

Supplemental Fig. S1. A⁶Sm⁶As identification.

(A) MeRIP-qPCR of m⁶A peaks at heterozygous sites at genes in Kidney-

2. The IP versus input of the peak was normalized to *GAPDH*.

(B) Pipeline for A⁶Sm⁶As identification from GLORI-seq and m⁶A-SAC-seq.

(C) Representative examples showing the A⁶Sm⁶As identified.

Fig. S1. ASm⁶As identification.

