

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: Raw mass spectrometry data for tandem affinity purified samples. Release 8.0 of T. brucei TREU927 database at <http://tritrypdb.org> was used for protein identification. Typical contaminations found in purified trypanosomal mitochondrial complexes have been removed. Raw data are provided in a separate sheet. Related to Figure 2.

File Name: Supplementary Data 2

Description: Raw mass spectrometry data for BioID experiments. Release 8.0 of T. brucei TREU927 database at <http://tritrypdb.org> was used for protein identification. Typical contaminations found in purified trypanosomal mitochondrial complexes have been removed. Raw data are provided in a separate sheet. Related to Figure 2.

File Name: Supplementary Data 3

Description: Sanger sequencing (96 well plate) of pre-edited RPS12 mRNA termini in KPAF4 RNAi cell line. RNA was isolated after 72 hours of RNAi induction. Thirteen representative sequences are listed.

File Name: Supplementary Data 4

Description: Mapping statistics for tails sequencing and CLAP experiments.

File Name: Supplementary Data 5

Description: Major components of KPAF4-WT and KPAF4-Mut complexes. Release 8.0 of T. brucei TREU927 database at <http://tritrypdb.org> was used for protein identification. Typical contaminations found in purified trypanosomal mitochondrial complexes have been removed. Raw data are provided in a separate sheet. Related to Figure 6

File Name: Supplementary Data 6

Description: DNA Oligonucleotides used in this study. Abbreviations: p, pre-edited; e, edited; fw, forward; rv, reverse.