

Supplementary Table Guide

Supplementary Table 1. Design of NuLibA.

Supplementary Table 2. NuLibA sequences and read counts in each sample.

Supplementary Table 3. Mapping statistics of NuLibA and NuLibB experiments.

Supplementary Table 4. Nuc/Cyt and WCE/Input ratios for NuLibA experiments with tile feature annotations.

Supplementary Table 5. Regions that led to nuclear enrichment identified in NuLibA.

Supplementary Table 6. Design of NuLibB.

Supplementary Table 7. NuLibB sequences and read counts in each sample.

Supplementary Table 8. Nuc/Cyt and WCE/Input ratios for NuLibB experiments with tile feature annotations.

Supplementary Table 9. Regions that led to nuclear enrichment identified in NuLibB.

Supplementary Table 10. Enrichments of binding of different proteins in the SIRLOIN region of Alu, based on eCLIP data from the ENCODE project.

Supplementary Table 11. Correlations between numbers of eCLIP clusters and Nuc/Cyt ratios in HepG2 and K562 cells.

Supplementary Table 12. Sequences of primers used in this study.