**The evolutionary fate of *rpl32* and *rps16* losses in the *Euphorbia schimperi* ( Euphorbiaceae) plastome**

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**Figure S1**

Chart

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**Figure S2**

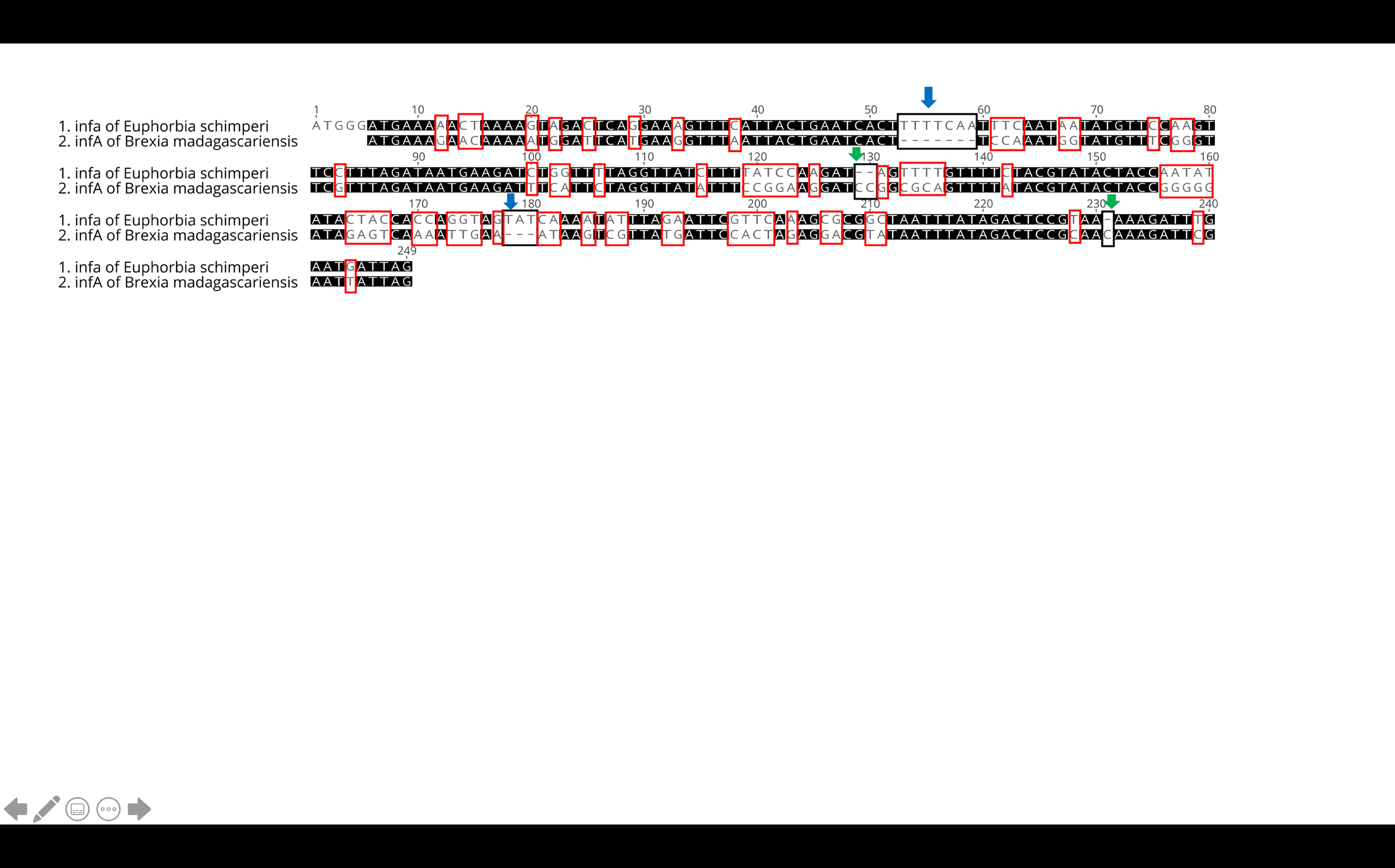
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**Figure S3**



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TP

**Figure 4**: Multiple alignments of nuclear **RPL32** of ***E.schimperi*** and **plastid** **RPL32** of other Euphorbiaceae (*E. marginata, Jatropha curcas, Hevea brasiliensis, Manihot esculenta, Ricinus communis, Vernicia fordii), Arabidopsis thaliana* and *Nicotiana tabacum.* Blue boxes indicate plastid transit peptide (TP) predicted using TargetP. Red box indicates a conserved domain of RPL32.

**ribosomal protein L32**