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Morphogenetic traits of the egg parasitoid *Trichogramma* for controlling certain date palms lepidopteran insect pests in the New Valley Governorate

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Abstract

Egg parasitoid *Trichogramma* strain (TNV) received from Kharga Oasis, New Valley Governorate, Egypt, is used in controlling certain date palm lepidopteran insect pests. The morphological features of adult wasps and genetically identification of the strain was carried out. Scanning electron microscopy (SEM) and light microscope were used for studying morphology of male and female adults and measurements. Data revealed that body length of male was shorter than female. However, adult parasitoid body length of TNV was longer than the species, *Trichogramma evanescens* West. studied by several authors. In addition, external morphology of antennae, wings, and male genitalia were differed in TNV *Trichogramma*. The strain TNV identification was based on the molecular genetic techniques using PCR amplification and sequencing of the internal transcribed spacer two (ITS2) region of the rDNA. The BLAST result and phylogenetic analysis of the ITS2 region of rDNA sequence revealed that the strain TNV is closely related to *Trichogramma turkestanica*. Obtained results confirmed that the *Trichogramma* sp., used as a biological control in the New Valley, is *T. turkestanica* and not *T. evanescens*.

Keywords: *Trichogramma turkestanica, T. evanescens,* Date palm, SEM, Phylogenetic analysis, Morphological and molecular identification

Background

In Egypt, date palm (*Phoenix dactylifera* L.) plantations markedly increased all over the different governorates, especially in the desert new reclaimed lands (El-Assal 2004). Dates are strategic product for Egypt, which is the world's top producer of the crop, outranking Iran and Saudi Arabia. Egypt produces 1,465,030 tons of dates annually, or 17.7% of the global production of 7.5 million tons, according to the 2014 statistics of the Ministry of Agriculture and Land Reclamation (FAO 2017).

The New Valley in the western part of Egypt, from the Nile to the Libyan frontier ,is occupied by a vast desert plateau. In this "valley", various oases are located that are, from the North to the South: Siwa, Baharia, Farafra, Dakla, Kharga, and Fayoum. The total number of productive date palms is estimated around 700,000 trees. A large amount of them (about 50%) is grown from seeds. These date palms are very heterogeneous and, in average, of low quality. Beside the palms from seeds, mainly, one variety, "Saidi," has been propagated. It could represent nearly 50% of the total number of palms of this area. Because of its excellent quality and of the importance of its production, it was transferred to the Nile valley (Riad 1993).

Many arthropod species are known as pests of the date palm throughout the World. Around 50 species of insects and mites are reported as pests of date palms worldwide,

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from which, approximately 25% are insects and mites that considered serious pests (Haldhar et al. 2017).

The palm trees in Egypt are exposed to many insects and mites' attacks. The loss in date crops may reach about 25%. Insect pests can attack florescence and date, leaves, roots, trunk and storage date (Alhudaib 2007).

Previously, methyl bromide was used to control insect pests in stored dates as fumigant because it was effective in killing most insect stages; but this fumigant appeared to be ozone-depleting agents (Marcotte 1993). Therefore, it was prohibited since 2005 in developing countries and by 2015 in under developed countries according to Montreal Protocol (Al-Taweel et al. 2014). Therefore, alternative safe control methods are needed. One of these methods is using biological control agents specifically the egg parasitoid, Trichogramma evanescens West. (Hymenoptera: Trichogrammatidae) in date orchards to minimize the percentage of dates infestation with lepidopteran insect pests as the Greater date moth (Arenipses sabella Hampsm), Almond moth (Cadra cautella Walk), Indian meal moth Plodia interpunctella Hub., and pomegranate fruit butterfly (Virachola livia Klug).

The Trichogrammatidae is a family of tiny wasps in the superfamily Chalcidoidea that include some of the smallest of all insects, ranging in size from 0.2 to 1.5 mm within the genus *Trichogramma* (Pinto 2006 and Ghoneim 2014). All members of this family attack eggs of insects belonging to 11 orders, e.g., Hymenoptera, Neuroptera, Diptera, Coleoptera, and Hemiptera especially Lepidoptera (Flanders and Quednau 1960 and Ghoneim 2014).

Despite the advances made in their taxonomy, *Trichogramma* spp. remain notoriously difficult to identify because of their tiny size and the relatively low interspecific morphological diversity (Ksentini et al. 2010). Recently, egg parasitoid *Trichogramma* is used for controlling the serious lepidopteran insect pests of date in the New Valley governorate.

The present work was conducted to lighten the morphological features of the adult wasps received for application in the field, with the purpose to determine the precise morphological and molecular identification of this strain.

Materials and methods

Trichogramma collection and rearing

Egg parasitoid *Trichogramma* was supplied from *Trichogramma* Mass Rearing Lab., Kharga Oasis, New Valley Governorate, Egypt. Eggs of *Sitotroga cerealella* Oliv., were used as host to produce sufficient numbers of the egg parasitoid for two generations under $(23 \pm 2^{\circ}c, 75 \pm 5\%$ RH, L 16: D 8) at the Biological Control Lab., Plant Protection Department, Assiut University, Assiut, Northern Upper Egypt.

Morphological and molecular identification of Trichogramma Morphological identification

For light microscope Mounted *Trichogramma* adults on microscope slides were used as permanent specimens for species identification. Identification relies upon microscopic examination of male genitalia and other morphological characters. Dr. John Pinto, University of California at Riverside provided us with the method of Knutson (1998) for this purpose. Measurement was done using HDMI MULTI -OUTPUT HD (Toup Cam_120) CAMERA.

For scanning electron microscopy (SEM) The material was fixed in FAE (formaldehyde, acetic acid, and ethanol) and stored in 70% ethanol. Skeletal structures studied using a Jeol JSM-5400 LV scanning electron microscope following critical drying point (Hitachi HCP-2) and sputter coating of samples with gold (Giko IB-3) (Polilov 2017).





Measurements morphological characters The specialized terminology for the male genitalia was followed according to the modification Polaszek et al. (2012) to that of Pinto (1999). Linear measurements are reported as a mean \pm standard deviation (SD). Species were initially identified based on morphology, using a combination of the following published taxonomic accounts: Nagarkatti and Nagaraja (1971), Pintureau and Babault (1988), Pinto (1999), Pintureau (2008), and Del Pino et al. (2013).

Molecular genetic identification

Isolation of genomic DNA Total genomic DNA was extracted as described by Stouthamer et al. (1999).

PCR amplification of the *Trichogramma* **ITS2 region of rDNA** For identification, the ITS2 region of *Trichogramma* ribosomal DNA was amplified using the specific primers as the following: forward, 5–TGTGAACTG CAGGACACATG–3, located in the 5.8S rDNA, and



Table 1 Measured characters (μ m) of *Trichogramma* strain TNV male antenna

Criteria (N = 10)	Measured characters (µm)				
	Flagellar length FL	Flagellar width FW	Scape length ScL	Longest seta LS	
Min.	149.96	28.09	75.55	70.99	
Max.	171.60	36.83	96.05	82.76	
Average	162.95	33.11	82.27	75.63	
±SD	6.72	2.90	5.94	3.59	

reverse, 5–GTCTTGCCTGCTGTGAG–3, located in the 28S rDNA closer to the 3 ends of the ITS2 (Stouthamer et al. 1999). The PCR reaction was performed in a final volume of 50 μ L containing GoTaq (Promega, Madison, WI, USA) Green Master Mix, 1 μ L of DNA sample, and 1 μ L of each primer (at a concentration of 0.5 mM). The PCR conditions were as follows: initial denaturation at 95 °C for 5 min, then followed by 36 cycles at 94 °C for 1 min, 55 °C for 1 min, and 72 °C for 1 min, with a final extension at 72 °C for 10 min and holding at 4 °C.

PCR products purification and sequence determination To verify the presence of appropriate sized amplicons, the PCR product for *Trichogramma* strain TNV was subjected to electroph oresis using 1% agarose gel in TAE buffer according to the standard method. Product of the correct size was purified using a TaKaRa agarose Gel DNA Purification Kit Ver.2.0 and sequenced in both directions using an ABI 3730 automated sequencer.

Alignment and phylogenetic analysis The ITS2 sequences of *Trichogramma* strain TNV were aligned with known ITS2 sequences *Trichogramma* spp. of rDNA in the GenBank database using the basic local alignment search tool (BLAST) at the National Center for Biotechnology Information (https://blast.ncbi.nlm.nih.gov/Blast.cgi?PRO GRAM=blastn&PAGE_TYPE=BlastSearch&LINK_LOC= blasthome), and the percent homology scores were generated to identify the *Trichogramma* strain TNV.

Phylogenetic analysis The phylogenetic tree was constructed with MEGA version 4.0 using a neighbor-joining algorithm, and the Jukes-Cantor distance estimation method was performed with bootstrap analyses for 1000 replicates (Hesham et al. 2012).

GenBank accession number The partial of the ITS2 region of rDNA sequence of the strain TNV reported in this paper is deposited in the DDBJ (www.ddbj.nig.ac.jp/), EMBL (www.embl.de/), and GenBank nucleotide sequence databases (http://www.ncbi.nlm.nih.gov) under accession number MH432649.









Data processing

Measurements of morphological data were statistically analyzed by using SAS 9.1 2008 software program.

Results and discussion

Morphological identification

Using scanning electron microscopy (SEM), some morphological structures of male and female parasitoid were illustrated. By light microscopy, collected parasitoid is an extremely tiny wasp with pale yellow to brown color (Fig. 1).

Body length

Body length of adult *Trichogramma* strain TNV female was $472.37-608.63 \mu m$ with an average of $539.44 \pm 64.34 \mu m$. However, male measurements reviled an average of $518.85 \pm 57.97 \mu m$ and ranged from 456.47 to $576.84 \mu m$. Thus, the male body length is shorter than female. Statistical analysis of the differences between the female and male body length are non-significant (*T* calculated value = 0.609421 with n = 5 each).



Compound eyes The compound eyes cover a large amount of the head (Fig. 2a). Detail of one compound eye showed interfacetal hairs distributed all across the eye (Fig. 2b). Detail of an interfacetal hair in a corner between three facets is shown (Fig. 2c).

The head

Antennae Antenna of the strain TNV is of geniculate type, consisting of scape, pedicel, and flagellum. As shown in (Fig. 3a and b), the male hairs were more condensed and elongated than female hairs. Also, in male, segment of flagellum fused with club.

Male antennal measurements Antennal trichiation one of the morphological characters have been used to identify *Trichogramma* (Dang et al. 2005 and Ksentini et al. 2010). Measurements of male antennae include flagellar length (FL), flagellar width (FW), scape length (ScL), and

exv. inv stl longest seta (LS). Data presented in Table 1 based on 10 male's antennae of TNV *Trichogramma* strain.

Mouthparts SEM photographs of TNV indicated that the mouthparts structure is typical of chewing type. Mouthparts consist of well-developed mandibles, maxillae, and labium (Fig. 4a and b). However, labrum was weakly developed. Mandibles with undulate margin and spines on internal surface are shown (Fig. 4c). Maxillae connected with labium by membranous septum into labiomaxillary complex (Fig. 4b). Maxillae consist of cardo, stipes, fused galea and lacinia, and maxillary palp. Maxillary palp 1-segmented strongly reduced.

The thorax

Wings As shown in light micrograph for *Trichogramma* strain TNV, wings with strongly depleted venation (Fig. 5). Three veins preserved in forewing: submarginal, marginal, and stigmal, fused into one arch near anterior wing margin (Fig. 5a). Hindwing narrower than forewing (Fig. 5b). Forewing rather wide, covered with small hairs, with longer hairs forming fringe on perimeter of wing. Hindwing bearing fringe of long setae along posterior margin.

Legs

Legs in *Trichogramma* strain TNV are slender and ambulatorial. Each leg consists of coxa, 2 segmented trochanter, femur, tibia, and 3 segmented tarsi. Tibiae of legs apically bear well-developed branched spurs. Apical tarsomere bears two claws and well-developed arolium (Fig. 6a).

The abdomen

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The lateral view of abdomen in TNV strain was illustrated (Fig. 7). The petiole not pronounced, mesosoma

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10µm



15kU







and metasoma broadly joined. Metasoma consists of six or seven visible tergites.

Measurements of female abdomen Length of female abdomen in *Trichogramma* strain TNV from the frons to the tip of the abdomen measured $247.086 \pm 10.682 \ \mu m$ (231.59–258.55 μm). The average abdominal area reached $42,923.65 \pm 5381.776 \ \mu m^2$ (35,365.42– 50,411.75 μm^2).

Ovipositor *Trichogramma* strain TNV female ventral view (Fig. 8a) illustrates the ovipositor plates and stylet. SEM of TNV ovipositor showed its external and internal valves with long setae and stylet (Fig. 8b).

External male genitalia Male genitalia in strain TNV is represented by simple aedeagus, phallobase, and parameres as shown in Fig. 9a and b. Measurements of male genitalia include general length (GL), general width (GW), apical distance (AD), basal distance (BD), and apical width (AW), which are presented in Table 2 based on 10 male genitalia of TNV *Trichogramma* strain.

Molecular identification of *Trichogramma* strain TNV using ITS-2 region of rDNA sequencing and phylogenetic analysis

To identify and determine the correct phylogenetic position of the *Trichogramma* strain TNV, molecular

Table 2 Measured characters (µm) of TNV male genitalia

Criteria $(N = 10)$	Measured characters (µm)					
	General length GL	General width GW	Apical distance AD	Basal distance BD	Apical width AW	
Min.	124.9	44.8	25.44	85.42	25.44	
Max.	138.19	58.85	44.42	101.75	34.55	
Average	126.46	50.31	33.94	92.64	30.23	
± SD	5.94	4.45	6.28	6.11	2.64	

genetic identifications were performed. As shown in (Fig. 10), the size of the amplified ITS2 region of rDNA was approximately 500 bp, which is the expected size of ITS2 region of rDNA for *Trichogramma* species.

The alignment and comparisons of the ITS2 sequences of rDNA of the strain TNV, to the published ITS2 sequences of rDNA in the GenBank database by BLAST search, were determined. The sequence results of the ITS2 region of rDNA of the TNV are shown to be highly homologous to *Trichogramma turkestanica* with 100% similarity. To confirm the position of TNV strain in the phylogeny, several sequences representing *Trichogramma* spp. were selected from the GenBank database to construct a phylogenetic tree. The phylogenetic analysis was performed for the TNV sequence and 10 reference sequences. As shown in Fig. 11, the phylogenetic tree indicated that TNV strain and *Trichogramma turkestanica* shared one clade. Therefore, the TNV strain was identified as *T. turkestanica*. Recently, molecular



Fig. 10 Amplified DNA of the ITS-2 region of rDNA with specific primers: lane 1, 100-bp DNA markers; lane 2 to lane 5, *Trichogramma* sp. The size of the amplified ITS2 region of rDNA was approximately 500 bp





genetic tools including PCR amplification and the sequencing of the internally transcribed spacer 2 (ITS-2) of the rRNA gene were applied for the accurate identification of many *Trichogramma* species (Ksentini et al. 2010; Sayed et al. 2011; Polaszek et al. 2012; Pino et al. 2013; Nasir et al. 2013; Almeida and Stouthamer 2015; Wu et al. 2016 and Liu et al. 2017).

The wildly use of egg parasitoid *Trichogramma* encouraged biological control specialists to give extensive and intensive attention for selecting the adequate species of the parasitoid. Egg parasitoid *Trichogramma* (strain TNV) released in Kharga Oasis, New Valley Governorate Egypt for controlling certain date palm lepidopteran insect pests is usually recognized as *Trichogramma evanescens*. Results in this work discussed and supported by literature serve as good informative background. Our morphological data was compared with other author's data for *T. evanescens*, i.e., Fischer et al. (2011), Polaszek et al. (2012) and Polilov (2016).

Concerning body length of adult TNV female, data showed an average of $539.44 \pm 64.34 \ \mu\text{m}$ and male measurements reviled an average of $518.85 \pm 57.97 \ \mu\text{m}$ with an average length of both adult sex of $529.145 \pm$

Table 3 Measured characters (μm) of strain TNV and T. evanescens male antennae

Taxon	Measured characters (µm)			
	Male antennal (average \pm SD)			
	FL/FW	FL/ScL	LS/FW	
TNV (<i>n</i> = 10)	4.95 ± 0.396	1.987 ± 0.12	2.299 ± 0.22	
T. evanescens	$5.6 \pm 0.1 \ (n = 4)$	$2.1 \pm 0.1 \ (n = 3)$	$3.1 \pm 0.2 \ (n = 4)$	

58.75 μ m (*n* = 10). However, the average body length of adult *T. evanescens* was 390 μ m (*n* = 10) as mentioned by Polilov (2016).

Comparing finds of TNV male antennae with *T. evanescens*, results of Polaszek et al. (2012) indicated merely adequate measurements FL/FW (not including setae). In addition, as shown in Table 3, it was inadequate for FL/ScL and LS/FW. Polaszek et al. 2012 and Polilov (2016) described the morphological features of *T. evanescens* wings. TNV differed in stigmal vein than *T. evanescens* (Fig. 12).

Several authors reported that male genitalia can be used to differentiate between egg parasitoid *Trichogramma* spp. (Polaszek et al. 2012 and Del Pino et al. 2013). Comparing TNV male genitalia with that of *T. evanescens*, results of Polaszek et al. (2012) indicated merely adequate measurements AW/GW. Also, as shown in Table 4, it was inadequate for GL/GW and AD/GL.



Fig. 12 Light micrograph of TNV forewing: (sv stigmal vein, mv marginal vein, and smv submarginal vein)

Taxon	Measured characters (µm)			
	Male genitalia (average \pm SD)			
	GL/GW	AD/GL	AW/GW	
TNV (n = 10)	2.52 ± 0.17	0.27 ± 0.04	0.60 ± 0.06	
T. evanescens $(n = 4)$	2.8 ± 0.29	0.3	0.6	

Table 4 Measured characters (μ m) of TNV and *T. evanescens* male genitalia

Conclusion

In conclusion, it is evident that the *Trichogramma* strain TNV was differed morphologically than *T. evanescens*, which is in use for biological control of lepidopteran date palm insect pests in the Egyptian New Valley Oasis. Also, molecular identification of the strain TNV, using ITS-2 region of rDNA and phylogenetic analysis, showed that it is highly homologous to *T. turkestanica* with 100% similarity.

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