

Supplementary caption:

Figure S1. The mitogenome maps of *Agrilus sichuanus*, *Coraebus cloueti*, *Coraebus diminutus*, *Meliboeus sinae*, and *Sambus femoralis*.

Figure S2. The secondary cloverleaf structure for the tRNAs of *Agrilus sichuanus*.

Figure S3. The secondary cloverleaf structure for the tRNAs of *Coraebus cloueti*.

Figure S4. The secondary cloverleaf structure for the tRNAs of *Coraebus diminutus*.

Figure S5. The secondary cloverleaf structure for the tRNAs of *Meliboeus sinae*.

Figure S6. The secondary cloverleaf structure for the tRNAs of *Sambus femoralis*.

Figure S7. Heterogeneous sequence divergence within datasets 13 PCGs and 2 rRNAs of Buprestidae species.

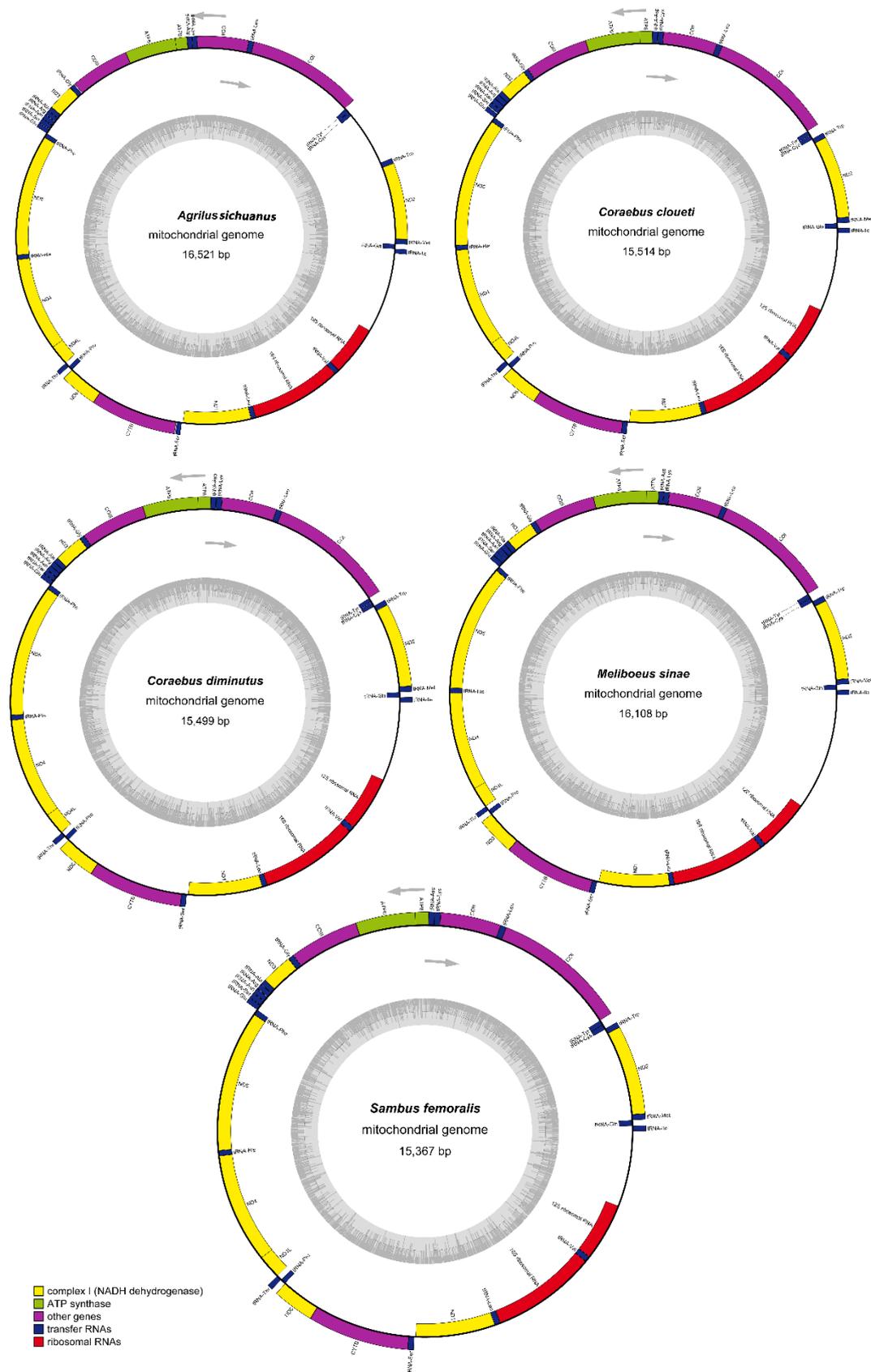


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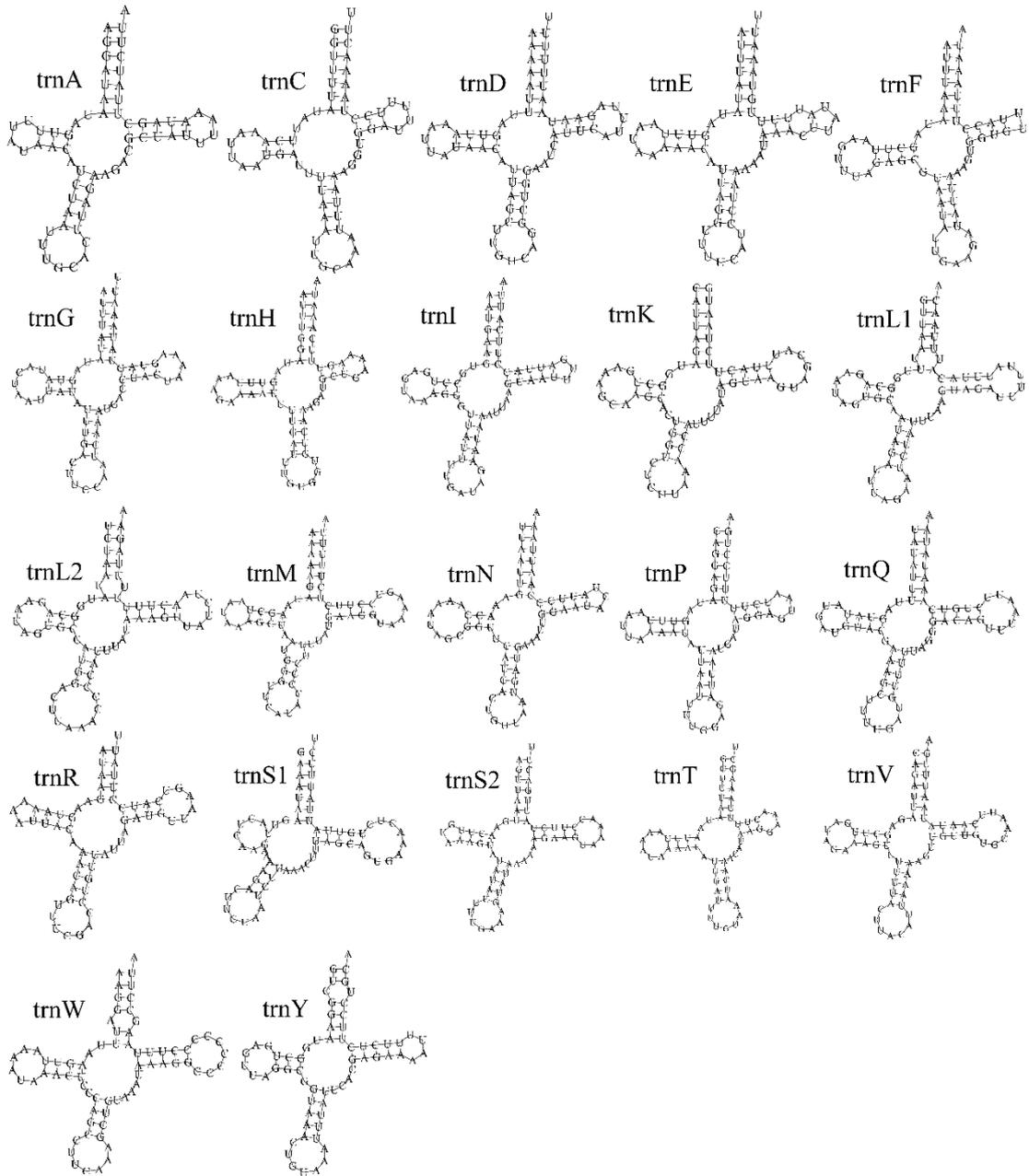


Figure S2. The secondary cloverleaf structure for the tRNAs of *Agrilus sichuanus*.

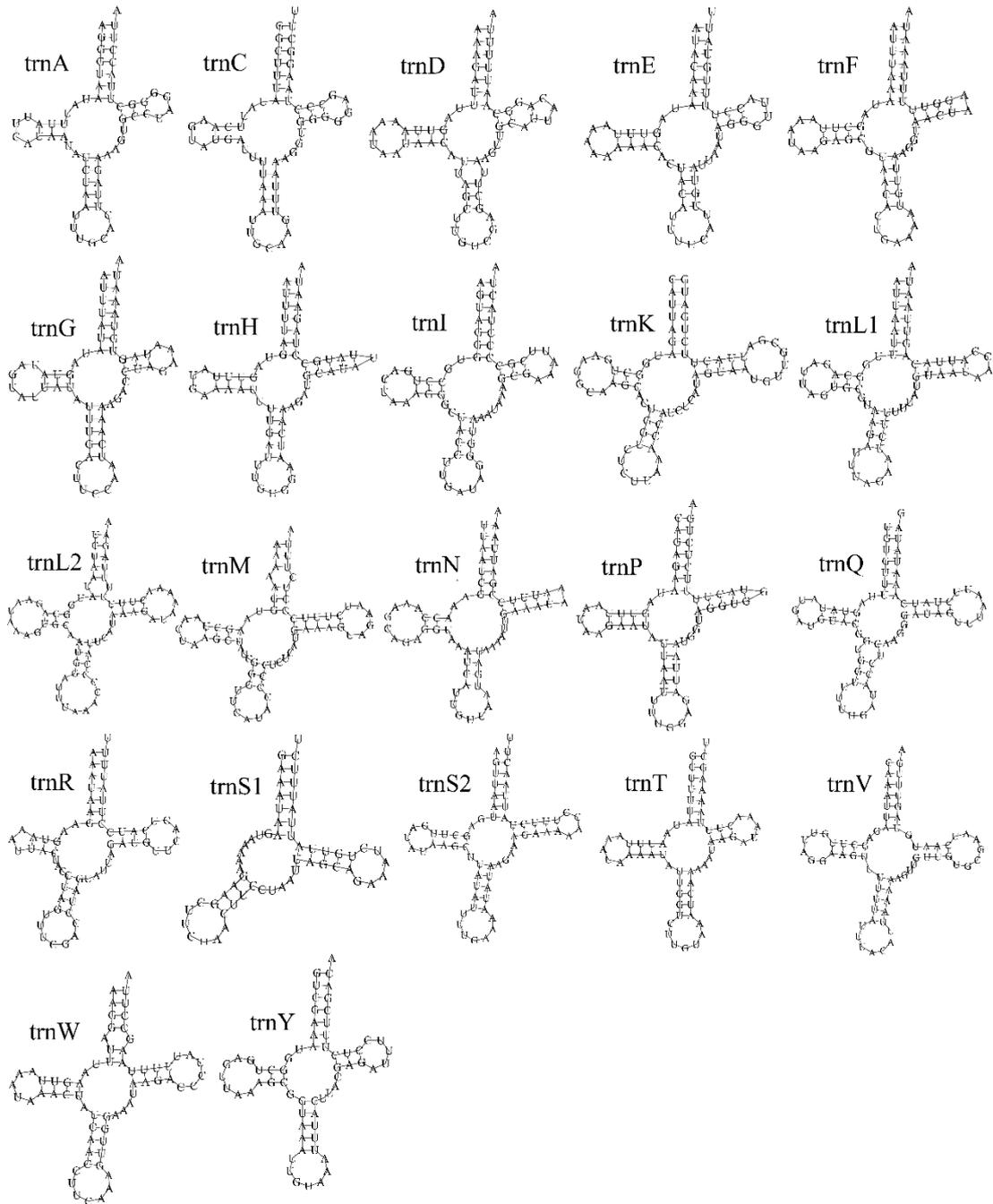


Figure S3. The secondary cloverleaf structure for the tRNAs of *Coraebus cloueti*.

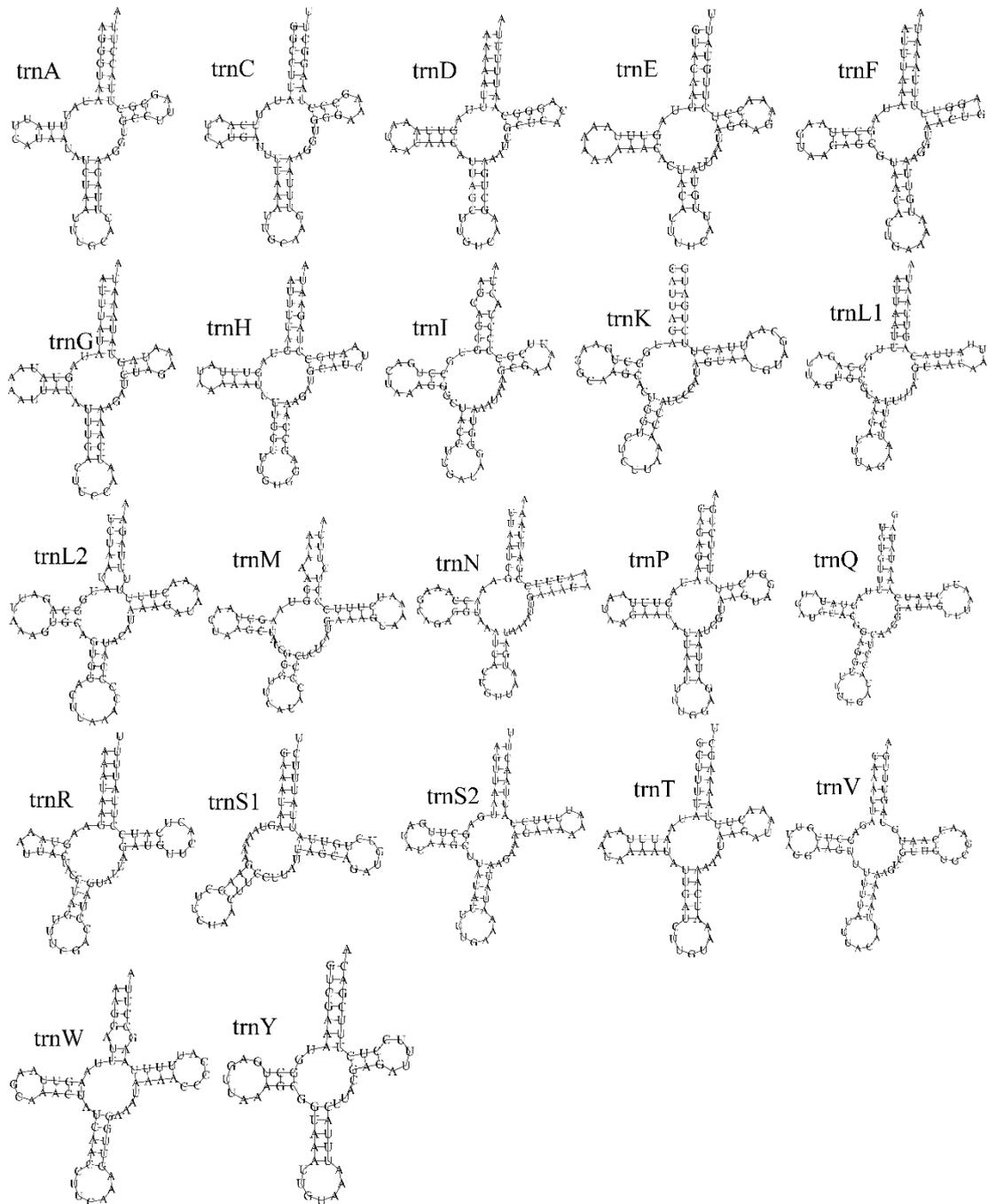


Figure S4. The secondary cloverleaf structure for the tRNAs of *Coraebus diminutus*.

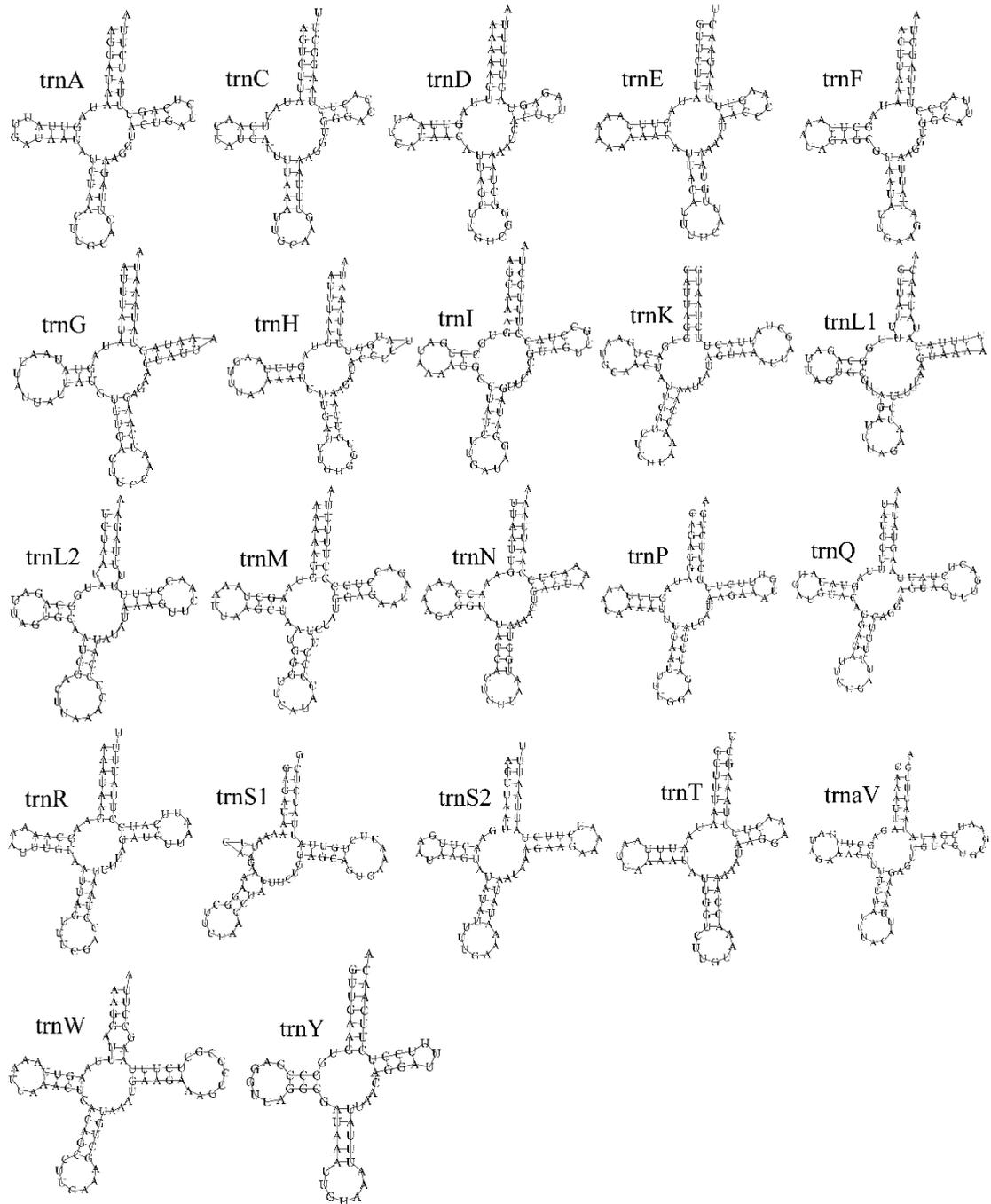


Figure S5. The secondary cloverleaf structure for the tRNAs of *Meliboeus siniae*.

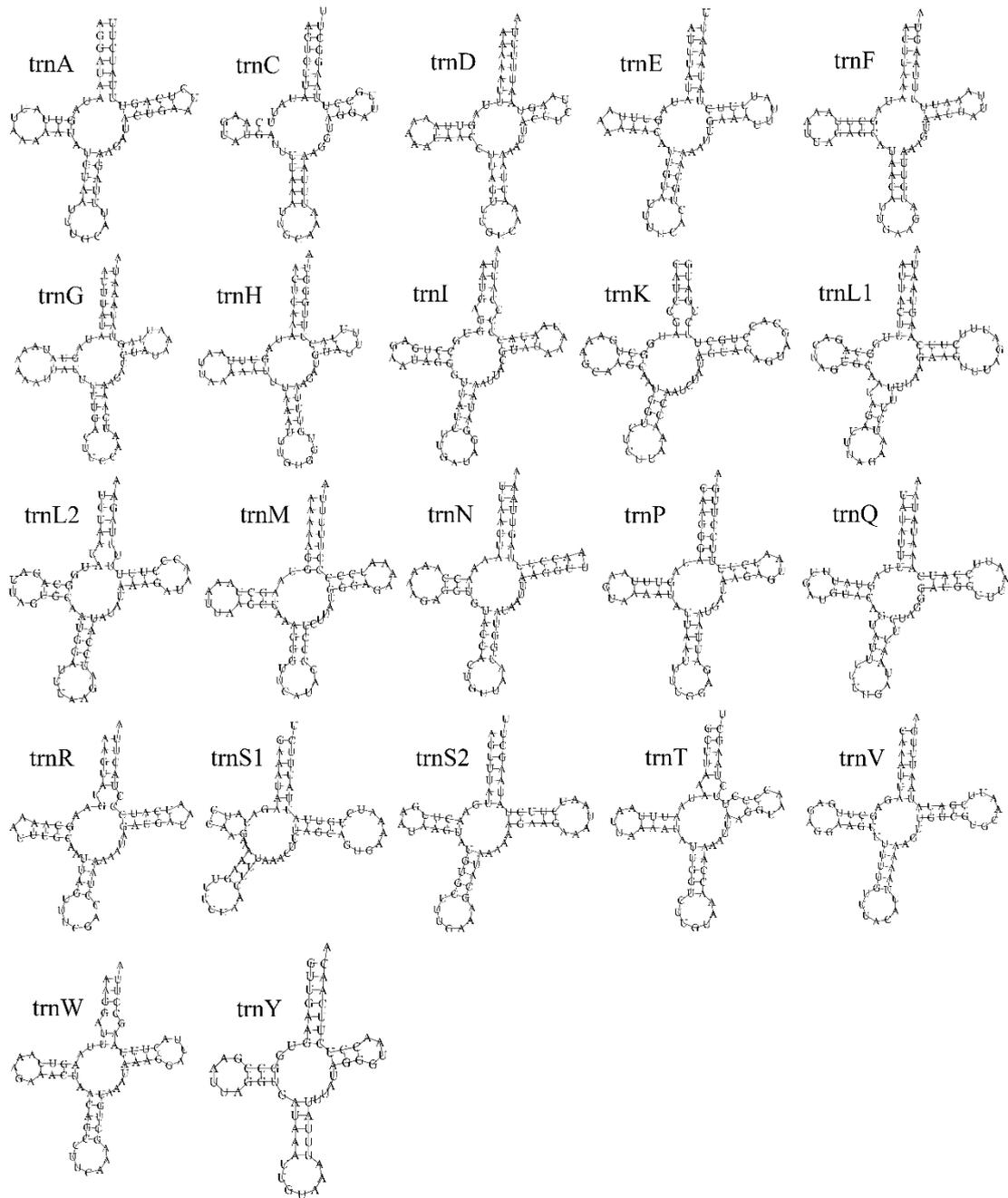


Figure S6. The secondary cloverleaf structure for the tRNAs of *Sambus femoralis*.

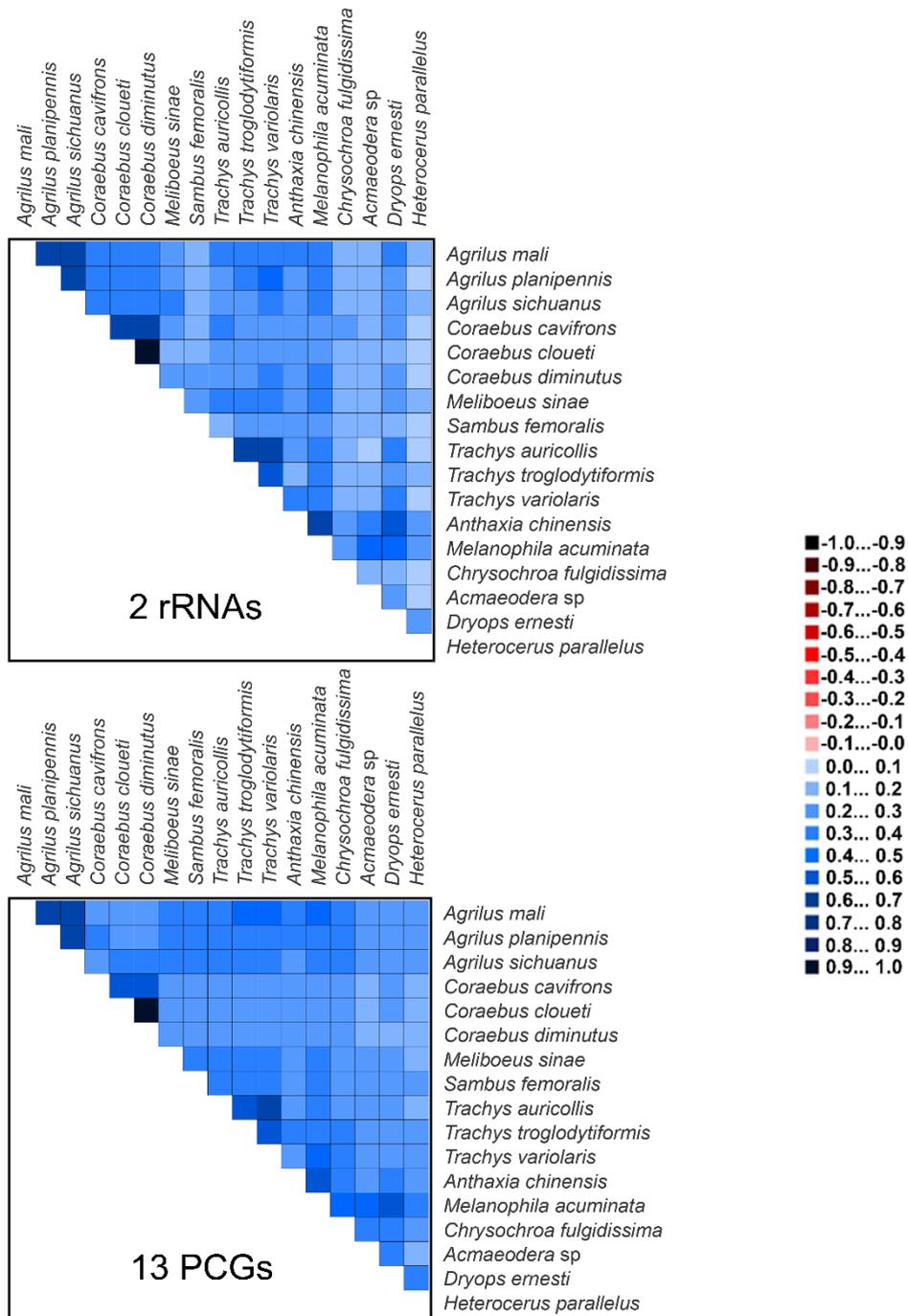


Figure S7. Heterogeneous sequence divergence within datasets 13 PCGs and 2 rRNAs of Buprestidae species