Naturalized *Dolichogenidea gelechiidivoris* Marsh (Hymenoptera: Braconidae) complement the

resident parasitoid complex of *Tuta absoluta* (Meyrick) (Lepidopera:Gelechiidae) in Spain

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Abstract

Our study aimed to assess the contribution of natural parasitism due to *Necremnus tutae* Ribes &

Bernardo (Hymenoptera: Eulophidae) to the biological control of *Tuta absoluta* (Meyrick)

(Lepidopera:Gelechiidae) in commercial plots where an IPM program based on the use of predatory mirid

bugs was implemented. During the samplings, the presence of another parasitoid was detected and,

therefore, a second part of our study intended to identify this species and to evaluate the importance of its

natural populations in the biological control of the pest. Leaflets with *T. absoluta* galleries were collected

during 2017–2020 from commercial tomato plots in the horticultural production area of Catalonia

(Northeast Spain), including greenhouses, open fields, and roof covered tunnels that lack side walls. In

the laboratory, *T. absoluta* larvae were classified as ectoparasitized, alive, or dead. Reared parasitoids

from ectoparasitized larvae were mostly morphologically identified as *Necremnus* sp. with parasitism

rates that peaked in summer months with values between 9 and 15%. Some of these ectoparasitized larvae

also yielded another parasitoid identified as *Dolichogenidea gelechiidivoris* Marsh (Hymenoptera:

Braconidae) by both morphological and molecular-DNA barcoding methods. In 2020, parasitism rates

due to *D. gelechiidivoris* that increased with season up to 22%. Our work reports for the first time in

Europe the presence of the neotropical species *D. gelechiidivoris* adding this biocontrol agent to the

resident parasitoid complex of *T. absoluta* in Spain.

Introduction

Tuta absoluta (Meyrick) (Lepidopera:Gelechiidae) is native to South America where has been considered

an important tomato pest since a long time (Larrain 1987). The first report outside its area of origin was

from Spain in 2006 (Urbaneja et al. 2007). From then onwards, the pest did spread over the

Mediterranean basin, and then quickly colonized Africa and Asia (Desneux et al. 2010; Desneux et al.

2011), threatening tomato production (Biondi et al. 2018).

Although insecticides still remain the main control tool in many world areas, many efforts have targeted

sustainable biological control methods (Biondi et al. 2018). In Spain, the positive role of the predatory

mirid bugs *Macrolophus pygmaeus* (Rambur) and *Nesidiocoris tenuis* (Reuter) (Hemiptera: Miridae) was

soon acknowledged (Arnó et al. 2009; Urbaneja et al. 2009). Successful IPM programs based on both

predators already in use at the time of invasion did greatly contribute to the control of *T. absoluta*

(Urbaneja et al. 2012). These two bugs remain as cornerstones for the biological control of several pests

in the area (Arnó et al. 2018).

Surveys of parasitoids that could complement the poor predator action on *T. absoluta* larvae were soon

undertaken in the Mediterranean (Zappalà et al. 2013; Gabarra et al. 2014), and several larval parasitoids

of *T. absoluta* within the Eulophidae, Braconidae, Chalcididae, Ichneumonidae and Pteromalidae were

recorded (Biondi et al. 2018; Mansour et al. 2018). Out of these species, *Necremnus tutae* Ribes &

Bernardo (Hymenoptera: Eulophidae), first identified as *Necremnus* nr. *artynes*, was found consistently

parasitizing *T. absoluta* (Gebiola et al. 2015), and several studies have recognized the contribution of

natural populations to the biological control of the pest (Abbes et al. 2014; Crisol-Martínez and van der

Blom 2019; Arnó et al. in press). This parasitoid attracted the attention of the biocontrol industry, and for

some time it was commercially available, but the high host-killing rate was a serious drawback for

successful mass rearing (Calvo at al. 2016) and production has been discontinued.

To further assess the contribution of *Necremnus tutae* to the biological control of *T. absoluta*, our first

goal was to estimate the natural parasitism in commercial plots where the IPM program based on the use

of predatory mirid bugs was implemented. However, since we observed the presence of another

parasitoid, the second part of our study aimed to identify this species and to evaluate the importance of its

natural populations in the biological control of *T. absoluta*.

Materials and Methods

Samples were collected during 2017–2020 from commercial tomato plots in the horticultural production

area of Catalonia (Northeast Spain), including greenhouses, open fields, and roof covered tunnels that

lack side walls (Table 1). Each plot was sampled up to 21 times depending on crop duration and

T. absoluta infestation levels. Plots were sampled by randomly walking in the plot and collecting leaflets

that had galleries big enough to host a second to third instar larva of *T. absoluta*. Sampling terminated

after 20 minutes or a maximum of 25 leaflets, whichever was reached first, in order to minimize sampling

time and costs, particularly when infestation levels were low (Naranjo 2008).

Leaflets were taken to the laboratory and inspected under a stereomicroscope. *Tuta absoluta* larvae were

classified as "ectoparasitized" (with pupae, larvae or eggs of a parasitoid on the *T. absoluta* larva), "alive"

(not ectoparasitized and able to crawl when touched with a fine brush), or "dead" (not ectoparasitized and

unable to crawl when touched with a fine brush), and the number of each category was recorded. As our

initial interest was on the contribution of *N.tutae* to the control of *T. absoluta*, during 2017, 2018 and

2019, only the ectoparasitized larvae were placed inside Petri dishes (maximum of 3 larvae per dish), and

kept at room temperature for at least 2-3 weeks.

During 2019, the recurrent presence in the samples, albeit at a low rate, of a parasitoid belonging to the

Braconidae was observed. Therefore, additional samples were collected from nine tomato fields in

September. From these samples, alive, dead and ectoparasitized larvae were individualized in Petri dishes

and kept in the climatic chamber (25°C) for a maximum of 42 days until the emergence of *T. absoluta* or

of adult parasitoids.

From the samples collected in 2020, the number of alive, dead, and ectoparasitized *T. absoluta* larvae was

also recorded as before, but only larvae that were alive, with no clearly visible ectoparasitoids, were kept

in order to determine the parasitization rate by endoparasitoids. Larvae were placed in a Petri dish

(maximum of 3 larvae per dish) together with the leaflet and stored at room temperature for a maximum

of 3 weeks until the pupation of *T. absoluta* or the emergence of a parasitoid. Emerged parasitoids from

the four years samples were collected and stored in 70% alcohol for further identification.

Due to the high irregularity of infestation levels in the plots, the number of leaflets collected from each

plot and date (a sample) was very variable, and furthermore, not all the galleries had a *T. absoluta* larvae

inside. Consequently, the total number of larvae (alive, dead or ectoparasitized) collected in each plot was

very variable, and sometimes very low. Leaflets with only empty galleries and no *T. absoluta* larvae were

discarded. To summarize the levels of parasitism, samples were pooled for each month and year. The

percentage of monthly ectoparasitism was calculated by dividing the number of ectoparasitized larvae by

the total number of larvae recorded each month. Additionally, for the 2020 samples of alive *T. absoluta*

larvae, monthly parasitism due to endoparasitoids was calculated dividing the number of emerged

parasitoids by the total number of larvae recorded each month.

Adult parasitoids were first identified to family and sub-family level using the keys by Grissell and

Schauff (1990) and Hanson and Gauld (2006). Eulophidae were further identified to genus using the keys

by Askew (1968) and Gebiola et al. (2015). Microgastrinae (Hymenoptera: Braconidae) were first

identified to genus following the descriptions of Fernandez-Triana et al. (2020), and then as

Dolichogenidea gelechiidivoris Marsh (=*Apanteles gelechiidivoris*) using the description of Marsh

(1975). Additional material examined for morphological identification were 10 ♂♂ and 10 ♀♀, from a

laboratory rearing started with adults that emerged from *T. absoluta* larvae collected in 2019 from several

tomato fields in El Maresme county (31TDF49 to 31TDG92, Catalonia, Spain). These voucher specimens

were prepared and deposited in Naturalis Biodiversity Center (Leiden, The Netherlands).

To confirm the morphological identification of *D. gelechiidivoris*, 11 specimens were sequenced for DNA

barcoding identification. The specimens emerged from samples collected in different locations from

eastern Catalonia along a transect of 100 Km (from 31TDF17 to 31TDG84 and 31TEG03, in the

municipalities of Viladecans (2018), Mataró (2019, 2020), Argentona (2018), Santa Susana (2017, 2019,

2020), Blanes (2017), Fornells de la Selva (2019), and Calonge (2018). An additional specimen from a

previous study (Arnó et al. in press) collected in 2016 in Malgrat de Mar was also DNA barcoded.

Total genomic DNA was extracted from whole insects by using SpeedTools Tissue DNA Extraction Kit

(Biotools, Madrid, Spain) following the manufacturer protocol and eluted in 100 µl of BBE buffer

provided by the manufacturer and stored at -20°C. A 658-bp region of the CO1 gene was amplified with

the following protocol using primers LepF1 5-ATTCAACCAATCATAAAGATATTGG-3 and LepR1 5-

TAAACTTCTGGATGTCCAAAAAATCA-3 (Smith *et al*. 2006). PCR reaction volumes (20 µl)

contained 2 µl of resuspended DNA, 10 µl of Master Mix (Biotools, Madrid, Spain) and 0.4 µl of each

primer [10 µM]. Samples were amplified in a 2720 thermal cycler (Applied Biosystems, CA, USA) using

the thermocycling profile of one cycle of 2 min at 94°C; five cycles of 40 sec at 94°C, 40 sec at 45°C, and

1 min at 72°C; followed by 35 cycles of 40 sec at 94°C, 40 sec at 51°C, and 1 min at 72°C, with a final

step of 5 min at 72°C (Smith *et al*. 2006). PCR products were analysed by electrophoresis in 2.4%

agarose gels stained with GelRed® (Biotium, Hayward, CA) and visualized under UV light. They were

purified with QIAquick PCR Purification kit (Qiagen) and bidirectionally sequenced by using BIGDYE

118 3.1 on an ABI 3730 DNA Analyzer (Applied Biosystems) at the Genomics Unit of the Barcelona Science

119 Park (University of Barcelona). Obtained sequences were compared against the reference database

120 *Barcode of Life Data System* (BOLD, http://www.boldsystems.org/), to find the matching species.

122 **Results**

123 DNA barcoding successfully confirmed the initial identification based on morphological characters of the

124 11 analyzed specimens as *D. gelechiidivoris,* regardless of location and year of collection. Obtained

125 similarity percentages ranged from 100 to 99,48% compared with the 13 *D. gelechiidivoris* available

126 sequences at the time of the analysis (February 2021) in the GenBank database (Accession codes:

127 KX443088, HQ558975-HQ558977, JN282071-JN282078 and JQ849955). . Obtained sequences were

128 also deposited to the GenBank database (Accession codes: MZ298974-MZ298984).

129 Monthly levels of ectoparasitized larvae over the four sampling years are displayed in Table 2. They were

130 recorded from April to November, with levels ranging from 0.1% (May 2017) to 35.7% (November

131 2020). Apart from this exceptionally high value, each year the parasitism peaked during the summer

132 months of August and September with values between 9 and 15%.

133 Most parasitoids that emerged from ectoparasitized larvae from 2017 to 2019 were Eulophids (87%) and

134 a smaller percentage (11%) belonged to Braconidae (Table 3). Among the Eulophidae, eight genera were

135 recorded: *Necremnus* (162 individuals), *Pnigalio* (7), *Neochrysocaris* (5), *Diglyphus* (4), *Stenomesius* (4),

136 *Aprostocetus* (3), *Cirrospilus* (2), and *Sympiesis* (1). Out of the 25 individuals belonging to Braconidae

137 that emerged from larvae that had an ectoparasitoid on them (Table 3), 23 were morphologically

138 identified as *D. gelechiidivoris* same as the four individuals collected in 2016.

139 The additional samples collected in September 2019 yielded 170 *T. absoluta* larvae. From 21

140 ectoparasitized *T. absoluta* larvae emerged 11 *Necremnus* sp., 2 *Diglyphus* sp., 1 *Neochrysocharis* sp.*,*

141 and 1 unidentified Eulophid. From 114 dead *T. absoluta* larvae emerged 6 *Necremnus* sp., and from the

142 35 alive larvae emerged 13 *D. gelechiidivoris.*

143 In 2020, 1,872 alive larvae were collected from 228 samples from 32 plots. From these larvae, we

144 obtained a total of 264 parasitoids that emerged from 92 different samples collected from 20 plots. A

145 subsample of 165 were morphologically identified as *D. gelechiidivoris*, together with one

146 *Neochrysocharis* sp., and one from the subfamily Alysiinae. As can be observed in Figure 1, the

percentage of endoparasitism steadily increased from May until October. No endoparasitoids were

recorded in March, April and November.

Discussion

Results of our four-years field survey confirmed the relevance of Eulophidae as larval parasitoids of *T.*

absoluta in the Mediterranean basin. All genera except *Aprostocetus* had been previously reported

(Biondi et al. 2018). We reared three individuals of this genus from a single sample collected in

September 2017. *Aprostocetus* are parasitic or hyperparasitic in Lepidoptera (Sakaltaş and Tüzün 2014),

and Mirchev et al. (2001) refer to an *Aprostocetus* sp. as an hyperparasitoid of a Braconid parasitizing a

Gelechiid.

Among Eulophidae, our results corroborate the importance of *Necremnus* sp. as the most widespread

ectoparasitoid of *T. absoluta* larvae in the Mediterranean, as reported before (Ferracini et al. 2012;

Zappalà et al. 2013; Gabarra et al. 2014; Gebiola et al. 2015; Biondi et al. 2018). In fact, *N. tutae*, which

is by far the predominant *Necremnus* species in the area of our survey (Arnó et al. in press), was

considered a promising parasitoid to be released for *T. absoluta* control (Calvo et al. 2013; Chailleux et

al. 2014; Bodino et al. 2016; Calvo et al. 2016), although currently it is not commercially available. Rates

of ectoparasitized larvae found in the present study, mostly due to *Necremnus* sp., were similar to those

recorded in other field studies in the Mediterranean basin. In the same area of the present study,

parasitism rates by *N. tutae* in sentinel plants were close to 20% (Arnó et al. in press), and in Tunisia it

was of 26% in sentinel plants, and between 11 and 15% when sampling the crop (Abbes et al. 2014).

Much higher rates, up to 73%, were recorded in tomato greenhouses in the southeast of Spain (Crisol-

Martínez and van der Blom 2019).

Contribution of *N. tutae* to the control of *T. absoluta* goes further than only parasitization. As many

Eulophidae, it kills more larvae than parasitizes in order to obtain nutrients that have a strong positive

effect on its reproduction (Ferracini et al. 2012; Balzan and Wäckers 2013; Calvo et al. 2013; Chailleux et

al. 2014; Calvo et al. 2016; Bodino et al. 2019). In the present study, although the number of dead larvae

was recorded, we could not determine the exact causes of mortality because even all sampled plots were

managed according to an IPM program based on predatory mirids, insecticide applications were also

occasionally required.

An important outcome of our study was the detection of *D. gelechiidivoris* in field collected samples from

2017 to 2020, and also in one sample collected in 2016. *Dolichogenidea gelechiidivoris* is native of South

America, where it is considered as an important agent of biological control (Larrain 1987, Agudelo and

Kaimowitz 1997; Vallejo 1999; Salas-Gervasio et al. 2019). In Colombia, mass rearing protocols were

developed to release this parasitoid for *T. absoluta* control (Morales et al. 2013). Furthermore, in 2017 it

was imported to Kenia from Peru to contribute to the control of *T. absoluta* in Africa (Aigbedion-Atalor

et al. 2020). Another species, *Dolichogenidea appellator* (Telenga) (= *Dolichogenidea litae* (Nixon)), was

occasionally found parasitizing *T. absoluta* in the same area as the present study (Gabarra et al., 2014),

and was found also associated with this pest in Sudan (Mansour et al. 2018).

To our knowledge, this is the first report of *D. gelechiidivoris* naturally occurring outside its area of

origin. Since there is no record of intentional introduction of *D. gelechiidivoris* into Europe, and the

importation to Africa took place in 2017, one year after our first detection of this species in 2016, the

results of our survey suggest that this parasitoid was unintentionally introduced from the Neotropics

together with the pest. Accidental introductions of natural enemies in new territories are not strange. Roy

et al (2011) stated that most of the alien arthropod predator and parasite species in Europe arrived

accidentally, as part of worldwide movement of invasive pests that is facilitated by global trade. Trade of

infested fruits has been pointed out as the most probable cause of the arrival of *T. absoluta* to Spain from

South America and the quick spread of the pest (Desneux et al. 2010).

As all Microgastrinae, *D. gelechiidivoris* is a koinobiont solitary larval endoparasitoid (Fernandez-Triana

et al. 2020), and the host remains alive until the end of the parasitoid development. Of the 2017–2019

samples of *T. absoluta* larvae, we only kept those that had an ectoparasitoid egg or larva on them, but

11.5% of the emerged parasitoids were *D. gelechiidivoris* (Table 3). This suggests that there is no clear

recognition of previous parasitism between ectoparasitoids (mainly *Necremnus* sp.), and *D.*

gelechiidivoris.

When we maintained all alive larvae to record parasitoid emergence (additional 2019 samples, and all

2020 samples), the endoparasitism rate was 7.6% in September 2019 and increased up to 22% from May

to October 2020 (Fig. 1). However, the real rate of parasitism by *D. gelechiidivoris* in 2020 was probably

underestimated since about 11% of ectoparasitized larvae collected between 2017 and 2019 yielded *D.*

gelechiidivoris, and ectoparasitized larvae of 2020 had been discarded.

In this scenario of coexistence of several natural enemies, the interactions between *D. gelechiidivoris* and *Necremnus sp.* will be of great importance. For example, young females of the ectoparasitoid *Dineulophus phthorimaeae* De Santis (Hymenoptera: Eulophidae) avoided multiparasitism on the microgastrinid endoparasitoid *Pseudapanteles dignus* (Muesebeck) (Hymenoptera: Braconidae), but older females did not discriminate heterospecific parasitized *T. absoluta* larvae, and joint action of both parasitoids exerted an important control of *T. absoluta* (up to 80% of host larvae mortality) (Salas-Gervasio et al., 2019). The outcome of competition between parasitoids attacking the same host depends on many factors that may explain the dominance of one parasitoid over another, e.g. where the venom of idiobiont ectoparasitoids has little or no effect on the development of endoparasitic koinobionts (Harvey 2013), although in multiple parasitisms between an ectoparasitoid and an endoparasitoid, the former normally wins (Mitsunaga and Yano 2004). Furthermore, the interaction between *D. gelechiidivoris* and predatory mirids will be also of interest. These predators prefer to prey on eggs but may also feed on young *T. absoluta* larvae (Arnó et al. 2009; Urbaneja et al. 2009), which are the preferred host instar of the parasitoid (Aigbedion-Atalor et al. 2020). *Nesidiocoris tenuis* did not prey on nor did reduce the oviposition by *D. gelechiidivoris*, and the efficacy of both natural enemies together on *T. absoluta* was significantly higher than either natural enemy alone (Aigbedion-Atalor 2020). The outcome of the interactions among these biocontrol agents will be determinant for a more successful control of *T. absoluta* (Tarusikirwa et al. 2020).

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Table 1: Number of plots surveyed, number of samples that had leaflets with at least one *T. absoluta*

larva, and week of the year when first and last samples were taken. Each sample corresponds to a

maximum of 25 leaflets during up to 20 minutes from one crop on one date.

Year	Number of		Week of the year	
	Plots	Samples	First	Last
2017	32	235	17	41
2018	61	296	11	40
2019	37	186	12	39
2020	32	228	12	46

354 **Table 2.** Percentages of ectoparasitized *T. absoluta* larvae recorded from tomato leaflets collected in

355 commercial tomato plots. Percentages were calculated as the number of ectoparasitoized larvae over the

356 total number of larvae collected per month. In brackets are the number of samples that had leaflets with at

357 least one *T. absoluta* larvae.

Month	% ectoparasitized larvae				*T. absoluta* larvae (num. samples)			
	2017	2018	2019	2020	2017	2018	2019	2020
Mar	-	0	0	0	-	480 (25)	113 (12)	11 (3)
Apr	0	0.8	1.0	0	32 (5)	612 (47)	308 (22)	14 (6)
May	0.1	3.0	5.0	0	714 (50)	833 (52)	360 (25)	148 (19)
Jun	0.6	7.3	6.2	0.3	519 (52)	575 (43)	389 (27)	398 (48)
Jul	4.9	5.2	4.3	0.6	409 (53)	515 (55)	531 (45)	530 (43)
Aug	10.6	5.5	13.5	1.8	256 (31)	455 (48)	319 (36)	610 (47)
Sep	11.6	14.6	7.6	9.2	242 (27)	213 (21)	197 (19)	370 (40)
Oct	9.1	3.1	-	7.4	186 (16)	65 (5)	-	202 (17)
Nov	-	-	-	35.7	-	-	-	14 (5)

360 **Table 3**: Number of adult parasitoids of different Hymenoptera families reared from ectoparasitized *T.*

361 *absoluta* larvae in samples collected from commercial plots in the different years of sampling.

Hymenoptera family	Year of sampling		
	2017	2018	2019
Eulophidae	23	68	98
Braconidae	5	8	12
Torymidae	1	0	0
Platygastridae	1	0	0
Diapriidae	0	1	0
Aphelinidae	0	0	1

366 **Figure 1:** Monthly percentage of endoparasitized *T. absoluta* larvae in commercial tomato plots in 2020.

367 Percentages were calculated as the number of emerged parasitoids over the total number of larvae

368 recorded per month. Total number of *T. absoluta* larvae and number of samples are displayed in Table 2.