

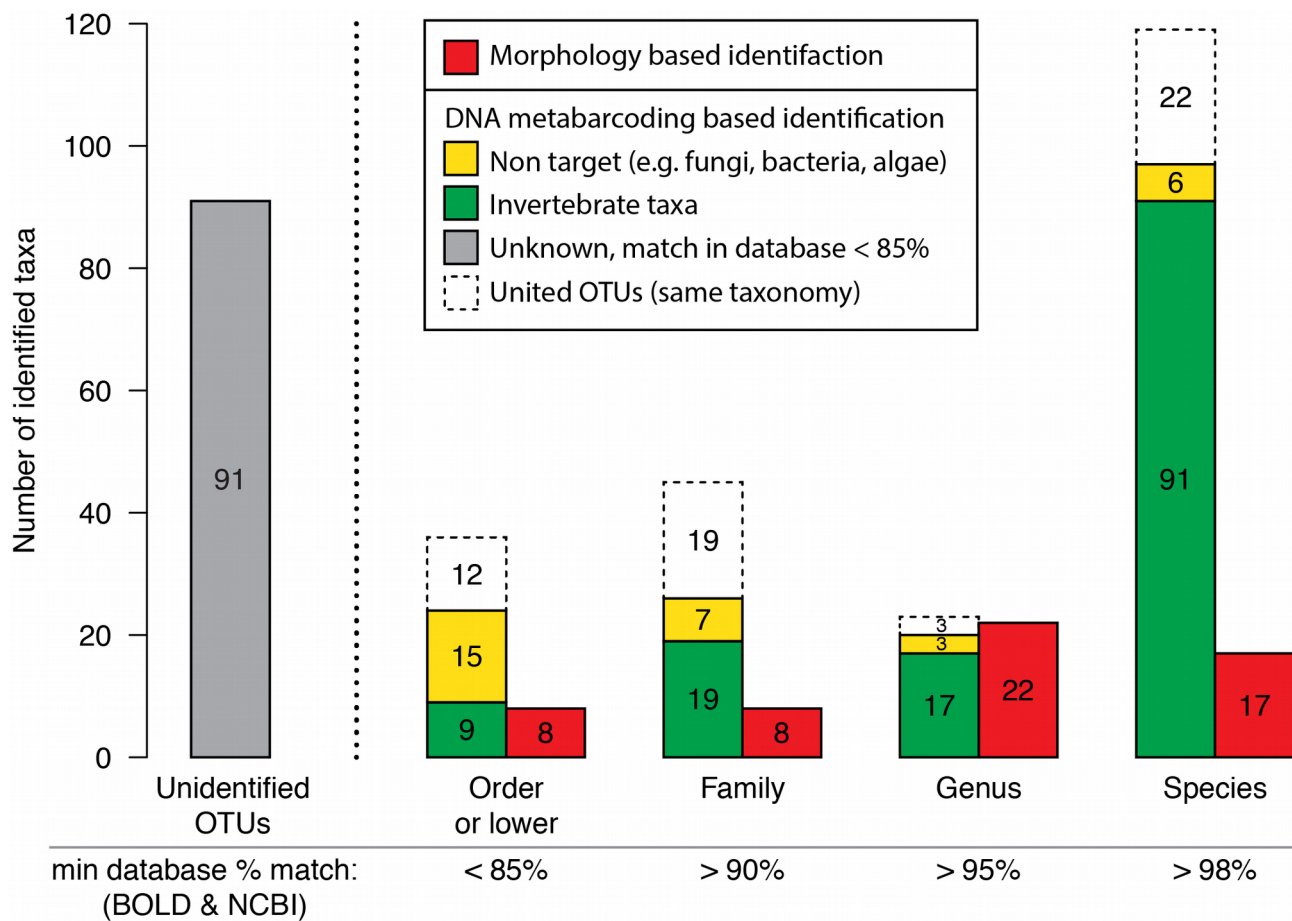


Figure S9: Comparison of taxa identified by DNA metabarcoding and morphological determination on different taxonomic levels. Below the bars the minimum required % match of the reference sequences required for sorting sequences into the respective category. OTUs with the same taxonomic hits in the reference database were merged into one single hit.