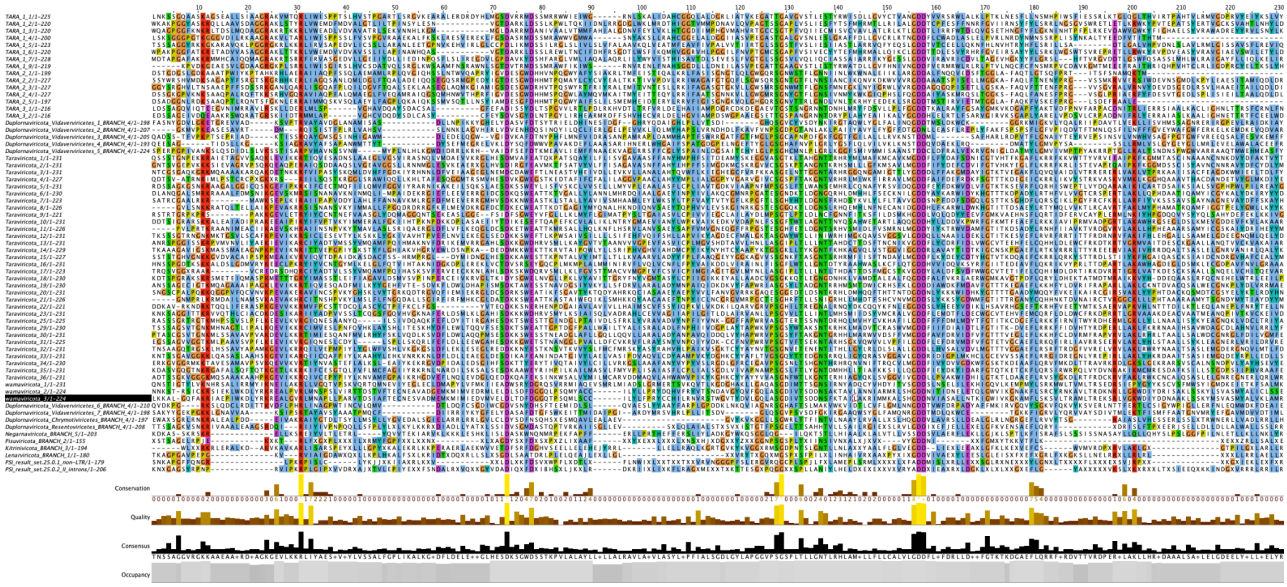Data S4: Sequence alignment used for the Global RdRp phylogenetic. Aligned sequences were derived from both individual megataxa and consensus sequences. The colors coding scheme represents amino acid properties per column in the alignment. The multiple sequence alignment and phylogenetic tree are available online for download (see **Data and materials availability** section).