

Figure S23

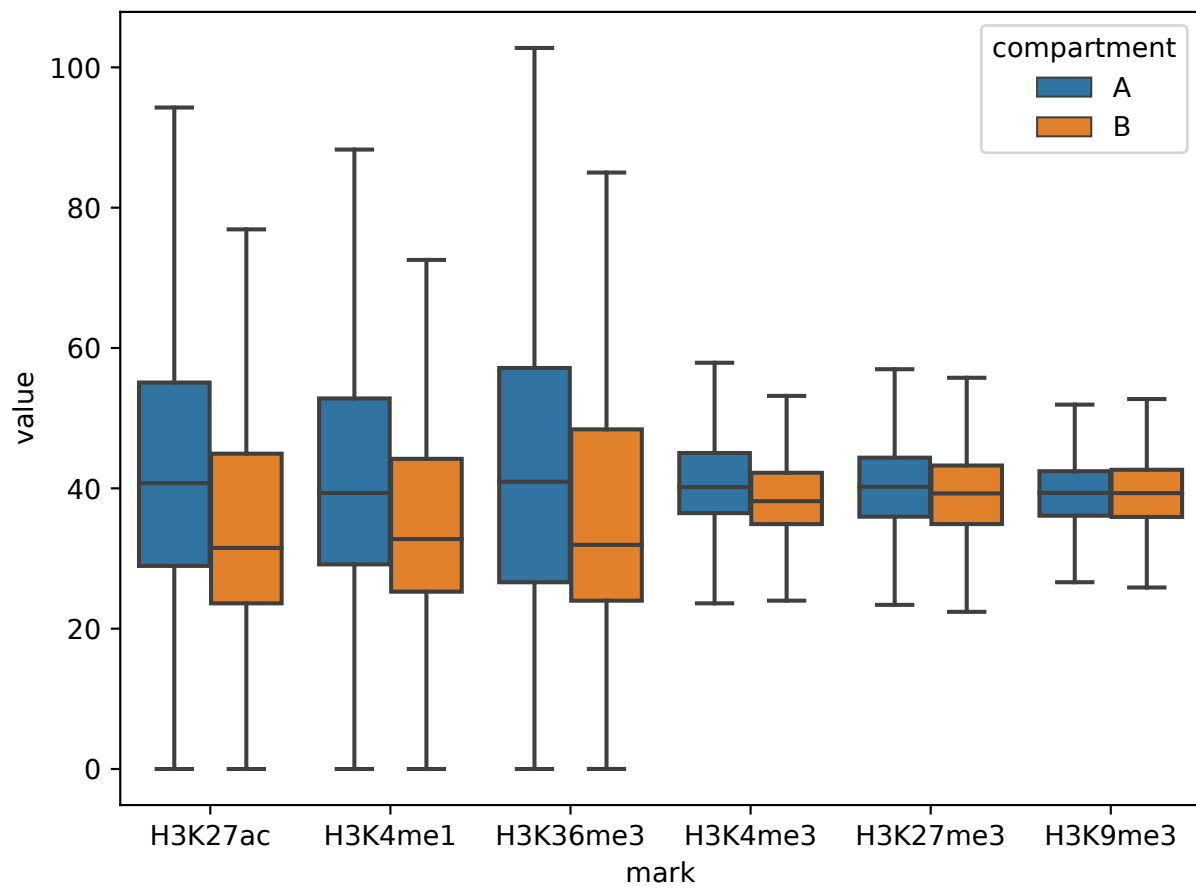


Figure S23. PCA-based A and B compartments (annotated based on GC content) and the distribution of various histone modification signals in each compartment (quantile-normalized mean signal).