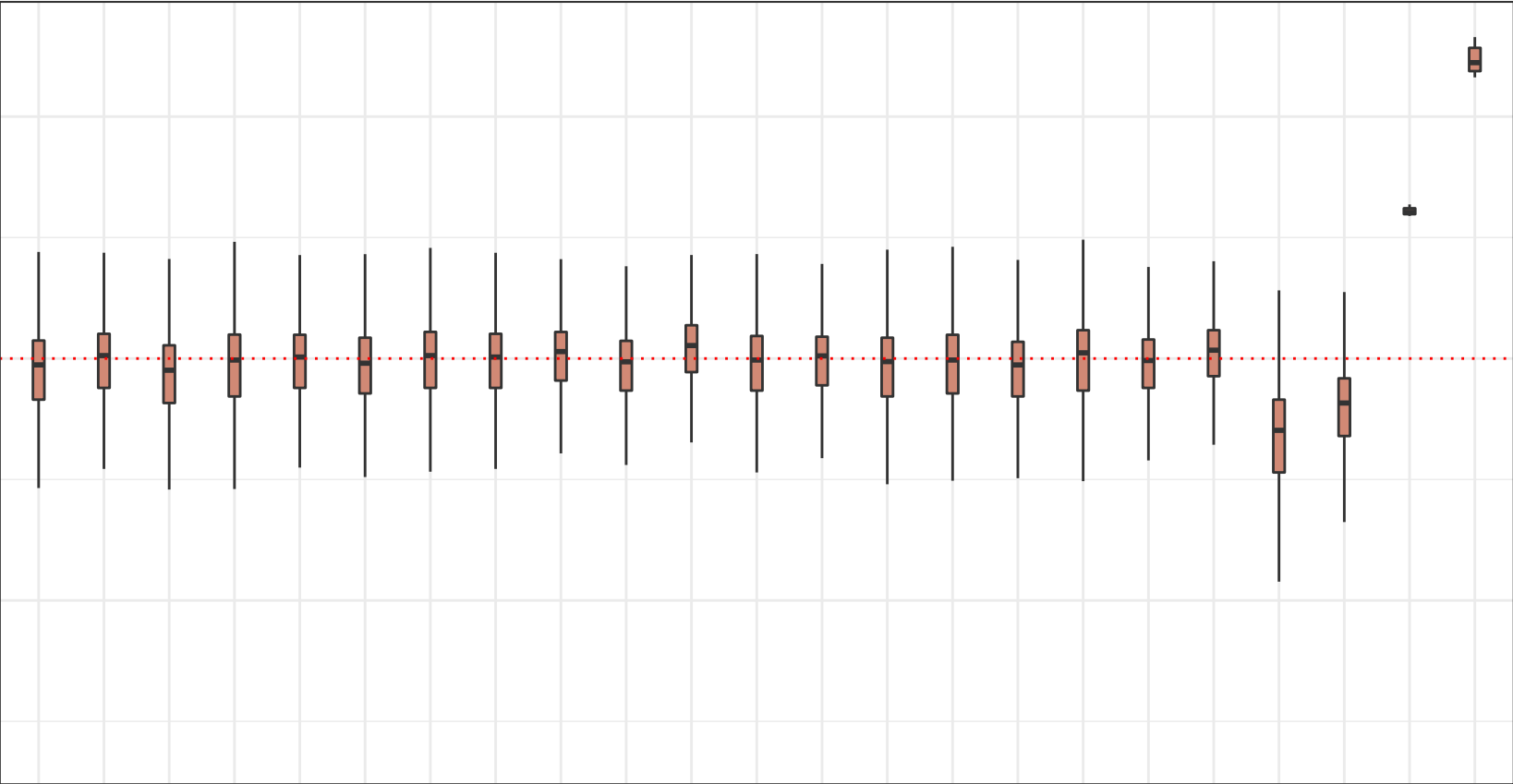


Normalised coverage per reference sequence per sample

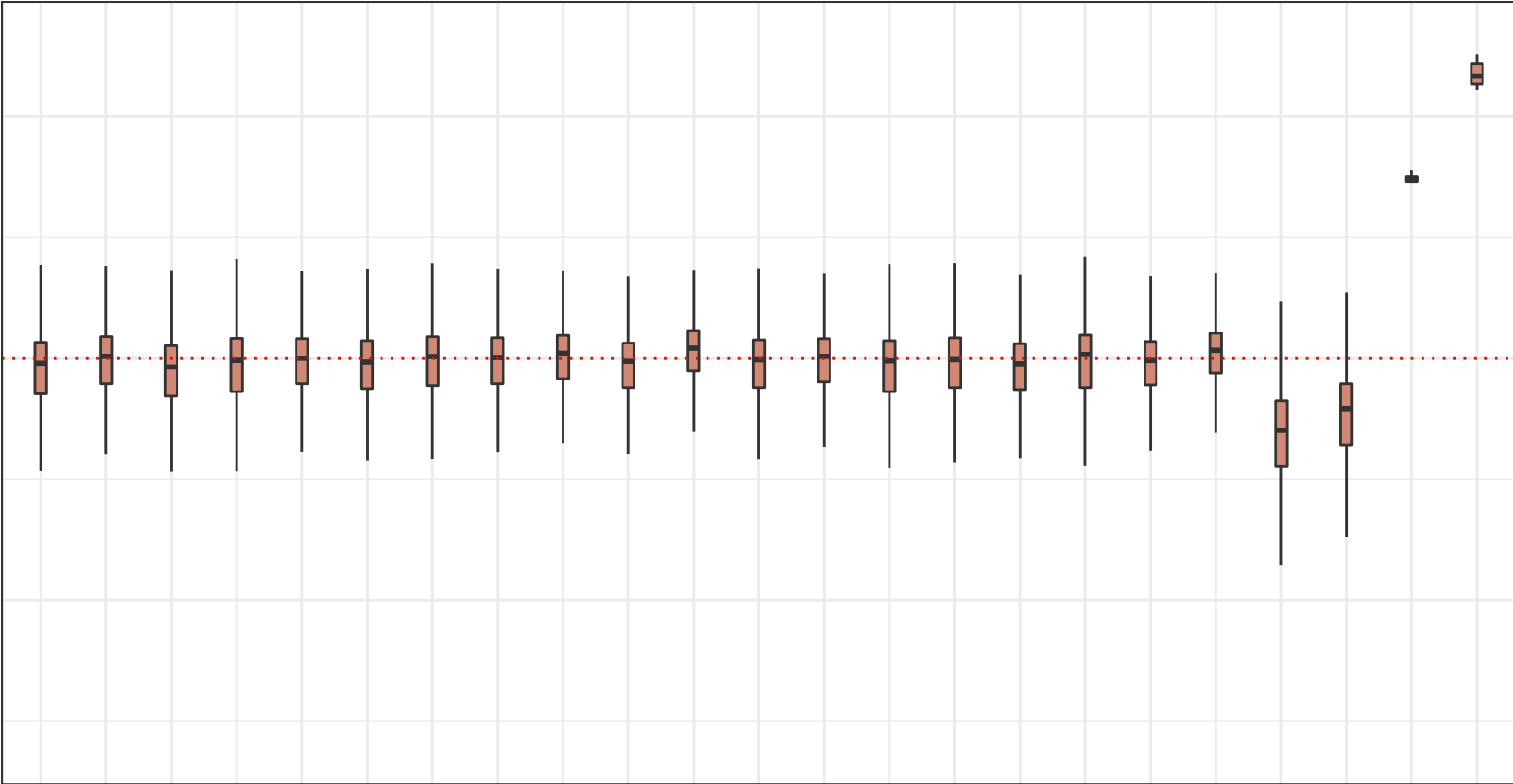
Note: Coverage (normalised to median autosomal genome coverage) is shown on a log10 scale

log10(Coverage normalised to median autosomal coverage)

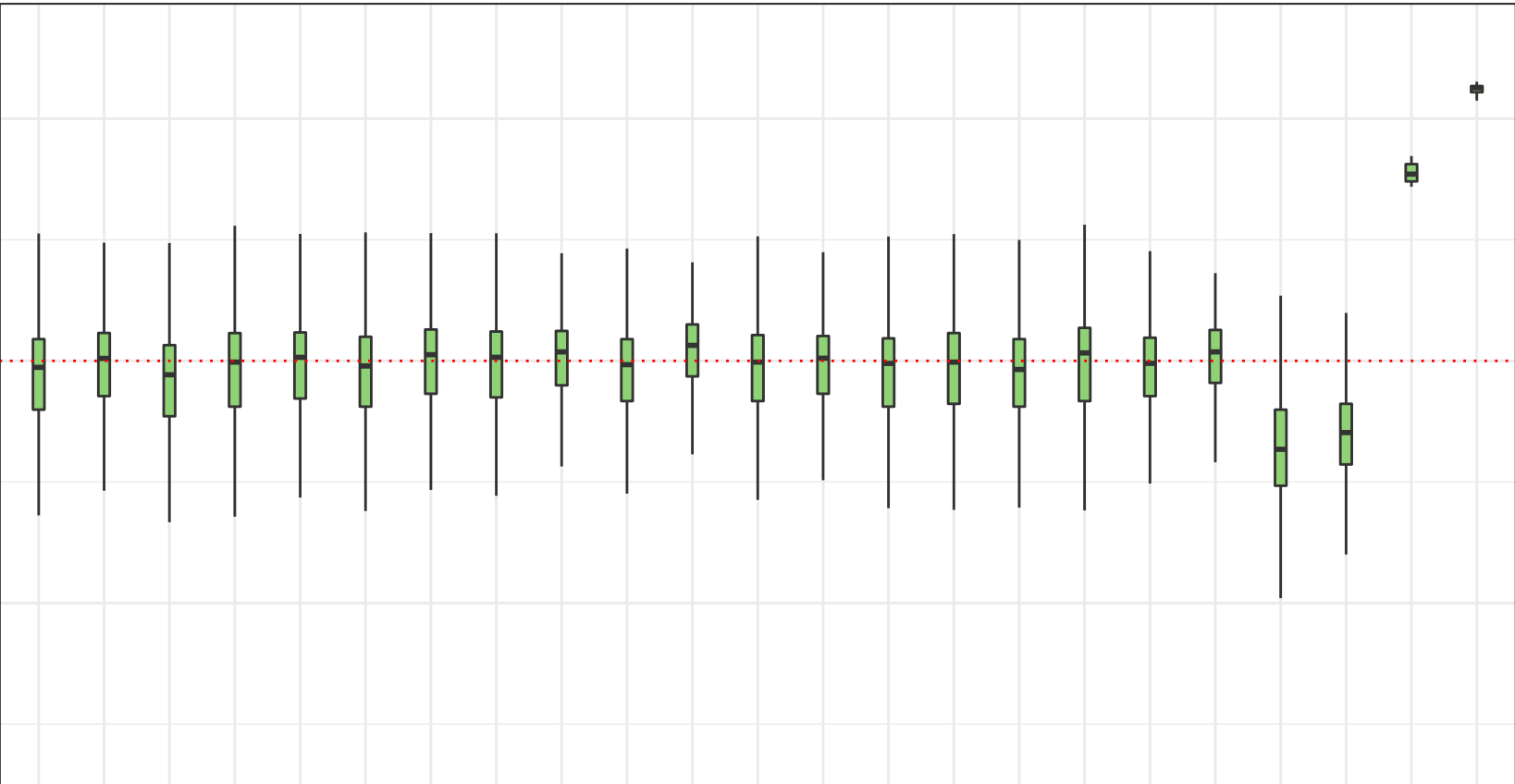
H3K27me3_brep1



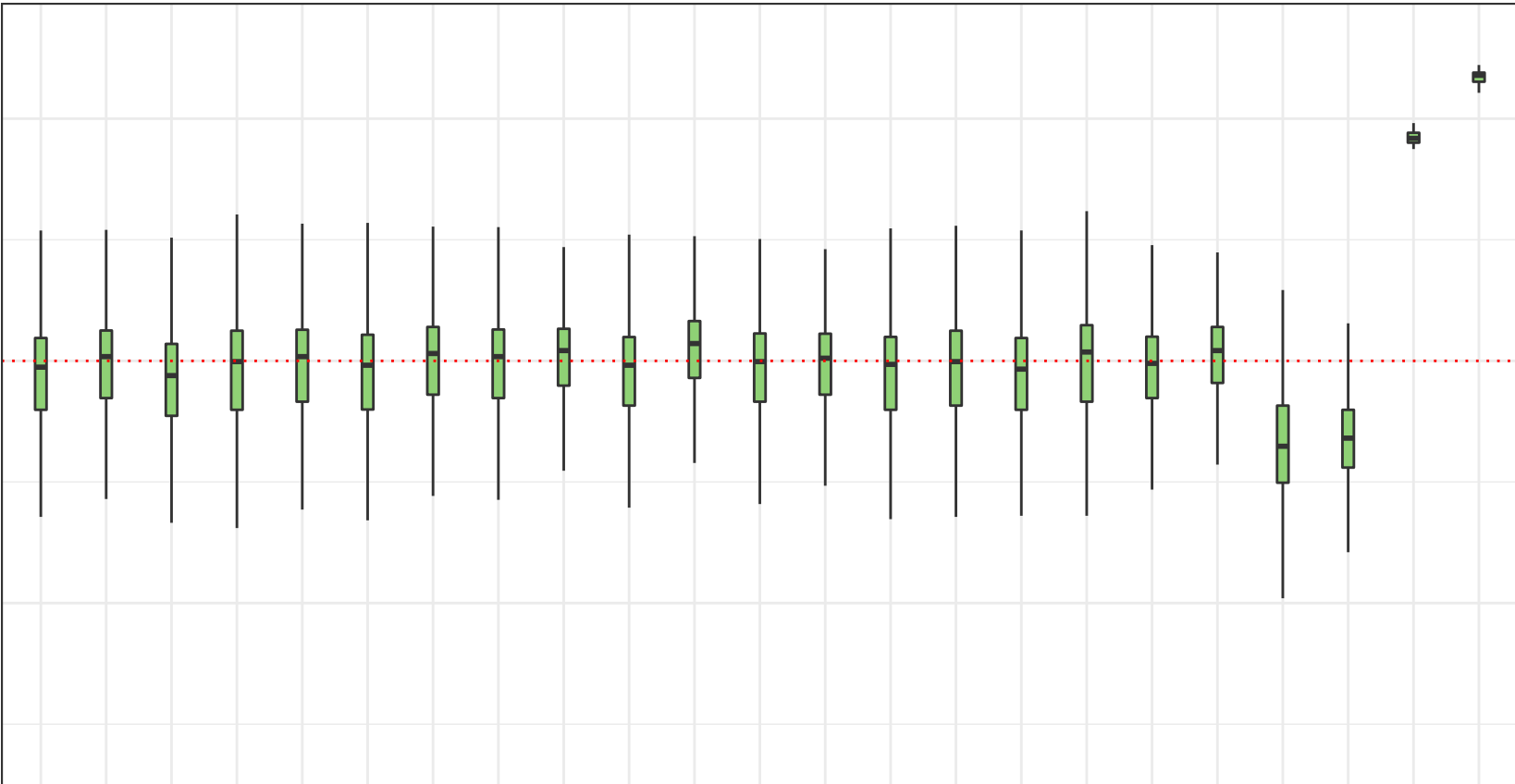
H3K27me3_brep2



input_brep1



input_brep2



Reference

Condition ChIP input