Classification of the Intertidal and Estuarine Upogebiid Shrimps (Crustacea: Thalassinidea), and Their Settlement in the Ryukyu Islands, Japan

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Abstract
Upogebiid shrimps (Crustacea: Thalassinidea) inhabit sandstones exposed on the intertidal zone of two of the islands of the Ryukyu Arc, Japan (islands located between 24°N, 123°E and 28°N, 129°E). The upogebiids here are morphologically similar; so, their classification is open to debate. In addition, when the upogebiids settled on the Ryukyu Arc has not been studied till date. Based on morphological features, we suspect that six stone upogebiid species are distributed in the Ryukyu Arc. This hypothesis was tested by molecular phylogenetic analysis of mitochondrial DNA markers. Furthermore, Bayesian evolutionary analysis, which was used to estimate TMRCA (time to most recent common ancestor) of each upogebiid, showed that stone upogebiids diverged 5-15 Ma (second half of the Miocene). Stone upogebiids may have occurred in the islands near the equator 5-15 Ma. On the contrary, Bayesian analysis indicated that the time of ‘intraspecific’ divergence varied from 1.15 Ma to 0 Ma, suggesting that stone upogebiids may have settled in the Ryukyu Arc from the Pleistocene to the present. In Iriomote Island, sedimentary layers suitable for burrowing occurred on the shores from the west to the northern part of the island, but not on the east. The Ryukyu Arc was a passive continental margin until the beginning of the Pleistocene about 1.55 Ma. With subduction of the Ryukyu Arc northward, sedimentary layers for burrowing may have appeared on the ‘front’ of the island (i.e. on the shores from the west to the north), thereby providing suitable habitats for the stone upogebiids to settle, since 1.15 Ma.

Keywords
Stone upogebiids; Iriomote Island; Classification; Habitat; Bayesian evolutionary analysis; 1.55 Ma; Subduction of Ryukyu Arc

Introduction
Muddy shores appear in intertidal areas protected from open ocean wave activity, i.e. in partially enclosed bays, lagoons, harbors, and estuaries, especially [1]. While sand beaches have larger grain size, muddy areas consist of finer-grained sediment particles [2,3]. The slope of muddy shores is much flatter than that of sandy beaches; so, these areas are often referred to as mudflats.

Mudflats show high standing stocks of phytoplankton, as determined by measuring the chlorophyll concentration [3,4]. Detritus is also abundant. The inhabitants of tidal mudflats (infauna) burrow in the sediment, and they feed on detritus in the sediment and water. Thus, a number of mudflat inhabitants may be deposit and suspension feeders [5]. At low tide, their presence may be shown by small holes on the flats, small sediment mounds topped by a hole, or piles of feces [3]. Among the large number of mudflat inhabitants, thalassinidean decapods (Crustacea) make elaborate burrows in the sediment [6-10]. They may feed on detritus that is filtered from water [4].

Mud shrimps (Upogebiidae) have been described from many of the world’s marine and estuarine shores [8,11-13]. The main bases for upogebiid identification are the morphological characteristics of their exoskeleton. Nine species of upogebiid were recorded from the mudflats on the coasts of the main islands of Japan (Kyuushu, Shikoku, Honshu, and Hokkaido), where the substrates were mud, muddy sand, gravels, or boulders [14].

Upogebiids are also found in partially enclosed bays and estuarine shores in the subtropical islands of Japan. A feature of some of these upogebiids is that they burrow into siltstones or sandstones lying on the shore. These upogebiids were first examined by Hirano and Sakai [15,16]. However, some important morphological features were not considered in their classification. A significant issue, therefore, is the uncertainty over the number of stone upogebiids inhabiting the subtropical islands of Japan, like the Ryukyu Islands [13,15-18]. To resolve some of these debates on classification, we collected stone upogebiids in the Ryukyu Island Arc and reclassified them. Based on morphological features, herein we propose a hypothesis on their classification (hereafter the ‘Hirano-Saigusa hypothesis’).

Genetic relatedness based on molecular phylogenetic studies may be available for the identification of similar species [19-24]. The Hirano-Saigusa hypothesis could be tested by using molecular phylogenetic studies, such as those employing mitochondria DNA (COI) markers; this was, therefore, our second task.

The third task we aimed to undertake was to determine the relation between the distribution and characteristics of stone upogebiid habitats. By collecting a number of stone upogebiids in Iriomote Island, we noticed that they inhabit the shores from the west to the northern parts (i.e. west side) of the island, and that no upogebiids burrow in the sedimentary layers exposed on the eastern shores. This observation led us to question why stone upogebiid distribution was limited to the western side of the island. To answer this question, we first collected samples of the substrate and analyzed their grain size. Then, we examined whether the sedimentary layers that are buried were exposed in the lower part of the intertidal zone. Distribution of the habitat of stone upogebiids might be strongly related to the subduction of Iriomote Island towards the northwest.

The Ryukyu Arc formed the eastern margin of the Eurasian continent between the Cretaceous and the Oligocene, and was...
Collection of stone upogebiids and identification of morphological species

Stone upogebiids inhabit the shores of the Okinawa and Iriomote islands. These upogebiids make U-shaped burrows in relatively soft stones or sedimentary layers. Soft stones and sedimentary layers are found at the banks of the estuaries (Figure 2a), or on shores along the partially enclosed bays when the tide recedes. Stone upogebiids were collected using a chisel and hammer at these locations when the water level was low (Figure 2b). Upogebiids were fixed in 70% ethanol, and identification was made under a stereomicroscope (Stemi 2000-C, Zeiss) in the laboratory. Collection was mostly carried out every day during two weeks in May 2004, November 2005, and May 2007, and every day for one month from April to May in 2017 (Table 1).

Reclassification of stone upogebiids

Stone upogebiids collected from the Ryukyu Islands have previously been identified by Hirano and Sakai [15,16]. However, some characteristics were not considered in their studies. Upogebiids were re-examined by employing some new standards, and a reconstructed classification (Hirano-Saigusa hypothesis) has been proposed in this study (Table 2). Other studies [11,17,31-33] were also referred to.

1) U. spinidactylus inhabits the estuarine shores and mudflats of the mangrove flots of Iriomote Island, as has been described by Sakai and Hirano [16].
2) U. saigusai inhabits the shores of partially enclosed bays on Iriomote Island, as has been described by Sakai et al. [15], but there the specimens were described as U. snelliusi.
3) U. iriomotensis inhabits the estuarine shores of Iriomote Island, as has been described by Sakai and Hirano [16].
4) U. neogenii (new species), inhabits the estuarine shores of Iriomote Island, as is first described in this study.
5) U. semicircula (new species), lives in the sedimentary layers on the shores of the partially enclosed bays of Iriomote Island, as is first described in this study.
Figure 2: Habitat of stone upogebiids and their morphological features: a) Habitat (dotted line) of *U. iriomotensis*, *U. neogenii*, and *U. spinidactylus* on an estuarine shore of Iriomote Island (site e in Figure 7A). HTL: high tide line. b) Collection of upogebiids in the mangrove flat (site g in Figure 7A). c) The whole body of *U. spinidactylus* (dorsal view). d) Lateral view of the head (the square shown in Figure 2c). es: eyestalk, st: assemblage of the seta, dp (a double arrow): depression between the hard spines (i and iii), sp: spine, ve: ventilation hole, an: antenna, ME: merus. Hard spines arranged along both sides of the head are shown by the numbers i-viii. a) Dorsal view of the head (the square indicated in Figure 2c). st: assemblage of the seta, es: eyestalk, gv: groove, iii: the third spine. f) The outer side of male right chela. PR: propodus, DA: dactylus, st: long setae borne on the segment. A small projection characterizing this species is shown by the white arrow. g) The inner side of male right chela. ME: merus, CA: carpus, PR: propodus, DA: dactylus. Scale is shown in each panel.

Table 1: Stone upogebiids employed for molecular philogenetic analysis.

<table>
<thead>
<tr>
<th>Species</th>
<th>Collection</th>
<th>Number*</th>
<th>Latitude (N)</th>
<th>Longitude (E)</th>
<th>Habitat</th>
<th>Distribution</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Upogebia saigusai</em></td>
<td>Iriomote Is.</td>
<td>12</td>
<td>24°35'-41'</td>
<td>123°73'-77'</td>
<td>Shores of partially enclosed bay</td>
<td>Iriomote Island</td>
</tr>
<tr>
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<td>12</td>
<td>24°35'-41'</td>
<td>123°73'-77'</td>
<td>Shores of partially enclosed bay</td>
<td>Iriomote Island</td>
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<td>Iriomote Is.</td>
<td>12</td>
<td>24°35'-41'</td>
<td>123°73'-77'</td>
<td>Estuarine shores</td>
<td>Iriomote Island</td>
</tr>
<tr>
<td><em>Upogebia iriomotensis</em></td>
<td>Iriomote Is.</td>
<td>10</td>
<td>24°34'-41'</td>
<td>123°73'-77'</td>
<td>Estuarine shores</td>
<td>Iriomote Island</td>
</tr>
<tr>
<td><em>Upogebia mioceni neogenii</em></td>
<td>Iriomote Is.</td>
<td>6</td>
<td>24°34'-41'</td>
<td>123°73'-77'</td>
<td>Estuarine shores</td>
<td>Iriomote Island</td>
</tr>
<tr>
<td><em>Upogebia rupicola</em></td>
<td>Okinawa Is.</td>
<td>12</td>
<td>26°38'</td>
<td>128°02'</td>
<td>Shores of partially enclosed bay</td>
<td>Okinawa Island</td>
</tr>
<tr>
<td><em>Laomedia astacina</em></td>
<td>Honshu</td>
<td>3</td>
<td>34°22'</td>
<td>133°31'</td>
<td>Mudflat in the estuary</td>
<td>Honshu and the Ryukyu Arc</td>
</tr>
</tbody>
</table>

*Number of sample sequenced in this study. **Employed for the out group.

Table 2: Gene Bank accession number of stone and mudflat upogebiids.

<table>
<thead>
<tr>
<th>Hirano-Saigusa hypothesis</th>
<th>Accession number</th>
<th>Upogebiids that have already been registered in Gene Bank</th>
</tr>
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<tr>
<td><em>Upogebia saigusai</em></td>
<td>AB71737</td>
<td>Upogebia snelliisi</td>
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<tr>
<td><em>Upogebia spinidactylus</em></td>
<td>AB71740</td>
<td>Upogebia spinidactylus</td>
</tr>
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<td><em>Upogebia semicircula</em></td>
<td>AB71736</td>
<td>Upogebia sp.</td>
</tr>
<tr>
<td><em>Upogebia iriomotensis</em></td>
<td>AB71738</td>
<td>Upogebia iriomotensis (Type 1)</td>
</tr>
<tr>
<td><em>Upogebia neogenii</em></td>
<td>AB71739</td>
<td>Upogebia iriomotensis (Type 2)</td>
</tr>
<tr>
<td><em>Upogebia rupicola</em></td>
<td>AB71735</td>
<td>Upogebia carinicauda</td>
</tr>
<tr>
<td><em>Laomedia astacina</em></td>
<td>AB71741</td>
<td>Laomedia astacina</td>
</tr>
</tbody>
</table>
6) *U. rupicola* inhabits the siltstone or sandstone layers on Okinawa Island, as described by Komai [18].

**Distribution of stone upogebiids in Iriomote and Okinawa Islands, and grain-size analysis of stone habitats**

We examined the soft stones burrowed into by the upogebiids along a number of shores of several islands. Stone upogebiids were only found in Iriomote Island and Okinawa Island (Figure 1). The distribution of stone upogebiids was investigated along the shores of these islands, especially Iriomote Island (Table 3).

Sedimentary stones were classified into three categories based on their grain size: puddingstone (more than 2 mm), sandstone (from 62 µm to 2 mm), and mudstone (less than 62 µm) [2]. Stones burrowed into by the upogebiids were collected from four locations on Iriomote-island, and were pulverized using a mortar and pestle. Grain size was estimated by sifting through screens of 10 mesh sizes, from 20-µm to 4-mm.

**DNA extraction, PCR, sequencing, sequence alignment, and phylogenetic analysis**

Genomic DNA was extracted from the chela muscle of the specimens fixed in 70% ethanol using a DNeasy Blood & Tissue kit (Qiagen, Hilden, Germany). Partial fragments of the mitochondrial DNA *Cytochrome c Oxidase Subunit I* (*COI*) were amplified by polymerase chain reaction (PCR) using genomic DNA as template and a primer set described by Folmer et al. [34]. PCR products were purified for direct sequencing using an ExoSAP-IT (Thermo Fisher Scientific, Waltham, MA, USA) according to the manufacturer's protocol. Sequencing reactions were performed using HCO2198 primer and a BigDye Terminator v3.1 Cycle Sequencing kit (Thermo Scientific, Waltham, MA, USA). Sequenced DNA was analyzed with an ABI 3100 automated sequencer (Applied Biosystems, Foster City, CA, USA).

**Estimation of divergence time**

Phylogenetic analyses were performed by Bayesian inference in MrBayes, version 3.2.6 [28] for a GTR+G+I substitution model. We estimated the time to the most recent common ancestor (TMRCA) of the different phylogenetic groupings, using the arthropod COI substitution rate of 0.0115 per site per million years in BEAST, version 1.8.4 [35,36]. The species tree prior was set to the default option of the Yule process. Markov chain Monte Carlo (MCMC) analyses were run for 1 billion steps, with trees sampled every 1,000 steps and the first 10% of the runs being discarded as burn-ins.

**Results**

**Morphological characteristics of upogebiids**

The morphological features of stone upogebi, *Upogebia spinidactylus* (dorsal view) revealed that its head (rostrum) is not sharp, but round (Figure 2c). Many hard spines are arranged along both sides of the head. In *U. spinidactylus*, eight projections (i-viii in Figure 2d) were distinguished along the head edge, and a groove (gv in Figure 2e) was formed inside the continuous hard spines.

Some species possessed a clear depression or space (dp in Figure 2d) between the hard spines. In *U. spinidactylus*, a depression was found between spines ii and iii, where the tip of the eyestalk is seen from the top. Others, e.g., *U. semicircula*, lacked this depression, and the eyestalk was not visible from the top.

In all species of Upogebiidae, including mudflat species, the eyestalk (es) was positioned beneath the hard parts of the head (Figure 2d). In *U. spinidactylus*, a sharp spine was borne along the edge of the exoskeleton around the base of the eyestalk (sp in Figure 2d). Some species (*U. saigusai*) bore several spines along the carapace edge, but others (*U. iriomotensis* and *U. neogenii*) lacked spines there. In stone upogebiids, the tip of the propodus (PR in Figures 2f and 2g) was not degenerated. The tips of the chela (propodus and dactylus) crossed, and this feature is common to many species of infaunal decapods, including the Axiidae, Laomediidae, and Callianassidae. Upogebiids may also be characterized by a number of long fine setae arranged on each segment of the pereiopods (st in Figures 2f and 2g).

**Comparison of morphology among stone upogebiids: the Hirano-Saigusa hypothesis**

A sketch of three stone upogebiids collected from the shores of Iriomote Island is shown in Figures 3a-3c. *Upogebia saigusai* (Figure 3a) is a typical example of a species that inhabits soft sandstone, which is distinguished by its thick head and long, thin setae. *Upogebia semicircula* (Figure 3b) is a species that inhabits hard sandstone, distinguished by its long head and thin eyestalk. *Upogebia iriomotensis* (Figure 3c) is a species that inhabits mudstone, distinguished by its short head and thick eyestalk.
3a) was most easily discriminated among the stone upogebiids because of the presence of a number of hard spines on its chelae. The head was round, and two hard spines are borne at the tip (Figure 3a-1). The tip of the eyestalk (es) was somewhat visible from the top. In lateral view (Figure 3a-2), a sharp spine (sp1) was borne from the base of the eyestalk. A large, sharp spine (sp 2) was also borne on the base of the antennae. A unique feature was a large spine (sp 3) positioned beneath the tip of the head (photo above Figure 3a-2). Five to six small spines were borne by the carapace edge (solid arrow without symbol in Figure 3a-2). Furthermore, U. saigusai was characterized by a number of small spines borne on the propodus (PR), carpus (CA), and merus (ME), as indicated by the solid arrows in Figure 3a-3. We could not find characteristics useful for identification on the abdomen and tail of this species (Figure 3a-4).

In combination with its habitat, U. semicircula was easily identified morphologically. The head of U. semicircula was round, and several small projections were borne on the head (Figure 3b-1). The eyestalk was not visible from the top. In lateral view, a spine was seen at the base of the eyestalk (solid arrow in Figure 3b-2). Small spines were also borne at edges of the carpus and merus (arrows without symbol in Figure 3b-3). No spine was borne under the propodus (Figure 3b-3). No clear characteristics were found on either the abdomen or tail (Figure 3b-4).

A feature of U. spinidactylus was a deep depression between two spines (ii and iii) at the head, from which the groove (gv) extended (Figures 2e and 3c-1). This depression was also visible in lateral view between spines ii and iii (Figure 3c-2). A spine was borne along the carapace edge under the head (solid arrow in Figure 3c-2). U. spinidactylus was characterized by a small projection borne on the dactylus (dotted arrows in Figure 3c). Several spines were borne on the edge of the pereiopods (solid arrows in Figure 3c-3). No clear characteristics were found on either the abdomen or tail (Figure 3c-4).

A small depression between the hard spines was also seen on the head of U. iriomotensis (Figure 4a-1). The tip of the eyestalk (e) was visible through the depression from the top. The groove (gv) on both sides of the head was weakly formed. In lateral view, no spine was borne near the base of the eyestalk (Figure 4a-2). Spines were borne at the frontal edge of the carapace, and at the lower end of the propodus (CA and PR in Figure 4a-3). No clear characteristics were found on either the abdomen or tail (Figure 4a-4).

Morphology of the carapace (cephalothorax) was much the same between U. iriomotensis (Figure 4a) and U. neogenii (Figure 4b). In U. neogenii, the eyestalk was slightly visible from the top (Figure 4b-1). No spines were borne at the base of the eyestalk (Figure 4b-2). The position and arrangement of spines were much the same as in U. iriomotensis (compare Figure 4b-3 with Figure 4a-3). The crossing of the dactylus (DA) and propodus (PR) at the tips was also the same as in other stone upogebiids. However, a unique feature of U. neogenii is the presence of long, brush-like setae borne on the dactylus and upper

![Figure 3: Comparison of morphological characteristics in stone upogebiids (3 species).](image-url)
propodus (Figure 4b-1, photo on the right; and Figure 4b-3). Brush-like setae are possibly due to setules borne on the seta.  

*U. rupicola* (Figure 4c) was only distributed in Okinawa Island (Figure 1). Several spines were borne along the top edge of the head. A groove was developed along both sides of the head (Figure 4c-1, left). The head (Figure 4c-1, photo on the right) resembled that of *U. semicircula* (Figure 3b-1, photo on the right). In lateral view (Figure 4c-2), a spine was borne along the carapace beneath the head (near the base of the eyestalk). Sexual dimorphism appeared a little stronger for this species than in the other five stone upogebiids examined (Figures 4c-3 and 4c-4). Several protrusions were arranged on the outer edge of the male propodus (PR) (photo above Figure 4c-3). A continuous arrangement of these protrusions was not obvious in the female. Many spines were borne by the edges of the propodus (PR), carpus (CA), and merus (ME) in both males and females (Figures 4c-3 and 4c-4).

Sexual dimorphism was not remarkable in these upogebiids. The only outstanding differences were those in the relative length of the merus (ME). ME length of females was comparatively greater than that of males (Figures 4c-3 and 4c-4).

**Molecular phylogeny of stone upogebiids**

A molecular phylogenetic tree was constructed based on 458 bp of mtDNA (COI) sequences (Figure 5). *U. spinidactylus* was estimated to have diverged about 13.85 Ma (middle Miocene, Tertiary), suggesting that reproductive isolation may have occurred very early among the stone upogebiids. The other group may have diverged about 12.34 Ma (middle Miocene). *U. saigusai* was separated from *U. semicircula* and *U. rupicola* about 10.03 Ma (late Miocene). *U. iriomotensis* was separated from *U. neogenii* about 5.32 Ma (end of Miocene).

As indicated in Figure 5, the TMRCA in *U. spinidactylus* (1.15 Ma) was the greatest among the six stone upogebiids tested (middle Pleistocene, Quaternary), and it was 0.61 Ma for *U. saigusai* (late Pleistocene). In *U. semicircula*, mtDNA (COI) sequence data of 10 individuals were identical, suggesting that *U. semicircula* settled in the Ryukyu Islands most recently among all the stone upogebiids. The TMRCA of each of the other three upogebiid species (*U. rupicola*, *U. iriomotensis*, and *U. neogenii*) was estimated to be about 0.6-0.7 Ma (late Pleistocene).

**Habitat of stone upogebiids in Iriomote and Okinawa Islands**

Stone upogebiids and their habitats are summarized in Figure 6. *U. saigusai* was found on the shores of the partially enclosed bays on Iriomote Island (Figures 6a and 6c). The habitat was exposed by the retreat of sea water during low tide. The burrow possessed two openings and was U-shaped, as observed for all the other upogebiids (Figure 6b). The burrow was dug not only in stones lying on the
shore, but also in the sedimentary layer exposed on the shores. The shores of the partially enclosed bays were also inhabited by *U. semicircula* (Figure 6d), and both *U. saigusai* and *U. semicircula* were also collected from the same habitat (Figure 6a).

*U. iriomotensis* (Figure 6h) and *U. neogenii* (Figure 6i) inhabited the burrows dug in stones lying on estuarine shores. Figure 6g indicates a stone wall made on a stream bank around 1940 (80 years ago). Burrows of *U. iriomotensis* and *U. neogenii* were found at the bottom of the stone wall. *U. iriomotensis* burrowed in not only soft stones but also in a decayed tree half-buried in the mudflat (Figure 6h). *U. neogenii* was collected from soft sandstones lying on the river bed. Soft sandstone strata exposed along the shores of estuaries were also inhabited by *U. spinidactylus* (Figures 6e and 6f). *U. spinidactylus* was distributed as far as inside the estuary.

*U. rupicola* (Figure 6j) was only collected from the shores of Okinawa Island (Figure 1). The habitat (26°6’N 127°39’E) was the soft siltstone-strata exposed at the time of low tide (Figure 6k). These strata were 300 m distant from the shore (low tide). The siltstone strata extended 10 km along the shore (figure not shown). A second habitat (26°40’N 128°0’E) was found on the shore of a small island (Yagachi Island) 100 km north from the first habitat (figure not shown). Yagachi Island was also positioned on the west side of Okinawa Island.

**Topography of Iriomote island and distribution of stone upogebiids**

Five species of stone upogebiids were identified on Iriomote Island (Tables 1 and 2). *U. saigusai* and *U. semicircula* were found from shores in partially enclosed bays. *U. iriomotensis* and *U. neogenii* inhabited the sedimentary layers in estuaries. *U. spinidactylus* was found as far as inside the estuary. A noteworthy feature of their distributions was that they were limited to the western and northern shores of the island (Figure 7a and Table 3). The seacoast in the south
Figure 6: Stone upogebiids and their habitat. a) Habitat of *U. saigusai* and *U. semicircula* (enclosed by a white line). Sedimentary layers and stones on the shore of a bay that is partially enclosed (site b in Figure 7A). Photographs (a,e,g,k) were taken at the time of low water. b) U-shaped burrow of *U. saigusai* with two openings (open arrow). Scale, 1cm. c) *U. saigusai*, d) *U. semicircula*. Scale (c, d, f), 1 cm. e) Major habitat of *U. spinidactylus* (enclosed by a white line) in an estuarine shore (site f in Figure 7A). g) Habitat of *U. iriomotensis* and *U. neogenii* in an estuarine shore (site e in Figure 7A). The stone wall was about 1 m high on the mudflat. h) *U. iriomotensis* burrowed in a decayed tree half buried in the mud. Scale, 1cm. i) *U. neogenii*. j) Habitat of *U. rupicola* on the shore (Okinawa Island). k) *U. rupicola* burrowing into a siltstone layer. Scale, 1 cm.

was made up of steep cliffs, and was directly exposed to strong waves because of less developed coral reefs (figure not shown). Southern shores consisted of sand and hard rocks (limestone); so, there was no habitat for upogebiids on the shore. In the east of the island, there are extensive coral rubble flats (reef flats), and there is a distance of more than 1 km between the land and reef crest. Sandstones and sedimentary layers were found in the estuary there, but they were hard to crack by a chisel and hammer.

Grain size analysis

Stones inhabited by upogebiids were collected from the shores at four sites of Iriomote Island: Mitara, Hoshidate, Urauchi River, and Funaura Bay (sites k, j, g, and m in Figure 7a, respectively). Grain size was estimated by sifting through screens of different mesh sizes (Figure 7b). Stones inhabited by upogebiids were composed of fine grain sizes of 60 µm on average. Stones with this grain size were classified as sandstone; more precisely, they were intermediate between fine-grained (125-62 µm) and ultrafine-grained (< 62 µm).

Discussion

Speciation and morphological relatedness

The time required for speciation (in terms of reproductive isolation) may be highly variable [37]. Three hundred species of cichlids in Lake Victoria may have evolved from one ancestor within the last 12,400 years [38]. At least 24 species of *Drosophila* may have originated on the ‘big island’ of Hawaii that formed less than 0.8 Ma. Among the snapping shrimps (*Alpheus*), geological evidence (i.e. the closure of the Panama seaway), combined with estimates of divergence time based on molecular data, suggests that sister species in the Caribbean and eastern Pacific achieved varying degrees of reproductive incompatibility around 3.0 Ma to 3.5 Ma [19].

The Ryukyu Arc (Figure 1) has been isolated from the Eurasian continent since 1.55 Ma [26]. Reproductive isolation between the Ryukyu Arc and the continent produced a number of endemic species in the Ryukyu Islands, including amphibians, reptiles [39], and mammals [40].
Based on morphology of the cephalothorax, *Upogebia neogenii* (Figure 4b-1) was very similar to *U. iriomotensis* (Figure 4a-1). *U. neogenii* was, however, characterized by an assemblage of long, brush-like setae borne on the chelae (Figure 4b-3). In addition, molecular clock data (Figure 5) suggested that *U. neogenii* had branched off from *U. iriomotensis* at 5.3 Ma, i.e. at the end of the Miocene period in the Neogene. *U. semicircula* (Figure 3b-1) may resemble *U. rupicola* in overall shape (Figure 4c-1), but molecular data (Figure 5) suggested that reproductive isolation had occurred between these two species as long ago as 5.3 Ma (late Miocene).

*U. rupicola* was described by Komai [18]. *U. rupicola* may also resemble *U. carinicauda*, which is widely distributed in the Indo-West Pacific [13]. In *U. rupicola*, intraspecific genetic divergence may have started 0.67 Ma, possibly in the Ryukyu Arc. If this is true, *U. rupicola* can certainly be regarded as a species different from *U. carinicauda*. *U. saigusai* was first described by Sakai et al. [15]. Morphologically, *U. saigusai* may be similar to *U. snelliusi* [17], but intraspecific genetic variation (Figure 5) did not support the possibility of both species dwelling on the same shores. The intraspecific divergence time (possibly in the Ryukyu Arc) was 0.61 Ma (Figure 5). In addition, *U. snelliusi* was collected from 5-6 m deep layers [13,17], showing a different habitat from that of *U. saigusai* (Figure 6a). Thus, *U. saigusai* should be regarded as a species different from *U. snelliusi*.

Figure 7: Distribution of stone upogebiids and grain size analysis. a) Distribution of five stone upogebiids in Iriomote Island. The figure illustrates data collected up to 2007. (*U. neogenii* had not yet been discriminated from *U. iriomotensis*.) See Table 3 for the number collected in each habitat. b) Grain size distribution of stones inhabited by upogebiids. Collection site: Urauchi River (site *g*), Mitara (site *h*), Funaura Bay (site *m*), and Hoshidate (site *i*) in Figure 7A.
Origin of the stones burrowed into by the upogebiids

Stones burrowed into by the upogebiids in Iriomote Island were composed of fine grain sizes (mean=60 µm) (Figure 7B). This grain size is just intermediate between fine-grained sandstone (125-62 µm: sandstone) and ultra-fine grained sandstone (<62 µm: siltstone) [2]. Mean grain size of the stones in Okinawa Island may be a little smaller than that of Iriomote Island (data not shown).

The uppermost layer of the Ryukyu Arc region (Figure 1) consists of paralic sediments with sandstone-rich facies, and contains some coal-measures [29,30]. Depositional facies analysis indicated four depositional cycles in this island, when the region was on the eastern margin of the Eurasian continent [41-43]. Calcareous nannofossil biostratigraphic data suggested that the Iriomote Formation was deposited at the lower shoreface to the inner tidal flat between 14.91 Ma and 13.53 Ma in the middle Miocene [25,44,45].

As shown in Figure 8, terrain characteristics are different between the western and eastern sides of Iriomote Island. While stratified sandstone facies are exposed on shores in the west, shores in the east are characterized by sedimentation of mud, sand, and coral rubbles. A number of stones are lying along the shores from the western to the northern parts, and these stones and the depositional layer are burrowed into by the stone upogebiids (Figures 6a, 6e, 6g and Figure 7A). Tidal flats composed of sedimentation of mud and coral rubbles were not used by the stone upogebiids. Thus, the distribution of stone upogebiids may be strongly influenced by difference in terrain characteristics between the western and eastern shores of the island.

Divergence time and settlement of stone upogebiids in the Ryukyu Arc

Global wind patterns and the Coriolis effect produce large, circular systems of surface. In the west Pacific Ocean, the Kuroshio current carries warm water from the equator to the north.

The Ryukyu Arc region (Figure 1) was originally on the eastern margin of the Eurasian continent during the Miocene (24-5 Ma). The Iriomote Formation was formed through depositional cycles between the tidal flat (shallow) and the inner shelf (deep) in the middle Miocene, and the uppermost interval was deposited from 14.91 Ma to 13.53 Ma [25]. Thus, a vast shallow sea (paleo-East China Sea) may have spread on the paleo-Ryukyu Arc region in the middle Miocene, and this region may have appeared above the sea later than 13.53 Ma. The warm, Kuroshio current may have originally flowed along the ‘east’ edge of this region. In the Quaternary, however, with the back-arc spreading of the Okinawa Trough [46], the paleo-Ryukyu Arc region was separated from the continental margin, and then started to move northeastwards. The arc transition from the continental to island changed the course of the Kuroshio current from the east to the west of the Ryukyu Arc, and it started flowing the Okinawa...
Trough about 1.55 Ma (figure 2 in Osozawa et al. [26]). The Ryukyu Arc reached 500 km east about 1.0 Ma, and the coral reefs developed around the islands because of the influence of the Kuroshio current [26].

The back-arc spreading of the Okinawa Trough converted the Ryukyu Arch from a passive to active margin (Figure 8A). The Ryukyu Arch region may now be subducting toward the northwest (Figure 8C). Subduction suggests that depositional strata suitable for burrowing by the stone upogebiids emerge only on the shores on the western side of the islands, and habitats would not be expected on the eastern shores (Figure 8B). The new shores formed along the Eurasian continent would have become the passive margin after 1.55 Ma. Deposition of sand and mud would not produce the habitat of stone upogebiids there either.

Bayesian inference (Figure 5) showed that stone upogebiids diverged 5-15 Ma (second half of the Miocene). Since the Ryukyu Arc had not yet been isolated from the continental margin in the Miocene (24-5 Ma) and Pliocene (5-1.8 Ma), we speculate that each species occurred in many islands near the equator 5-15 Ma. In contrast, intraspecific divergence time differed from 1.15 Ma (Pleistocene) to 0 Ma (present). With enlargement of the Okinawa Trough (back arc basin- rift valley), the Ryukyu Arc was separated from the continental margin, and the warm current (Kuroshio) isolated the Ryukyu Arc at 1.55 Ma (Pleistocene). The Ryukyu Arc has been moving towards the northwestern after that, and sedimentary strata for burrowing may have been exposed on the shorefront, i.e., western shores of the island.

Stone upogebiids would have settled in the Ryukyu Arc at different times (1.15-0 Ma) since 1.55 Ma. Intraspecific genetic variation of *U. spinidactylus* (Figures 6e and 6f) indicated 1.15 Ma (Figure 5), suggesting that *U. spinidactylus* settled in the Ryukyu Arc shortly after isolation of the Ryukyu Arc from the continent. The beginning of intraspecific genetic variation in the other upogebiids (*U. saigusai, U. rupicola, U. iriomotensis, and U. neogenii*) was between 0.61 Ma (*U. saigusai*) and 0.71Ma (*U. iriomotensis*). These upogebiids may have settled in the Ryukyu Arc during the second half of the Pleistocene, after which genetic variation in them appeared. No intraspecific genetic variation was shown in *U. semicirculara* (Figure 5). *U. semicirculara* may have settled in the Ryukyu Arc most recently among the six stone upogebiids examined.

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