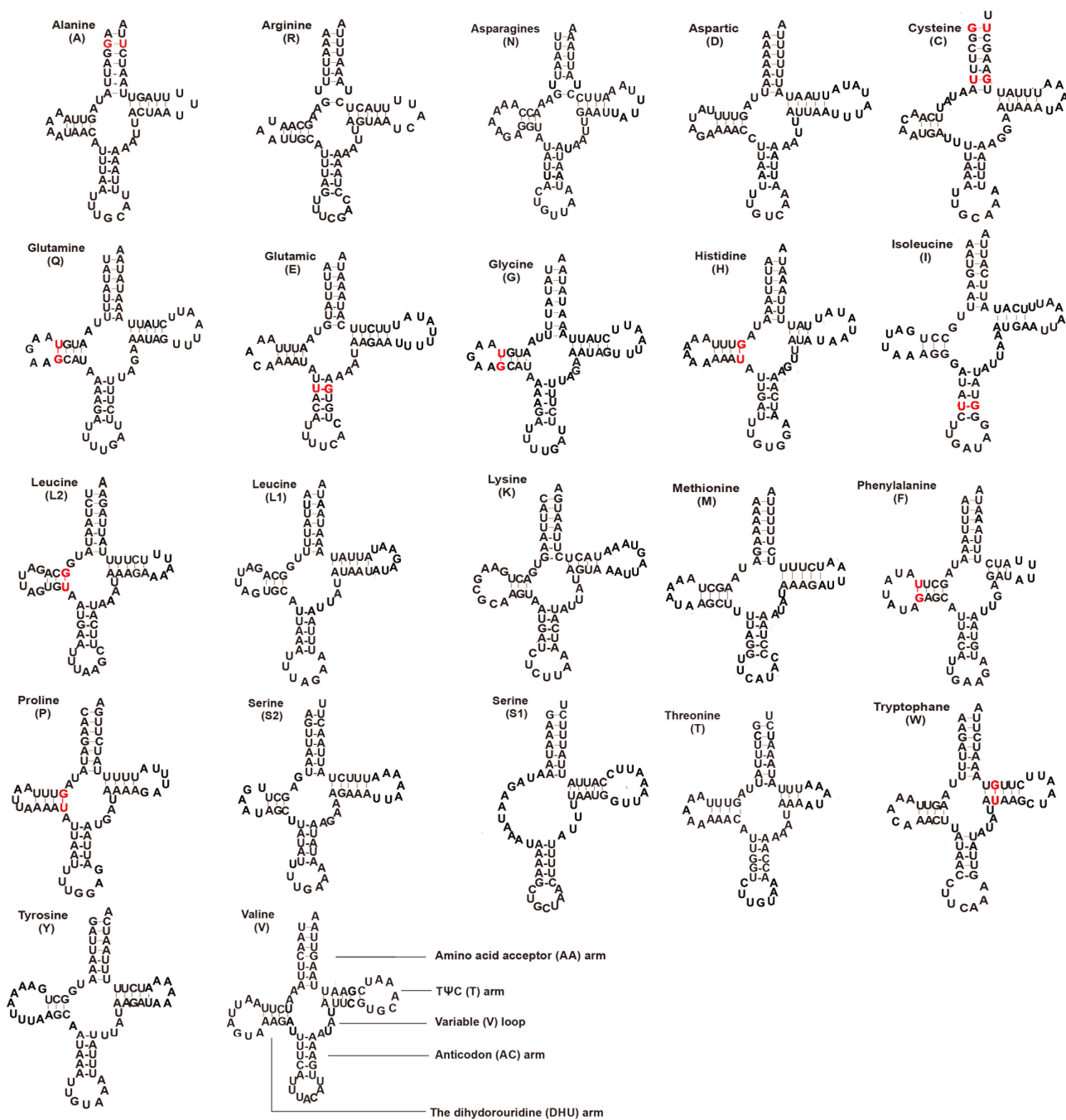


Figure S1. Predicted secondary structures for 22 tRNAs in *Penthetria simplicioides*

(A), and *Plecia hardyi* (B) mtgenomes. Nucleotides highlighted in red indicate mismatch.

A



B

