

Table S1. Primers sequences and target region of rRNA used in the metabarcoding analysis of olive xylem sap.

PCR	Primers Pairs	Primer Forward	Primer Reverse	Region	Size (pb)	References
PCR1	799F+1062R	AACMGGATTAGATACCKG	CTCACRRCACGAGCTGAC	V5-V6	270	(Chelius and Triplett, 2001; Diskin et al., 2017)
PCR2	799F+1115R	AACMGGATTAGATACCKG	AGGGTTGCGCTCGTTG	V5-V6	320	(Laforest-Lapointe et al., 2017)
PCR3	967F+1391R	CAACGCGAAGAACCTTACC	GACGGGCGGTGWGTRCA	V6- V8	430	(Callahan et al., 2016; Sogin et al., 2006; Walker and Pace, 2007)
PCR4	799F+1193R	AACMGGATTAGATACCKG	ACGTCATCCCCACCTTCC	V5- V7	400	(Beckers et al., 2016; Bodenhausen et al., 2013; Walker and Pace, 2007)