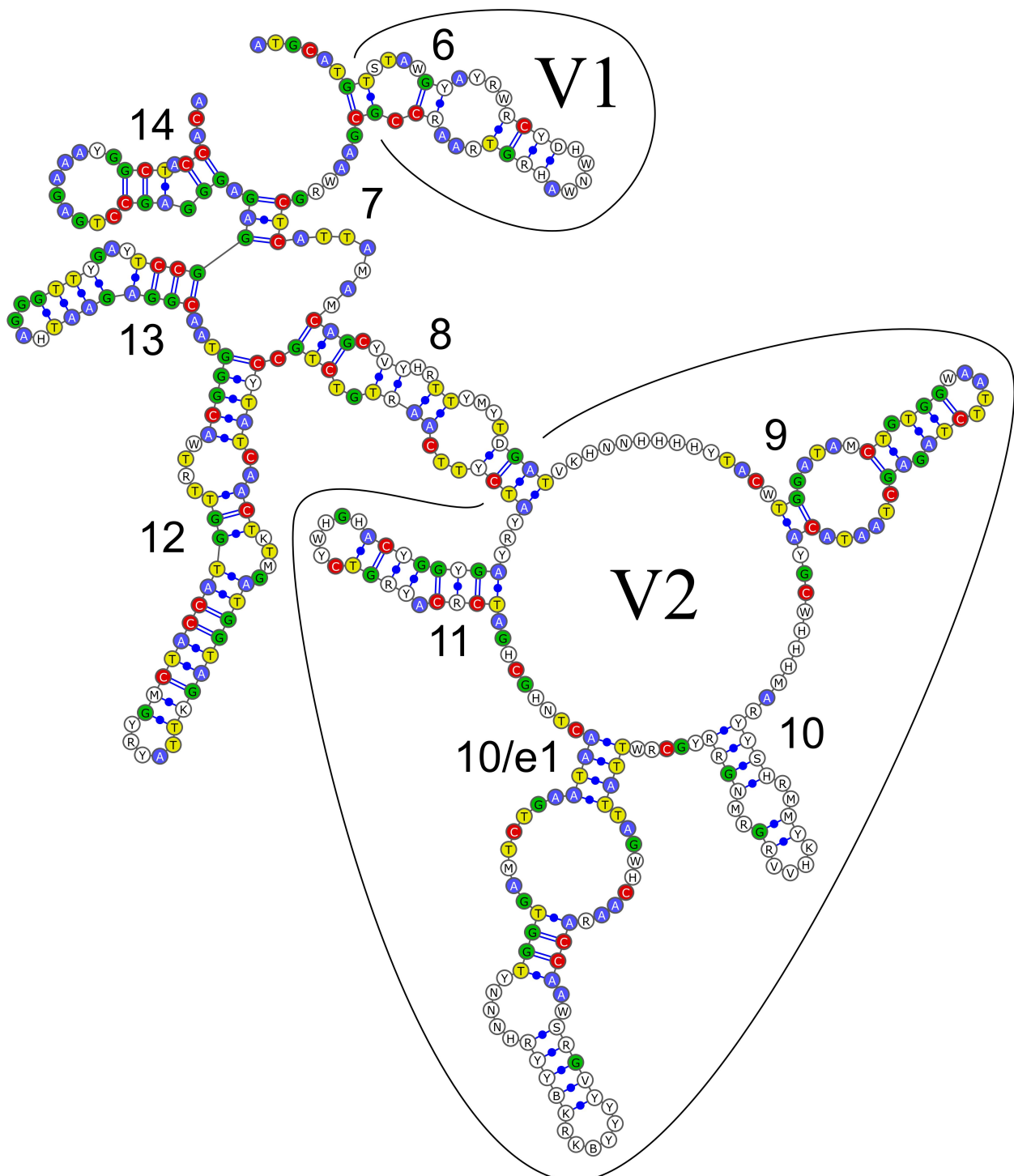


**Supplementary file for the article:**

Holovachov O, Haenel Q, Bourlat SJ, Jondelius U. Taxonomy assignment approach determines the efficiency of identification of OTUs in marine nematodes. *Royal Society Open Science*.

**Supplementary Figure 1.** Barcoding region used in this study – generalized secondary structure model of the 90% consensus sequence (excluding rare insertions) based on combined data from secondary-structure-based alignment of reference sequences and mothur-based alignment of OTU sequences (444 sequences in total). Helices (6-10, 10/e1, 11-14) are according to [63]. Variable regions V1 and V2 are labelled according to [64].



## References

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