

**MOLECULAR PHYLOGENY OF GALL-FORMING APHIDS TRIBE
FORDINI (HOMOPTERA, APHIDIDAE, ERIOSOMATINAE) ON *PISTACIA*
TREES FROM KURDISTAN REGION NORTH OF IRAQ**

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ABSTRACT

Four aphid species belonging to the Tribe Fordini (Homoptera, Aphididae, Eriosomatinae) induce galls on *Pistacia* trees in North Iraq (Kurdistan) were recorded in this study. Molecular phylogeny of Iraq Fordini species was constructed for These species (*Forda riccobonii*, *Paraclsetus cimiciformis*, *Baizongia pistaciae* and *Forda marginata*), based on sequences of mitochondrial Cytochrome Oxidase I (COI) gene of these aphid species to detect the relationship and taxonomy of Iraqi species for tribe Fordini. Our results confirmed that each of *Forda* and *Paraclsetus* are a sister genus, or monophyletic group by designing in same clade, while *B. pistaciae* is separate from other clades, supporting the taxonomic division of the tribe Fordini into two subtribes (Fordina and Baizongina).

Key words: fordini, molecular phylogeny, (COI) gene, gall aphids, *Pistacia*

علي وآخرون

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علاقات النشوء الجزيئية لحشرات المن المكونة للعقد قبيلة Fordini (Homoptera, Aphididae, Eriosomatinae)

على أشجار الفستق من منطقة كردستان شمال العراق

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المستخلص

اربعة انواع من حشرات المن المكون للعقد على اشجار الفستق تعود لقبيلة Fordini (Homoptera, Aphididae, Eriosomatinae) قد سجلت في هذه الدراسة. علاقات النشوء الجزيئية لانواع هذه القبيلة في العراق تمت دراستها في الانواع (*Forda riccobonii*, *Paraclsetus cimiciformis*, *Baizongia pistaciae* and *Forda marginata*) اعتمادا على تتابعات الجين mitochondrial Cytochrome Oxidase I (COI) للتحري عن علاقات الانواع العراقية لقبيلة Fordini وتصنيفها. اكدت نتائج هذه الدراسة على ان كل من الجنس *Forda* و *Paraclsetus* هي اجناس شقيقة عن طريق انضمامها في مجموعة وحيدة الاصل (clade) في حين انفراد النوع *B. pistaciae* في مجموعة اخرى مما يدعم تصنيف هذه القبيلة الى تحت قبيلتين هما (Fordina and Baizongina).

الكلمات المفتاحية: قبيلة Fordini ، العلاقات الجزيئية الجينية، جين COI ، المن المكون للعقد، الفستق.

INTRODUCTION

The genus *Pistacia* belongs to a cosmopolitan family “Anacardiaceae, this family include about 70 genera and over 600 species (24). The species of the genus *Pistacia* are dioecious trees or shrubs and characterized as xerophytic trees which and growing up to 8–10m height. In Iraq AL-Saghir and Porter (4) listed three *Pistacia* species, *Pistacia atlantica* Desf., *Pistacia eurycarpa* Yalt. and *Pistacia khinjuk* Stocks. The subfamily Eriosomatinae has recently been placed with family the Aphididae (16, 17). It was previously placed in the family Pemphigidae (= Eriosomatidae). There are about 19 aphid genera with 107 valid species belong to the tribe Fordini (Subfamily Eriosomatinae) (5, 6, 28), according to Inbar, (19) about 30 of these species are gall-inducing aphids belong to this tribe. The division of subtribes in Fordini is confusing. Some believe that Fordini includes two subtribes, Fordina and Baizongina (14, 37), others suggest it is divided into Fordina and Melaphidina (8, 18, 34). Almost all the species in Fordini are gall-forming and stimulate species-specific galls on their primary hosts, whereas the primary hosts for all species are full within two genera, *Pistacia* and *Rhus*, which belong to Anacardiaceae. Galls are the characteristic of aphids (29). Their position, morphology and structure of these galls are important aphid appearing. Among the whole Aphididae, only Eriosomatinae, Hormaphidinae and some species of Aphidinae produce galls (7), Galls of Fordini are significantly variable in morphology and position, nevertheless fascinatingly, are species-specific (32), since the gall founder, fundatrix forms a unique-shaped gall in a species-specific manner, the galling aphid species can usually be identified exclusively based on the gall morphology (21). Martinez, (23) and Wool, (33) reported about 15 species of aphids on *Pistacia* trees induce species-specific gall. The more common galls are those of *Slavum wertheimae* [H.R.L.], *Smynthuodes betae* West, *Forda riccobonii* [Stefani] and *Geoica* sp..The geographic distribution of these aphid species everywhere in the Mediterranean Basin, from Morocco and Spain to east Afghanistan and Iran, agrees with the distribution of host

Pistacia sp. trees (9, 14, 36). In India Chakrabarti, (10, 11) listed four species of aphids on *Pistacia* trees included *Forda riccobonii*; *F. marginata*; *Baizongia pistaciae* and *Geoica urticularia*. In Turkey 7 aphid species were Identified forming galls on *Pistacia* species. *Forda formicaria*, *Smynthuodes betae*, *Aploneura lentisci*, *Geoica urticularia*, *Baizongia pistaciae*, *F. Magenta* and *Rectinasus buxtoni*. (12) Ortiz-Rivas *et al.* (26) conducted a survey to explain the phylogenetic relationships among 10 Fordina species present in the Iberian Peninsula and the Canary Islands when they studied the implications for the taxonomy of genera *Forda* and *Paraclsetus* Mróz and Depa, (25) proved the usefulness of molecular analysis for solving taxonomic problems in aphids when they studied the DNA sequences of Mediterranean populations of *Paraclsetus cimiciformis* in aphid specimens morphologically identical with *Forda rotunda* Theobald, 1914. Li *et al.* (22) analyzed the molecular phylogeny of the Subfamily Eriosomatinae (also called Pemphiginae) which traditionally consists of 3 tribes, Eriosomatini, Pemphigini, and Fordini based on morphological and biological evidence. In Iraq, Aphid fauna induce species-specific gall almost poorly known, they are known mainly by insufficient surveillance lists, three aphids forming galls on *Pistacia khinjuk* where listed by Daoud and El-Haidari, (13); Al-Ali, (1) in North of Iraq the species are *Baizongia pistaciae*, *Geoica articular* and *Forda formicary*. The present study aimed to determine forming galls aphid in the North of Iraq, and clarifying the evolutionary relationships among the Tribe Fordini, and further investigating the relationships within and between *Forda*, *Paraclsetus* and *Baizongia*. This study represented the first phylogenetic study to this group in Iraq, preceded by few studies on other groups such as: Ali *et al.*, (2) on the widow spider in Iraq; Al-Saad and Aletby, (3) on the red palm weevil; Faraj *et al.*, (15) on the protozoan parasites causing Equine babesiosis in Baghdad city and few others.

MATERIALS AND METHODS

Sampling

Specimens for DNA extraction and amplification of mitochondrial Cytochrome

Oxidase I (*COI*) gene were collected from recent years, mostly from different types of galls on the infected primary hosts *Pistacia* trees (figure 1). *COI* gene was applied widely in the molecular phylogenetic studies of aphids and are easy to use in PCR. (22). All specimens were collected from different locations in Duhok, Erbil and Sulaimaniya provinces/ Kurdistan region North of Iraq during Summer seasons 2012-2013. Collection information of these specimens, including locations, sample numbers, and collection dates, (shown in table 1). Specimens were stored at (95–100)% ethanol at 4°C. (25). All

specimens were deposited in the Iraq National History Research center and Museum, Department of Entomology, University of Baghdad. The specimens were identified according to their main morphological diagnostic features (7, 31). The extraction of total genomic DNA using the Geneaid DNA Mini extraction kit for (Tissue), according to the standard protocol recommended by the manufacturer, with some modification. Extracted DNA was being stored at -20°C until use. the whole aphid individual was selected for molecular experiments in 10 replicates for each per specimen.

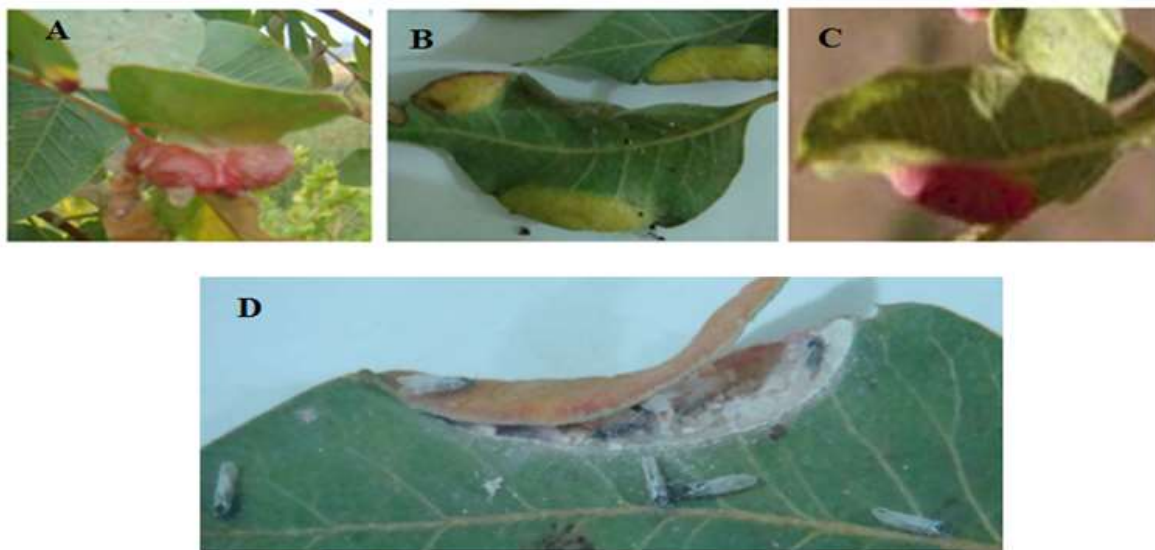


Figure 1. Shape of *Pistacia* trees-feeding aphids galls. (A,B and C), Pea/Margin galls on the blade of leaves by: A, *Forda riccobonii* from Erbil and Duhok provinces; B, *Paracletus cimiciformis* from Erbil, Sulaimaniya and Duhok provinces; and C, *Forda marginata* from Duhok province); D) Alate form of *Paracletus cimiciformis* from opening matured gall.

Table 1. Summary of collecting data relative to species (*Ford riccobonii*, *Paracletus cimiciformis*, *Baizongia pistaciae* and *F. marginata*) included in the present study.

Species	Host	Locality, Province	Number of collecting galls	date
<i>Forda riccobonii</i>	<i>Pistacia khinjuk</i>	Erbil and Duhok	6	14/05/2012
			Erbil (1) and Duhok (5)	10/6/2013
<i>Forda marginata</i>	<i>Pistacia</i> Sp.	Duhok	3	12/6/2013
				15/5/2012
<i>Paracletus cimiciformis</i>	<i>Pistacia khinjuk</i>	Erbil and Duhok	12	13/6/2013
			Erbil (4) and Duhok (8)	14/6/2013
				10/5/2012
<i>Paracletus cimiciformis</i>	<i>Pistacia</i> sp.	Sulaimaniya	2	11/5/2012
<i>Baizongia pistaciae</i>	<i>Pistacia khinjuk</i>	Duhok	3	16/5/2012

Sequencing

The classical method of CTAB (hexadecyltrimethylammonium bromide) extraction was reliable to obtain whole-genome DNA from each sample. The gene *COI* was amplified, and sequenced directly with an automated sequencer. The gene was amplified with LepF (5'ATTCAACCAATCATAAAGATATTGG-

3') (forward primer) and LepR (5' - TAAACTTCTGGATGTCCAAAAATCA-5') (reverse primer) (22). Complete sequences were deposited in GenBank under accession numbers given in table 2, together with the accession numbers of the sequences used for data comparison from the NCBI databases from different countries.

RESULTS AND DISCUSSION

The problems of morphological taxonomy of the gall-forming species identity, is confusing since they very closely related species, which

are usually morphologically very similar and thus, making it difficult for identifying these species. (25, 27, 30). So the present study was done to avoid these problems.

Table 2. Summary of data relative to specimens analyzed and accession numbers for the mitochondrial DNA Cytochrome Oxidase I (COI) gene sequences

Organsims	Accession no. local samples	Accession no. of reference samples of GenBank	Nucleitid no. of GenBank	Score/Identities
<i>Forda riccobonii</i>	MK111622.1	EU701669.1	41-637	588/(99%)
<i>Forda marginata</i>	MK111626.1	EU701668.1	50-637	576/(99%)
* <i>Paracletus cimiciformis</i>	MK111623.1	JQ241782.1	30-623	585/(99%)
** <i>Paracletus cimiciformis</i>	MK111625.1	JQ241782.1	43-620	440/(92%)
<i>Baizongia pistaciae</i>	MK111624.1	KY841873.1	46-449	377/(98%)

* specimens from Erbil and Duhok provinces.

** specimens from Sulaimaniya province.

The comparison of the studied sequence specimens with deposits (COI) gene copies of the same species in the NCBI database showed high identity (98-99%) with the reference species with accession number (table 2) (*Forda riccobonii* EU701669.1- Israel, *Forda marginata* EU701668.1 Canada, **Paracletus cimiciformis* JQ241782.1- Poland and *Baizongia pistaciae* KY841873.1- Canada), while the specimen ** *Paracletus cimiciformis* which collected from Sulaimaniya Province showed identity (92%) which differ from specimen * *Paracletus cimiciformis* this probably because of the polymorphism of their nucleotide as proofed by the alignment matrix, thus GenBank providing this specimen in different accession number. The results of phylogenetic analysis for a segment of 513 bp of the mitochondrial barcoding gene cytochrome oxidase I (COI) obtained with both applied markers indicated that sequences of studied specimens with those deposited in the data bank belonging to species (*Ford riccobonii*, *Paracletus cimiciformis*, *Baizongia pistaciae* and *F. marginata*). The present phylogenetic analyses support the taxonomic division of the tribe Fordini into two subtribes (Fordina and Baizongina) by creating two clades, the first one represents two sister groups, the first one includes *Ford riccobonii* and *F. marginata* and the second includes *P. cimiciform* which represents a sister group to the genus *Forda* sp. or

monophyletic group, while the species *B. pistaciae* individually with the second clade. Comparing our molecular phylogenetic analysis with previous studies our results are in agreement with the phylogeny of Ortiz-Rivas *et al.* (26) which presented that each of *Forda* and *Paracletus* are a sister genera when they studied Molecular phylogeny of Iberian Fordini, they phylogeny analysis suggests that the group can be divided into two main clades. One clade is composed of aphids inducing small, low-capacity galls including genera *Forda*, *Paracletus* and others, while the second clade is composed of species inducing larger galls such as *B. pistaciae* and other species, although they are detecting others genes. Similar results are shown by Mróz and Depa, (25). Although Zhang and Qiao, (35) suggest Fordini consisting of two subtribes, Fordina and Melaphidina. They concluded that in Fordina, two groups: one is represented by *Forda*, and the other is represented by *Baizongia*. The second clade included the species (*B. pistaciae* and *Slavum wertheimae*), the molecular relationships among them would agree with previous literatures (21, 26, 35). The present study concludes that each of *Forda* and *Paracletus* are a sister genus, or monophyletic group by designing in the same clade while, *B. pistaciae* separately with other clades, supporting the taxonomic division of the tribe Fordini into two subtribes (Fordina and Baizongina).

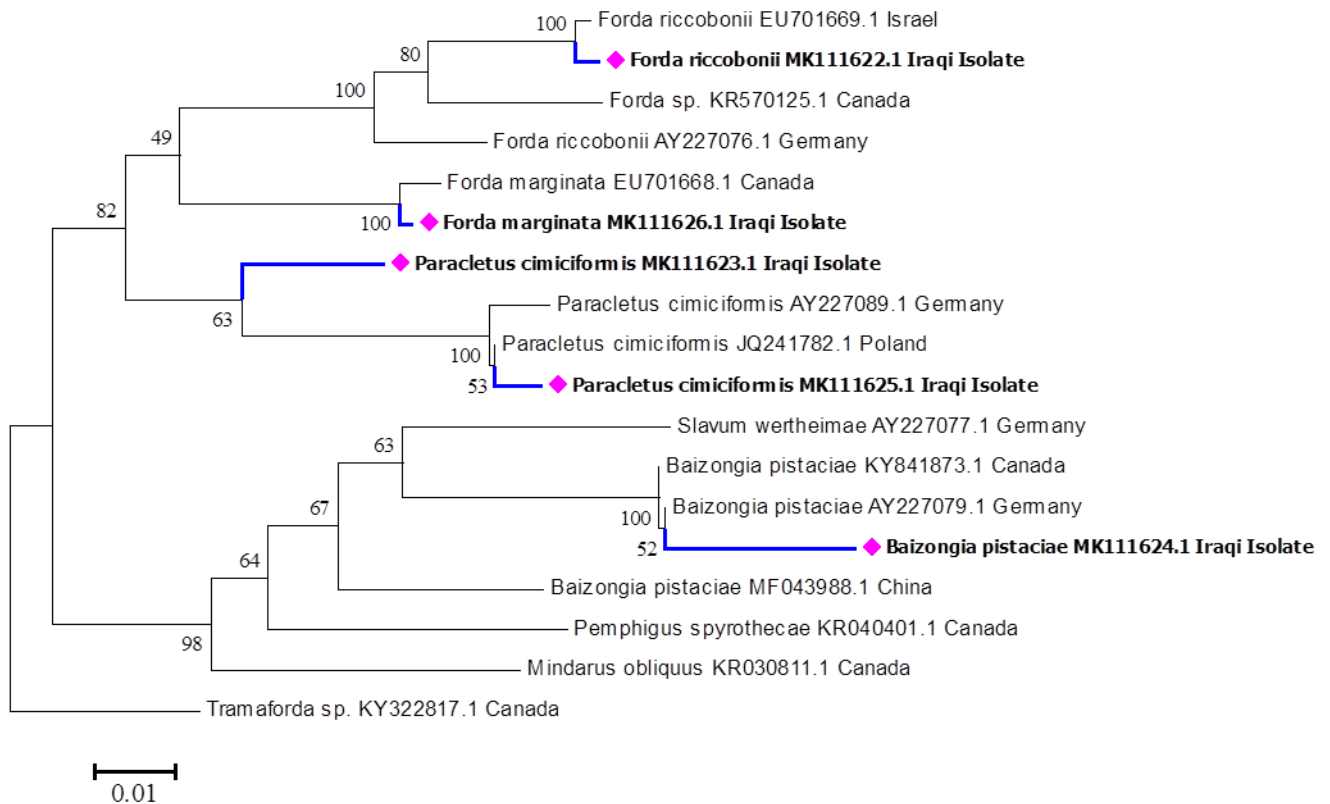


Figure 2. Neighbour joining phylogenetic tree constructed of Cytochrome Oxidase I gene sequences from galls forming aphids collected from different locations in Duhok, Erbil and Sulaimaniya, with their accession numbers; local samples in Bold, reference samples of GenBank from different countries. A measure of distance between taxa is indicated by the scale bar below the tree

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