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TAXONOMIC AND MOLECULAR STUDY OF THE WIDOW SPIDER GENUS *LATRODECTUS* WALCKENAER, 1805 (ARANEAE: THERIDIIDAE) IN IRAQ

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ABSTRACT

The widow spider, *Latrodectus*, (Araneae: Theridiidae) comprise about 31 currently recognized species, some species are very hard to distinct; in this study morphological and molecular method has been used to confirm the diagnosis of the spiders. Phylogenetic tree was constructed using the neighbor-joining of some other *Latrodectus* sp. sampled, including taxa occurring in the Middle East, Australia, New Zealand, North and South America and Europe, and with the two genera *Steatoda* Sundevall, 1833 and *Asagena* Sundevall, 1833 outgroup. The results of the analysis show the highest identity (90%) for *Latrodectus thoracicus* Nicolet 1849 represented by three specimens and for *L. pallidus* O.P.-Cambridge, 1872 represented by one specimen in addition to (85%) for the false widow *Asagena phalerata* (Panzer, 1801) represented by one specimen, these three species is recorded in Iraq for the first time.

Keywords: Latrodectus, Taxonomy, Iraq Spider Fauna, Widow Spiders

INTRODUCTION

The genus *Latrodectus* Walckenaer, 1805 (Araneae: Theridiidae) is a worldwide distribute genus (Graudins *et al.*, 2001), it includes a group of species commonly referred to as widow spiders. It's considered a taxonomically complex genus as the status of several forms had not been properly evaluated and specific boundaries are not well defined or understood (Levi, 1959; 1967; Garb *et al.*, 2001), therefore, in multiple cases, populations has been uncritically referred to as different taxa. Discriminating between *Latrodectus* species using morphology has always been problematic (Levi, 1983), it is difficult taxonomically and readily separated from members of other Theridiid genera (Mirshamsi, 2005).The Genus *Asagena* Sundevall, 1833 was revalidated by Wunderlich (2008), this genus was earlier considered as a junior synonym of *Steatoda* Sundevall, 1833.

The information and knowledge about the widow spiders in Iraq are very limited; in general spider in Iraq was neglected and rarely studied until the last few years, recently the interest in studying of this group was increased because of the emergence of several cases of bites, which was supposed to be caused by black widow spiders, researchers have revealed the existence of three *Latrodectus* species in Iraq, *L. scelio*

(Abdul-Rassoul *et al.*, 2012), *L. hasselti* (Al-Hadlag and Najim, 2015), while Zamani and El-Hennawy (2016) rejected these two species from the list of Iraqi spiders fauna and attributed them to *L. renivulvatus*. The difficulties of diagnosis underlines the importance of molecular characters in creating a valid phylogeny for this genus, this study characterizes the first phylogenetic relationship to the *Latrodectus* genus in Iraq and is made by sequence of the partial mt DNA-COI gene. The aim of the present study is to provide a taxonomic status of the widow spider species based on molecular analysis for the first time in Iraq.

MATERIALS AND METHODS

Spiders collection

Widow spiders were collected using forceps, preserved in 70-80 % ethanol, ten individual spider specimens obtained from different locations in Baghdad, Najaf and Babylon provinces. All the examined specimens were deposited in the Iraq Natural History Research Center and Museum, University of Baghdad. Identification was carried out by using morphological characteristics according to several diagnostic keys (Mirshamsi, 2005; Levi and Randolph, 1975; Sutton, *et al.*, 2006).

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DNA extraction

DNA was extracted from (1-2) legs removed with clean forceps from each specimen with 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.

Amplification for COI Gene

The Cytochrome (C) Oxidase I gene was amplified with LepF (5'ATTCAACCAATCATAAAGATATTGG-3') (forward primer) a n d LepR (5'-TAAACTTCTGGATGTCCAAAAAATCA-5') (reverse primer) (Brandon-Mong, et al., 2015). The DNA fragments to be analyzed were amplified using AccuPower PCR PreMix (BIONEER, Corp., Daejeon, Korea) in 50µl reaction mixtures containing 0.4 µM each primer and 100-200 ng of genomic DNA template. The PCR amplification protocol was performed according to the following procedure: initial denaturation at 95 °C for 5 min, followed by 35 cycles of 95 °C for 60 s; annealing temperature 46.7°C for 30 s; and extension at 72°C for 60 s and final extension at 72°C for 5 min.

RESULTS AND DISCUSSION

The results of PCR tests on 10 spider's specimens showed that LepF and LepR primers amplify one part of the mtDNA-COI gene with the length about 710 bp. only, 5 PCR products of studying spiders were sequenced. The alignment of acquired sequences with submitted sequences in Gene Bank showed that these sequenced samples belong to *Latrodectus thoracicus* for (BAB-1, KUF-3 and BAG-5) specimens and *L. pallidus* for KUF-2 specimen with Identities (90%), and belongs to *Asagena phalerata* for BAB-4 specimen with Identities (85%).

The nucleotide sequences of (BAB-1) submitted to Gene Bank under accession number MG645011 for *L. thoracicus*, nucleotide sequences of KUF-2 submitted under accession number MG645012 for *L. pallidus*, and nucleotide sequences of BAB-4 submitted under accession number MG645013 for *A. phalerata*. Sequencing ID in Gene Bank, score, expect and compatibility of Cytochrome Oxidase Subunit 1 (COI) gene, partial cds; mitochondrial sequences of the isolates spiders were summarized as in Table 1.

Table 1

Sequencing ID in GenBank, score, expect and compatibility of Cytochrome Oxidase Subunit 1 (COI) gene, partial cds; mitochondrial sequences.

*isolates	Organisms	Sequence ID	Score	Expect	Identities	Position in Gene Bank
BAB-1, KUF-3, BAG-5	Latrodectus thoracicus	GU112105.1	826	0.0	90%	1 to 639
KUF-2	Latrodectus pallidus	KY007713.1	821	0.0	90%	1 to 624
BAB-4	Asagena phalerata	KY269726.1	593	2e-165	85%	71 to 658

Table 2

Sequence Identity Matrix of the partial mtDNA-COI gene.

Seq<-	BAB-1	BAB-4	BAG-5	KUF-2	KUF-3
BAB-1	ID				
BAB-4	•. ٧٩٦	ID			
BAG-5	۱.۰۰۰	• ٧٩٦	ID		
KUF-2		•.975	·. 9V£	ID	
KUF-3	1	• <u>.</u> ٧٩٦	1	•.975	ID

Phylogenetic Analysis

Phylogenetic relationship of these widow spiders was analyzed (Fig.1). The presented phylogenetic analysis distinct two clades (*"Latrodectus"* clade and *"Steatoda"* clade) all *Latrodectus* species in this phylogeny due to the *"mactans"* clade as resolved by the phylogenetic tree of Garb *et al.* (2004). From the Sequence Identity Matrix Table 2 results support the convergence between *L. thoracicus* and *L. pallidus*, similarly, that appears as sister group, the phylogenetic tree which presented by Mollaiizadeh *et al.* (2017) distinct this convergence, although they didn't present as sisters, Garb *et al.* (2004) interpreted the relationships and why *L. pallidus* is distinct from many other *Latrodectus* spp. in the *"mactans"* clade.



Fig.1

Neighbor-joining phylogenetic analysis of mtDNA-COI gene in widow spiders, (BAB-1, KUF-3 and BAG-5) for *Latrodectus thoracicus* Iraq, (KUF-2) for *Latrodectus pallidus* Iraq, and (BAB-4) for *Asagena phalerata* Iraq

Morphological Analysis

Key to widow spiders in this study:

Latrodectus pallidus O.P.-Cambridge, 1872 (white widow spider)

Material examined: 1^{\bigcirc} and 1 juvenile Babylon province, Al-Nael, June, 2016; 1^{\bigcirc} Al-Najaf province, Kufa, Aug, 2016; 1^{\bigcirc} Baghdad province, Abu Ghraib, Feb. 2017.

Diagnosis: Female: body length 10.0-12.5 mm, cephalothorax: carapace light brown to orange brown; legs very light brown with broad dark annular stripes; abdomen pale cream to white with several dark dots on dorsal side.

Global distribution: Cape Verde Islands, Libya to Central Asia (Platnick, 2014).

Latrodectus thoracicus Nicolet, 1849

Material examined: 1 \bigcirc Babylon province, Al-Neel district, Jun. 2016; 1 \bigcirc Al-Najaf Province, Kufa, Aug. 2016; 1 \bigcirc and 1 juvenile Baghdad province, Abu Ghraib, Feb. 2017.

Diagnosis: Female: body length 10.2-11.4 mm cephalothorax: carapace and legs shiny black; abdomen black with large dorsal red marking from the middle of central line to the end of the abdomen; ventrally with red hourglass marking small in size.

Male: smaller than female, body length 2.8 mm; cephalothorax: carapace dull brown wider than longer; abdomen white with creamy parts, with several dark dorsal bands irregular in size and shape; legs longer than in female; color black to dark brown with broad white annular stripes.

Global distribution: Central and Southern parts of Chile (Aguilera *et al.*, 2009) and other region.

Asagena phalerata (Panzer, 1801)

It's a synonym to *Steatoda phalerata* (Panzer, 1801) (Le Peru, 2011).

Material examined: 1^{\bigcirc} and 1^{\bigcirc} and several specimens juveniles collected from Babylon province, Al-Neel district, Aug. 2016

Diagnosis: Female: body length 4.8 mm; cephalothorax: carapace shiny dark brown to black with numerous tiny tubercles; legs reddish brown with broad dark annular stripes. Palps reddish brown to brown; abdomen shiny dark brown with a pattern of yellow or white spots. Male: body length 4.2 mm. cephalothorax: carapace shiny dark reddish brown with many very fine tubercles on whole surface, length longer than wide; femora of legs I and II darker brown, than III and IV, all the other segments yellowish brown with broad dark annular stripes; abdomen uniformly black with pairs of definite yellowish white horizontal patterns, with small dots at the posterior part; ventral surface dark brown with no pattern.

Habitat: arid grasslands, agricultural areas, forests, and wetlands (Vona-Túri, *et al.*, 2016).

Global distribution: wide distribution from Europe and N Africa to Asia Minor, Karakorum and Trans-Baikal area (Namkung *et al.*, 1996).

CONCLUSION

This study characterizes the first phylogenetic relationship to the *Latrodectus* genus in Iraq and is made by the sequence of the partial mtDNA-COI gene; also phylogenetic tree was constructed. The results of the analysis show the highest identity for *L. thoracicus* and for *L. pallidus*, in addition to the false widow *A. phalerata* these three species are recorded in Iraq for the first time.

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AUTHORS' CONTRIBUTION

Hayder Badry Ali prepared the experimental design and wrote the manuscript. Hula Younis Fadhil Performed the DNA extraction, amplification for COI Gene, analyzed the bioinformatics data and approved the final manuscript; while Ishraq M. Baker collected the specimens.

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