LEGENDS FOR SUPPLEMMENTARY TABLES

Table S1. List of the 607 NS1-bound RNAs (genes), summarized from the fragments of peaks enriched compared to the controls. For each gene, the sum of fold changes, average fold change, and the number of enriched peaks are listed by the different comparisons between experimental vs controls. WN1 & WN2 are duplicate experimental (WSN infected cells and immunoprecipitation performed with anti-NS1 antibody) and duplicate controls: FN1, FN2 (cells non-infected and immunoprecipitation performed with anti-NS1 antibody), WG1 and WG2 (WSN infected cells and immunoprecipitation performed with IgG). Table S2. Transcriptome analysis of A549 cells mock-infected or infected for 6h with WSN at MOI 5. Comparative ratios and fold changes (FC) are shown of RNAs genome-wide or of RNAs that are up-regulated and/or are bound by NS1 (marked in yellow). Table S3. Transcriptome analysis of A549 cells mock-infected or infected for 10 h with WSN or WSNΔNS1 at MOI 5. Comparative ratios and fold changes are shown.