

**The First Report for the invasion of *Artemia franciscana* Kellogg 1906 in Tashk and Bakhtegan Lakes, Iran**Sepideh Shafaie<sup>1</sup>, Samad Zare<sup>2</sup>, Ramin Manaffar<sup>3</sup> and Afagh Falahati<sup>4</sup>

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**Abstract:** *Artemia*, a small crustacean, with high commercial value is a valuable model organism for researchers. This creature by tolerating extreme range of different environmental conditions was dispersed to more than 600 and 18 sites over the world and Iran, respectively. Tashk and Bakhtegan Lakes are one of the natural parthenogenetic *Artemia* habitats in Iran. Due to occurrence of an unknown bisexual *Artemia* in Tashk Lake, the species of this endemic *Artemia* was inferred. In this regard, four different molecular markers as Na/K ATP-ase, 12S-16S by PCR-RFLP technique and COI and HSP26 by sequencing and subsequent Genbank data were studied. The conducted analyses with emphasizing to ability of molecular techniques for identifying unknown species characterized the new population as *A. franciscana* in these two lakes. These analyses also revealed a molecular diversity between the sequenced genes with the data found in the Genbank.

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**1. Introduction**

Brine shrimp *Artemia* is one of the important zooplanktons in aquaculture. Having special physiological adaptabilities such as the capability of producing resistant cysts, a very effective osmotic pressure regulation system, *Artemia* has been adapted to the life in salty and very salty water. In fact, among Metazoan, *Artemia* is the only creature that can endure high degrees (up to 300 gr per liter) of salinity (Browne, 1992). In addition to its great nutritional value (fatty acids and necessary proteins), *Artemia* is an appropriate research model in molecular and evolutionary experiments. This genus harbors two bisexual (comprising of six species) and parthenogenetic strains (Van Stappen, 2008). Each of these populations has been adapted to different climates and habitats based on their molecular characteristics and physiologic. In 1995, the first official paper has referred to only 80 areas as *Artemia* habitats (Abonyi, 1915), while around 600 geographical regions has been introduced as *Artemia* habitats in the latest list of the year 2002 (Van Stappen et al, 2002). The fast dispersion of *Artemia* around the earth and also the discovery of new regions through developments in specific researches is the reason for the change in dispersion list of *Artemia*. Likewise, the extinction of *Artemia* in some regions (such as Lamington region in England and Shurabil Lake in

Iran) can also make changes in the habitat and dispersion list of *Artemia* (Van Stappen, 2008).

So far, 18 different sites of *Artemia* have been reported in Iran (Abatzopoulos et al, 2006; Asem et al, 2009). All of these sites, except Urmia Lake, have the endemic parthenogenetic *Artemia*. The geographical position of Bakhtegan Lake, the lake in Fars province of Iran, is 53° 50' N and 29° 40' E and it is located at a distance of 80 Km from the east of Shiraz. Tashk Lake, the other lake in Fars province, in geographical position of 53° 50' N and 29° 60' E is located at a distance of 50-160 Km from the east of Shiraz (Agh, 2007). Tashk lake was previously introduced as Brackish Lake that had connection to Bakhtegan Lake via a connective bridge. Therefore, the salinity of water has been increased which has made this region a biologically suitable place for parthenogenetic *Artemia* (Agh, 2007). The temperature of these lakes which has adaptability to thermo iso-plates of the region is between less than 5 degrees and more than 40 degrees and even reaches to 45°C in sloughs (Alamdari, 1987). There is no idea about the first report of parthenogenetic *Artemia* in these lakes, but several reports have been recorded about the existence of parthenogenetic *Artemia* in these lakes in 1980, 1984, 2002, and 2007 (Geddes, 1980; Browne et al, 1984; Van Stappen et al, 2002; Agh, 2007).

*A. franciscana*, as one of the very potatic bisexual species in the world (with great frequency 74

sites) is the dominant *Artemia* species in Great Salt Lake, U.S. (Van Stappen, 2002). In fact, the high and fast adaptability capacity of this species in new ecosystems has caused it to be selected for commercial Pond production industry in most of the countries (Amat et al, 2007). The molecular and physiological capacity of *A. franciscana* to a wide domain of ecological conditions and also the high growth velocity and reproduction potential of this *Artemia* has caused it to be easily dispersed around the world.

The studies have indicated that the molecular adaptability power of the *Artemia* (specially *A. franciscana*) is the main reason for this wide dispersion. It is inferred that some genes of heat shock proteins (HSPs) or mitochondrial genes can show these molecular changes quickly in a way that they have been considered as the successful adaptability criterion to new conditions. The researches have indicated that these chaperon proteins has a vital role in increasing physiological adaptability of the living creature facing with unpleasant biological conditions (Clegg et al, 2000). These proteins change a lot in *Artemia* which have experienced a successful period in a new and different habitat (Bossier et al, 2009). It has been proven that other proteins of this family also have significant role in stress tolerance and creating molecular adaptability (Federand and Hofmann, 1999; Prohaszka and Fust, 2004).

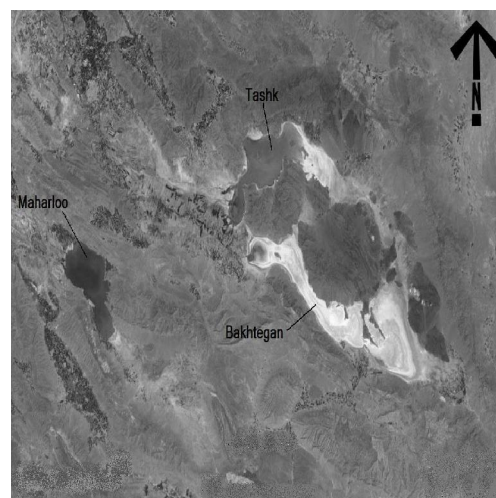
In order to investigate the species diversity of *Artemia*, different methods have been introduced including different types of morphometric and molecular techniques. Mitochondrial genome performance has been confirmed as an appropriate option for identification of indefinite taxon, analysis of species creation and even identification of *Artemia* population (Avisé, 2000; Bossier et al, 2004). Likewise, a molecular method has been presented recently in order to recognize the level of parthenogenetic from bisexuality level which can be used for the separation of these two levels (Manaffar et al, 2011).

Since *A. urmiana* is the only endemic bisexual *Artemia* population observing a huge population of bisexual *Artemia* in Tashk and Bakhtegan Lakes, the identification of level and species of the observed *Artemia* was considered as the main goal of the present study.

## 2. Material and Methods

*Artemia* cysts were collected from four different parts of Tashk and Bakhtegan Lakes which located in Fars province in July 2011 (Figure 1). The cysts were hatched after rinsing and purification in standard laboratory conditions in Urmia Lake water which was diluted to the saltness of 35 gr per liter, a temperature of 27°, pH=8, and equipped with aeration systems and

enough light (Lavens and Sorgeloos, 1996). Instar I were nauplii transferred into one liter bottles containing 80 gr per liter saline water in 4 repetitions and reached for 20 days with mixture of enriched yeast with fatty acid, and unicellular Alga *Dunaliella salina* (Coutteau et al, 1992).



**Figure1- Location of Tashk and Bakhtegan Lakes which are near the Maharloo Lake.**

### 2.1. Molecular analysis

The DNA were extracted from cyst individuals of cyst using Chelex method (Estoup et al, 1996). In order to extract DNAs of the mature *Artemia* samples, the CTAB method was used (Doyle and Doyle, 1990).

In order to investigate the species diversity of unendemic of *Artemia* population in Tashk and Bakhtegan Lakes, 20 samples (cyst and mature *Artemia*) were used. To this end, genetic parts of Na/K ATPase (a part of core genome for identification of bisexually or parthenogenetic) (Manaffaret al, 2011) and 12s-16s (a part of mitochondrial genome for identification of *Artemia* species) (Bossier et al, 2004) were used.

PCR program and also used primers have been summarized in Table 1. The PCR product were analyzed in all experiments using the electrophoresis of 2% Agarose gel and photographed by Gene Flash gel. Documentation system after approving the quality of PCR product in RFLP technique, in order to characterize strain and species of *Artemia* the exon-7 fragment of Na- K ATPase gene and the 12s-16s gene fragment were digested by *TruI* and *Hpa* II restriction digestion enzymes and they were analyzed on 2% Agarose gel. Fragments of HSP26 and COI genes were also sequenced by Sina Gene Company. (Folmer et al, 2006).

Table 1. primers and PCR program

|             | Forward and reverse primers  | programPCR   |
|-------------|--|--|
| Na/K ATPase | -cca-aac-gta-tgg-ctt-c-3'<br>5'-cag<br>-agc-acg-act-gca-aga-3'<br>5'-gaa-ttc             | 94°C 2 Min<br>35 cycle (94°C 2 Min, 56°C 25 Sec, 72°C 1 Min)<br>72°C 3 Min   |
| COI         | -atc-ata-aag-ata-tgt-g-3'<br>5'-ggg-aca<br>-tga-cca-aaa-aat-ca -3'<br>5'-taa-act-tca-ggg | 95°C 3 Min<br>33 cycle (95°C 1Min, 50°C 1 Min, 72°C 1.20 Min)<br>72°C 10 Min |
| 12S-16S     | -cca-aac-gta-tgg-ctt-c-3'<br>5'-cag<br>-agc-acg-act-gca-aag-3'<br>5'-gaa-ttc             | 95°C 2 Min<br>34 cycle (94°C 1.15 Min, 52°C 1 Min, 72°C 2 Min)<br>72°C 4 Min |
| HSP 26      | -gga-gaa-gaa-tga-gaa-g-3'<br>5'<br>-tgg-acg-tgt-cca-tat-tc-3'<br>5'-tct-ctt              | 94°C 2 Min<br>35 cycle (94°C 15 Sec, 54°C 25 Sec, 72°C 30 sec)<br>72°C 4 Min |

### 3. Results

The analysis of the implied a 700 bp which can be related to cytochrome oxidase gene and 217 bp fragment which can be related to heat shock gene of HSP 26 indicated that the sequence of the PCR product also checked in Genbank data base by Blast Run. Enzymatic cutting of piece number 280 bp of core genome which is produced by *HpaII* showed that the created cutting pattern in *Artemia* samples of Tashk and Bakhtegan Lakes, are exactly similar to the pattern of Bisexual *Artemia* (Figure 2).

In order to analyze the species of *Artemia*, the enzymatic cutting of piece number 1500 bp of 12S-16S mitochondrial genome was used. Enzyme cutting by the limitative *HpaII* enzyme created the profile figure of *A. franciscana* (Figure 3). Analysis of the sequence results using Genbank internet search by Blast software emphasized that the above bisexual *Artemia* belongs to the bisexual American *Artemia* in molecular structure of this gene in comparison to genes recorded in gene bank.

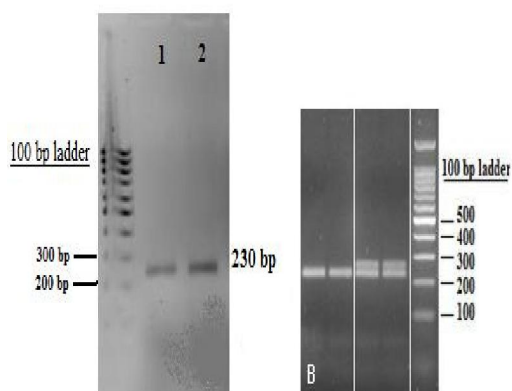


Figure 2- Agarose gel of restriction enzyme fragment of 280 bp region Image A): 230 bp band was produced from Brine Shrimp samples (1: Tashk Lake, 2: Bakhtegan Lake)

Image B): taken from the reference gel. Manaffar et al, 2011 pattern, cut two bands of parthenogenetic *Artemia* and single-band pattern is indicative of bisexual *Artemia*.

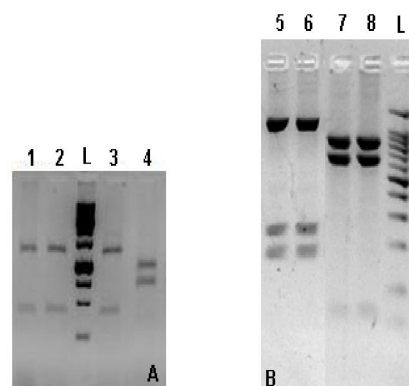


Figure 3. Agarose gel electrophoresis of 12S-16S fragment of 1500 bp region of the enzyme restriction by the enzyme *HpaII*. Images were: Image A) Restriction enzyme fragment above the exotic *Artemia* in Tashk (sample 1) Bakhtegan (sample 2) control samples (3. *A. franciscana* and 4. *A. sinica*) with 1 Kb marker. Reference gel from (Bossier et al, 2004) samples 5 and 6 related to *A. franciscana*, samples 7 and 8 related to *A. sinica* with 100 bp ladder.

### 4. Discussion

The present study is considered as the first report about the observation of *A. franciscana* in Tashk and Bakhtegan Lakes, the lakes of Fars province. Some scientific analysis had already confirmed the existence of *A. franciscana* in Iran Plateau (natural habitat of *Artemia*) (Manaffar et al, 2008). In present study, it was attempted to use new technique which forecast genetic distinction patterns and extend diversity in several creatures with little error (Chow et al, 2006). The merits of the used molecular methods for strain and species identification of *Artemia* was already approved. But, setting sequences of cytochrome oxidase and also small heat shock protein and the analysis of its sequence with samples of gene bank indicates a genetic difference of more than 30% between tested *Artemia* and existed samples from the above-mentioned website. Of course, the effect of other phenomenon such as Founder effect and genetic drift should not be neglected. Regardless it should be noted that this new population have managed to adapt themselves to Tashk and Bakhtegan Lakes successfully.

However, with regard to the probable method of transferring this species of *Artemia* to Tashk and Bakhtegan Lakes, it should be mentioned that although

the *Artemia* egg is generally transferred by wind and aquatic birds (Persoone and Sorgeloos, 1980), but from 1970s until now, humans have been responsible for dispersion of *Artemia*, specially *A. franciscana*. Regarding to existence of *A. franciscana* in Maharloo Lake and also close distance of these three lakes, it is suggested that *A. franciscana* have been transferred from Maharloo Lake to these Lakes by birds.

*Artemia* has shown the highest level of phenotypical and genetic flexibility and with a very high reproduction speed, quick adaption to difficult conditions and molecular adaptability with environment it has been dispersed successfully in Asia, Europe, and America and has often caused the elimination of local *Artemia* (Browne et al., 1988; Kappas et al, 2004; Pogge, 2004). With the transfer of this species in 1970s to the island of Pacific Ocean and Brazil, it was announced that the above species will probably substitute other species including *A. Salina* (Van Stappen et al, 2002). However, the first report about the offensive power of *A. franciscana* is related to Camara in 2001 who has reported that this species is located in Rio Grand do Norte in northern Brazil. Other similar report have been recorded in Portugal (Amat et al, 2005), France (Thiery, 1992), Egypt (Triantaphyllidis et al, 1998), Italy (Mura et al, 2004), Spain and Morocco (Amat et al, 2007). Research done by Kappas in 2004 on non-local *A. franciscana* in Vietnam indicated that there are significant differences between local American *A. franciscana* and commercial Vietnamese *Artemia* which had been transferred to Vietnam 10 years ago.

It should be noted that the permanent settlement of non-local *Artemia* population and the development in dispersion of *A. franciscana* around the world have been one of the note worthy issues in recent years (Abatzopoulos et al, 2006; Amat et al, 2005, green et al, 2005; Mura et al, 2006). At present, *A. franciscana* is considered as the dominant population in the west of Mediterranean Sea, Slat mines in Portugal, Mediterranean beach of France, and the Cadiz Gulf in Spain (Amat et al, 2005). This study has proved that *A. franciscana* has been able to eliminate local population within a few years (Amat et al, 2005; Green et al, 2005; Amat et al, 2007). Researches in Iran have also indicated that *A. franciscana* has managed to become a dominant population in Nogh Pool of Rafsanjan (which was a natural habitat for parthenogenetic *Artemia*) in a contest with parthenogenetic *Artemia* (Abatzopoulos et al, 2006).

According to the results of the present study the found genetic diversity in *A. franciscana* and also its potential capability, it is expected that this bisexual species may be able to completely eliminate local parthenogenetic *Artemia* population in the future.

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