Genes with hypomethylated promoters in our study

12 32 31

Escape genes or mostly escape genes

in the previous

study

**Figure S3. The number of escape genes.** The pink circle indicates the number of genes with hypomethylated promoters in our study. The blue circle represents the number of escape genes or mostly escape genes in the previous study (1). The overlapped area shows the genes which we evaluated in the patients in this study.

## Reference

1. Balaton BP, Cotton AM, Brown CJ. Derivation of consensus inactivation status for X-linked genes from genome-wide studies. Biol Sex Differ 2015;30:6:35.