Supplementary material

How does marker choice affect your diet analysis: comparing genetic markers and digestion levels for diet metabarcoding of tropical-reef piscivores

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Fig. S1. Schematic overview of the assembly pipeline, starting from raw forward (F) and reverse (R) reads. A detailed description is given in the ‘Materials and methods’ section. Numbers indicate the respective amount of sequences or blast results.
Fig. S2. Species richness of four low- and four high-digestion samples, compared across the four markers before and after blast-quality filtering.
Fig. S3. Efficiency of host amplification with host-blocking components for COI. Proportion of the number of host sequences over total number of sequences per sample is contrasted against the host species and digestion stage. Error bars represent standard errors.