

Newly Collected Specimens of the Sleeper *Eleotris acanthopoma* (Teleostei: Eleotridae) from French Polynesia Indicate a Wide and Panmictic Distribution in the West and South Pacific¹

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Abstract: The morphology of *Eleotris acanthopoma* collected from Moorea in French Polynesia is described. This is the first record of this species from French Polynesia, greatly expanding the known range, which was previously only considered to extend from southern Japan to New Caledonia. Nucleotide sequences of the mitochondrial ND5 gene of several *Eleotris* species and related genera indicate that *E. acanthopoma* from Moorea belongs to the same lineage as *E. acanthopoma* from Japan and the Philippines. Despite being separated by a distance of approximately 10,000 km, two of the specimens from Moorea and one from the Philippines had identical nucleotide sequences. Results of this study indicate that extensive dispersal occurs during the pelagic larval stage of this species.

FISHES OF THE genus *Eleotris* have a global distribution, primarily inhabiting estuaries and freshwater streams of tropical and subtropical regions (Pezold and Cage 2002). In all of the species studied to date, newly hatched larvae are very small and undeveloped (Lindstrom 1999, Dotsu et al. 2004, Maeda et al. 2008) and spend a relatively long time as pelagic larvae (2–5.5 months) before becoming established in their adult habitats (Bell et al. 1995, Maeda and Tachihara 2005, Maeda et al. 2007). During this pelagic stage, the larvae are often transported considerable

distances from their natal habitats (Maeda et al. 2007). Consequently, several *Eleotris* species have been reported to have extensive geographical distributions. In particular, *Eleotris fusca* may be the most widely distributed amphidromous fish, with a range extending from South Africa to Japan and French Polynesia (Pusey et al. 2004). However, because the taxonomy of this genus has not yet been sufficiently clarified, determining species distribution ranges is difficult. It is therefore necessary to accurately identify and record all occurrences of *Eleotris* species to increase our understanding of their distribution.

Although *E. fusca* has been recorded on most of the mountainous islands in the Marquesas, Austral, Gambier, and Society islands (Fowler 1932, Herre 1931, 1932, Marquet and Galzin 1992, Resh et al. 1999, Englund 2003, 2004), other *Eleotris* species have not yet been reported in French Polynesia. However, here we describe the morphologies of five *Eleotris acanthopoma* specimens collected on Moorea and in so doing establish a record of this species in French Polynesia. To discuss the phylogeography of this species, nucleotide sequences of the mitochondrial ND5 gene of these specimens were compared with those of other *Eleotris* species, including *E. acanthopoma* and related genera mainly from the Northwest Pacific.

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MATERIALS AND METHODS

Five *Eleotris acanthopoma* specimens (URM-P 45009–45013; 25.3–45.7 mm SL; three adult males, one adult female, and one juvenile) were collected on the coast of Papetoai, Moorea, French Polynesia (17° 29' S, 149° 52' W) on 4 December 2006 by K.M. with a hand net. Tissue samples obtained from the right pectoral fins of four of the five specimens were fixed in 99% ethyl alcohol for DNA analysis. Specimens, including the fish not used for tissue samples, were fixed in 10% formalin and preserved in 70% ethyl alcohol for morphological analysis.

Morphological features were described following the methods given in Nakabo (2002), with the following exceptions: scale counts followed Akihito (1967), notation for the pattern of interdigitation of the dorsal fin pterygiophores between the neural spines (P-V) followed Akihito (1984), and body depth was measured at the base of the pectoral fin and at the origin of the anal fin. Measurements were taken from the left side using a dial caliper and micrometer to the nearest 0.1 mm. The number of vertebrae was counted from radiographs. Cephalic sensory papillae were observed by staining the specimens with cyanine blue.

Total genomic DNA of four *E. acanthopoma* specimens was extracted from the right pectoral fin samples. DNA extraction, gene amplification, and sequencing of a part of the mitochondrial NADH dehydrogenase subunit 5 (ND5) gene (ca. 1 kbp) were performed following the methods in Mukai et al. (2005). The nucleotide sequences obtained were

compared with the same gene region in *E. acanthopoma* specimens collected in Japan ($n = 7$) and the Philippines ($n = 2$); *E. fusca* collected in Japan ($n = 3$), the Philippines ($n = 11$), and Moorea, French Polynesia ($n = 1$); *Eleotris melanosoma* collected in Japan ($n = 1$); *Eleotris oxycephala* collected in Japan ($n = 2$); *Bunaka gyrinoides* collected in Japan ($n = 2$); and *Belobranchius belobranchus* collected in the Philippines ($n = 1$). The nucleotide sequences of these species were obtained using the same protocols as those used to analyze the French Polynesian *E. acanthopoma* samples, and all of the sequences were deposited in the GenBank/EMBL/DDBJ databases. In addition, a published nucleotide sequence of *E. acanthopoma* collected from Wakayama Prefecture, Japan (GenBank/EMBL/DDBJ accession number: AP004455 [Miya et al. 2003]) was also used for the analysis. The catalog numbers of the voucher specimens and accession numbers of the nucleotide sequences are provided in the Appendix.

RESULTS

Morphological Description

A photograph of *Eleotris acanthopoma* is shown in Figure 1. Dorsal fin rays VI–I,8; anal fin rays I,8; pectoral fin rays 16; pelvic fin rays I,5; caudal fin rays $i + 13 + i$ ($n = 3$) or $i + 14$ ($n = 2$); vertebrae $10 + 15 = 25$ ($n = 4$) or $10 + 16 = 26$ ($n = 1$); P-V 3/I II II I/8.

Scales in longitudinal row 50–54; scales in transverse row 13–15; scales in transverse series in the caudal peduncle 11–13. Ctenoid scales covering the lateral sides of the trunk



FIGURE 1. *Eleotris acanthopoma* (URM-P 45009, female, 45.3 mm in standard length) collected at Papetoai, Moorea, French Polynesia, on 4 December 2006.

TABLE 1

Morphometric Measurements of *Eleotris acanthopoma*
Collected in Moorea, French Polynesia

Parameter	Adults	Juvenile
Number of specimens examined	4	1
Standard length (mm)	33.3–45.7	25.3
Total length (mm)	42.5–58.3	32.7
In % of standard length		
Body depth at pectoral-fin base	18.6–20.8	17.8
Body depth at anal-fin origin	18.0–19.4	15.8
Head length	34.5–39.1	35.2
Snout length	8.1–11.0	9.5
Eye diameter	7.2–8.0	7.9
Postorbital length of head	19.8–21.8	20.6
Upper jaw length	11.7–13.8	13.0
Length of caudal peduncle	25.2–27.6	25.3
Depth of caudal peduncle	12.9–13.5	11.9
Predorsal length	41.0–43.8	43.5
Pre second dorsal-fin length	59.5–62.5	61.7
Preanal length	53.5–60.0	55.7
Length of first dorsal-fin base	15.5–17.7	16.6
Length of longest first dorsal-fin ray	13.2–15.0	15.4
Length of second dorsal-fin base	14.5–16.4	15.8
Length of longest second dorsal-fin ray	15.9–17.3	16.6
Length of anal-fin base	14.6–15.5	13.8
Length of longest anal-fin ray	15.9–17.1	16.5
Length of pectoral-fin base	9.3–9.9	8.7
Length of pectoral fin	24.9–27.0	22.5
Length of longest pectoral-fin ray	23.8–25.2	22.1
Length of pelvic fin	18.6–22.1	22.5

and almost the entire tail. Cycloid scales covering the dorsal trunk and head extending anteriorly to the interorbital area, upper opercle, belly, thorax, pectoral-fin base, and caudal-fin base.

Morphometric measurements are given in Table 1. Body elongate and cylindrical anteriorly and somewhat compressed posteriorly. Head somewhat depressed. Anterior nostril tubular; posterior nostril not tubular. Mouth oblique. Upper jaw extending below middle of pupil. Tip of lower jaw preceding tip of upper jaw. Anterior end of gill opening aligned vertically with the posterior margin of the preopercle. Second dorsal fin slightly higher

than first dorsal fin, and of nearly equal height as the anal fin. Posterior end of first dorsal fin base separate or nearly touching second dorsal fin origin. Spines of first dorsal fin not elongated. First dorsal fin, 2nd, 3rd, or 4th spines longest. Anal fin origin aligned vertically with the base of 1st or 2nd soft rays of second dorsal fin. Posterior end of anal fin base aligned vertically with the base of 7th or 8th soft rays of second dorsal fin. All soft rays of second dorsal fin and anal fin branched; second dorsal fin, 4th, 5th, or 6th soft rays longest; anal fin, 5th or 6th soft rays longest. Posterior tips of second dorsal fin and anal fin extending to middle of caudal peduncle when depressed. All pectoral fin rays branched, except for uppermost 1 and lowermost 0–1 rays. Pectoral fin without free ray, slightly exceeding posteriorly to vertical of base of 6th spine of first dorsal fin. Pelvic fin separated without frenum; all segmented rays of pelvic fin branched; 4th segmented ray longest. Caudal fin rounded; all segmented rays of caudal fin branched, except for uppermost 1 and lowermost 0–1 rays.

Urogenital papilla of adult specimens exhibiting sexual dimorphism, slender in male and broad and rounded in female.

Patterns of cephalic sensory systems are shown in Figure 2. Head lacking canal and canal pore. Posterior ends of two longitudinal rows of sensory papillae on opercle well separated from each other; infraocular area with five transverse rows of sensory papillae, 2nd and 4th of which extended beyond longitudinal infraocular row, 3rd of which just attached to the longitudinal row; lower margin of preopercle with five transverse rows of sensory papillae.

Color in preservation: Dorsal and dorso-lateral sides of body and head dark brown; ventral and ventrolateral sides of body and head creamy white background. Ventrolateral side of tail with six obscure dark brown transverse bands. Rather dark patches at 3, 4, 5, and 9 o'clock around eye and both upper and lower parts of pectoral-fin base. First dorsal fin translucent with dark brown proximal area and a longitudinal bar across middle part of fin; second dorsal fin translucent with dark brown proximal area and three to seven longitudinal bars; anal fin grayish brown without

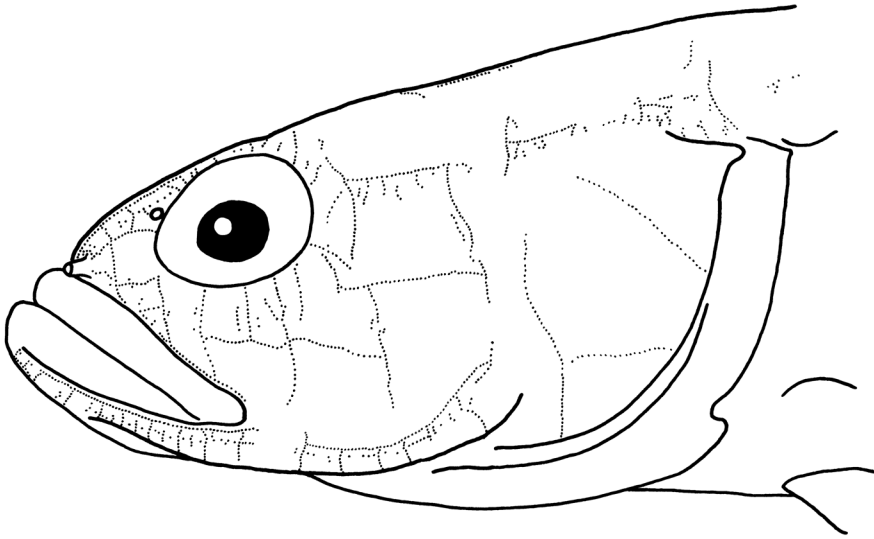


FIGURE 2. Head of *Eleotris acanthopoma* (URM-P 45009, 45.3 mm in standard length) showing arrangement of the cephalic sensory papillae.

distinct patterns; pectoral and pelvic fins uniformly pale grayish brown without distinct patterns; caudal fin translucent with eight to 10 dark brown transverse bars.

Ecological Notes

Eleotris acanthopoma was found together with juveniles of blacktail snapper, *Lutjanus fulvus*, in the vicinity of stones or wooden debris in shallow-water areas (5–20 cm in water depth) with muddy bottoms near the shore in north-western Moorea.

Mitochondrial DNA Phylogeny

A total of 995 bp of the mitochondrial ND5 gene sequence was determined or cited from 34 specimens of four *Eleotris* species and two species of related genera. The 995 characters included no indels (insertion and/or deletion events) and were perfectly aligned. A neighbor-joining tree of nucleotide sequences (haplotypes) from *Eleotris* and related genera (Figure 3) clearly grouped the four *E. acanthopoma* specimens from Moorea into the same lineage as *E. acanthopoma* specimens from Japan and the Philippines. Two individuals from Moorea and one from the Philippines had identical nucleotide sequences.

DISCUSSION

Akihito (1967) examined the morphology of 72 *Eleotris acanthopoma* specimens mainly from southern Japan and Taiwan, as well as Bleeker's holotype from Sumatra. The morphology of *E. acanthopoma* collected in Moorea is identical to that described by Akihito (1967) for all of the characters examined, including the arrangements of the cephalic sensory papillae and meristic characters. It is therefore suggested that there is no difference in morphology between *E. acanthopoma* from Moorea and those from the Northwest Pacific.

Eleotris acanthopoma was described originally by Bleeker (1853) based on specimens from western Sumatra. Records of this species are not commonly known; however, this is likely due to *E. acanthopoma* being considered a junior synonym of the congeneric *E. melanostoma* until Akihito (1967) demonstrated that they are two independent species. *Eleotris acanthopoma* has been reported from southern Japan, including Honshu, Shikoku, Kyushu, the Ryukyu Archipelago, and the Ogasawara Islands (Akihito 1967, Nakabo 2002); Taiwan (Akihito 1967); Hong Kong (Ni and Kwok 1999); Cebu, Philippines (this study); Lombok, Indonesia (Harada and Suharti 2000);

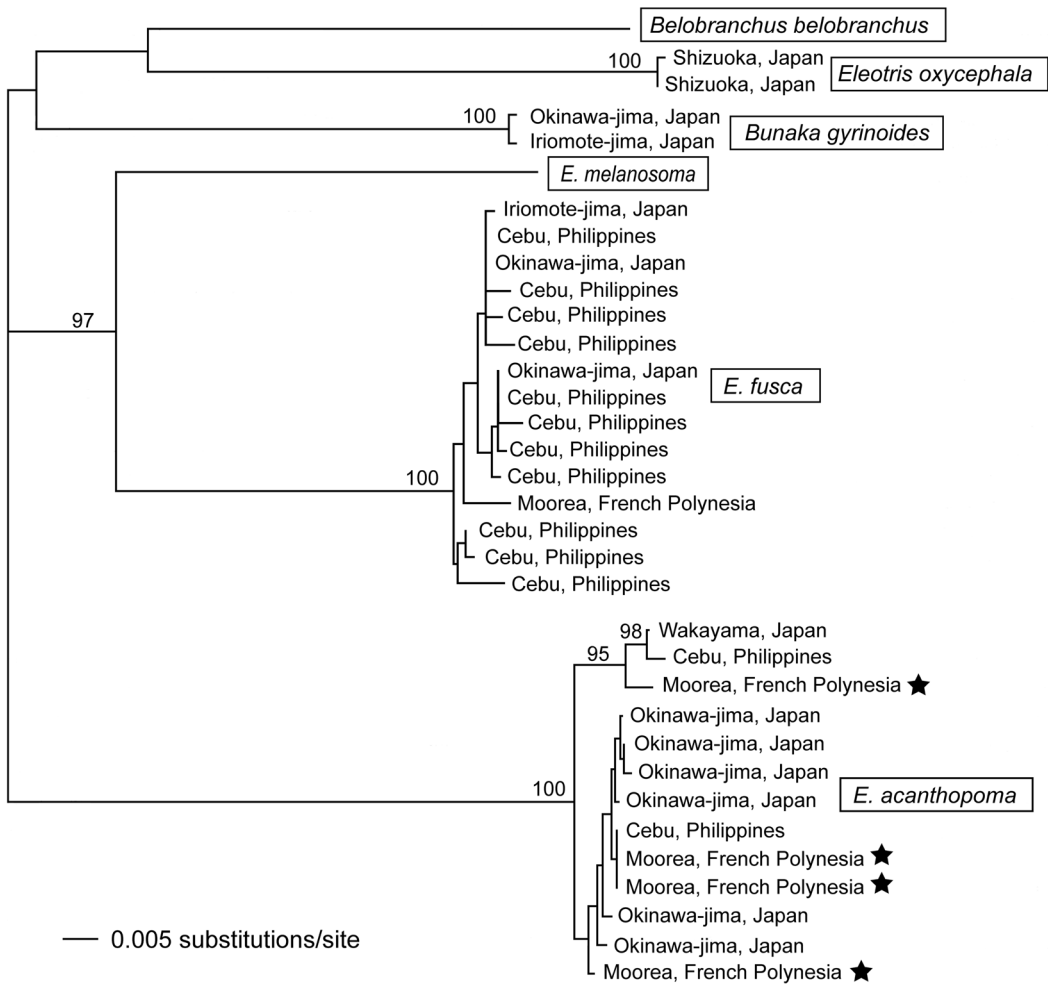


FIGURE 3. Neighbor-joining tree based on genetic distances estimated from partial mitochondrial NADH dehydrogenase subunit 5 (ND5) gene sequences (995 bp) of four *Eleotris* species including French Polynesian samples and two related genera. Distances were based on Kimura’s two-parameter model. Numbers adjacent to internal branches indicate bootstrap probabilities (>90%) based on 1,000 pseudoreplicates. Solid stars represent *E. acanthopoma* from French Polynesia.

the Mariana Islands (Donaldson and Myers 2002); northeastern Australia (Allen et al. 2002); the Solomon Islands (Blaber and Milton 1990); New Caledonia (Marquet et al. 2003), and Moorea (this study) (Figure 4). Because Moorea is located approximately 4,500 km from New Caledonia, the closest previously known locality of *E. acanthopoma*, this discovery markedly extends the known range of this species.

These findings therefore suggest that *E. acanthopoma* has a range that is considerably

larger than was previously thought, extending from the subtropical Northwest Pacific to the east-central tropical South Pacific. Maeda et al. (2007, 2008) indicated that *E. acanthopoma* larvae have a considerable capacity for dispersal. The mitochondrial DNA phylogeny presented here suggests that the panmictic *E. acanthopoma* populations in the tropical Pacific Ocean are widely distributed by the North and South Equatorial Currents and the Equatorial Counter Current during the pelagic larval stage.

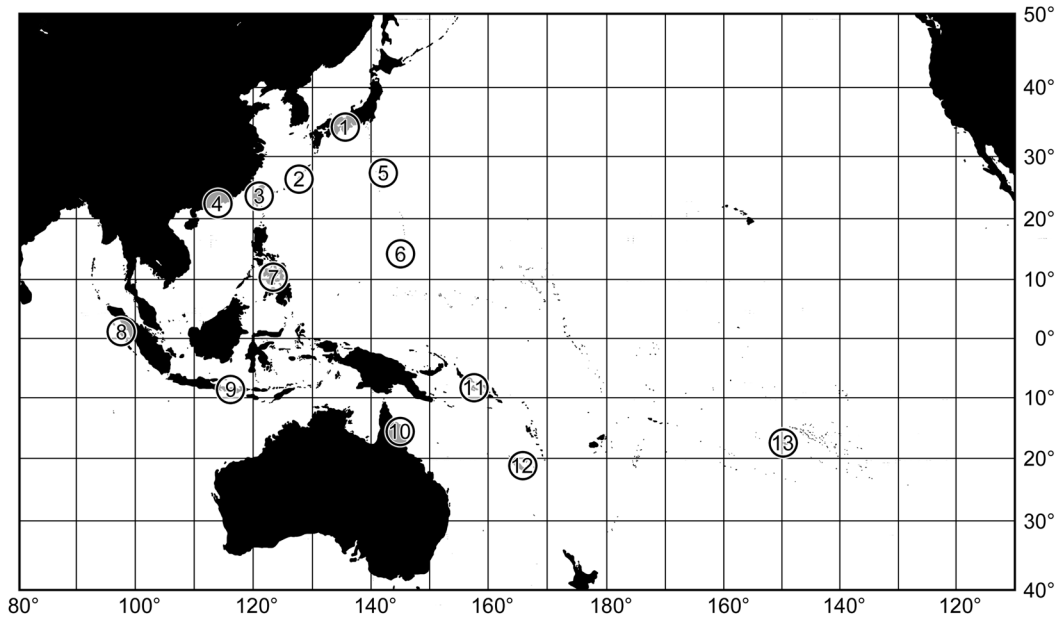


FIGURE 4. Geographical distribution of *Eleotris acanthopoma*: 1, Honshu, Shikoku, and Kyushu; 2, Ryukyu Archipelago; 3, Taiwan; 4, Hong Kong; 5, Ogasawara Islands; 6, Mariana Islands; 7, Cebu; 8, western Sumatra; 9, Lombok; 10, northeastern Australia; 11, Solomon Islands; 12, New Caledonia; 13, Moorea.

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Appendix
Samples for DNA Analysis

Species	Locality	Catalog No. of Voucher Specimen	Acc. No.
<i>Belobranchus belobranchus</i>	Danao, Cebu, Philippines	URM-P 45064	AB499299
<i>Bunaka gyrimoides</i>	Teima, Okinawa-jima, Japan	URM-P 45134	AB544426
	Uehara, Iriomote-jima, Japan	No voucher	AB499300
<i>Eleotris oxycephala</i>	Yoshida, Shizuoka, Japan	No voucher	AB499301
	Shizuoka, Shizuoka, Japan	KPM-NI 25014	AB544427
<i>Eleotris melanosoma</i>	Taiho, Okinawa-jima, Japan	KPM-NI 23490	AB499302
<i>Eleotris fusca</i>	Uehara, Iriomote-jima, Japan	No voucher	AB499303
	Yona, Okinawa-jima, Japan	URM-P 45150	AB544428
	Yona, Okinawa-jima, Japan	URM-P 45631	AB544429
	Panalipan, Cebu, Philippines	URM-P 45051	AB544430
	Panalipan, Cebu, Philippines	URM-P 45052	AB544431
	Panalipan, Cebu, Philippines	URM-P 45053	AB544432
	Panalipan, Cebu, Philippines	URM-P 45054	AB544429
	Panalipan, Cebu, Philippines	URM-P 45055	AB544433
	Panalipan, Cebu, Philippines	URM-P 45056	AB544434
	Panalipan, Cebu, Philippines	URM-P 45057	AB544428
	Panalipan, Cebu, Philippines	URM-P 45058	AB544435
	Panalipan, Cebu, Philippines	URM-P 45059	AB544436
	Panalipan, Cebu, Philippines	URM-P 45060	AB544437
	Danao, Cebu, Philippines	URM-P 45062	AB544438
	Moorea, French Polynesia	URM-P 45014	AB499304
<i>Eleotris acanthopoma</i>	Kirime, Wakayama, Japan	No voucher	AP004455
	Aritsu, Okinawa-jima, Japan	No voucher	AB499305
	Yona, Okinawa-jima, Japan	URM-P 45632	AB544439
	Yona, Okinawa-jima, Japan	URM-P 45633	AB544440
	Yona, Okinawa-jima, Japan	URM-P 45634	AB544441
	Yona, Okinawa-jima, Japan	URM-P 45635	AB544442
	Yona, Okinawa-jima, Japan	URM-P 45636	AB544443
	Dalaguete, Cebu, Philippines	URM-P 45047	AB499306
	Dalaguete, Cebu, Philippines	URM-P 45049	AB499307
	Moorea, French Polynesia	URM-P 45010	AB499307
	Moorea, French Polynesia	URM-P 45011	AB499308
	Moorea, French Polynesia	URM-P 45012	AB499307
	Moorea, French Polynesia	URM-P 45013	AB499309