Supplementary Figure

Fig. S1. ClustalW-generated tree showing sequence distances between putative PPR proteins from *T. brucei* and *L. major*. The branch-lengths are drawn to scale and bootstrap support (from 1000 trials) is indicated at each node. To the right of the tree, each protein is presented schematically (approximately to scale) with predicted PPR motifs indicated by arrows. Predotar or Mitoprot strongly predict a mitochondrial localization for almost all of these proteins. The exceptions are the clusters related to Tb927.6.4190, Tb927.8.6040, Tb11.01.7210 and Tb927.2.4460. The 8 putative PPR proteins of *T. brucei*, termed TbPPR1-8, that were selected for experimental studies are indicated by numbers.