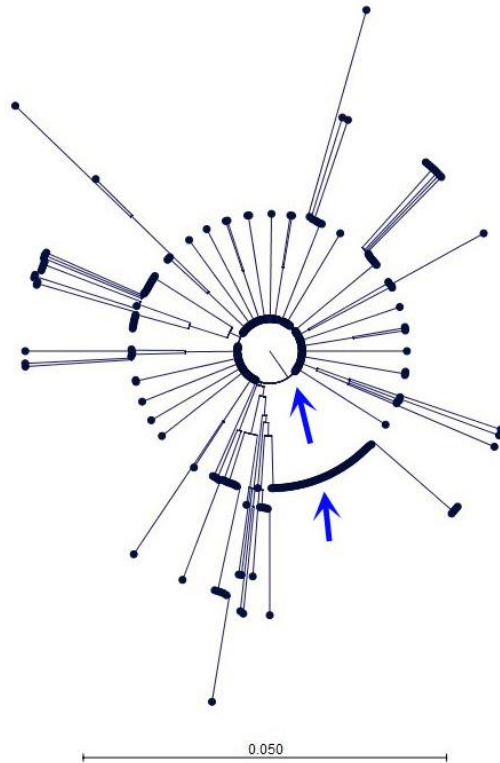


Figure S4. Phylogenetic NJ trees constructed from 500 aligned Illumina reads derived from the 5S genic part (A) and an intergenic spacer (B). Well supported branches (bootstrap >60%) are indicated by blue arrows. (C) A sequence of the 5S clone b from *E. lucius* (GenBank KX965717.1) with highlighted subregions used in the phylogenetic analysis.

A



B

