

Supplementary Material

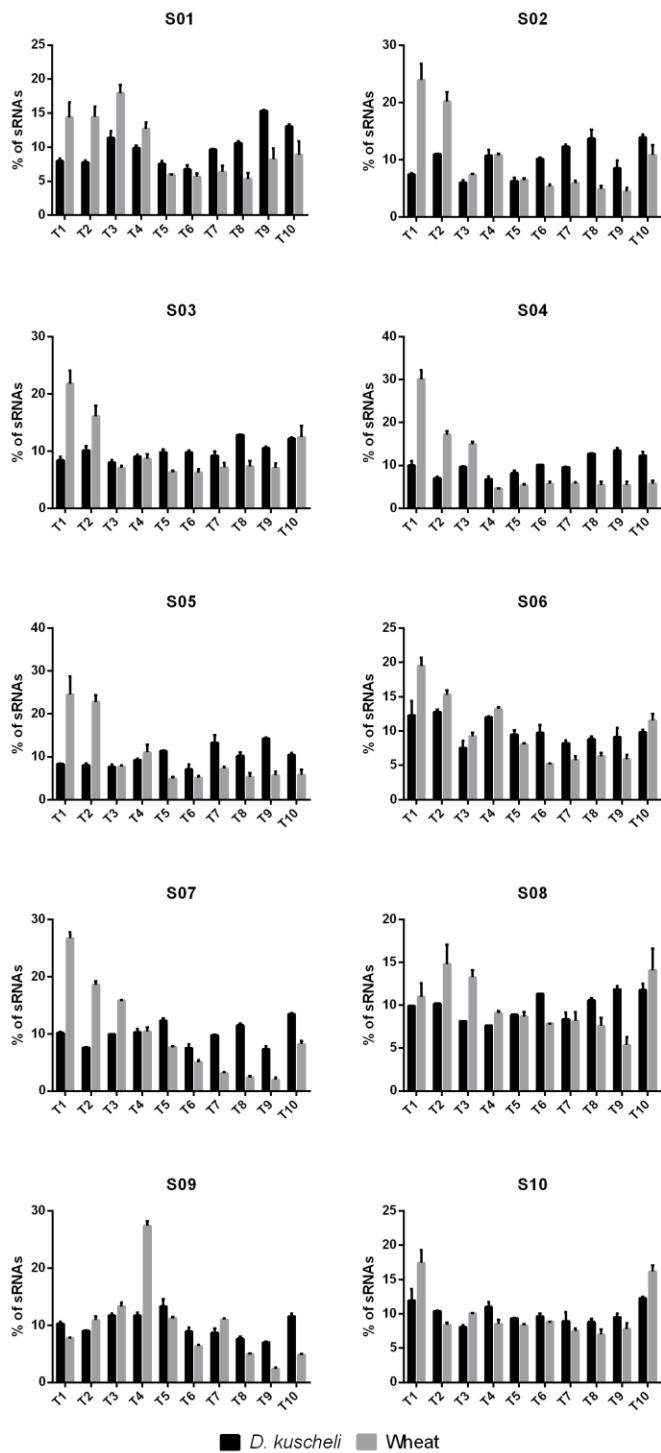
Mal de Río Cuarto virus infection triggers the production of distinctive viral-derived siRNAs profiles in wheat and its planthopper vector

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1 **Supplementary Figure S1**



Supplementary Figure S1. Distribution of vsiRNAs from *D. kuscheli* and wheat within each MRCV genome segment (S1 to S10). All MRCV genome segments were divided in ten equal fractions (T1 to T10). Percentage bar graphs represent the mean number of total vsiRNAs that mapped to the ten fractions within each segment.