

Morphological And Genetic Features of Oxyurichthys Ophthalmonema Distributed in Central Vietnam

Nguyen Ty¹•, Nguyen Thi Nhu Y¹•, Nguyen Thi Lan¹, Nguyen Van Thuan¹, Pham Thanh¹ and Tran Van Giang¹*

¹University of Education, Hue University, Hue city, 49000, Vietnam

ABSTRACT:

Gobius (Oxyurichthys ophthalmonema Bleeker, 1856) is a species of the Gobiidae family with high commercial and nutritional value, making it a promising candidate for development in Vietnam. A total of 408 fish samples were collected at 9 sites in 5 study areas in central Vietnam from October 2019 to September 2020. The research results showed that the average fish size ranges from 67.70 - 190.20 mm, corresponds to a mass of 4.70 - 29.25g, and the correlation between standard length and body mass is proportional to each other. Compared to 23 morphological criteria, it was shown that the farther the distribution distance, the greater the difference in morphological criteria (p<0.05). 19 total DNA samples of O. ophthalmonema in 5 study areas were extracted, in which 14 samples were amplified and sequenced for the COI gene region; the results obtained the sequences that had differences at different positions compared to the COI gene region of O. ophthalmonema. The phylogenetic tree construction of 14 fish samples with COI sequences of O. ophthalmonema and 5 species of the genus Oxyurichthys showed that the fish samples belonged to O. ophthalmonema species with a 100% bootstrap value. However, their genetic distances were far apart, and there were also differences between the samples.

KEYWORDS: Central region, Genetic diversity, Gobius, Morphological characteristics, O. ophthalmonema.

INTRODUCTION

Oxyurichthys ophthalmonema (Bleeker, 1856) belongs to the family Gobiidae, in the order Gobiiformes, which is mainly distributed in lagoons and estuaries of coastal areas. This fish has good resistance, health and high nutritional value, so it is favored by many people. Currently, O. ophthalmonema is one of the fish species with high consumption. However, the main supply source of this fish still comes from small-scale fishing with rudimentary tools such as small mesh nets, gill nets, etc., making the fish species currently declining in numbers and decreasing day by day. Therefore, the cultivation of this fish is an urgent issue to reduce the pressure of overfishing and provide enough for the market's demand. However, studies of this species around the world, as well as in our country, are very limited. In the world, the most remarkable research is the work of Remys Mohan & Sherly Williams (2018) on the nutritional and reproductive characteristics of this species in Lake Ashtamudi, India. In Vietnam, only a few studies have been recorded at Tam Giang - Cau Hai Lagoon in Hue, Rao Cai river in Ha Tinh and Mai Giang river in Nghe An (Thai & Dang 2015, Nguyen, 2005). Nowadays, there is very little research on this species, and there is no connection between the previous studies. Therefore, we conducted a study on morphological and genetic characteristics of Goby (Oxyurichthys ophthalmonema (Bleeker, 1856)) to provide scientific data for research and teaching at universities, as well as to regulate fishing and to conserve the existence of this species in central Vietnam.

MATERIALS AND METHODS

Sample collection locations

A total of 408 samples at 9 sites in 5 study areas in central Vietnam were collected from the beginning of October 2019 to the end of September 2020 (table 1). Samples were randomly collected and repeated at the site by direct fishing together with local people, or ordering samples from fishermen in the study area.

Table 1. Collection sample

No.	Location	Collection time	Number of samples
1	Water area in Quang An commune, Quang Dien district, Thua Thien Hue province	Oct, 2019 – Jan, 2020	51
2	Water area of Thuan An town, Thua Thien Hue province	Jan, 2019 – May, 2020	50
3	The brackish water area of Quang Ngan commune, Quang Dien district, Thua Thien Hue province	Sept, 16 th , 2020	11
4	The brackish water area of Dien Hai commune, Quang Dien district, Thua Thien Hue province	Aug, 30 th , 2020	1
5	Trieu An commune, Trieu Phong district, Quang Tri province (at the market)	Feb, 22 th , 2020	39
6	Trieu Phuoc Commune, Trieu Phong District, Quang Tri Province (at the market)	July, 22 th , 2020	71
7	Nhat Le River Estuary, Dong Hoi City, Quang Binh province	March, 2020 – Sept, 2020	115
8	River door Sot, Nghen river, Ha Tinh province	Sept, 2020	69
9	Ben Thuy river, Nghe An province	Sept, 2020	1

Morphological study: samples were treated when the fish was still fresh by fixing them with a 40% formol solution; After imaging, the sample was transferred to a 4% formol solution. After that, the fish samples underwent morphological characterization using the fish morphological description guidelines provided by Pravdin (1973) and Nguyen (2005).

Genetic study: fish samples were cut for pectoral fin samples and stored in eppendorf tubes containing 96% alcohol, and stored at -20°C until analysis.

Table 2. Total DNA extraction fish samples

Province	Sample number	Sample symbol				
Thua Thien Hue	4	TTHTG002 (MZ798204), TTHTG006 (MZ798208), TTHTG033 (MZ798216), TTHTG034 (MZ798217)				
Quang Tri	3	QT004 (MZ798206), QT007 (MZ798209), QT008 (MZ798210)				
Quang Binh	7	QB005 (MZ798207), QB009 (MZ798211), QB010 (MZ798212), QB036, QB037, QB041, QB042				
Ha Tinh	4	HT003 (MZ798205), HT011 (MZ798213), HT012 (MZ798214), HT009				

Nghe An 1 NA013 (MZ798215)

DNA Barcoding: DNA extraction, PCR amplification and decoding. Total DNA was extracted from the pectoral fins of each individual fish using the "GeneJET Genomic DNA Purification Kit DNA Extraction Kit" according to the manufacturer's instructions. The 683 bp mitochondrial *CO*I was amplified with the primer pairs Fish F1 (5'- TCAACCAACCACAAAGACATTGGCAC-3') and Fish R1 (5'-TAGACTTCTGGGTGGCCAAAGAATCA-3'). PCR reactions were performed with a total volume of 60 μ I solution consisting of: 2 μ I of total DNA (~50ng), 30 μ I of Go Taq (2X), 3 μ I of each primer (10pmol/ μ I) and 22 μ I of Kit distilled water. The PCR machine (Esco) was run according to the following temperature program: DNA denaturation: 95°C/2 mins; (2). Amplification: 35 cycles: (94°C/3 mins; 54°C/3 mins; 72°C/1 mins); final 72°C/10 mins. PCR products were electrophoresed on a 0.8% agarose geI and the DNA bands were visualized under a UV projector. The results were recorded using the GeIDoc image analysis system.

DNA barcoding and phylogenetic analysis: sequences were initially aligned using the BioEdit sequence editing program. After that, the research sequences were compared with similar sequences on Genbank using BLAST (https://blast.ncbi.nlm.nih.gov). The phylogenetic trees were built by Clustal W. and constructed by the Test Maximum Likelihood method using MEGA X software (Kumar *et al.* 2018) with the bootstrap values repeated 1000 times for the sample.

RESULTS AND DISCUSSION

Morphological characteristics of Oxyurichthys ophthalmonema

External appearance

From morphological characteristic analysis combined with the COI gene barcoding results, the goby species has been identified as having the scientific name Oxyurichthys ophthalmonema (Bleeker, 1856). The study samples have the Vietnamese names "goby" or "goby van eye fish".



Figure 1. Morphology of the Goby Oxyurichthys ophthalmonema

Based on 25 measurement criteria and 5 count criteria, 408 samples collected in 5 research areas in central Vietnam (only one sample in Nghe An) have been identified as a fish species named Oxyurichthys ophthalmonema (Bleeker, 1856) with the following main features: antennae, upper lip retraction at the anterior jaw joint, upper jaw has a single row of teeth, nape has a thin membrane (crest), has 2 dorsal fins, 1st dorsal fin with filamentous, not elongated spines and 2nd dorsal fin with 1 starting spine and posterior rays, with the last ray branching at the base (Figs 1 and 2). According to Betancur *et al.* (2017), this species belongs to the genus *Oxyurichthys*, family Gobiidae, order Gobiiformes, class Actinopterygii, phylum Chordata.

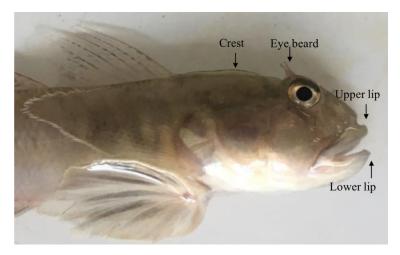


Figure 2. Head part of O. ophthalmonema

The results showed that this species has an average size ranging from 67.70 - 190.20 mm, corresponding to a weight of 4.70-29.25g; the number of spines and dorsal fin rays 1 is 6.0; the number of spines and dorsal fin rays 2 is 1-2, 12-14; the number of spines and pectoral fin rays is 1-5, 14-22; the number of spines and ventral rays is 2.10; the number of spines and pelvic fin rays is 2-11, 12-19; the number of spines and rays of anal fin is 1.13–14.

The external morphological characteristics of the population of *O. ophthalmonema* goby fish distributed in central Vietnam have many similar characteristics, which can be distinguished by the naked eye. They have a diamond-shaped body; their bodies are elongated and flat, covered with comb scales; the back is straight; the belly is slightly curved; the head is short; the mouth of the fish is inclined at an angle of about 45° to its body axis; the mouth is at the tip of the snout, facing forward; the arc is quite wide; the muzzle is long and slightly protruding; the developed muzzle covers the upper lip; the upper lip constricts; a wide gill membrane is attached to the gill waist (Figure 2). The fish's eyes are slightly convex laterally, favoring the top of the head and forming a crest on the dorsal surface. The distance between the eyes is narrow, and there is especially the presence of an eye antenna extending above the back of the eye.

The color of the body varies from ventral to dorsal; dark gray back, white gray belly. When the fish is alive, its body color is slightly pinkish gray, and its pelvic and anal fins are opaque gray. The dorsal, pectoral, and anal fins are quite clear. The tips of the caudal and dorsal fins are red. The pelvic and pectoral fins are bordered with thin white edges.

Size and body mass of O. ophthalmonema

The standard length and body mass of O. ophthalmonema in different regions were carefully studied. The results showed that both the largest mass and the longest standard length were from the samples collected in Quang Binh province (table 3).

Table 3. Body mass and standard length of *O. ophthalmonema* (Bleeker, 1856) distributed in the central Vietnam

Targets	Location	Thua Thien Hue	Quang Tri	Quang Binh	Ha Tinh	Medium
1. Weight	Min- Max	8.60-29.25	4.70-20.80	7.60-24.35	5.39-11.01	4.70-29.25
(g)	M± SD	15.05±4.04	9.54±3.42	16.55±4.59	8.07±1.45	12.80±5.10

۷.	Min-	80.00-	67.70-	82.00-	70.11-	67.70-
	Max	109.65	190.20	180.11	84.52	190.20
d length (mm)	M± SD	93.83±7.65	84.45±13.9 1	101.81±13.3 0	77.66±4.2 6	90.82±14.0 8

The morphological parameters of *O. ophthalmonema* in the central area were carefully studied, indicates that: the length of fish in Thua Thien Hue ranged from 80.00-109.65 mm, average 93.83± 7.65, Quang Tri fish ranged from 67.70-190.20 mm, average 84.45±13.91, Quang Binh fish ranged from 82.00-180.11 mm, average 101.81± 13.30 mm and Ha Tinh fish from 70.11 to 84.52 mm, average 77.66±4.26. Measures have variation among individuals in the same province. There are significant differences between provinces in a number of criteria.

The lengths of other body parts relative to the head length were measured. The results showed that the head width reaches 57.04% of the head length, while the figures for the height of the head, the distance between the eyes, the eye diameter and the width of the mouth are 68.57%, 18.81%, 23.79% and 48.96% respectively (table 4).

Location		Thus Thiss					
Compared to HL		Thua Thien Hue	Quang Tri	Quang Binh	Ha Tinh	Medium	
1. Head	Min- Max	45.45 - 70.83	47.16 - 71.43	34.13 - 68.04	50.14 - 56.42	34.13 - 71.43	
Width	M ± SD	56.73 ± 4.53	58.38 ± 4.38	57.44 ± 4.24	54.74 ± 1.14	57.04 ± 4.18	
2.	Min- Max	42.31 - 80	43.04 - 82.00	39.02 - 78.95	58.78 - 67.03	39.02 - 82.00	
Head Height	M ± SD	70.25 ± 4.96	68.88 ± 4.95	69.72 ± 5.19	63.37 ± 2.59	68.57 ± 5.29	
3. Distan	Min- Max	13.64 - 27.89	11.76 - 28.12	12.00 - 26.02	16.62 - 21.78	11.76 - 28.12	
ce betwe en eyes	M ± SD	21.08 ± 3.04	18.43 ± 3.09	17.56 ± 3.04	17.87 ± 1.43	18.81 ± 3.18	
4. Eye Diame	Min- Max	14.29 - 27.81	15.00 - 38.59	14.75 - 30.8	21.49 - 26.65	14.29 - 38.59	
ter	M ± SD	22.76 ± 3.33	23.66 ± 3.69	25.19 ± 2.49	23.39 ± 1.7	23.79 ± 3.13	
5.	Min- Max	23.69 - 74.20	32.41 - 95.83	27.74 - 58.58	23.78 - 45.66	23.69 - 95.83	
Mouth width	M ± SD	54.68 ± 9.62	48.56 ± 9.43	47.60 ± 5.30	42.53 ± 3.78	48.96 ± 8.73	

Table 4. The lengths of other body	parts relative to the l	nead length (%HL)

Besides the measurement criteria, the count criteria for the species have also been studied. The analysis and comparison for counting indicators found that fish species in the central area of Vietnam has the following characteristics: number of spines and dorsal fin rays 1 is 6 and 0; number of spines and dorsal fin rays 2 is 1 - 2 and 12 - 14; number of spines and pectoral fin rays is 1 - 5 and 14 - 22; number of spines and ventral rays is 2 and 10; number of spines and pelvic fin rays is 2 - 11 and 12 - 19; number of spines and rays of anal fin is 1 and 13 - 14 (table 5).

Location Targets		Thua Thien Hue	Quang Tri	Quang Binh	Ha Tinh
1. Number of	Min-Max	1 – 5	1-4	2 - 4	3
pectoral fin spines (P)	M ± SD	2.55 ± 0.83	2.67 ± 0.62	2.75 ± 0.62	3 ± 0
2. Number of	Min-Max	17 – 22	14 – 20	18 - 20	18 - 20
pectoral fin rays	M ± SD	18.62 ± 1.21	18.64 ± 0.89	18.66 ± 0.87	19.07 ± 0.9
3. Number of	Min-Max	2 – 10	4 - 11	5 - 10	7 - 8
caudal fin spines	M ± SD	6.56 ± 2.88	7.93 ± 1.13	6.98 ± 1.16	7.17 ± 0.38
4. Number of	Min-Max	13 – 17	12 – 19	13 - 18	14 - 14
caudal fin rays	M ± SD	14.45 ± 0.78	14.68 ± 1.26	14.34 ± 0.77	14.0

Table 5. Comparison of the counting parameters of O. ophthalmonema

Through the analysis of measurements and counts, the average size of *O. ophthalmonema* ranges from 67.70 - 190.20 mm; average 90.82 \pm 14.08 mm; the head length is about 1/5 of the standard length while body height is about 1/4 of the standard length; the average weight is 12.80g \pm 5.10 g. These results are similar to the previous descriptions of Pezold and Larson (2015) and Nguyen and associates (2020) on goby fish collected in Thua Thien Hue province.

Table 6. Mean values (± standard deviation) of parameters compared to standard length (SL) of *Oxyrichthys ophthalmonema*

Compared to SL (%)	Thua Thien Hue	Quang Tri	Quang Binh	Ha Tinh
1. Head length	24,18 ± 0,71 ^a	24,38 ± 1,74 ^a	24,66 ± 2,00 ^a	26,04 ± 0,55 ^b
2. Body height	20,15 ±1,37 ^{ac}	19,14 ± 1,73 ^b	19,66 ± 1,88 ^{ab}	20,33 ±0,95 ^{ac}
3. Tail stalk length	9,7 ± 1,36ª	10,1 ± 1,80 ^{ab}	9,91 ± 1,75ª	10,43 ± 0,35 ^b
4. Tail stalk height	10,9 ± 0,75ª	10,12 ± 0,67 ^b	10,67 ± 1,49 ^{ac}	10,57 ± 0,55°
5. Dorsal fin length 1	19,37 ± 1,98ª	19,02 ± 2,08ª	19,14 ± 2,25ª	19,33 ± 3,28ª
6. Dorsal fin length 2	39,08 ± 3,85ª	39,82 ± 12,90 ^{ab}	39,94 ± 6,57 ^{ab}	40,74 ± 1,02 ^b
7. Pectoral fin length	28,90 ± 2,84ª	29,83 ±3,66 ^{abc}	30,79 ± 3,43 ^{bc}	30,52 ± 1,61°
8. Pelvic fin length	22,98 ± 1,63ª	23,06 ± 2,12ª	23,59 ± 2,32 ^{ab}	23,97 ± 1,07 ^b

9. Tail fin length	45,45 ± 4,17ª	49,2 ± 7,46 ^b	47,74 ± 4,77 ^b	44,56 ± 3,34ª
10. Anal fin length	41,29 ± 1,66ª	39,53 ± 2,68 ^b	40 ± 2,98 ^b	39,87 ± 1,01 ^b
11. Dorsal fin height 1	14,19 ± 3,91ª	11,1 ± 2,44 ^b	11,1 ± 2,44 ^b 12,56 ± 1,82 ^c	
12. Dorsal fin height 2	13,55 ± 5,20ª	10,37 ± 2,22 ^b	10,98 ± 3,35 ^b	9,34 ± 1,79°
13. Pectoral fin height	18,05 ± 1,56ª	17,33 ± 2,43ª	18,96 ± 2,26 ^b	15,81 ± 0,97°
14. Pelvic fin height	18,47 ± 2,70 ^a	18,32 ± 2,59ª	19,8 ± 2,55 ^b	18,28 ± 0,76ª
15. Tail fin height	21,06 ± 2,82ª	21,14 ± 3,4ª	23,16 ± 3,6 ^b	18,21 ± 1,47°
16. Anal fin height	11,02 ± 3,51ª	8,91 ± 1,52 ^{bc}	9,49 ± 3,37 ^b	8,53 ± 0,94°
17. Head length behind eyes	12,32 ± 1,70ª	13,13 ± 3,16 ^{ab}	13,01 ± 3,10 ^{ab}	13,6 ± 0,41 ^b
18. Muzzle length	11,7 ± 0,70 ^ª	11,65 ± 1,15ª	12,56 ± 1,89 ^b	12,43 ± 0,39 ^b

Note: Values in the same row with the same letter are not significantly different (p>0.05).

Table 7. Mean values (± standard deviation) of parameters compared to head length (HL) of *Oxyrichthys ophthalmonema*

Compared to HL (%)	Thua Thien Hue	e Quang Tri	Quang Binh	Ha Tinh
1. Head Width	56,73 ± 4,53ª	58,38 ± 4,38 ^b	57,44 ± 4,24 ^{ab}	54,74 ± 1,14 ^c
2. Head Height	70,25 ± 4,96ª	68,88 ± 4,95ª	69,72 ± 5,19ª	63,37 ± 2,59 ^b
3. Distance between eyes	21,08 ± 3,04ª	18,43 ± 3,09 ^b	17,56 ± 3,04 ^b	17,87 ± 1,43 ^b
4. Eye Diameter	22,76 ± 3,33ª	23,66 ± 3,69ª	25,19 ± 2,49 ^b	23,39 ± 1,7ª
5. Mouth width	54,68 ± 9,62ª	48,56 ± 9,43 ^b	47,6 ± 5,3 ^b	42,53 ± 3,78 ^c

Note: Values in the same row with the same letter are not significantly different (p>0.05).

Among 23 surveyed criteria, the ratio of criteria has a significant difference (p<0.05) between fish populations distributed in 4 provinces as follows: Thua Thien Hue and Quang Tri only differ in 11 criteria (48%), Thua Thien Hue and Quang Binh differ in 14 criteria (61%), Thua Thien Hue and Ha Tinh differ in 18 criteria (78%), Quang Tri and Quang Binh differ in 7 criteria (28%), Quang Tri and Ha Tinh differ in 12 criteria (57%), Quang Binh and Ha Tinh differ in 12 criteria (52%).

The research results indicated that if *O. ophthalmonema* in the provinces are close to each other, the average values of the measured criteria change little. However, if they are far apart, the mean values of those criteria have more changes between the two criteria. Among all the fish samples collected in Nghe An, we conducted analysis and determined that only one of them was *O. ophthalmonema* species, with morphological characteristics similar to those of Goby in the remaining 4 study areas. However, in such small quantities, it is not possible to compare the measurement and count critieria with other study areas.

Genetic diversity of goby fish

Total DNA extraction, PCR performance

Total DNA samples were extracted with good quality, 0.8% gel electrophoresis, and the electrophoresis images showed clear dark bands (Figure 4A).

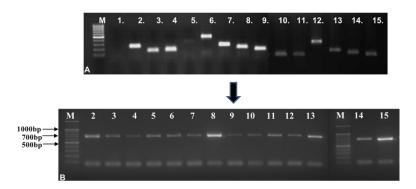


Figure 4. Total DNA electrophoresis (A): (M, Marker 1, HT009; 2, HT003 (MZ798205); 3, HT011 (MZ798213); 4, HTT012 (MZ798214); 5, QT004 (MZ798206); 6, QT007 (MZ798209); 7, QT008 (MZ798210); 8, NA013 (MZ798215); 9, TTHTG002 (MZ798204); 10, TTHTG006 (MZ798208); 11, QB005 (MZ798207); 12, QB009 (MZ798211); 13, QB010 (MZ798212), 14, TTHTG033 (MZ798216); 15, TTHTG034 (MZ798217). Results of PCR product electrophoresis (B): (M, Marker 2, HT003 (MZ798205); 3, HT011 (MZ798213); 4, HTT012 (MZ798214); 5, QT004 (MZ798206); 6, QT007 (MZ798209); 7, QT008 (MZ798210); 8, NA013 (MZ798215); 9, TTHTG002 (MZ798204); 10, TTHTG006 (MZ798208); 11, QB005 (MZ798207); 12, QB009 (MZ798211); 13, QB010 (MZ798212), 14, TTHTG033 (MZ798216); 15, TTHTG034 (MZ798217)

The 14 genomic DNA samples were amplified by PCR reaction using 2 pairs of primers FishF1 & FishR1. The PCR products were analyzed using a 0.8% agarose gel and the results showed that their sizes were about 650 - 750 bp (Figure 4B). The size of the PCR products of 14 fish samples in 5 areas is consistent with the theoretical size of the target *CO*I gene fragment.

COI gene sequence analysis results

After sequencing and editing with MEGA X software, all research samples were compared with similar sequences on Genbank using BLAST. As a result, the fish samples had a high similarity with other species (not *O. ophthalmonema*) while the remaining 14 samples had a high similarity rate with the *O. ophthalmonema* species.

The nucleotide sequence of each fish sample (14 samples) was compared with the COI gene sequence of *O. ophthalmonema* on the Genbank. The sequence with code MZ798204 was selected for detailed analysis for nucleotide differences, other sequences not shown here. The fish sample MZ798204 has a 93.43% similarity with *O. Ophthalmonema (JX193749.1)*, with 58 nucleotide positions difference (Figure 5). Nucleotide frequencies of 14 samples were T 32.3%; C 25.3%; A 24.1%; G 8.2%.

JX193749.1 O. ophthalmonema MZ798204	10 20 30 40 50 60 70
JX193749.1 O. ophthalmonema MZ798204	80 90 100 110 120 130 140 AATTCGAGCTGAACTAAGTCAGCCAGGAGCTCTTTTTGGTGACGACCAAATCTACAATGTCATTGTTACT TACC.TT.
JX193749.1 O. ophthalmonema MZ798204	150 160 170 180 190 200 210 GCACATGCCTTTGTAATAATTTTCTTTATAGTAATGCCAATTATGATTGGAGGGGTTTGGAAACTGACTTA C
JX193749.1 O. ophthalmonema MZ798204	220 230 240 250 260 270 280 TCCCACTAATGACTTGGCGCGCGCGCTGATATGGCCTTCCTCGGATGAATAATATGAGTTTTTGACTTCTTCC T.AC.
JX193749.1 O. ophthalmonema MZ798204	250 300 310 320 330 340 350 ACCTTCTTTCCTCCTCTTAGCATCCTCAGGGGTAGAGGCAGGGGGGGG
JX193749.1 O. ophthalmonema MZ798204	360 370 380 390 400 410 420 CCCCCACTAGCGGGCAATCTAGCACATGCTGGGGGCTTCTGTAGACTTAACAATTTTTTCTCTTCATTTGG
JX193749.1 O. ophthalmonema MZ798204	430 440 450 460 470 480 490
JX193749.1 O. ophthalmonema MZ798204	500 510 520 530 540 550 560 TTCACAGTACCAAACACCTCTCTTTGTTTGAGCTGTACTAATCACTGCCGTTCTTTTATTACTATCCCTC G. C. T. T. T.
JX193749.1 O. ophthalmonema MZ798204	570 580 590 600 610 620 630 CCTGTTCTTGCTGCTGGCATTACAATGCTTCTTACAGATCGCAACCTCAACACAAACCTTCTTTGACCCTT ACT
JX193749.1 O. ophthalmonema MZ798204	640 650 660 670 680 CTGGAGGAGGTGACCCAATCCTTTACCAACATCTATTC

Figure 5. Results comparing nucleotide sequences of fish sample MZ798204 with similar sequences in Genbank

The detailed analysis of the remaining sequences shows that fish samples named MZ798206 and MZ798213 have 93.65% similarity with *O. ophthalmonema* (JN242585.1), in which there is a difference in 41 nucleotide positions. However, both samples differed in 40 similar positions, and 1 nucleotide position difference from the COI gene sequence of *O. ophthalmonema* was different. Fish samples MZ798205, MZ798214, MZ798216, MZ798217, MZ798208, MZ798210, MZ798209, MZ798207, MZ798211, MZ798212 and MZ798215 have 93.81% similarity with *O. ophthalmonema* (JN242585.1), with a difference in 40 nucleotide positions. In these samples, the MZ798207 pattern at the 348th different nucleotide position (T-G) is different from the rest (T-A). The most similarity to Favonigobius gymnauchen (KJ013043.1) was 99.84%, with only 1 nucleotide position could not be identified, and the similarity to *O. microlepis* (MK777350.1) was 99.67%, with 2 nucleotide positions difference, but only 495 nucleotide sizes were compared.

Results comparing gene distance and similarity between *CO*I gene fragments of 14 samples of *O*. *ophthalmonema* are shown in table 8.

Table 8. The similarity of COI gene of 14 samples of O. ophthalmonema in central Vietnam

	MZ	MZ	MZ	MZ	MZ	MZ								
	798	798	798	798	798	798	798	798	798	798	798	798	798	798
	204	216	217	208	206	209	210	207	211	212	205	213	214	215
MZ														
798	***	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
204	**	089	089	029	044	029	029	044	029	029	029	044	029	029
MZ														
798	0.9	***	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
216	91	**	000	059	074	059	059	074	059	059	059	074	059	059
MZ														
798	0.9	1,0	***	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
217	91	00	**	059	074	059	059	074	059	059	059	074	059	059
MZ														
798	0.9	0.9	0.9	***	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
208	97	94	94	**	015	000	000	015	000	000	000	015	000	000
MZ														
798	0.9	0.9	0.9	0.9	***	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
206	95	92	92	98	**	015	015	029	015	015	015	029	015	015
MZ														
798	0.9	0.9	0.9	1,0	0.9	***	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
209	97	94	94	00	98	**	000	015	000	000	000	015	000	000
MZ 798	0.9	0.9	0.9	1,0	0.9	1,0	***	0.0	0.0	0.0	0.0	0.0	0.0	0.0
210	0.9 97	0.9 94	0.9 94	00	98	00	**	0.0	0.0	0.0	0.0	0.0	0.0	0.0
	57	51	51	00	50	00		010	000	000	000	015	000	000
MZ	0.0	0.0	0.0	0.0	0.0	0.0	0.0	***	0.0	0.0	0.0	0.0	0.0	0.0
798 207	0.9 95	0.9 92	0.9 92	0.9 98	0.9 97	0.9 98	0.9 98	**	0.0 015	0.0 015	0.0 015	0.0 029	0.0 015	0.0 015
	55	52	52	50	57	50	50		015	015	015	025	015	015
MZ		0.0		1.0	0.0	1.0	1.0	0.0	***		0.0			
798 211	0.9 97	0.9 94	0.9 94	1,0 00	0.9 98	1,0 00	1,0 00	0.9 98	**	0.0 000	0.0 000	0.0 015	0.0 000	0.0 000
	97	94	94	00	90	00	00	90		000	000	015	000	000
MZ														
798	0.9	0.9	0.9	1,0	0.9	1,0	1,0	0.9	1,0	***	0.0	0.0	0.0	0.0
212	97	94	94	00	98	00	00	98	00	<u>ት</u> ት	000	015	000	000
MZ														
798	0.9	0.9	0.9	1,0	0.9	1,0	1,0	0.9	1,0	1,0	***	0.0	0.0	0.0
205	97	94	94	00	98	00	00	98	00	00	* *	015	000	000
MZ														
798	0.9	0.9	0.9	0.9	0.9	0.9	0.9	0.9	0.9	0.9	0.9	***	0.0	0.0
213	95	92	92	98	97	98	98	97	98	98	98	**	015	015
MZ														
798	0.9	0.9	0.9	1,0	0.9	1,0	1,0	0.9	1,0	1,0	1,0	0.9	***	0.0
214	97	94	94	00	98	00	00	98	00	00	00	98	**	000

1		1	1	1	l	I	1	1	1	1	l	I	I	I	I I
	MZ														
	798	0.9	0.9	0.9	1,0	0.9	1,0	1,0	0.9	1,0	1,0	1,0	0.9	1,0	* * *
	215	97	94	94	00	98	00	00	98	00	00	00	98	00	**

Genetic relationships of Goby populations distributed in central Vietnam

Development of a phylogenetic tree of 14 fish samples with 10 *CO*I sequences of 5 species of the genus *Oxyurichthys* (sequences from Genbank) is shown in figure 6. The results indicated that all fish samples belong to the same clade as *O. ophthalmonema* with a bootstrap value of 64%.

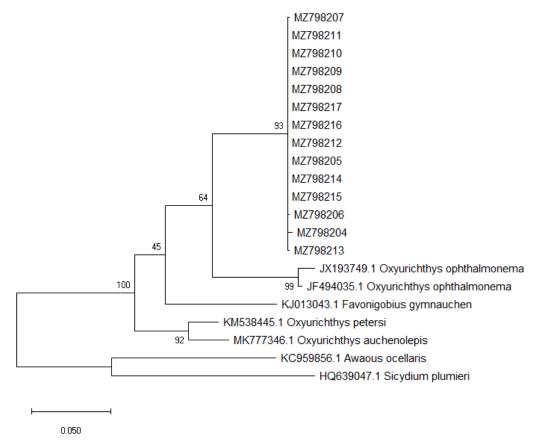


Figure 6. Relative relationship of Goby in central Vietnam with 5 species of genus Oxyurichthys

Through the results of sequence analysis and pedigree building of 14 fish samples identified as *O. ophthalmonema* species, the following results were obtained: the Goby samples in the same province or between the study areas have differences in the number of nucleotides and in the sequence of one or several nucleotides (MZ798204, MZ798206, MZ798207, MZ798215, MZ798213, MZ798216, MZ798217). However, the remaining fish samples in the study areas did not show any significant difference (MZ798205, MZ798214, MZ798209, MZ798210, MZ798208, MZ798211, MZ798212). The difference between the *COI* gene sequence of *O. ophthalmonema* (Genbank) and the fish samples is quite high. This difference was expressed in the positions of 40 to 43 nucleotide, the similarity is not high. This can be explained because the *COI* gene sequence region is prone to mutations, thus leading to errors in the pairing process as well as efficient use of primers in PCR.

In summary, when studying the morphological characteristics of all samples distributed in different areas in the Central region, the obtained results showed that there were differences in the morphology of the research subject in each analysis criterion. Although they were only small differences, they were also significant in specific features for identification. When studying the COI gene sequencing and analyzing the genetic tree, it was found that the Goby populations had genetic diversity showing differences in nucleotide sequences in the analyzed genes. However, there were still some differences in nucleotide sequences. The genetic results firmly confirmed the morphological taxonomic criteria. Therefore, for fish species whose morphology is still uncertain, gene sequencing is required to identify not only the *CO*I gene but also many other genes in the fish's mitochondria. For 1 out of 15 samples collected from Nghe An, when analyzing genetics, the similarity with the bootstrap value was still 100%. However, for this area, it is necessary to collect more samples and conduct genetic analysis to get accurate data.

CONCLUSION

Species *O. ophthalmonema* had an average size ranging from 67.70 - 190.20 mm, corresponding to a weight of 4.70 - 29.25g. It was found that there was a correlation between body length and body mass. Among 23 surveyed criteria, the ratio of criteria has significant difference (p<0.05) between fish populations distributed in 4 study areas as follows: Thua Thien Hue and Quang Tri are different in 11 criteria (48%), Thua Thien Hue and Quang Binh differ in 14 criteria (61%), Thua Thien Hue and Ha Tinh differ in 18 criteria (78%), Quang Tri and Quang Binh differ in 7 criteria (28%), Quang Tri and Ha Tinh differ in 13 criteria (57%), Quang Binh and Ha Tinh differ in 12 criteria (52%).

19 total DNA samples of Goby in 5 areas of central Vietnam were extracted, in which 14 samples were amplified and sequenced for *CO*I gene region; the results obtained the sequences that had differences at different positions in comparison to the COI gene region of *O. ophthalmonema*. The phylogenetic tree construction for 14 fish samples with COI sequences of *O. ophthalmonema* and 5 species of the genus *Oxyurichthys* showed that the fish samples belonged to *O. ophthalmonema* species with a 64% bootstrap value. However, their genetic distances were far apart and there were also some differences between the researched fish samples.

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