

Description of Additional Supplementary Files

Supplementary Data 1 (related to Fig. 2): Analysis of sequential RIP-sequencing (seqRIP-seq) data.

Shown are the features reproducibly enriched upon sequential immunoprecipitations of Mmi1 and Mei2 from biological duplicates.

Supplementary Data 2 (related to Fig. 3): Analysis of RNA-sequencing data.

Shown are the up-regulated and down-regulated genes in *mot2Δ*, *mamRNAΔ* and *mot2Δ mamRNAΔ* cells relative to the *wt* strain, and the associated *p*-values.