

Figure S1. Calibration curves for the quantifications of A in mRNA. The amounts of A ranged $0.1 - 2.5 \mu$ mol.

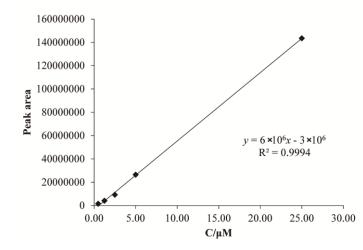


Figure S2. Calibration curves for the quantifications of C in mRNA. The amounts of C ranged 0.5 - 25 μ mol.

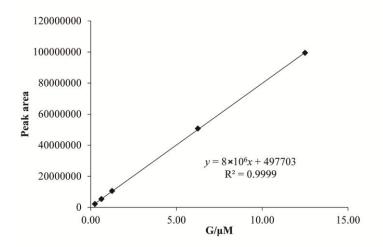


Figure S3. Calibration curves for the quantifications of G in mRNA. The amounts of G ranged $0.25 - 12.5 \mu mol$.

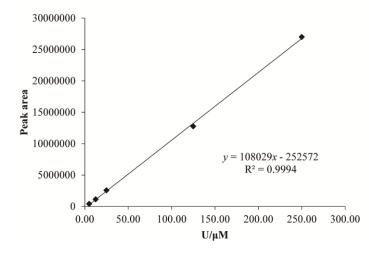


Figure S4. Calibration curves for the quantifications of U in mRNA. The amounts of U ranged 5 - 250 μ mol.

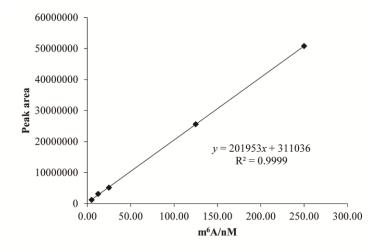


Figure S5. Calibration curves for the quantifications of m^6A in mRNA. The amounts of m^6A ranged $0.005 - 0.250 \ \mu mol$.

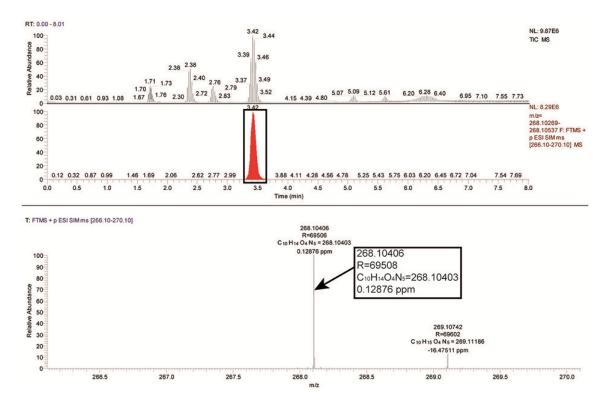


Figure S6. Representative UHPLC-HR-MS/MS results for the quantifications of A in N. tabacum K-326 with selected-ion chromatograms.

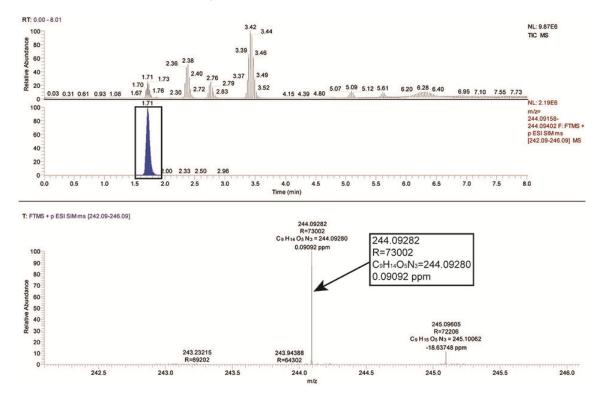


Figure S7. Representative UHPLC-HR-MS/MS results for the quantifications of C in N. tabacum K-326 with selected-ion chromatograms.

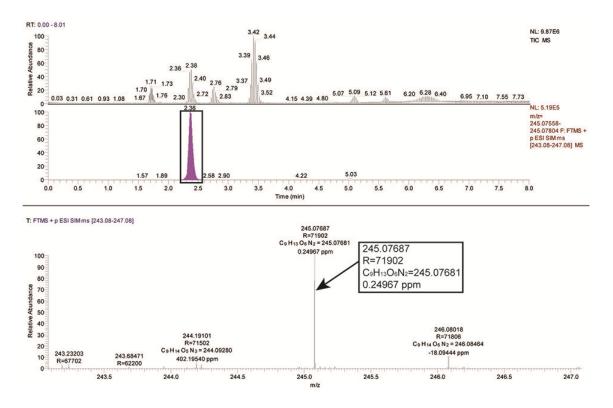


Figure S8. Representative UHPLC-HR-MS/MS results for the quantifications of U in N. tabacum K-326 with selected-ion chromatograms.

Table S1. Primer sequences for RT-qPCR analysis

Gene name	Primer sequences (5'-3')	Amplicon length (bp)
XM_009766347	F: ACGCTCATTTTGCTTCCAGT	171
	R: ATCCCCTTCCCAGTCGTCAT	
XM_009766348	F: TTGGCCGTAGAAGGCTTGAG	114
	R: GTGTAGGCCTGAACTCAGCA	
XM_009775897	F: GGCAAACGGAAGATGAAGCC	172
	R: CTGGTGCTGGTGTGATGGAT	
XM_009770927	F: TAGGCCGACACGTACTGAGA	163
	R: CGGCGGCATATTGAAACAGG	
XM_009801708	F: AGGTTTATGTCCCCACTGCG	114
	R: TCTGCAGAACTTTTTGTGACATT	
XM_009801663	F: GTTCCGCCACTTTTGTCGAG	148
	R: AGATGGCACCTCTGGGCATA	
β -actin	F: GACATGAAGGAGGAGCTTGC	178
	R: ATCATGGATGGCTGGAAGAG	

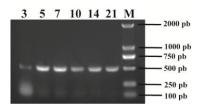


Figure S9. TMV detection by the PCR method. M: DL2000 maker, Lanes 3, 5, 7, 10, 14 and 21: The

TMV coat protein gene in *N. tabacum* K326 at 3, 5, 7, 10, 14 and 21 days post infection TMV.