

Supplementary information, Fig. S9. The C-terminal of VIRMA plays a vital role in the process of m⁶A methylation. The N-terminal region (1-333) before the visible region and C-terminal region (1587-1812) after the Halter domain were truncated for RNA binding and methylation assay. **a** The binding affinities of $HWVZ^{(1106-1668)}$, $HWV^{(1-1586)}Z^{(1106-1668)}$, and $HWV^{(334-1586)}Z^{(1106-1668)}$ complex with FAM-labeled actb-1 RNA from EMSA experiments. Binding curves were generated using the mean of three biologically independent experiments, with standard deviation (SD) values indicated by error bars (n = 3). **b** Actb-1 RNA N6-adenosine methylation activity of MAC, MAC/HWVZ^{(1106-1668)}, MAC/HWV^{(1-1586)}Z^{(1106-1668)}, and MAC/HWV^{(334-1586)}Z^{(1106-1668)} complex. The bar chart was generated using the mean of three biologically independent experiments, with standard deviation (SD) values indicated by error bars (n = 3).