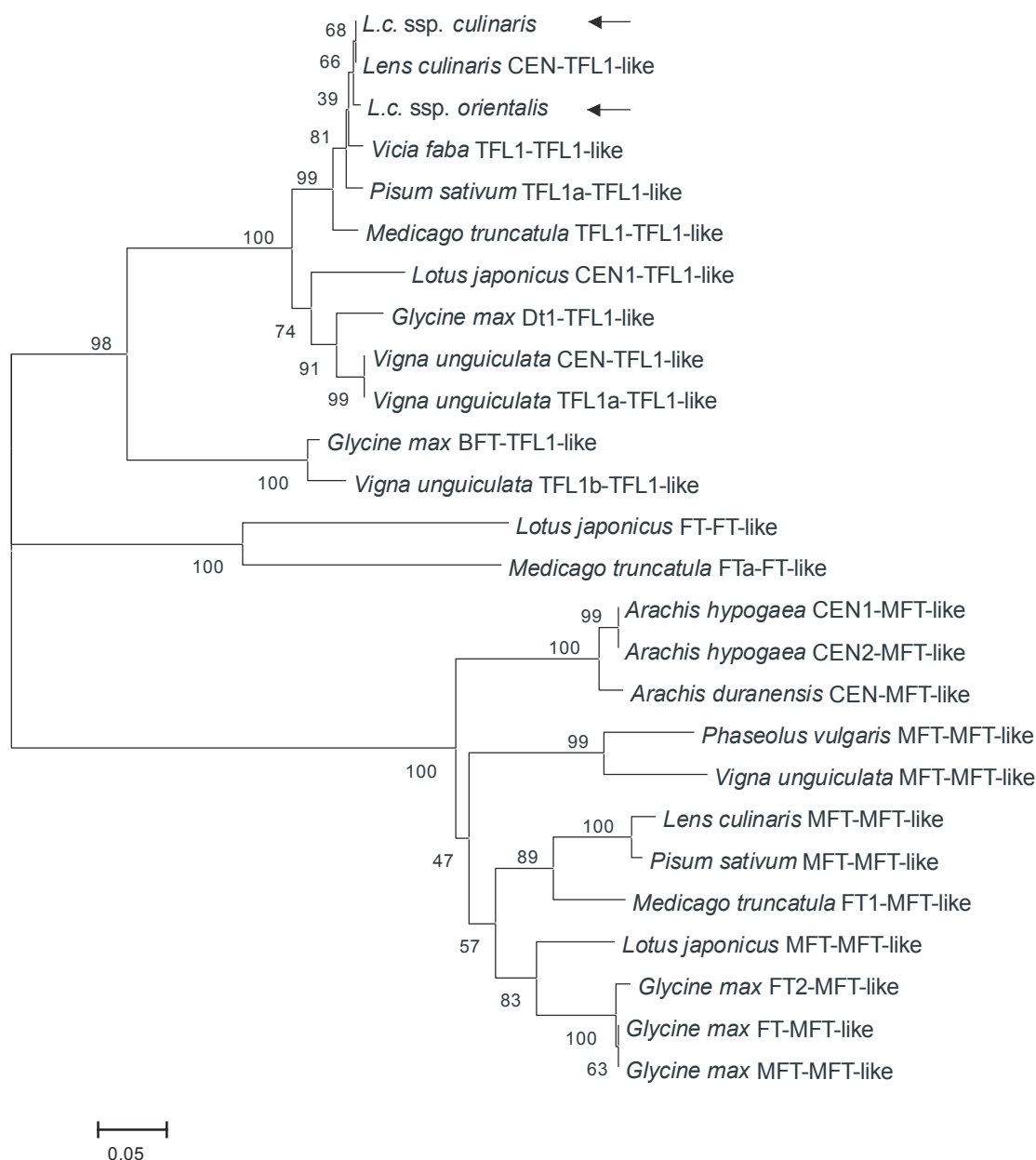




Supplementary Figure 1. Unrooted dendrogram showing the relationships between the *Lens* TFL1a amino acid sequences obtained in this work (arrows) in relation to other legume sequences available in data bases. The nomenclature from databases has been maintained. The distance dendrogram was obtained from amino acid sequences using the Jones *et al.* (1992) method and the Neighbor-Joining clustering procedure as implemented in the software MEGA 5.0 (Tamura *et al.*, 2011). Figures indicate the node bootstrap (500 replicates) supporting values.