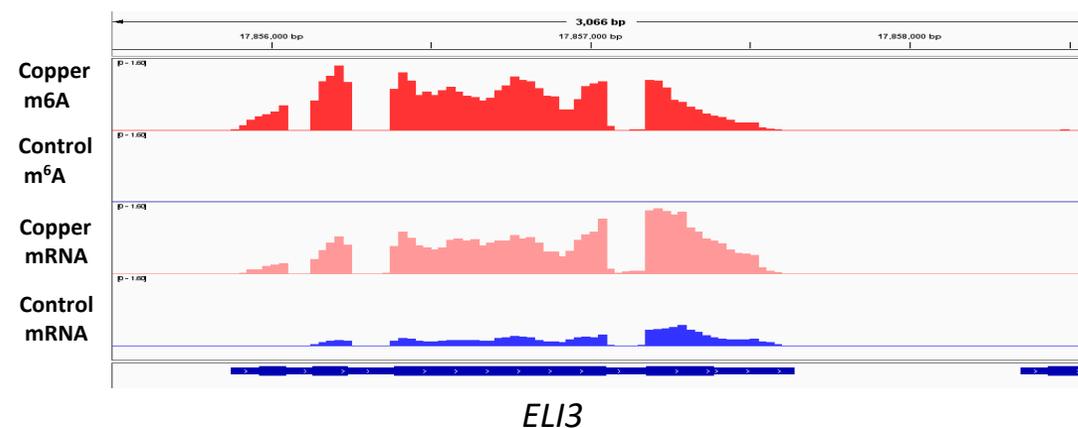
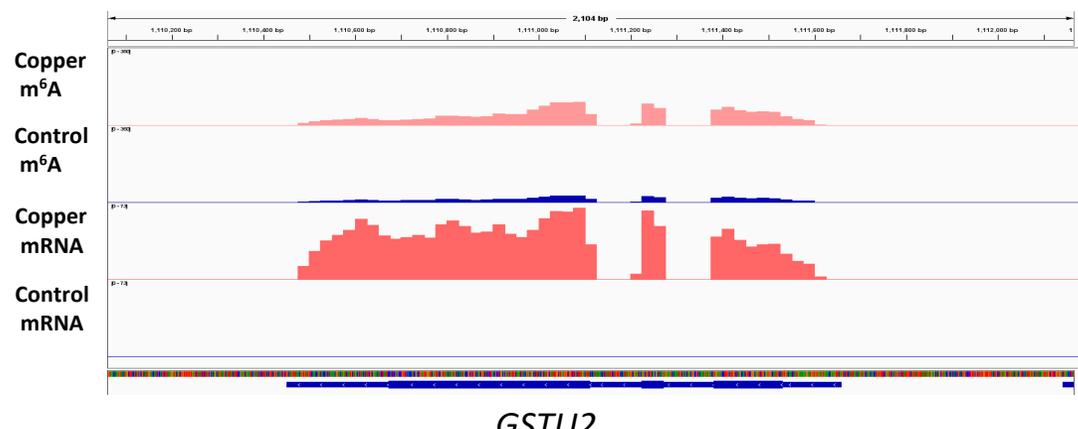


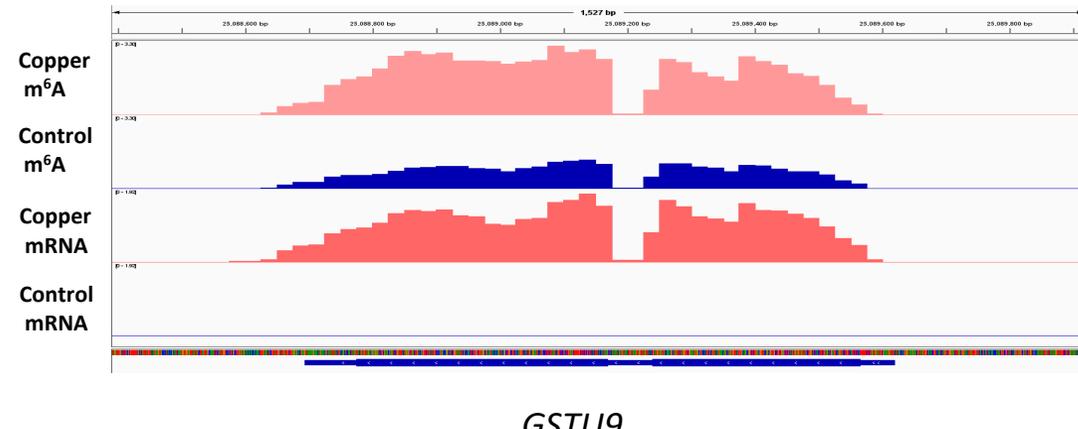
CRK11



ELI3

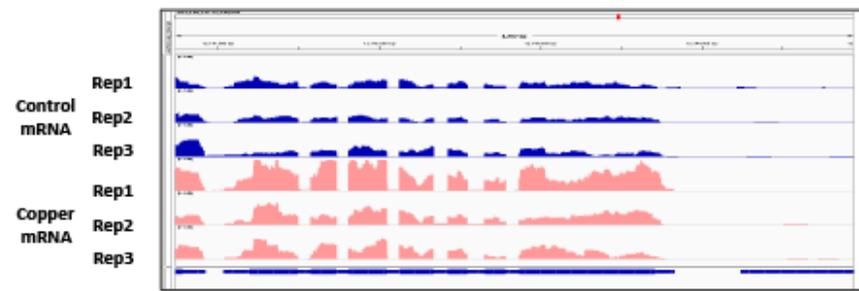


GSTU2

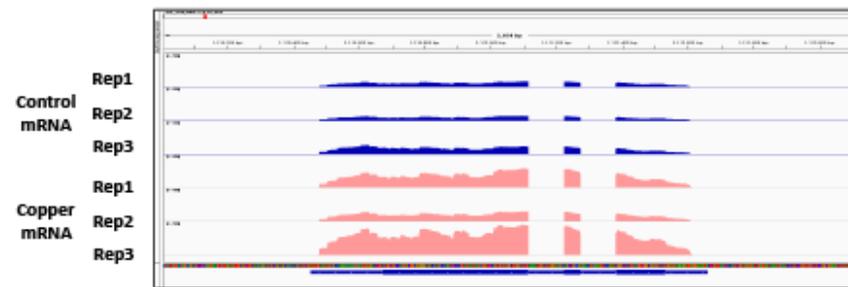


GSTU9

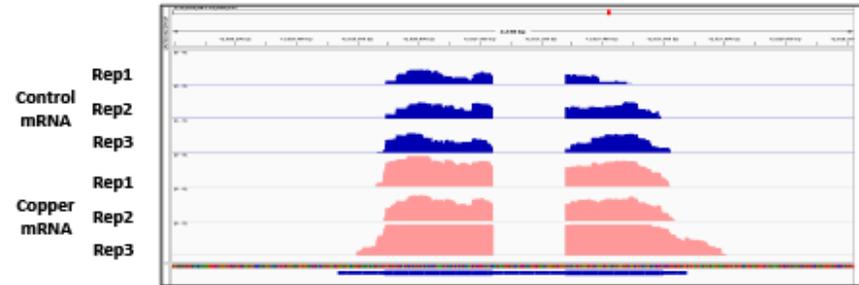
Supplemental Figure S1: Example transcripts displaying copper-induced oxidative stress-specific m⁶A increases that correspond with increases in overall mRNA abundance of those transcripts. The transcripts shown here are *CRK11*, *ELI3*, *GSTU2*, and *GSTU9* as indicated.



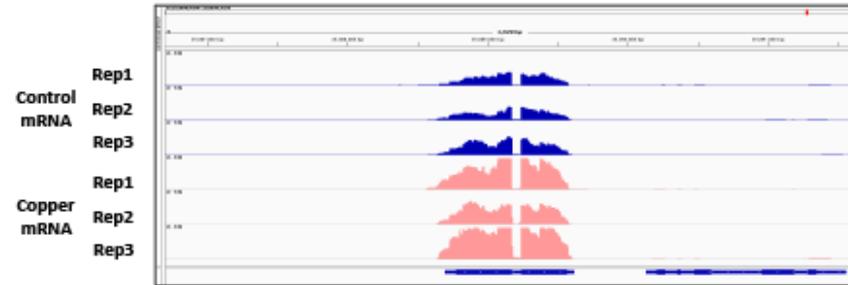
CRK11



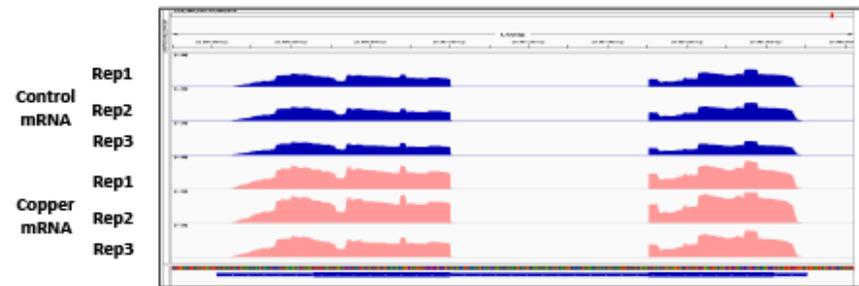
GSTU2



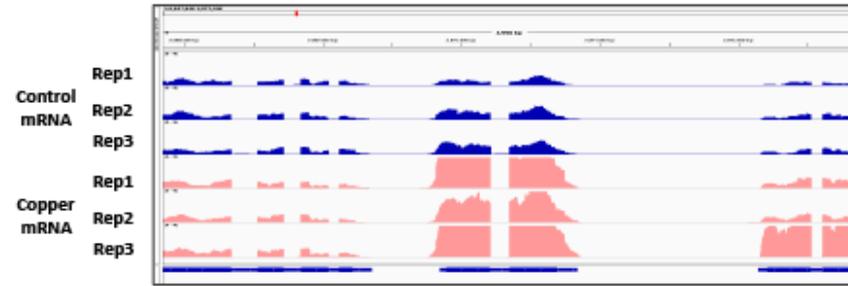
GSTU4



GSTU9



GSTU19



GSTU24

Supplemental Figure S2: Example transcripts displaying copper-induced oxidative stress-specific increases in overall mRNA abundance of those transcripts. The transcripts shown here are *CRK11*, *GSTU2*, *GSTU4*, *GSTU9*, *GSTU19*, and *GSTU24* as indicated.