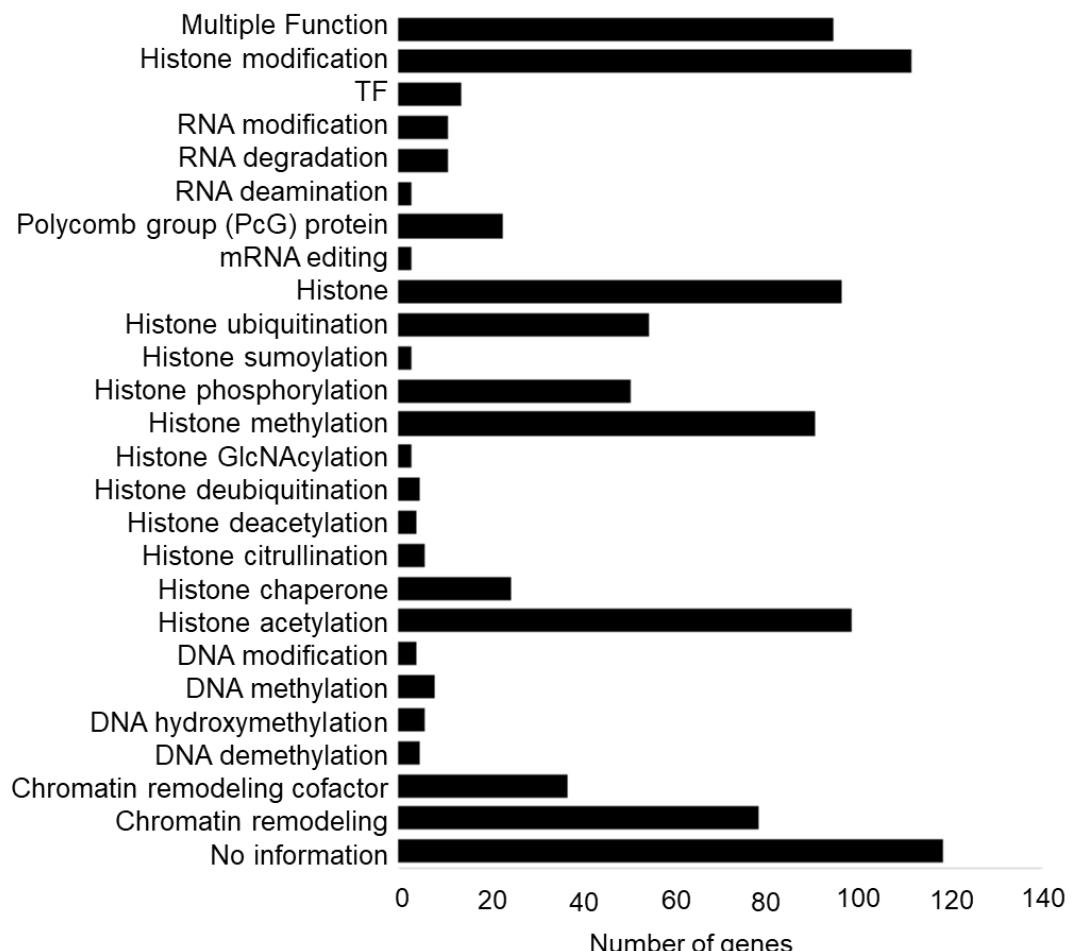
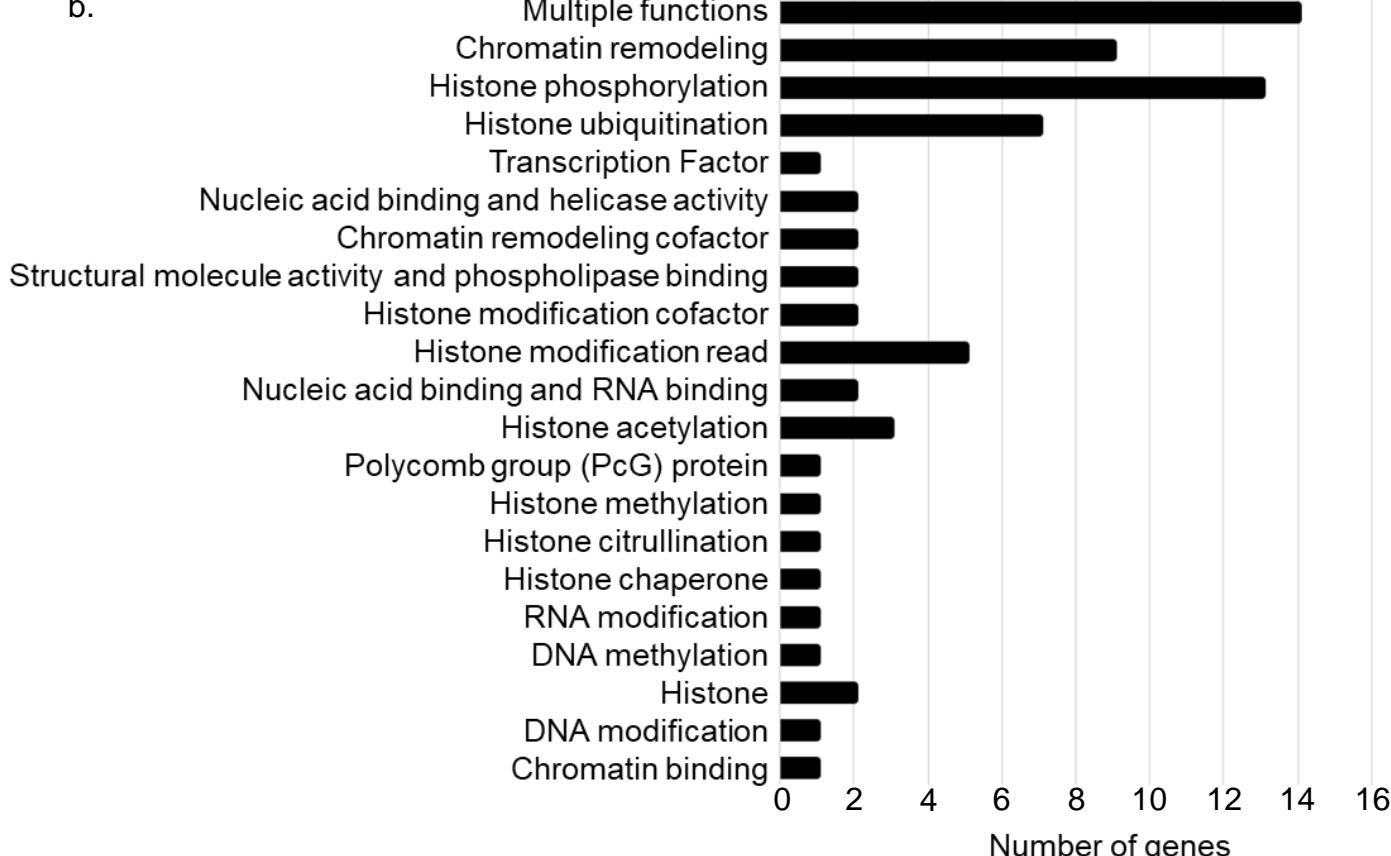


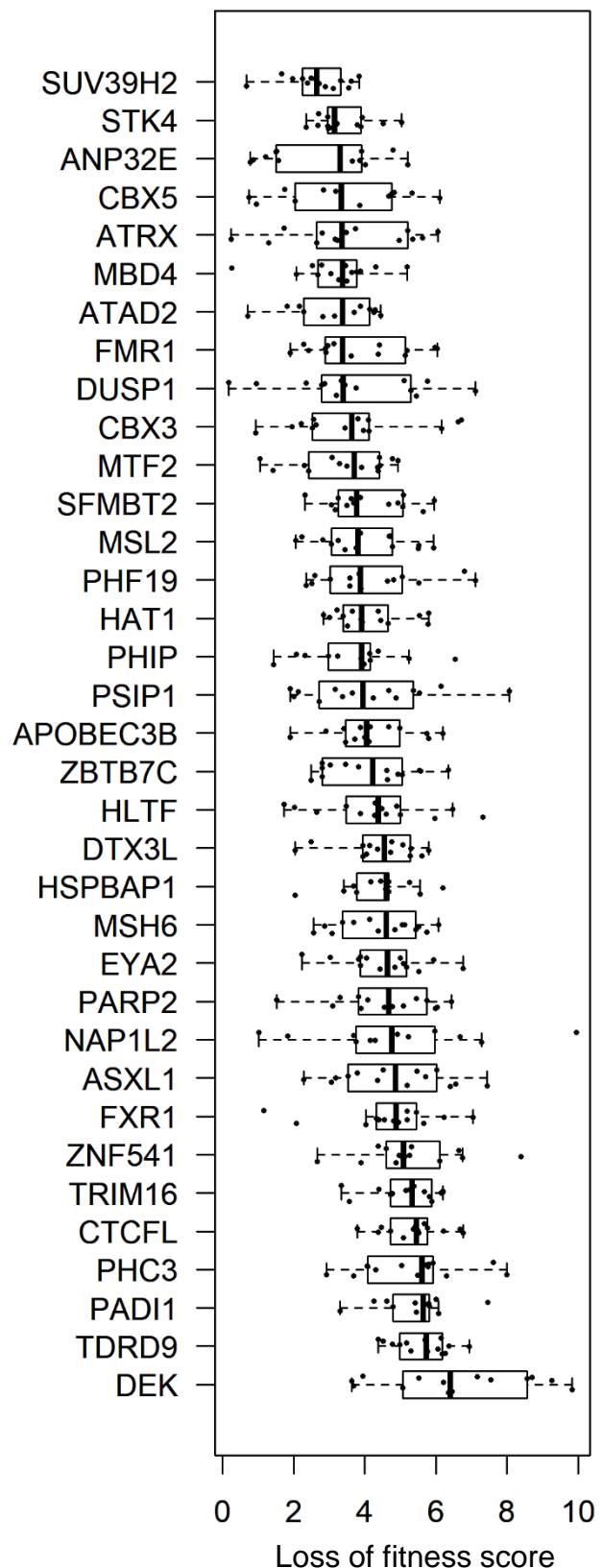
a.



b.



Supplementary figure S1: Epigenomic and chromatin remodeling regulators. a, Classes of molecules curated in the present study. Y-axis represents the functional classification and X-axis represents the number of genes in each functional group, and b, Functional sub-types of 73 differentially expressed epigenomic and chromatin regulators.



Supplementary figure S2: Genes with poor fitness dependency values in cervical cancer cell lines.