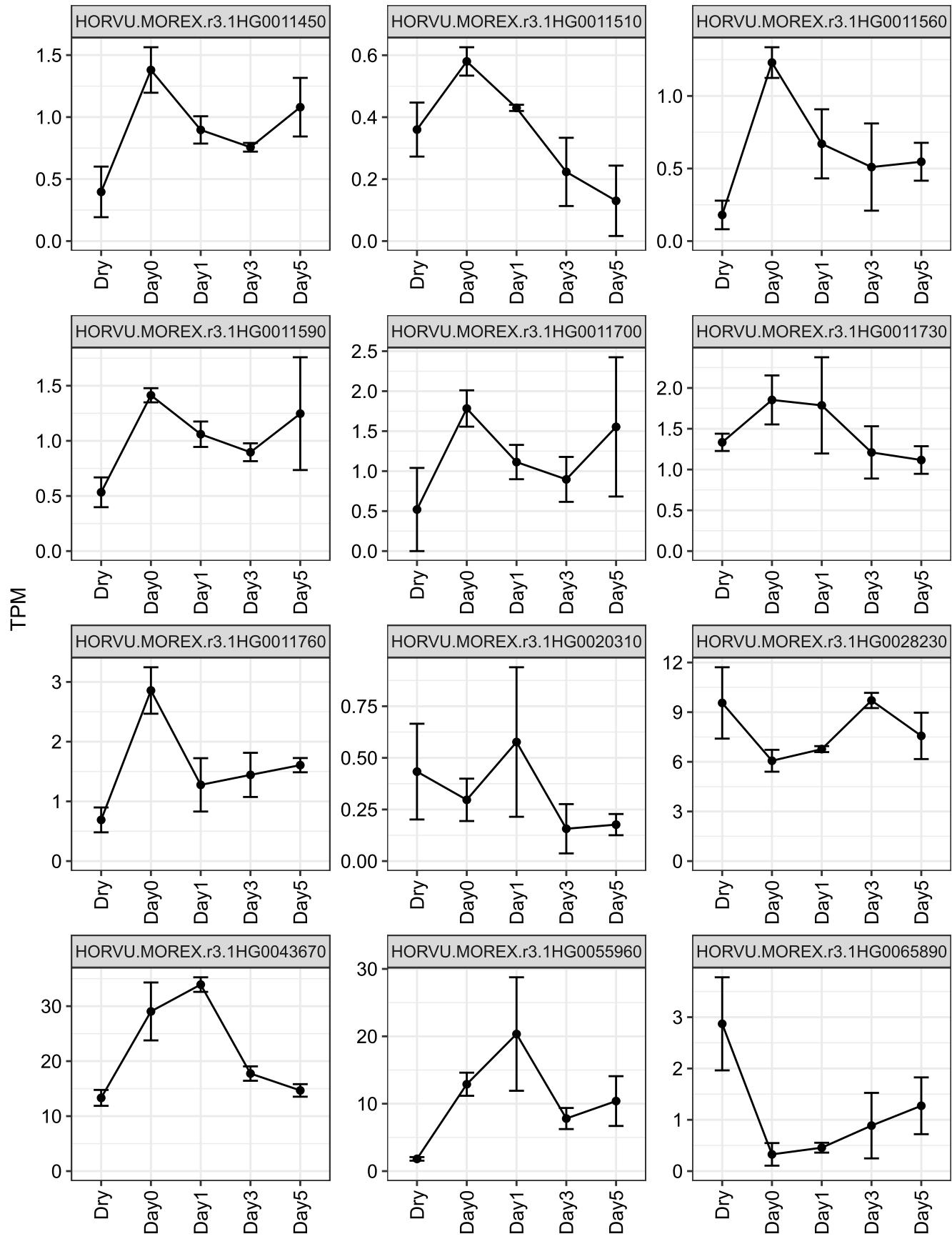


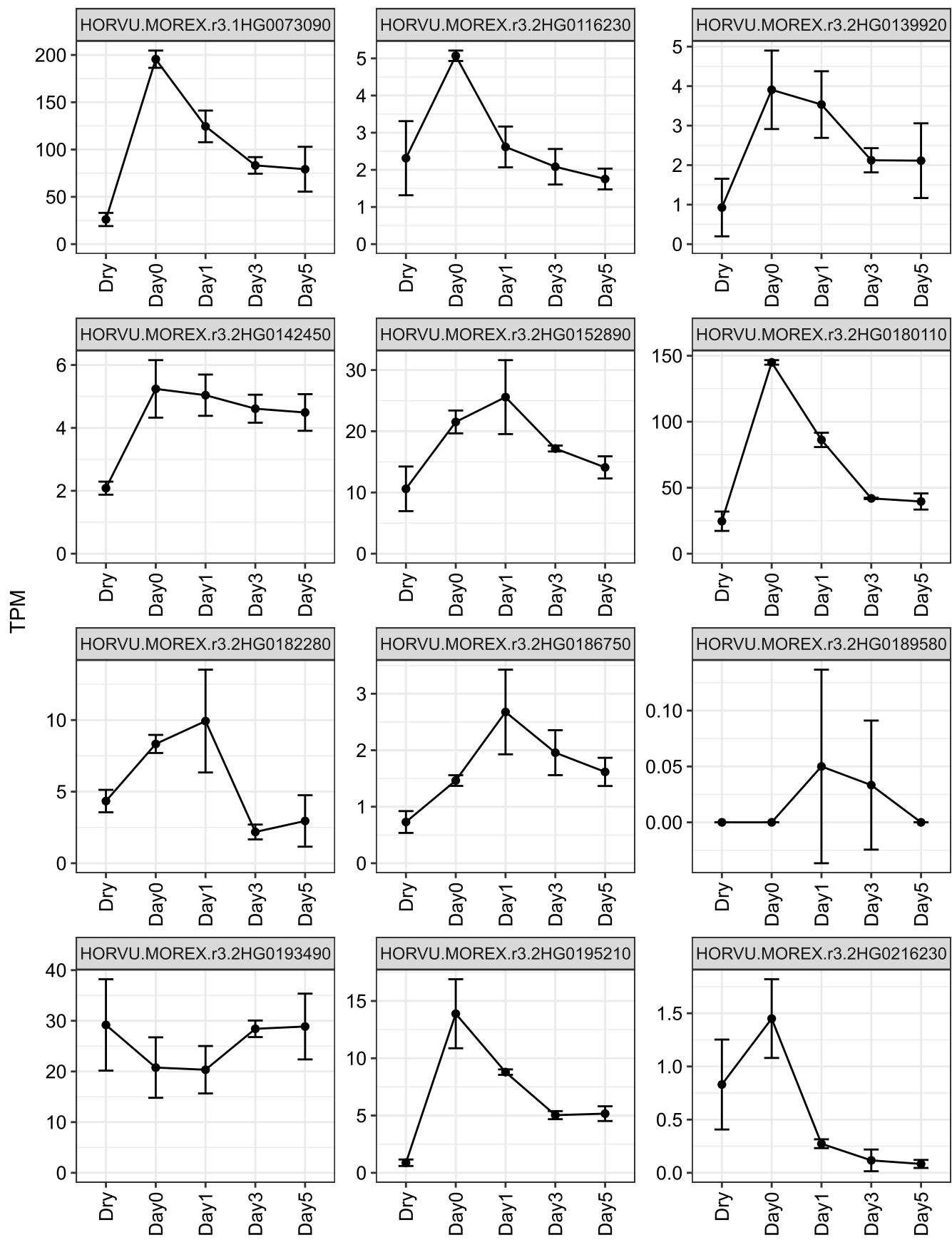
Figure S6

Expression across the five malting stages of the 64 genes with CleaveLand4 category 0 predicted slice sites. Expression is represented as the mean transcripts per million (TPM) +/- standard deviation, n = 3. The normalized read count data are publicly available at NCBI's Gene Expression Omnibus through GEO Series accession number GSE 295574 (<https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE295574>).

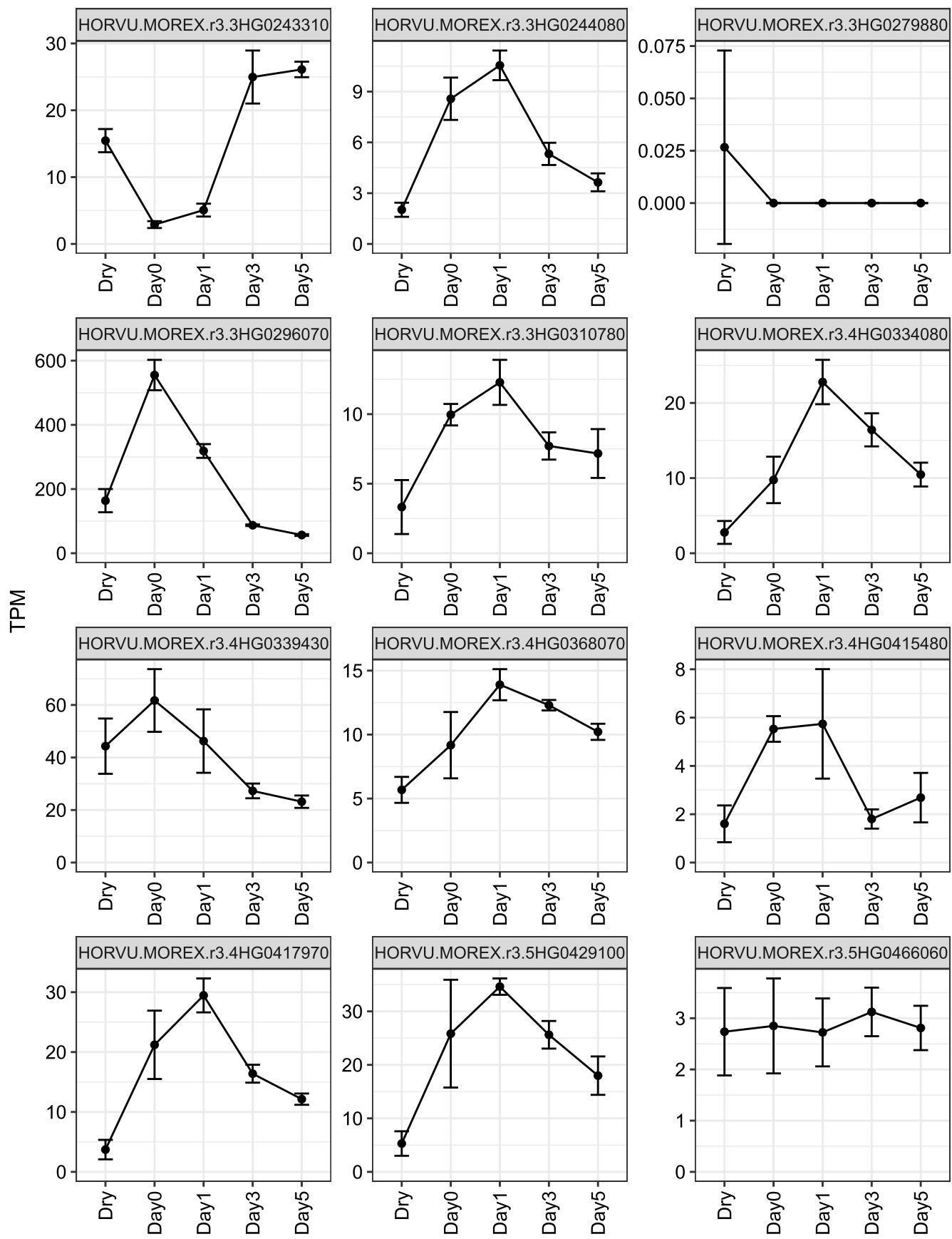
expression of CleaveLand4 category 0 predicted targets of miRNA-mediated slicing
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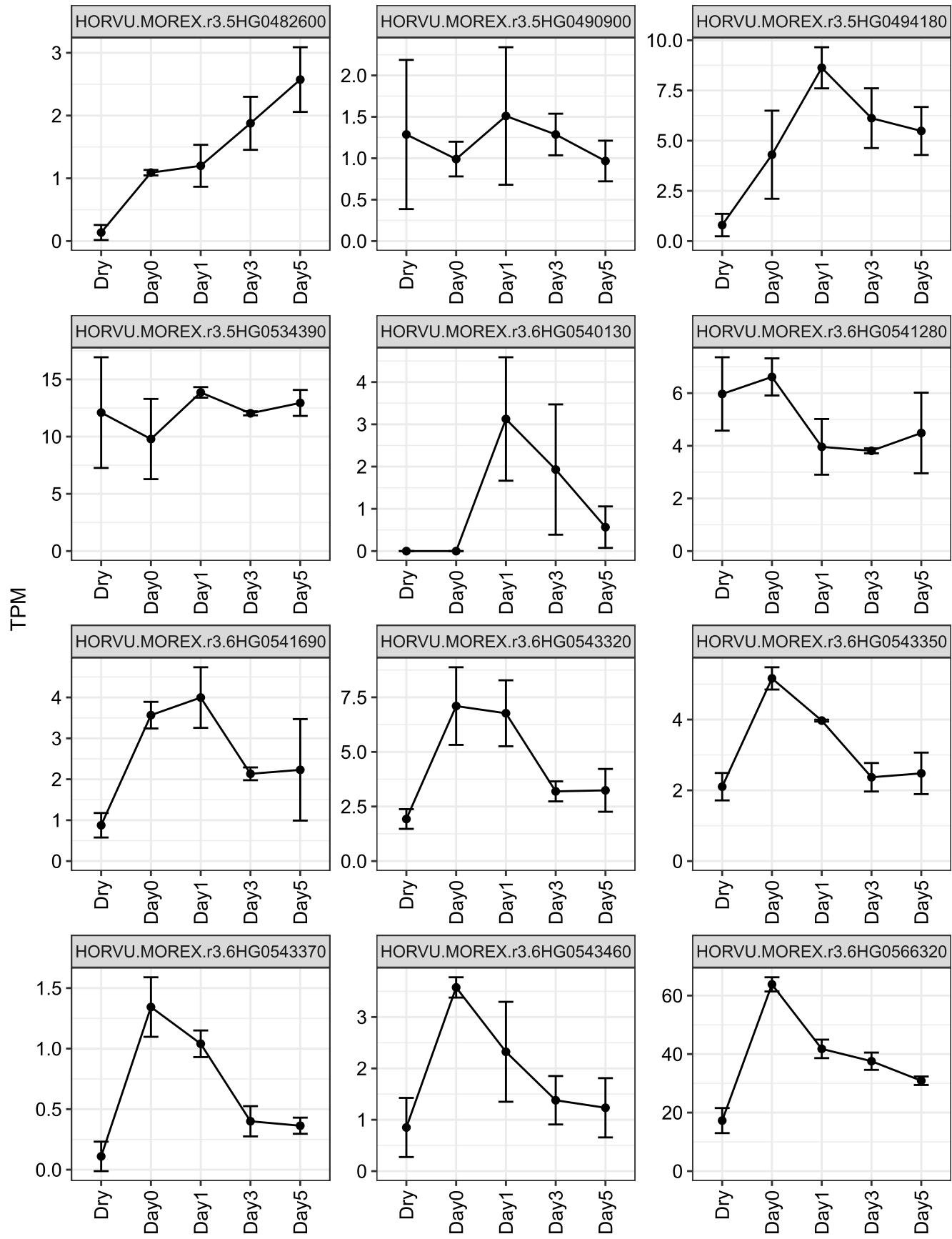
expression of CleaveLand4 category 0 predicted targets of miRNA-mediated slicing
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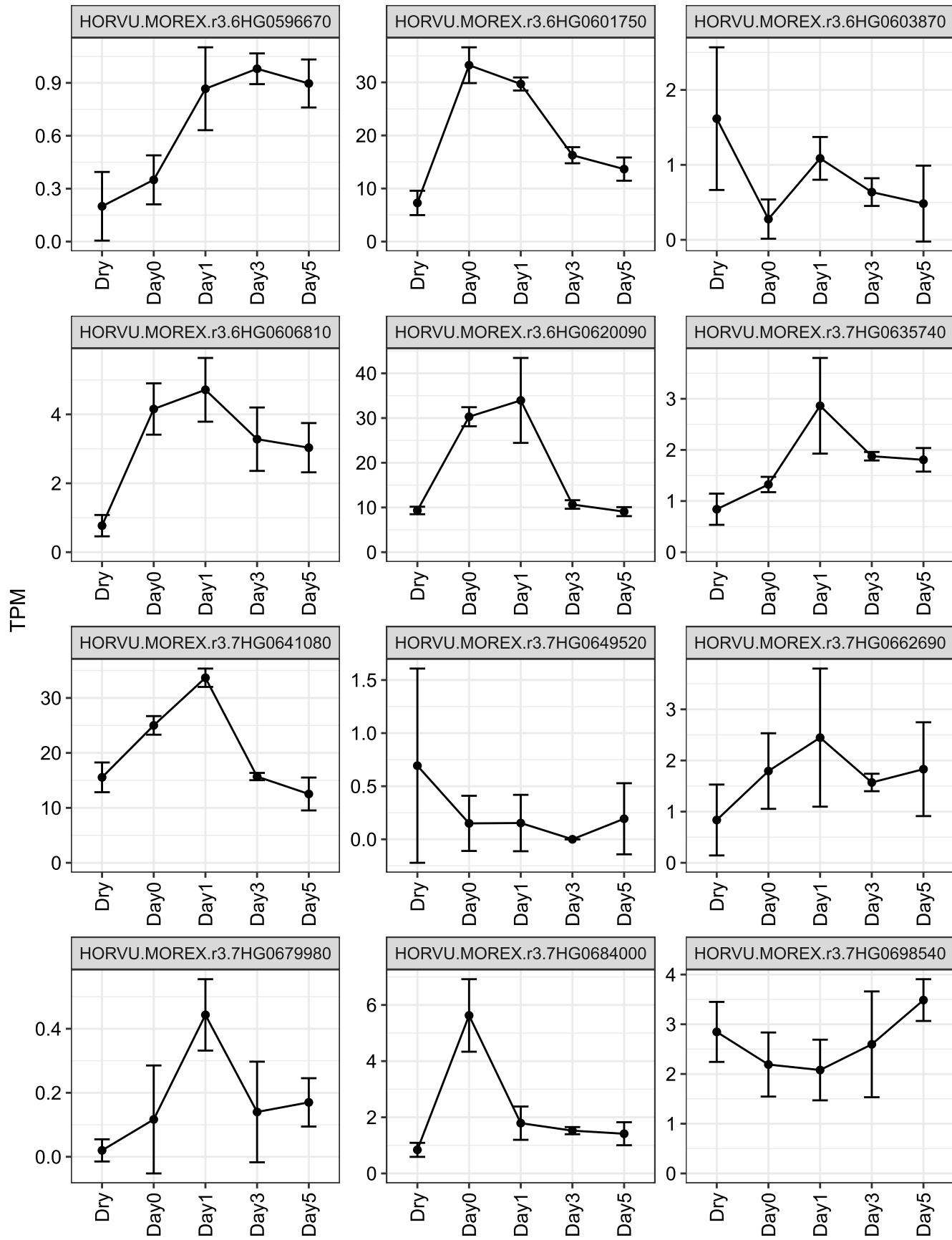
expression of CleaveLand4 category 0 predicted targets of miRNA-mediated slicing
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