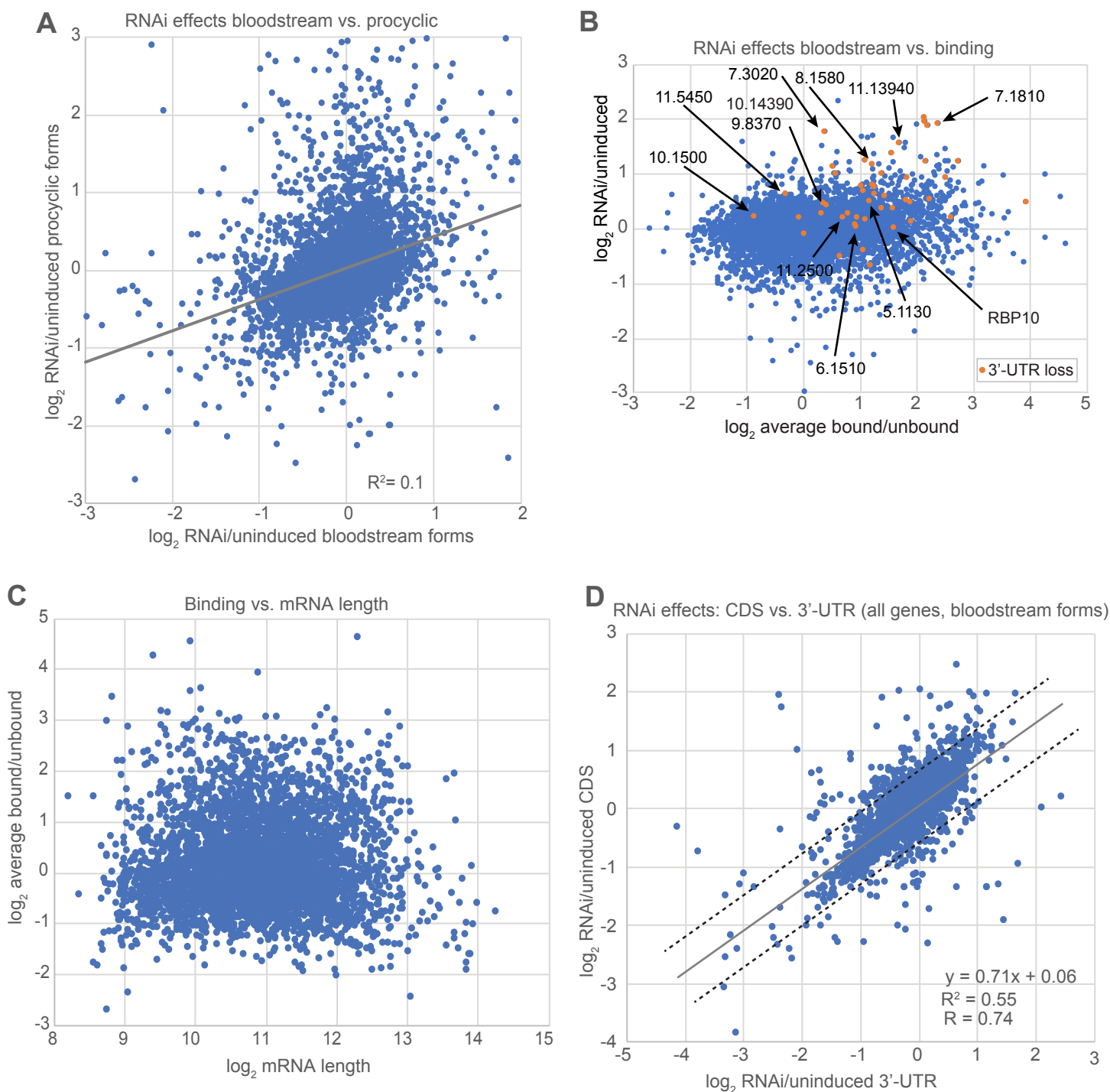




Supplementary Figure S3. DRBD18 depletion and binding analysis

A. The effect of *DRBD18* RNAi in procyclic forms (y-axis, published values) was plotted against the effect of *DRBD18* RNAi in bloodstream forms (x-axis). Results are for coding regions.

B. Lack of correlation between effects of *DRBD18* depletion in bloodstream forms (y-axis) and *DRBD18* binding (x-axis). For repeated genes only one representative is shown. Results for mRNAs that showed loss of 3'-UTR reads are in orange and those studied in detail are indicated.

C. Lack of correlation between mRNA length (x-axis) and RNA binding by *DRBD18* (y-axis).

D. Correlation between the effect of *DRBD18* depletion in bloodstream forms on coding sequences (y-axis) vs 3'-UTRs (x-axis). Dotted lines indicate $\pm 1.5x$.