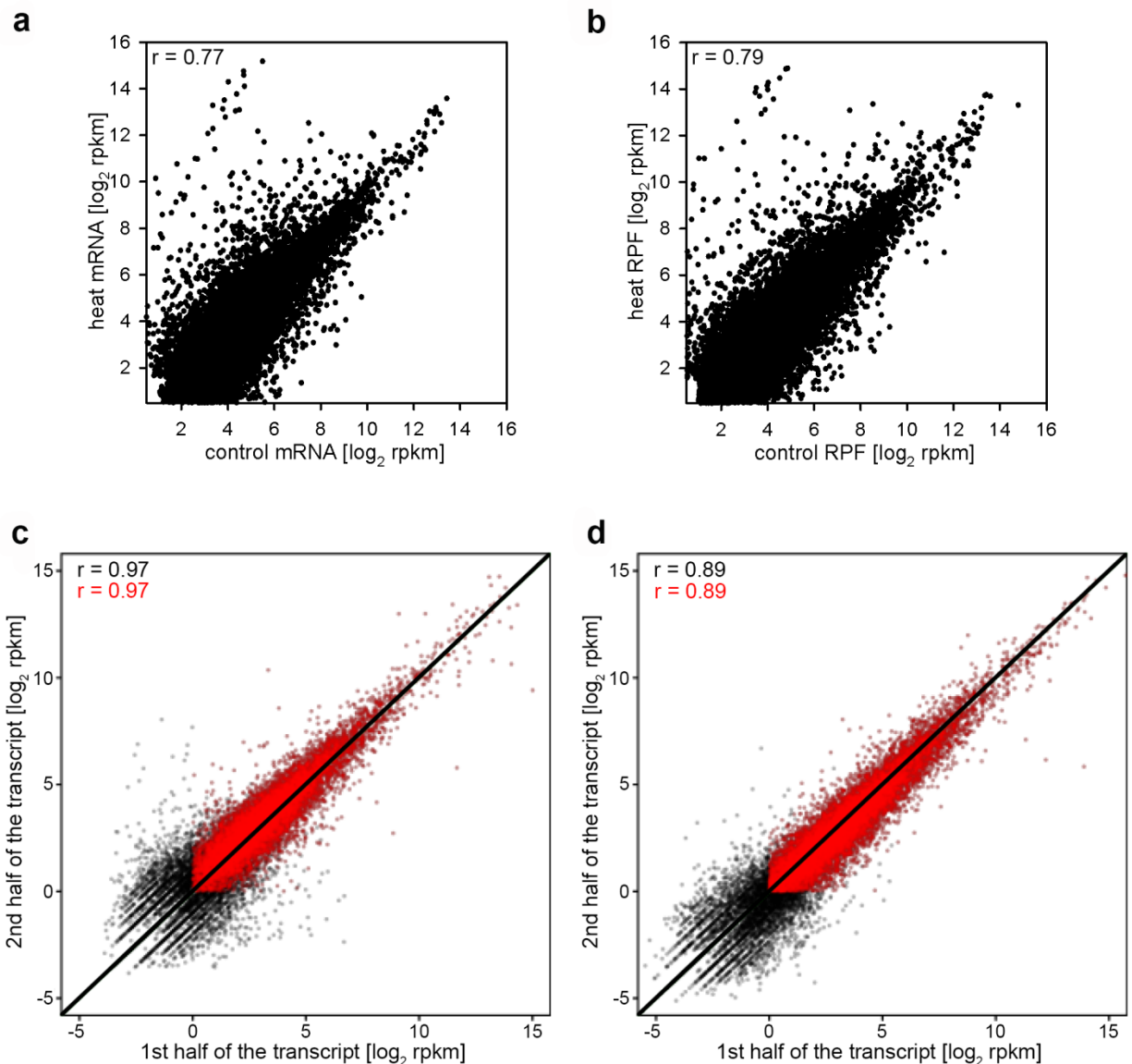




Additional File 2. Comparison of transcriptional and translational features of all protein-coding genes between control plants and those exposed to thermal stress. **(a, b)** Correlation of the normalized RPF (a) and randomly fragmented mRNA reads (b) for each gene from control plants or plants subjected to heat stress. **(c, d)** Symmetric distribution of the mRNA (black) and RPF (red) reads between the first and second halves of the CDS of each transcript for control (c) and heat stress (d). r , Pearson correlation coefficients.