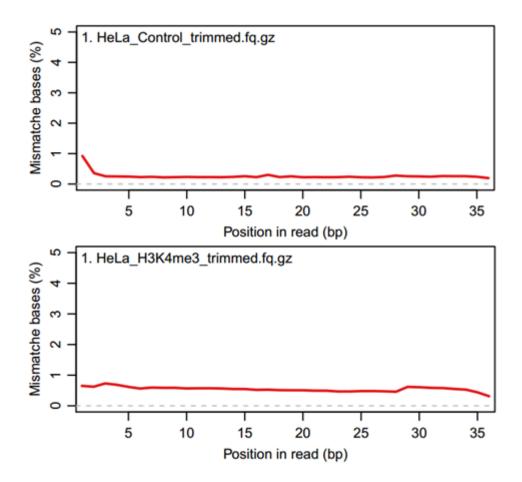
SUPPLEMENTARY INFORMATION

A ChIP-Seq Data Analysis Pipeline Based on Bioconductor Packages

Seung-Jin Park^{1,2}, Jong-Hwan Kim^{1,2}, Byung-Ha Yoon^{1,2}, Seon-Young Kim^{1,2}*

¹Personalized Genomic Medicine Research Center, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Daejeon 34141, Korea, ²Department of Functional Genomics, University of Science and Technology (UST), Daejeon 34113, Korea



Supplementary Fig. 5. Mismatch frequency. It shows the frequency and position of mismatches in the alignments against the reference genome.