

Title: **A molecular characterization of the invasive fig weevil *Aclees taiwanensis* in Italy**

Authors: **Rodolfo BERNARDI, Lucia GROSSI, Andrea CAVALLINI, Flavia MASCAGNI, Massimo MEREGALLI, Erika Carla PIERATTINI, Pier Luigi SCARAMOZZINO, Andrea LUCCHI, Barbara CONTI**

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A.taiwanensis_2LU_B_MK478005 -----CCAGGAGTG-TGGTGCATGGCCGTTCTTAGTTGGTGGAGCGATTTG
A.taiwanensis_4MS_B_MK478007 -----CCAGGAGTG-TGGTGCATGGCCGTTCTTAGTTGGTGGAGCGATTTG
A.hirayamai_11SIB_B_MK478004 -----CCAGGAGTG-TGGTGCATGGCCGTTCTTAGTTGGTGGAGCGATTTG
A.hirayamai_7SUB_B_MK478009 -----CCAGGAGTG-TGGTGCATGGCCGTTCTTAGTTGGTGGAGCGATTTG
A.hirayamai_11SIB_A_MK478003 GCTCTTTCTTGATTCCGGTGGGTGGTGGTGCATGGCCGTTCTTAGTTGGTGGAGCGATTTG
A.taiwanensis_4MS_A_MK478008 GCTCTTTCTTGATTCCGGTGGGTGGTGGTGCATGGCCGTTCTTAGTTGGTGGAGCGATTTG
A.taiwanensis_2LU_A_MK478006 GCTCTTTCTTGATTCCGGTGGGTGGTGGTGCATGGCCGTTCTTAGTTGGTGGAGCGATTTG
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A.taiwanensis_2LU_B_MK478005 TCTGGTTAATTCGGATAACGAAACGAGACTCTAGCCTGCTAACTAGGCGTATTTTGACATC
A.taiwanensis_4MS_B_MK478007 TCTGGTTAATTCGGATAACGAAACGAGACTCTAGCCTGCTAACTAGGCGTATTTTGACATC
A.hirayamai_11SIB_B_MK478004 TCTGGTTAATTCGGATAACGAAACGAGACTCTAGCCTGCTAACTAGGCGTATTTTGACATC
A.hirayamai_7SUB_B_MK478009 TCTGGTTAATTCGGATAACGAAACGAGACTCTAGCCTGCTAACTAGGCGTATTTTGACATC
A.hirayamai_11SIB_A_MK478003 TCTGGTTAATTCGGATAACGAAACGAGACTCTAGCCTGCTAACTAGGCGTATTTTGACATC
A.taiwanensis_4MS_A_MK478008 TCTGGTTAATTCGGATAACGAAACGAGACTCTAGCCTGCTAACTAGGCGTATTTTGACATC
A.taiwanensis_2LU_A_MK478006 TCTGGTTAATTCGGATAACGAAACGAGACTCTAGCCTGCTAACTAGGCGTATTTTGACATC
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A.taiwanensis_2LU_B_MK478005 CTAAAGGCCCGCCGGCTTGGGCTTCGCCTTTAGTCGTGCGCGGTTTTTACTGTC-----
A.taiwanensis_4MS_B_MK478007 CTAAAGGCCCGCCGGCTTGGGCTTCGCCTTTAGTCGTGCGCGGTTTTTACTGTC-----
A.hirayamai_11SIB_B_MK478004 CTAAAGGCCCGCCGGCTTGGGCTTCGCCTTTAGTCGTGCGCGGTTTTTACTGTC-----
A.hirayamai_7SUB_B_MK478009 CTAAAGGCCCGCCGGCTTGGGCTTCGCCTTTAGTCGTGCGCGGTTTTTACTGTC-----
A.hirayamai_11SIB_A_MK478003 CTAAAGGCCCGCCGGCTTGGGCTTCGCCTTTAGTCGTGCGCGGTTTTTACTGTCGGCGTA
A.taiwanensis_4MS_A_MK478008 CTAAAGGCCCGCCGGCTTGGGCTTCGCCTTTAGTCGTGCGCGGTTTTTACTGTCGGCGTA
A.taiwanensis_2LU_A_MK478006 CTAAAGGCCCGCCGGCTTGGGCTTCGCCTTTAGTCGTGCGCGGTTTTTACTGTCGGCGTA
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A.taiwanensis_2LU_B_MK478005 -----
A.taiwanensis_4MS_B_MK478007 -----
A.hirayamai_11SIB_B_MK478004 -----
A.hirayamai_7SUB_B_MK478009 -----
A.hirayamai_11SIB_A_MK478003 CAAACAATTCTTCTTAGAGGGACAGGCGGCTTC
A.taiwanensis_4MS_A_MK478008 CAAACAATTCTTCTTAGAGGGACAGGCGGCTTC
A.taiwanensis_2LU_A_MK478006 CAAACAATTCTTCTTAGAGGGACAGGCGGCTTC

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Figure S1. Multialignment of partial 18S rDNA sequences of *Aclees taiwanensis* and *Aclees hirayamai*. The organism, sample, haplotype and accession number in gene bank are reported (e.g., *A. taiwanensis*_2LU_B_MK478005 is organisms ‘*Aclees taiwanensis*’, the sample ‘2LU’, haplotype ‘B’ and accession number in gene bank ‘MK478005’). Conserved sites are marked by an asterisk.