

TRANSMISSION OF ALFALFA MOSAIC VIRUS THROUGH *SUBCOCCINELLA VIGINTIQUATORPUNCTATA* (L.) (COLEOPTERA: COCCINELLIDAE) IN ALFALFA GROWING AREAS IN TÜRKİYE

A. Barış^{1,*}, A. F. Morca¹ and M. Alkan²

¹Directorate of Plant Protection Central Research Institute, Ankara, Türkiye

²Yozgat Bozok University, Faculty of Agriculture, Department of Plant Protection, Erdogan Akdag Campus, Türkiye

*Corresponding author's e-mail: aydemirbaris01@gmail.com

ABSTRACT

Alfalfa *Medicago sativa* L. (Fabaceae: Leguminosae) is an important fodder crop due to its highly nutritious animal feed and potential to adapt various environmental conditions. The average alfalfa yield in Türkiye is considerably higher than the world's average. Several diseases and pests exert negative impacts on alfalfa yield. This study investigated the status of the alfalfa mosaic virus (AMV) and the presence of insect species that could be new vectors for AMV in the alfalfa growing areas of Zonguldak and Bartın provinces in Türkiye. AMV is non-persistently transmitted by aphid species, but it is also known to be transmitted by pollen and seeds. The most prevalent harmful insect species in the studied area were *Subcoccinella vigintiquatorpunctata* (L.) (Coleoptera: Coccinellidae), and *Gonioctena fornicata* (Brüggeman) (Coleoptera: Chrysomelidae). The AMV was prevalent in the tested plants at a rate of 85.00%, with infection rates of 80.55% in mature larvae of *S. vigintiquatorpunctata*, 80.00% in pupae, and 96.77% in adults. When looking at the relationship of AMV with common pest species, AMV was not detected in all life stages of *G. fornicata*. In contrast, AMV was detected in all life stages of *S. vigintiquatorpunctata* except eggs. This study is the first in the world to detect the presence of AMV in *S. vigintiquatorpunctata*. However, further studies are needed to determine whether *S. vigintiquatorpunctata* transmits AMV persistently, semi-persistently, or non-persistently.

Keywords: Forage crops, 24-spot ladybird, Molecular detection, Insect Vector, Prevalence

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INTRODUCTION

Alfalfa (*Medicago sativa*) is a perennial forage crop known as the queen of fodders. It is cultivated on >30 million hectares' temperate regions of the world (Acharya *et al.*, 2020; Li *et al.*, 2020; Tuck *et al.*, 2020). It is regarded as a superior forage crop and contains high amount of protein, minerals, vitamins, and a balanced amino acid composition (Wang *et al.*, 2015). Alfalfa is a natural host for different plant viruses, like several other crops (Samac *et al.*, 2015).

Thirty-one viruses are known to infect alfalfa crops (Usta and Güller, 2020). Natural plant openings or wounds are required for the pathogens to penetrate the plant. Insects are part of the disease complex because feeding wounds constitute the entry point for plant pathogens (Willsey *et al.*, 2017). Several modes of virus transmission exist, including plant contact, vegetative reproduction, and vector-mediated transmission. Insects acquire and spread viruses by feeding on infected plants, vomiting sap and virus particles, and depositing them in chewing wounds; the efficiency and duration of virus retention varies among insect species. Insects are vital

virus vectors and more than 70 species are known to transmit economically important crop viruses. They come from families such as Chrysomelidae, Coccinellidae, Curculionidae, and Meloidae. Both adult insects and larvae can effectively transmit plant viruses thanks to their feeding behavior and modes of transmission that facilitate the spread of the virus within plants (Wielkopolan *et al.*, 2021).

Alfalfa mosaic virus (AMV) reduces the alfalfa crop's production, quality durability, and causes significant economic damage. The AMV has the largest range of hosts among plant viruses as it infects 698 species belonging to 167 genera in 71 families (Hull, 1969; Edwardson and Christie, 1997). However, the main hosts for the AMV are *M. sativa*, *Phaseolus vulgaris* L., *Pisum sativum* L., and *Nicotiana tabacum* L. (Mangeli *et al.*, 2019). The AMV infection in alfalfa causes serious crop losses with mosaic spots, yellowing in the form of oak leaves, vein banding, and different deformities in addition to severe stunting. Several studies have indicated that AMV causes 30 to 100% yield loss in alfalfa, depending on ecological conditions and the strain of the virus (Zschau and Janke, 1962; Jaspars and Bos, 1980;

Van Regenmortel and Pinck, 1981). The AMV is a multipartite virus composed of three linear RNAs (RNA1, RNA2, and RNA3) and a non-infective subgenomic RNA (sgRNA4). The four viral RNAs are encased in individual bacilliform particles. The RNA1 and RNA2 are viral replicas subunits. These contain a single open reading frame (ORF) encoding (P1 and P2, respectively). The RNA3 contains two ORFs encoding movement protein and coat protein (CP). (Scott *et al.*, 1998; Moradi and Mehrvar, 2021).

The virus can spread rapidly from one plant to another. It is known that the AMV is non-persistently transmitted by at least fourteen aphid species. The prevalent aphid species in Zonguldak and Bartın province of Türkiye are *Acyrtosiphon pisum* (Harris), *Aphis gossypii* Glover, and *Therioaphis (Pterocallidium) trifolii* (Monell) (Hemiptera: Aphididae) (Barış *et al.*, 2017). Apart from the aphid species, the AMV has also been reported to transmitted by pollen, *Cuscuta* spp. and seeds (Frosheiser, 1974; Hemmati and McLean, 1977).

Larvae and adults of 24-spot ladybird, *Subcoccinella vigintiquatuorpunctata* (L.) (Coleoptera: Coccinellidae), feed on the leaves of host plants. They eat the lower epidermis and parenchyma of the leaves, whereas the upper epidermis remains in the form of a membrane (Barış, 2021; Barış, 2022). The damaged leaves appear in the form of lace, which can cover the whole plant under severe infestation. Alfalfa forage yield can be reduced by 40 to 60% as a result of the 24-spot ladybird infestation (Keresi and Sekulic, 2005). The ladybird adults overwinter in the soil near host plants (Marriner, 1927; Tanasijevic, 1958; Richards *et al.*, 1976; Wheeler and Henry, 1981; Baldwin, 1988). The other significant pest alfalfa leaf beetle *Gonioctena fornicata* (Brüggemann) (Coleoptera: Chrysomelidae) is substantial pests of alfalfa crop. It is a widespread species in Asia, Siberia, Europe, North Africa, Russia, and Western Asia (Horion, 1961). It has more than 70 hosts, especially in alfalfa (Richards *et al.*, 1976). Bodenheimer (1958) reported that its presence in Türkiye for the first time and named it as 'Clover beetle'. It causes significant crop losses, especially in legumes. The larvae and adults of *G. fornicata* cause significant yield losses in alfalfa. Barış (2021) reported that larvae and adults feed on young shoots, leaf buds, flowers, leaves and the tips of alfalfa stems. The alfalfa leaf beetle was firstly reported by Alkan (1946) in Türkiye. Afterwards, Bodenheimer (1958) regarded it alfalfa leaf beetle and stated that there is an epidemic risk of the pest which would exert significant economic damages.

This study, it was aimed to determine the infection rate of AMV in alfalfa plants, which is a problem in alfalfa cultivation areas in Türkiye and causes yield losses, as well as the detection of pest species in alfalfa cultivation areas and to test these detected pest species for the possible presence of AMV virus.

MATERIALS AND METHODS

Survey studies: Field sampling was carried out in alfalfa fields in the Western Black Sea Region of Türkiye (Bartın and Zonguldak provinces) every two weeks in April, May, and June in 2013-2014 (Table 1). Eggs were collected in April, mature larvae in May, pupae in May-June, and adults in June. The altitude of the surveyed alfalfa fields varied between 55 and 575 m. The green parts of the alfalfa plant were carefully examined with macro-observations, and the presence of any life stage of the harmful species (Çalışkaner *et al.*, 1989; Tamer *et al.*, 1997) and the symptoms caused by AMV in the plant were recorded. As a result of macro-observation, plant parts and insects including, eggs, larvae, pupae and adults of the pests were taken in a paper bag and brought to the plant pest laboratory of the Plant Protection Central Research Institute. The insects were separated according to their stages and taken into tubes containing 99% ethanol in the laboratory. These tubes were then labeled and stored in the +4°C refrigerator, and the plant parts were stored in the deep freezer at -20°C for further analysis.

Eggs, mature larvae, and adult stages of the 24-spot ladybeetle observed on alfalfa plants during the current study are shown in Fig. 1.

Different life stages of (egg, mature larvae, and adult) the harmful insect species were recorded and collected with the infected plant parts. The collected samples were transferred to eppendorf tubes containing 99% ethanol. Afterward, these tubes were brought to the laboratory by placing them in polyethylene bags tagged with the collection date, location, and the number of host plants. The pupae of the pest were cultured in breeding cages at 25±2°C temperature, 16:8 light: dark period, and 65±5% relative humidity under laboratory conditions. The pupae were observed for 8 every hour intervals and taken into 99% ethyl alcohol for analysis before they started feeding.

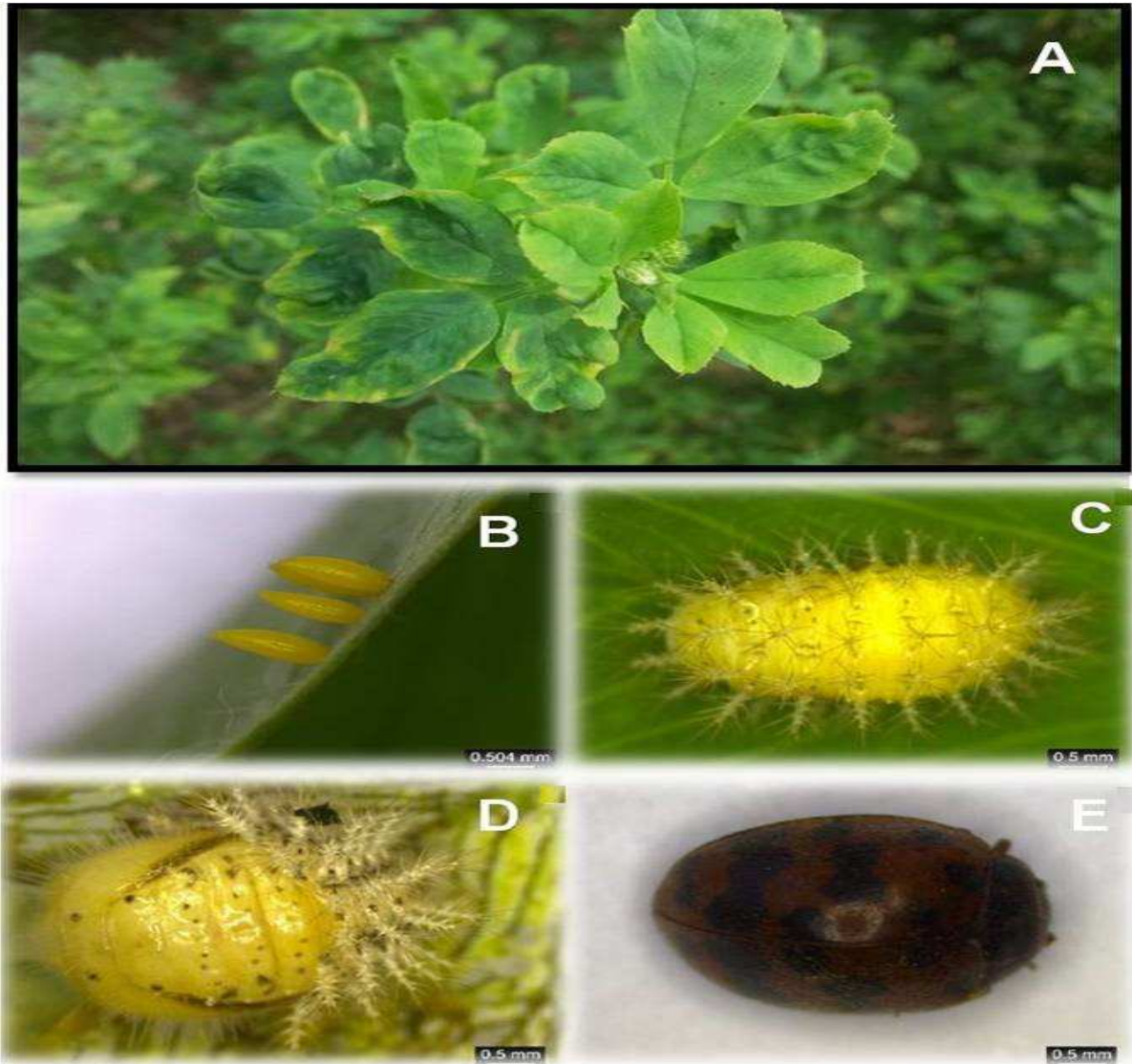
The prevalence of pests was recorded, and the infection rate calculated by Bora and Karaca, (1970).

$$\text{Prevalence rate} = \frac{\text{Sampled area}}{\text{Total area}} \times 100$$

Total RNA extraction: The alfalfa plant samples and different life stages (eggs, mature larvae, pupae, and adults) of the harmful insect species collected during surveys were analyzed simultaneously. In insects, each individual was considered separately, and in eggs, each cluster was considered a single sample. RNA isolations for all samples were made according to the silica gel method (Foissac *et al.*, 2001). The concentrations of the obtained total RNAs were adjusted using Nanodrop (Thermo Scientific, USA).

Table 1. Information on the geographical coordinates of the province and districts where the materials were obtained.

Bartın province	Zonguldak province
Merkez 41°32'41.0"N 32°20'38.0"E 95 m 41°30'33.0"N 32°21'02.0"E 88 m 41°33'14.8"N 32°20'09.3"E 55 m 41°28'28.5"N 32°17'53.1"E 76 m 41°28'51.6"N 32°18'13.7"E 84 m	Eceler 41°20'39.4"N 32°01'30.1"E 115 m 41°20'30.8"N 32°00'48.7"E 153 m 41°20'03.5"N 32°00'32.1"E 191 m
Ulus 41°32'52.8"N 32°36'21.3"E 126 m 41°31'56.2"N 32°35'37.1"E 175 m 41°32'21.6"N 32°38'00.8"E 197 m 41°37'58.9"N 32°38'44.6"E 457 m 41°38'56.4"N 32°41'03.0"E 575 m	Kayıkçılar 41°22'44.2"N 32°05'35.5"E 66 m 41°23'06.3"N 32°05'43.6"E 70 m 41°23'30.8"N 32°05'38.0"E 57 m
Amasra 41°43'22.8"N 32°26'47.3"E 182 m 41°43'20.8"N 32°26'56.4"E 187 m	Nebiöğlü 41°26'08.9"N 32°15'09.3"E 102 m 41°26'33.8"N 32°15'21.6"E 110 m 41°26'37.1"N 32°14'39.5"E 93 m
Yılanlıca 41°16'07.7"N 31°58'32.7"E 142 m 41°16'14.2"N 31°58'17.7"E 133 m 41°16'12.5"N 31°57'57.7"E 153 m	
Yassören 41°17'44.5"N 31°56'09.7"E 443 m	
Bölücek 41°18'42.6"N 31°58'54.3"E 416 m 41°19'01.2"N 31°59'06.5"E 390 m	
Karaman 41°21'34.0"N 31°58'57.7"E 240 m 41°21'25.9"N 32°00'43.1"E 141 m 41°21'09.8"N 32°00'56.6"E 131 m	

**Figure 1. Alfalfa plant with AMV infection symptoms (a), eggs (b), larva (c), pupa (d) and adult (e) of *Subcoccinella vigintiquatuor punctata***

RT-PCR (Reverse Transcription Polymerase Chain Reaction): The resulting total RNAs were analyzed in a single-stepped RT-PCR (Xu and Nie, 2006). The primer pairs (AMV-F 5' - CCATCATGAGTTCTTCACAAAAG-3' and AMV-R 5' -TCGTCACGTCATCAGTGAGAC-3') were used for the amplification of the coat protein genome region of AMV. The RT-PCR was performed according to following conditions: total volume of 25 µl PCR mix: 8 µl 5X GoTaq Flexi buffer, 1.2 µl MgCl₂ (25 mM), 0.7 µl dNTP (10 mM), 0.25 µl GoTaq polymerase enzyme (5 U µl), 1 µl forward primer (10 µM), 1 µl Reverse primer (10 µM), 0.2 µl Reverse-transcriptase enzyme (200 U/µl), 0.2 µl RNase inhibitor (5000 U/ml), 2 µl RNA prepared to total volume with sterile water without nuclease. The reaction conditions for the one-step RT-PCR were; 60 min at 37 °C, 5 min at 94 °C, 30 s at 94 °C, 30 s at 57 °C, 45 s at 72 °C, and 10 minutes at 72 °C. The amplicons obtained with the primer pair were run on a 1.5% agarose

gel prepared with Pronosafe (Conda, Madrid, Spain) DNA dye at 80 V for 60 minutes and visualized under UV transilluminator.

RESULTS AND DISCUSSION

During the research, two different harmful species, *S. vigintiquatuor punctata* and *G. fornicata*, were detected in alfalfa fields. The prevalence rate of *G. fornicata*, and *S. vigintiquatuor punctata* was 100%. Similarly, the AMV prevalence rate in the alfalfa samples was 85.00% (Table 2).

Due to the high level of AMV infection in the same fields, the relationship between the pest and the virus was evaluated. As a result of RT-PCR studies, while AMV was not detected in all life stages of *G. fornicata*, the presence of AMV was detected in all life stages of *S. vigintiquatuor punctata* except egg (Fig. 2).

Table 2. The RT-PCR testing of harmful insect species recorded from alfalfa fields in Zonguldak and Bartın provinces during 2013-2014

Locations	<i>Suboccinella vigintiquatuor punctata</i>				<i>Gonioctena fornicata</i>			Alfalfa	
	Number of								
	Eggs	Larvae	Pupae	Adults	Eggs	Larvae	Adults	Plant samples	
Zonguldak	Yassiören	75	15	18	25	60	18	14	15
	Bölücek	69	18	22	34	82	12	18	12
	Karaman	82	14	16	32	58	10	22	17
	Eceler	60	12	14	38	54	22	20	13
	Kayıkcılar	36	20	12	33	72	20	19	18
Bartın	Nebioğlu	62	16	18	42	64	17	15	14
	Merkez	68	17	20	40	66	14	14	8
	Ulus	74	14	16	37	48	11	18	13
	Amasra	34	18	19	29	42	10	12	10
	Total*	560	144	155	310	546	134	152	120
AMV-positive samples	0	116	124	300	0	0	0	102	
AMV infection rate (%)	0.00	80.55	80.00	96.77	0.00	0.00	0.00	85.00	

*Total number of eggs, insects and alfalfa plants collected during the surveys

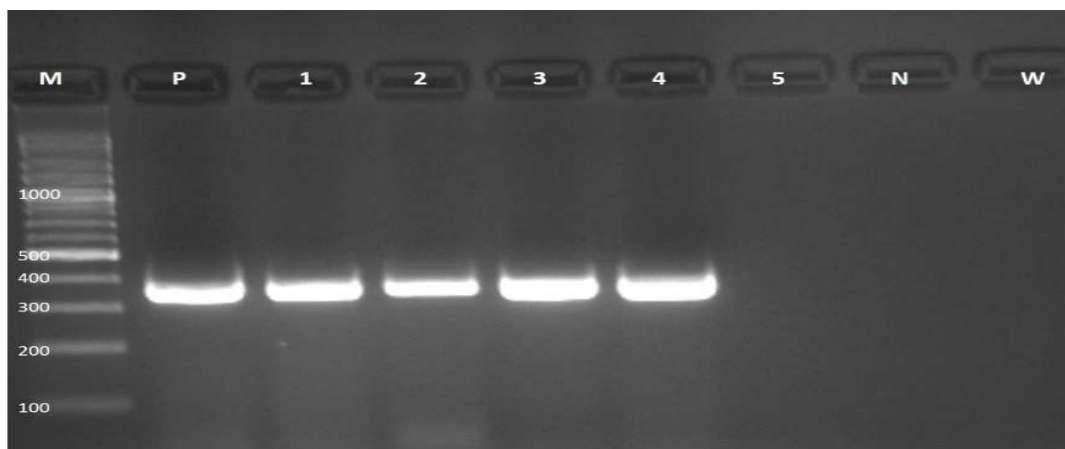


Figure 2. Agarose gel electrophoresis of RT-PCR amplification product for *Suboccinella vigintiquatuor punctata* using AMV-F/R primer pairs. In the upper figure panel; M=DNA 100 bp ladder, P= positive control, 1=mature larvae, 2= alfalfa plant sample, 3=pupae, 4=adult, 5=egg, N= negative control and W= water control.

The RT-PCR confirmed the presence of AMV in 80.55% of the tested *S. vigintiquatuorpunctata* mature larvae, 80.00% of pupae and 96.77% of the adults. The highest AMV presence in adults can be linked to adult flights and feeding from different plants. Previous research has demonstrated that an insect can obtain a virus from plant tissue through a single bite. However, the effectiveness of virus acquisition rises as the insect engages in more extensive feeding, as noted by Fulton and Scott in 1977. Moreover, insects are capable of ingesting and transmitting the virus within a matter of seconds, and the duration of virus retention varies between 1 and 10 days, contingent upon the insect species, according to Agrios (2008).

On the other hand, no natural enemies of *S. vigintiquatuorpunctata* were found in alfalfa fields in the present study. However, the larval parasitoid *Tetrastichus* sp. (Hymenoptera: Eulophidae) has been reported from other parts of the world (Wheeler and Henry, 1981). The presence of AMV in all locations where *S. vigintiquatuorpunctata* was discovered is believed to be attributed, in part, to the absence of natural predators for this insect within the examined fields. Analysis of *S. vigintiquatuorpunctata* larvae for the presence of AMV virus showed that the virus was also present in larvae. In parallel with this study, similar studies on the presence of viruses in larvae have been recorded in the literature. Nault (1978) found that larvae of *Oulemamelanopus* (Chrysomelidae) transmit maize chlorotic mottle virus (MCMV, Tombusviridae) more efficiently than adults. Additionally, *O. melanopus* and *O. gallaeciana* can effectively transmit cocksfoot mottle virus (CfMV, Solemoviridae) up to 15 days after acquisition (Catherall, 1987). Furthermore, Musser *et al.*, (2003) noted that *Epilachna varivestis* (Coccinellidae) exhibits a preference for feeding on plants that have undergone visual changes due to infections by bean pod mottle virus (BPMV, Secoviridae) and southern bean mosaic virus (SBMV, Solemoviridae). The observations of the current study suggest that the pest ingests the virus through diet during feeding. The non-infected eggs indicate that AMV is not transmitted from one generation to the other in *S. vigintiquatuorpunctata*.

In the observations made for *G. fornicata*, it was determined that only adult individuals were fed with alfalfa, which caused yield losses (Bariş *et al.*, 2021). However, the AMV was detected in *G. fornicata* feeding plants, but the virus was not found in any stage of this pest. The ability of an insect to serve as a vector has been reported to depend on several factors, including specific molecular interactions between the virus and the insect host (Chen *et al.*, 2015). As stated in Chen *et al.*, (2015), the interaction between the non-structural protein of a virus and the cytoplasmic actin of leafhoppers is related to insect vector specificity. In this study, we did not find

AMV in *G. fornicata*, which is thought to be due to molecular incompatibility between host and pathogen.

It is known that aphids are the only vectors for transmission of AMV. There are >20 aphid species that can transmit AMV. Among these species, *Aphis pisum*, *A. craccivora*, *A. fabae*, and *Myzus persicae* are the most important and transmit AMV non-persistently (Edwardson and Christie, 1997). Although aphid species were not prioritized in this study, observations showed that the density of some aphid species was very low in samples collected from alfalfa fields. It can be said that natural enemies suppress aphid populations. Aphid plays an important role in transmitting of AMV; however, no pest management strategies, the presence of natural enemies, and their activities lower the aphid population (Katis *et al.*, 2007). The prevalence of AMV in alfalfa causes significant damage. Several natural enemies, especially those in the Coccinellidae family, significantly suppress aphids population in Türkiye (Kök *et al.*, 2020). Kaygın and Kaptan (2017) reported that 14 Coccinellidae species from Bartın province and the species with the highest prevalence rates were; *Coccinella septempunctata* (L.), *Harmonia axyridis* (Pallas), *Scymnusquadri guttatus* (Capra), *Halyziasedecim guttata* (L.), *Oenopiacon globata* (L.), *Propyleaquatuordecim punctata* (L.), and *Adaliade punctata* (L). The *C. septempunctata* was commonly observed from both provinces during the surveys. In alfalfa fields, no pest management strategy was favored that was considered effective in promoting beneficial fauna and suppressing aphid populations.

In conclusion, this is the first report that the AMV was only recorded from *S. vigintiquatuorpunctata* in the current study. Insects can serve as vectors for the transmission of viruses, including plant viruses. The transmission of viruses by insect vectors can occur through circulative or non-circulative modes, depending on the specific interactions between the virus and the insect host. Transmission studies under controlled conditions are needed to confirm the vector status of the insect. Understanding these interactions is crucial for developing strategies to control the spread of plant diseases and protect plants from viral infections.

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