



## Parasitological and molecular study of nosemosis in migratory apiaries in Hormozgan Province, southern Iran

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**Abstract.** Nosemosis is a microsporidian disease caused by *Nosema ceranae* and *N. apis* and transmitted via oral-fecal and oral-oral routes. It is globally distributed among adult bees in honeybee colonies. Considering the health importance of nosemosis in honeybees, the study aims to determine the frequency of *Nosema* spp. infection in migratory apiaries in Hormozgan province by microscopic and molecular methods. In the present study, 20 bees from ten randomly selected hives in 84 migratory apiaries were collected. In the laboratory, the abdomen of the bees was separated from the rest of the body with entomological tweezers and scissors and then ground up in a mortar containing saline serum. The prepared suspension was filtered by passing through a sieve, then the prepared suspension was transferred to test tubes and centrifuged. The pellets were repeatedly washed by saline solution and centrifuged. Finally, the pellets were examined for spores of *Nosema* spp. by light microscopy and conventional PCR. In microscopy, 38.2 % of apiaries were positive for *Nosema* spp. spores. By PCR however, DNA of *N. ceranae* was detected in 39.2. % of apiaries with no samples positive for *N. apis*. Due to the considerable frequency of infection in migratory apiaries in Hormozgan province, it is necessary to carry out appropriate health measures such as screening of apiaries with appropriate diagnostic methods and training of beekeepers to disinfect hives in order to control *Nosema* infection in Iranian apiaries by the veterinary health officials.

**Keywords:** Insect pathology, characterization, honeybee, molecular detection

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## Introduction

Nosemosis is one of the most destructive diseases of adult honeybees around the world (Bailey & Ball, 1981). The causative agents are unicellular microsporidian parasites *Nosema cernea* and *Nosema apis* (WOAH, 2019). Adult honeybees become infected during consumption of food and water contaminated with spores, during cleaning contaminated combs, robbing contaminated hives, or by infected bees drifting to new hives (Galajda *et al.*, 2021). The wall of ingested spores is broken by the enzymes of honeybees' gut and then sporoplasm is injected into the cell mid-gut by polar tubes. *Nosema* spp. grows and produces several million spores and damages the intestinal cells. Honeybees infected with *N. apis* may be shown dysentery, but bees infected with *N. ceranae* show no symptoms. They often die away from the hive and only a few sick or dead bees may be found

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near the hive entrance (Galajda *et al.*, 2021). *Nosema apis* infection is more distributed among apiaries located in cold and temperate climates during spring and winter seasons, while *N. ceranae* infection is in tropical and subtropical climates (Fries *et al.*, 1996). The prevalence of nosemosis has been reported in the range of 23% to 53% in honeybee colonies from different climatic regions of Iran (Mohammadian *et al.*, 2018). Hormozgan province is the southernmost region of Iran with a tropical climate and mild weather which is suitable for the growth of flowering plants in the winter season. Every year, many honeybee colonies from northernmost regions migrate to Hormozgan province in winter. Considering the effect of nosemosis on the health of honeybee colonies, the study aimed to determine the rate of *Nosema* spp. in the migratory honeybee colonies in the Hormozgan province using microscopical and molecular tools.

## Materials and methods

### Study area

Hormozgan province is located in the South of Iran, the North of Hormoz Strait, with geographical coordinates between 25° 23' to 28° 57' N and 52° 41' and 59° and 15' E. The boundaries of the province lie on the Oman Sea from the East to the Persian Gulf on the West. More than 70% of the province is covered by mountains and hills. The province is bounded by Kerman province in the north and northeast, Fars and Bushehr provinces in the west and northwest, and Sistan and Baluchistan province in the east. Three types of climates exist in this province, i.e., forest, rangeland, and desert. The average temperature affected by humidity is moderate and rarely gets higher than 45 °C in summer. In the deserts, the temperature is about 0 °C but there is no frigid weather in winters. The annual rainfall is less than 250 mm and the relative humidity is more than 80% (Safa *et al.*, 2012)

### Sample collection

The sample size was calculated 840 hives. The sample size was calculated using a 98% confidence level with 4% desired absolute precision (Thursfield, 1986), based on the prevalence of *Nosema* spp. infection (54%) that was previously reported in humid climate (Mohammadian *et al.*, 2018). After visiting, we obtained the data such as the apiary address, the name of the owner, and bee population from the beekeeper. Then, the samples were collected from 10 seemingly healthy hives in each apiary, consisting of 20 old worker bees from peripheral frames of each hive (200 honeybees in each apiary) (WOAH, 2019). The collected bees were put in storage containers and transported immediately to the laboratory under cold conditions. From December 2022 to March 2023, 84 migratory apiaries were sampled one to two weeks after entering migratory apiaries in autumn and winter to Hormozgan province (Fig. 1) No clinical symptoms related to nosemosis were observed in hives of each apiary during sampling.

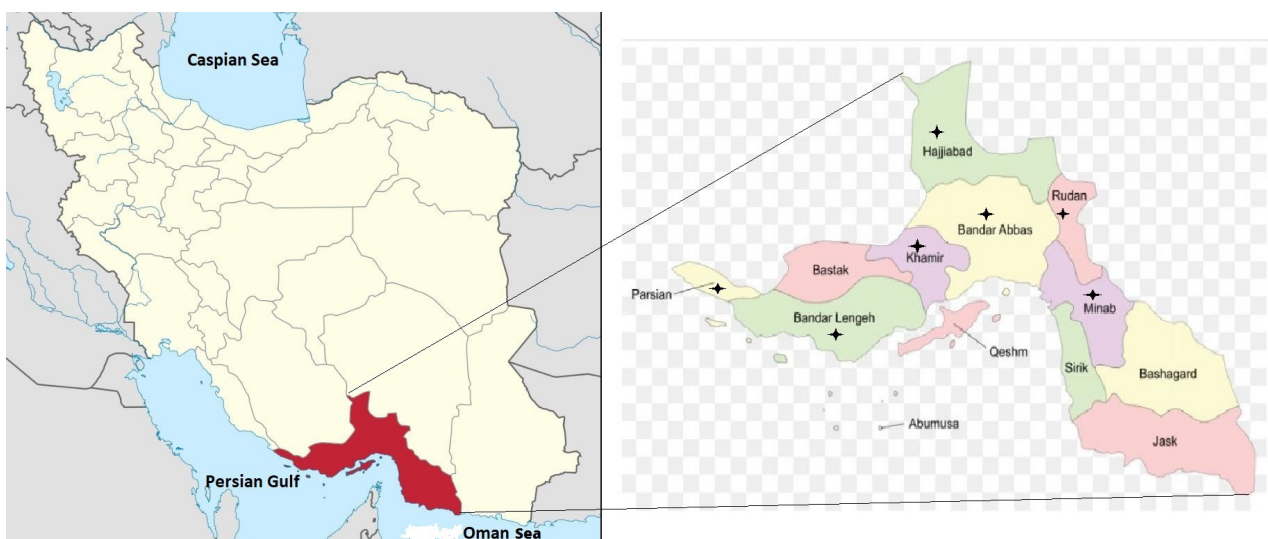


Fig. 1. Sampling areas in Hormozgan province.

## Microscopy Examination

The collected bees were put in storage containers and transported immediately to the laboratory under cold conditions. Abdomens of 20 honeybees from each hive were washed with normal saline solution and ground in 5 ml of this solution. The suspensions were then filtered through two layers of muslin to remove coarse bee parts and centrifuged at 2500 g for 5 min. The supernatants were removed and pellets were mixed with saturated saline solution and again centrifuged at 2500 g for 5 min. One milliliter of supernatants was taken, and the rest of the solution was discarded. The supernatants were washed three times with distilled water and each time they were centrifuged at 2500 g for 3 min and the upper parts were discarded. The final pellets were resuspended in 1.5 ml of distilled water. Three drops of the suspension were put on a microscopic slide covered with an 18×18 mm coverslip and examined by a light microscope at ×400 magnification (Shirzadi & Razmi, 2021). The rest of the homogenate was transferred to an Eppendorf tube and kept at -20°C until molecular examination.

## DNA extraction and PCR assay

Genomic DNA of samples was extracted with a commercial DNA extraction kit (MBST, Tehran, Iran) according to manufacturer's instructions. A multiplex PCR targeting 16S rRNA region was employed for simultaneous detection of two *Nosema* species in a single reaction (Martín-Hernández *et al.*, 2007) (Table 1). In positive samples, 321 bp product of *N. apis* and 218 bp product of *N. ceranae* could be detected.

Amplification was conducted in 25 µl reaction volumes (Accupower PCR premix kit, Bioneer, Seoul, South Korea) with a final concentration of each dNTP of 250 µM in 10 mM Tris-HCl pH 9.0, 30 mM KCl and 1.5 mM MgCl<sub>2</sub>, 1U Taq DNA polymerase (Takapouzist, Tehran, Iran) and 10 pmol of each PCR primer, then 1 µl of DNA template was added to each reaction. The remaining 25 µl reaction volume was filled with nuclease-free distilled water. The thermocycler program consisted of 94°C for 2 min, followed by 10 cycles of 15 s at 94°C, 30 s at 61.8°C, 45 s at 72°C, 20 cycles of 15 s at 94°C, 30 s at 61.8°C, and 50 s at 72°C plus an additional 5 s of elongation for each successive cycle and a final extension step at 72°C for 7 min. The PCR products were electrophoresed in a 2% agarose gel with TBE buffer and visualized using ethidium bromide. The positive DNA controls were obtained from a previous study (Shirzadi & Razmi, 2021) and nuclease-free distilled water was used as a negative control for each PCR amplification.

## Statistics analysis

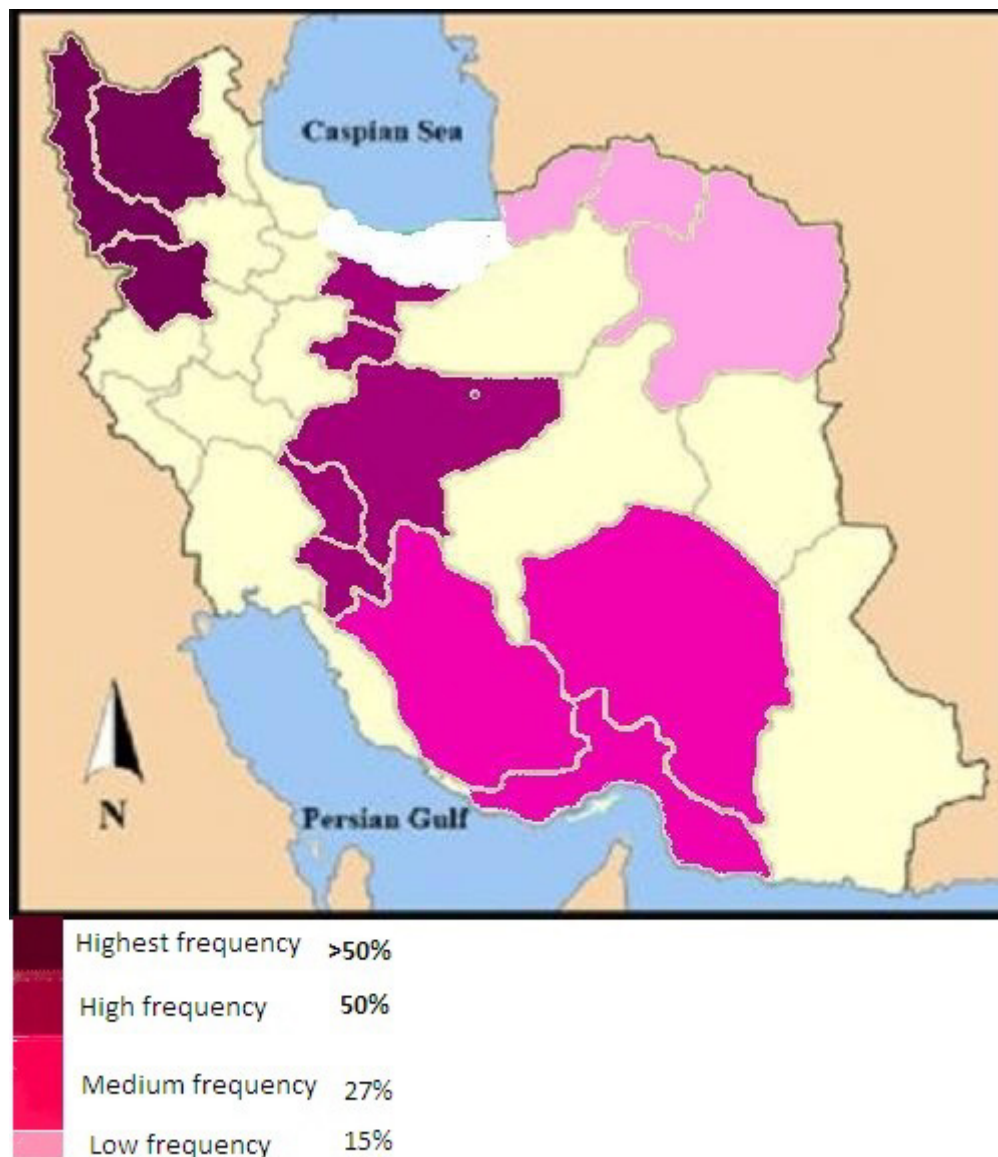
The relationship between the *Nosema* infection rate by molecular results and the region of each migratory apiary was analyzed by the Chi-square test. *P*-value below 0.05 was considered statistically significant. The agreement between the molecular and microscopic tests was shown as a Kappa-coefficient. The agreement is poor if Kappa-coefficient is between 0.2 and 0.4, moderate if between 0.4 and 0.6, substantial if 0.6 and 0.8, and good if it exceeds 0.8 and 1 (Petrie & Watson, 2006). Analyses were performed in SPSS ver. 18.0.

## Results

During the study, honeybees of sampled hives looked to be healthy. Microscopy positivity rate of *Nosema* spp. infection was detected in 38.2% (32/84) of migratory apiaries (Table 2). Molecular examination showed that 39.2% (33/84) of apiaries were infected with *N. ceranae* (Table 2). The highest prevalence of *Nosema* spp. infection was detected in the west and northwest regions while the lowest frequency was in the east and northeast regions in Iran (*P*<0.05) (Fig. 2) (Table 3). A good agreement was observed in the results between the microscopy and PCR methods (Fig. 3) (Table 4) (Kappa= 0.975).

**Table 1-** Primers selected for detection of *N. ceranae* and *N. apis* in Duplex PCR (Martín-Hernández *et al.*, 2007).

Primer	Sequences
<i>N. ceranae</i> forward	5'-GGCGACGATGTGATATGAAAATATTAA-3'
<i>N. ceranae</i> reverse	5'-CCCGGTCATTCTCAAACAAAAACCG-3'
<i>N. Apis</i> forward	5'-GGGGGCATGTCTTTGACGTACTIONTATGTA-3'
<i>N. Apis</i> reverse	5'-GGGGGGCGTTTTAAAATGTGAAACAACTATG -3'



The frequency of *Nosema* infection in different regions in Iran

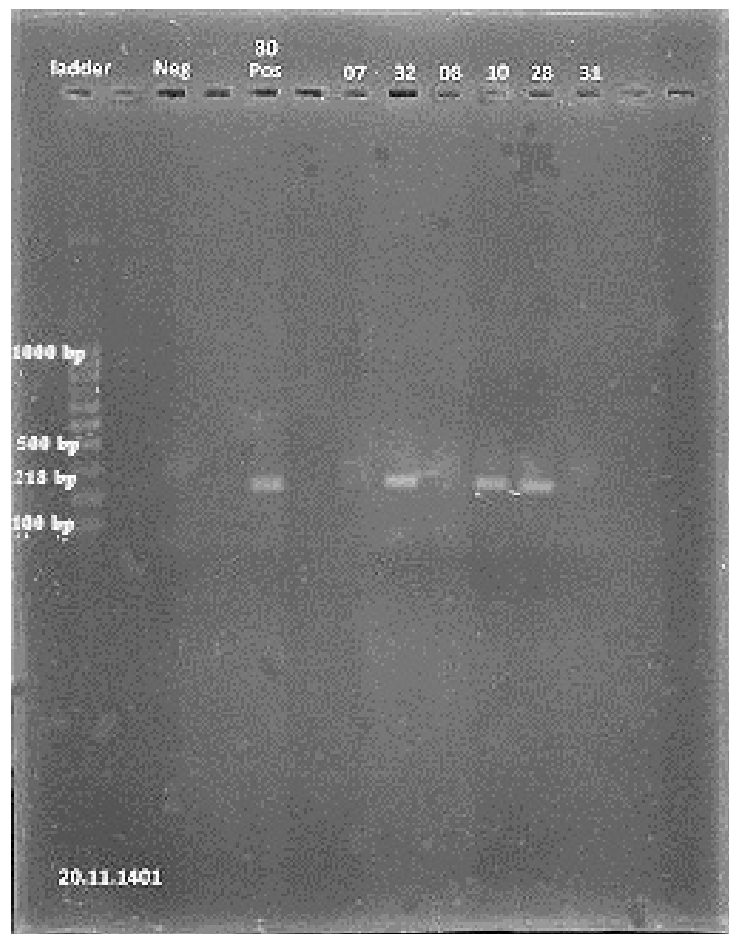
Fig. 2- The frequency of *Nosema* infection in different regions of Iran

## Discussion

The range of frequency of *Nosema* infection was determined from 15.7% to 57.14% in migratory apiaries in Hormozgan province. The highest frequency of *Nosema* spp. infection was reported in the north and northwest regions of Iran and the lowest frequency of *Nosema* spp. in the east and northeast regions. The frequency of *Nosema* spp. infection was determined in the range of 9.59% to 67% in the apiaries of different areas in East and West Azerbaijan (Lotfi *et al.*, 2009; Razmaraii, *et al.*, 2013; Moeini *et al.*, 2022). The prevalence of *Nosema* spp. infection was also reported in the range of 10-60% in different areas of Kurdistan province (Khezrei *et al.*, 2018). The reported frequency of *Nosema* spp. infection in the above studies is consistent with the reported frequency in the west and northwestern areas in this study. Mohammadian *et al.* (2018) showed a high frequency of *Nosema* spp. infection in humid and semi-humid climates and the lowest frequency in semi-arid and arid areas of Iran. A study showed that increasing the humidity and temperature caused high *N. ceranae* density and low *N. apis* density in honey bee colonies in Turkey (Özgör *et al.*, 2015).

**Table 2.** The frequency of microscopy and PCR positivity rate in migratory apiaries in Hormozgan province, Iran.

Pronince	PCR		Microscopy method	
	<i>n. examined</i>	<i>n. positive</i>	<i>n. positive</i>	<i>n. doubtful</i>
East Azerbaijan	3	1	1	
West Azerbaijan	27	17	16	2
Isfahan	5	1	1	1
Alborz	2	1	1	
Chaharmahal and Bakhtiari	3	1	1	
South Khorasan	3	-	-	1
Khorasan Razavi	15	3	3	4
North Khorasan	1	-	-	
Fars	5	3	3	
Qom	2	2	2	
Kerman	2	-	-	1
Kurdistan	5	2	2	
Golestan	1	-	-	
Lorestan	2	2	2	
Hormozgan	4	-	-	
Mazandaran	2	-	-	
No data	2	-	-	1
<b>Total</b>	<b>84</b>	<b>33</b>	<b>32</b>	<b>10</b>

**Fig. 3.** Electrophoresis results of 16 *SSUrRNA* gene with special primers, M: Marker, P: Positive control, N: negative control, *Nosema ceranea* positive samples (218bp).

**Table 3.** The frequency of PCR positivity of *Nosema* spp. based on the region of migratory apiaries in Hormozgan province (2021-2022).

Regions	Provinces	n positive (%)	Total
West and Northwest	West Azerbaijan, East Azerbaijan, Kurdistan	20 (57.14)	35
East and Southeast	Khorasan Razavi, South Khorasan, North Khorasan	3 (15.7)	19
Central	Alborz, Isfahan, Qom, Chaharmahal and Bakhtiari, Lorestan	7 (50)	14
South	Kerman, Fars, Hormozgan	3 (27.27)	11
<b>Total</b>		33	79

Other studies in China and Saudi Arabia reported a high prevalence of *N. ceranaea* in tropical wet and dry regions and a low prevalence in the regions with hot arid climates (Ansari *et al.*, 2017; Wang *et al.*, 2019). However, some studies showed that temperature and humidity do not have a positive or even negative effect on increasing the *N. ceranaea* incidence in four Mediterranean countries and Serbia (Jabal-Uriel *et al.*, 2022; Vejnovic *et al.*, 2017). A good agreement was obtained between detecting *Nosema* infection in the examined apiaries by microscopic and molecular methods. The result was in line with the results of the two studies that have reported substantial to good agreement between microscopy and PCR methods (Khezri *et al.* 2018; Papini *et al.*, 2017).

The low cost and ease of work of the microscopic method could be used to prove the *Nosema* infection in the apiaries, although detection of the *Nosema* species needs the molecular methods application. In this study, the samples were tested by Multiplex PCR method using two primer pairs of *N. apis* and *N. ceranaea* at the same time. All samples were positive for *N. ceranaea* infection. Our results are consistent with other molecular studies that determined *N. ceranaea* as the only causative agent of noseamosis in Iranian apiaries. (Nabian *et al.*, 2007; Razmaraii *et al.*, 2013; Modirrousta *et al.*, 2014; Aroee *et al.*, 2014; Shirzadi & Ramzi, 2021; Moeini *et al.*, 2022). The results of these studies have been summarized in Table 5. In the neighboring countries, a high prevalence of *N. ceranaea* infection has been reported in apiaries in Turkey (Ivgin Tunca *et al.*, 2016), Azerbaijan (Ütük *et al.*, 2019), Iraq (Kareem *et al.*, 2021), and Saudi Arabia (Ansari *et al.*, 2017). To date, *N. ceranaea* have been detected in honeybee colonies in different countries of all continents including Asia (Martín-Hernández *et al.*, 2018), Africa (Higes *et al.*, 2009), Europe (Higes *et al.*, 2006; Fries *et al.*, 2010), Australia (Giersch *et al.*, 2009), North America (Williams *et al.*, 2008; Huang *et al.*, 2015), and South America (Medici *et al.*, 2012; Teixeira *et al.*, 2013).

Diarrhea is the only detectable clinical sign in honey bees infected with *N. apis*. This symptom is not observed in hives infected with *N. ceranaea* (Huang *et al.*, 2015; Papini *et al.*, 2017). The main clinical symptom is decreasing honeybees' population with the progression of the disease (Huang *et al.*, 2015). It has also been reported that *N. ceranaea* is more virulent than *N. apis*; affects learning and homing behavior, causes higher energy costs and immune suppression, and affects queen health (Huang *et al.*, 2015). Despite the similar lifecycle of both *Nosema* species in midgut of honeybees, the reason for the difference in the symptoms is unknown (Huang *et al.*, 2015). However, no hives of *N. ceranaea*-positive apiaries in the present study showed any clinical sign at the time of sampling.

## Conclusion

*Nosema ceranaea* was the only species of *Nosema* in local and migratory apiaries in Hormozgan province. The major limitation of this study was in sampling of some apiaries that beekeepers were not willing to cooperate with us in taking bees samples. Due to the high frequency of infection in migratory apiaries in Hormozgan province, it is necessary to carry out appropriate health measures such as screening of apiaries with appropriate diagnostic methods and training of beekeepers to disinfect hives in order to control *Nosema* infection in Iranian apiaries by the veterinary health officials. However, more epidemiological studies are needed to determine the actual frequency of noseamosis in apiaries in different parts in Iran.

**Table 4.** The comparison of the results of *Nosema* spp. detection in apiaries by microscopy and PCR.

PCR Method	Microscopy Method				Total
	Results Number	Positive	Negative	Doubtful	
Positive		32	0	1	<b>33</b>
Negative		0	42	9	<b>51</b>
Doubtful		0	0	0	<b>0</b>
Total		32	42	10	<b>84</b>

**Table 5.** Studies on frequency of *Nosema* infection in different areas of Iran until October 2023.

Province	Study year	Method	n examined	n infected	<i>Nosema</i> species	References
West Azarbijan	2002–2003	Microscopy	478	138	<i>N. apis?</i>	Tavassoli et al., 2009
Ardabil	2008	Microscopy	294	59	<i>Nosema</i> spp.	Lotfi et al., 2009
North Khorasan	2011	Microscopy	54	12	<i>N. apis?</i>	Moshaverinai et al., 2012
Mazandaran	2011	PCR	6 microscopy positive	6	<i>N. ceranae</i>	Nabiean et al., 2011
East Azarbijan	2011	Microscopy	387	225	<i>N. ceranae</i>	Razmaraai et al., 2013
		PCR	387	260		
Alborz, East Azarbijan, Ghazvin, Gilan, Tehran	2013	PCR	41 microscopy positive	41	<i>N. ceranae</i>	Modirrosta et al., 2014
Chaharmahal and Bakhtiari, Isfahan, Fars	2017	Microscopy	180	-	<i>N. ceranae</i>	Aroee et al., 2017
		PCR	180	-		
Kurdistan	2018	Microscopy	100	29	<i>N. ceranae</i>	Khezeri et al., 2018
		PCR	100	32		
Different provinces	2019	Microscopy	183	85	<i>N. ceranae</i>	Mohammadian et al., 2018
		PCR	183	66		
Mazandarn	2020	Microscopy	320	250	<i>N. ceranae</i>	Shirzadi and Razmi, 2021
		PCR	320	278		
West Azarbijan	2022	Microscopy	840	269	<i>N. ceranae</i>	Moeini et al., 2022
		PCR	840	488		
East Azarbijan	2022	Microscopy	165	165	<i>Nosema</i> spp.	Imani and Hamidiam, 2022

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### چکیده

نوزموزیس یک بیماری میکروسپوریدیایی ناشی از *Nosema apis* و *Nosema ceranae* است که باعث آلودگی زنبوران بالغ در کلنی‌های زنبورعسل در سراسر جهان می‌گردد. انتقال اسپورهای نوزما از طریق مسیرهای مدفوعی-دهانی و دهانی-دهانی اتفاق می‌افتد. با توجه به اهمیت سلامت زنبورهای عسل، هدف از این مطالعه بررسی فراوانی آلودگی با نوزما در زنبوداری‌های مهاجر به استان هرمزگان با استفاده از روش‌های میکروسکوپی و ملکولی بود. بدین منظور از ۸۴ زنبورستان مهاجر در استان هرمزگان نمونه‌برداری شد، بدین صورت که از هر زنبورستان ۱۰ کندو انتخاب شده و از هر کندو ۲۰ تا ۳۰ زنبورعسل مسن به صورت تصادفی انتخاب شده و به آزمایشگاه منتقل شدند. در آزمایشگاه، شکم زنبورها با پنس و قیچی حشره‌شناسی از بقیه بدن جدا شده و در حاوی سرم نمکی آسیاب شد. سوسپانسیون تهیه شده با عبور از الک صاف شده و به لوله‌های آزمایش منتقل و سانتیفریوژ گردید. رسوب ته لوله‌های آزمایش به طور مکرر با افزودن سرم فیزیولوژی و سانتیفریوژ شسته شده و برای تشخیص عفونت نوزما با روش‌های میکروسکوپی و ملکولی مورد بررسی قرار گرفتند. در بررسی میکروسکوپی اسپورهای گونه‌های *Nosema* در ۲۸/۲ درصد زنبورستان‌ها مشاهده شدند. با روش PCR اما ۳۹/۲ درصد زنبورستان‌ها آلوده به دنای *N. ceranae* تشخیص داده شدند. هیچ موردی از آلودگی *N. apis* مشاهده نشد. با توجه به فراوانی قابل ملاحظه‌ی آلودگی نوزما در زنبورستان‌های مهاجر به استان هرمزگان، انجام اقدامات بهداشتی مناسب همچون غربالگری زنبورستان‌های کشتور با روش‌های تشخیصی مناسب و آموزش زنبورداران جهت ضدعفونی صحیح کندوها برای کنترل بیماری در زنبورستان‌های ایران توسط مسئولان بهداشتی دامپزشکی ضروری است.

**کلمات کلیدی:** بیماری شناسی حشرات، زنبور عسل، شناسایی ملکولی

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