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• Re: [biodiv] Editor Decision Yahoo/Sent ★

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To: Smujo Editors

Dear Smujo editors,  
I herewith send you the revised version of A-8563-Article Text-51369-1-4-20211026 (enclosed). I have made many corrections and necessary sentence restructure. I hope this has met your requirements. Thank you

Sincerely Yours,  
Silvester B. Pratasik

On Thursday, October 28, 2021, 06:04:52 PM GMT+8, Smujo Editors <smujo.id@gmail.com> wrote:

Silvester Benny Pratasik:

We have reached a decision regarding your submission to Biodiversitas Journal of Biological Diversity, "Deep Sea Squid in Sulawesi Sea, North Sulawesi Province, Indonesia".

Our decision is: Revisions Required

## Korespondensi

# Deep sea squid in Sulawesi Sea, North Sulawesi Province, Indonesia

Commented [JB1]: Make sure your title is specific. Is this about identifying the squid species?

**Abstract.** This study aimed to identify one of the deep sea squids caught in artisanal fisheries by traditional fishermen of Manado Bay, North Sulawesi, in Sulawesi Sea. Samples were collected using vertical hand line of traditional fishermen. The specimen was molecularly identified using Cytochrome c oxidase subunit 1 (CO1). For morphometric characteristics, all body parts were measured and each part was compared with mantle size. Results showed that the specimen was identified as purpleback flying squid *Sthenoteuthis oulaniensis* and had 99.35% similarity to *S. oulaniensis* (CO1) from China (NCBI - MN101944) with sufficient genetic diversity. Based on the body size, specimen recorded in the present study belongs to the dwarf form, the smallest form of *S. oulaniensis* groups. The species has the following morphometric characteristics: Head length is 0.28 of mantle length, tentacle length is 1.158 mantle length, and fin area is 0.5 mantle length. The fourth arm is the shortest and the second arm is the longest. *S. oulaniensis* recorded in the present study belonged to dwarf form living in equatorial waters. This finding has contributed to the list of fisheries marine resources, especially squid, in Indonesian waters.

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**Keywords:** species identification, CO1 gene, morphometric, traditional fishermen, hand line.

**Running title:** Deep sea squid identification

## INTRODUCTION

Squids is a cephalopod, mollusk, living in marine environment. This group is ecological opportunist adapted to exploit favorable environmental conditions (Rodhouse 2013). It is a favorite food type due to its highly nutritional content with high quality protein for human consumption and nearly all body parts are edible (Triharyuni and Puspasari 2012). Cephalopod production from fishing has continued to grow (Doubleday et al. 2016), with total commercial annual catches varying between 3.5 and 4.9 million tonnes in 2008–2017 (FAO 2019) and averagely supports about 15 and 20% of marine fishery landings and landed values, respectively (FAO 2019). Demand for this commodity has increased in both fresh and processed forms (Baskoro and Mustaruddin 2019). High market demand for squids as export commodity of Indonesia has made squids become one of major catches besides fish and lobster (Wulandari 2018). This trend will urge fishermen to conduct intensive fishing, while the squid production is still dependent upon the stock in the wild (Tresnati et al. 2012). The squids, in general, with the other coleoid cephalopods, are semelparous, have high reproductive rates and generally shortlived approximately one year with a single spawning event, then die (Anusha and Fleming 2014; Rodhouse et al. 2014). They also grow fast with high feeding rates and conversion efficiencies (Arkhipkin et al. 2015). These biological features make them be ecological opportunists in which they can quickly exploit favorable environmental conditions, but their abundance responds rapidly to poor conditions so that recruitment and abundance may be highly variable on annual time scales (Rodhouse et al 2014). Therefore, high dependence on the wild stock can lead to stock overfishing.

In Indonesia, squid production is far below the world market demand in spite of its increasing squid production, so that the export development is still wide opened (Triharyuni and Puspasari 2012; Hualalata et al. 2013). This condition is caused by uneven distribution of squid fisheries across the country. Only several regions do the squid fisheries, and the other do in very low scale for local consumption.

Information on this group is, so far, mostly related with commercial promotion as fisheries production. This group has been introduced together with other cephalopod groups as export commodity. Very few studies have been accomplished, so that people's knowledge on squids is very limited and they know the animal under a common

43 name "squid", but there are so many species belonging to this group. At fishermen community's level, squids have  
44 different names with their morphological characteristics. Nevertheless, several members of this group that usually  
45 occur in regular period of time have disappeared for longer time. It is evident that there is a major artisanal and  
46 small-scale inshore element to the world squid fishing fleet and that large volumes of loliginid squids caught in  
47 tropical and subtropical regions have high species diversity. Thus, species identification study on this group in  
48 Indonesian waters is needed. The taxonomy of the squid fauna generally is poorly understood. Correct species  
49 identification is basis of the ecological studies (Veijalainen 2011).

50 There are 290 species of squids and about 30–40 species have substantial commercial importance (Arkhipkin et  
51 al 2015) belonging to family Ommastrephidae, Loliginidae, Onychoteuthidae, and Gonatidae. Triharyuni and  
52 Puspasari (2012) have grouped family Loliginidae from Indonesian waters into several genera, *Afrololigo*,  
53 *Allotheuthis*, *Dorytheuthis*, *Heterololigo*, *Loliolus*, *Lollinguncula*, *Pickfordiateuthis*, *Sepioteuthis*, and *Urotheuthis*  
54 (Wulandari 2018), but there are many more species described (Jerep and Roper 2010). In Rembang waters, Central  
55 Java, four species were found, *Loligo chinensis*, *L. singhalensis*, *L. edulis*, and *L. duvaucelli*, in which *L. duvaucelli*  
56 is the most often caught species.

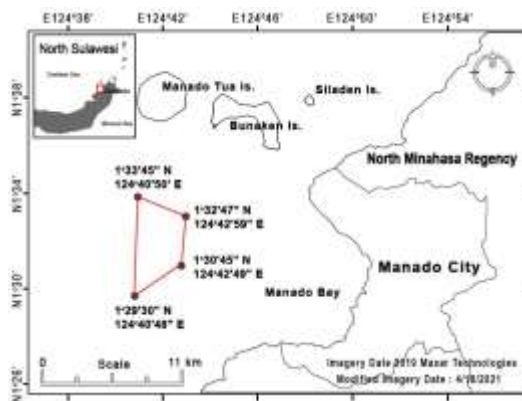
57 In North Sulawesi, Indonesia, there are no squid fisheries and the squid catches are bycatch of other fisheries, but  
58 squid production of different species is available every month. Squid catches are obtained at certain moon phase,  
59 especially during early new moon, when the large squid schools come near the surface. The local fishermen catch  
60 them using small mesh-sized seine, lift net or using scoop net. A small group of fishermen along Manado Bay,  
61 North Sulawesi, Indonesia, have benefitted deep sea squid as income source and food ingredients by relying on hand  
62 line fishing or jig fishing as practiced by foreign squid fishermen (Sundaram and Sawant 2014). Nevertheless, very  
63 few fishermen do squid fishing, the amount of catches is very low. This animal has become very important resource,  
64 so that its sustainability needs to be maintained, and thus, basic information on this resource needs to be provided.  
65 This study is intended to identify the deep sea squid caught by local fishermen of Manado Bay in Sulawesi waters.  
66 The findings may enrich the inventory of economically valuable squid species and develop squid fisheries in this  
67 area for future regional economic growth.

Commented [JB3]: obtained at during moon phases

68

## MATERIALS AND METHOD

69 Squid samples were obtained from fishermen's catch in Sulawesi Sea, North Sulawesi, about 5 miles from the  
70 shore of Manado Bay. In relation with this study, fishing operations were carried out at the geographic position  
71 covered by the area formed in the east border (1° 30' 45"N and 124° 42' 49"E to 1° 32' 47"N and 124° 42' 59"E) and  
72 the west one (1° 29' 30"N and 124° 40' 48"E to 1° 33' 45"N and 124° 40' 50"E) (Figure 1). Sample collection was  
73 done in May to June 2020.



74

75 **Figure 1.** Sampling location. Red lines point out the sampling area.

76 A total of 600 fresh squid samples were collected and measured in order to describe the species physical  
77 characteristics through comparison of mantle size and other morphometric characteristics, such as head length,  
78 tentacle length, arm length, fin length and fin width. For DNA identification, a piece of the squid arm was taken and

Commented [JB4]: As in from 600 individual squids? Were they all from the same species? Did you use any keys to identify species?

Commented [C5R4]: Yes, they are from the morphological characteristics

79 preserved in 95% ethanol solution before extraction to wash the sample from salt water and draw water from the  
80 cell. All samples were preserved in 95% ethanol and stored at room temperature before DNA extraction.

#### 81 **Extraction, PCR, and sequencing**

82 Genome DNA extraction of all samples used Innu PREP DNA Micro Kit (Analytic Jena). The CO1 gene was  
83 amplified applying universal primer pairs LCO1490: 5'-ggtcaacaaatcataaagatattgg-3' and HCO2198:  
84 5'taaacttcagggtagcaaaaaatca-3' (Folmer et al. 1994). Polymerase chain reaction (PCR) was carried out in 35 cycles  
85 at 95°C (30 sec.), 50°C (30 sec.), 72°C (50 sec.). The PCR product was visualized in 1% (b/v) agarose gel  
86 electrophoresis. Bi-directional sequencing was done by First Base CO (Malaysia) using Big Dye© terminator  
87 chemistry (PerkinElmer).

#### 88 **Data analysis**

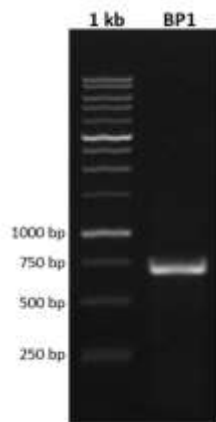
89 The specimen chromatogram was edited using Mega X v10.1 software (Kumar et al. 2018). The sequences were  
90 then compared with Gen Bank data using BLAST (BasicLocal Alignment Search Tools) method (Altschul et al.  
91 1997) and BOLD Systems (Ratnasingham and Hebert 2007). In this study, the sample sequence was compared with  
92 that of 7 specimens from South China Sea, one from Mid-Pacific, and one from Eastern Pacific. Besides, another  
93 specimen of the same genus *S. pteropus* from Spain was also used as an outgroup in order to strengthen the  
94 comparison.. The phylogenetic tree was built using Neighbor-Joining Method of MEGA X and the heterogeneity  
95 index was also estimated (Kumar et al 2018).

96

#### 97 **RESULTS AND DISCUSSION**

##### 98 **DNA characteristics**

99 Based on molecular identification using the Internal Transcribed Spacer (ITS) region, the DNA bands obtained  
100 were around 500-750bp, the success of PCR was detected by the presence of a single DNA band around 680 bp, the  
101 PCR results can be seen in Figure 2.



102

103 **Figure 1.** PCR product of the sample specimen

104 The sample was identified as purpleback flying squid *Sthenoteuthis oualaniensis* as shown in the BLAST results in  
105 the Gen Bank of the National Center for Biotechnology Information (NCBI). This species, according to WoRMs  
106 details, has synonymized names as *Loligo brevitentaculata* Quoy & Gaimard, 1832 (synonym), *L. oualaniensis*  
107 Lesson, 1830 (original combination), *L. vanikoriensis* Quoy & Gaimard, 1832 (synonym), and *Symplectoteuthis*  
108 *oualaniensis* (Lesson, 1830). This species belongs to class Cephalopod, subclass Coleoides, suborder  
109 Decapodiformes, order Oegopsida, superfamily Ommastrephidea, family Ommastrephidae, subfamily  
110 Ommastrephinae, and genus *Sthenoteuthis* (Jerep and Roper 2010). This finding is the first report on this species  
111 from Sulawesi Sea. To confirm the species status, the sample sequence was then compared to 9 *Sthenoteuthis*

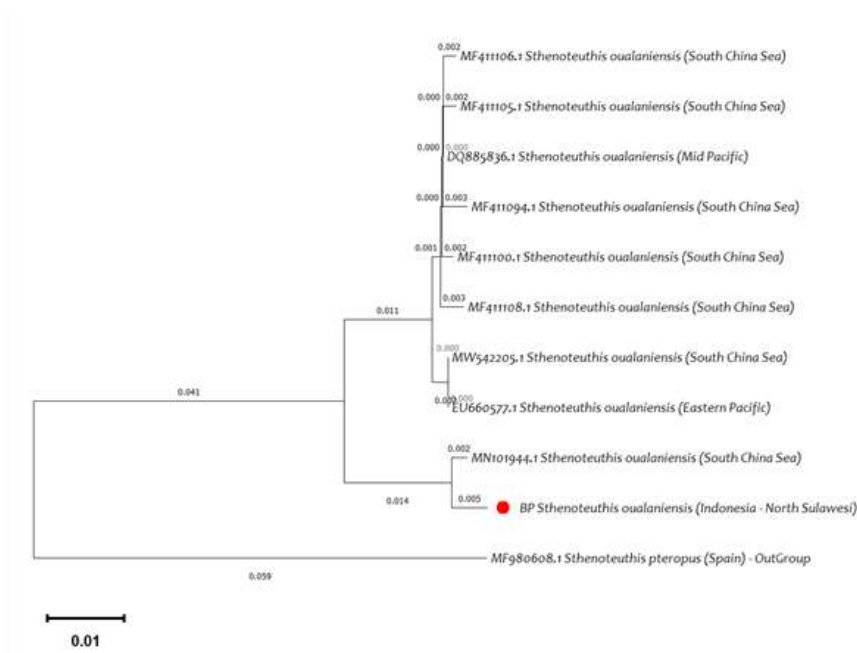
112 *oualaniensis* specimens from South China Sea, Eastern Pacific, and Central Pacific. The present study shows that *S.*  
 113 *oualaniensis* specimen from Sulawesi Sea has similarity rate of 99.3% to NCBI record (acc. no. MN101944.1) from  
 114 South China and 96.4-96.8% to 8 other NCBI records from South China Sea, Eastern Pacific, and Mid Pacific  
 115 (Table 1). Specimen of the same genus *Sthenoteuthis pteroptus* from Spain used as outgroup also clearly indicates  
 116 great difference from the specimen and supports the kinship status of the specimen on study.

117 **Table 1.** Similarity rate of *S. oualaniensis* from the Sulawesi Sea and those from the Gen Bank DNA sequence database.

118

No	Sequence	1	2	3	4	5	6	7	8	9	10
1	MN101944.1_ <i>S. oualaniensis</i> (South China Sea)										
2	MW542205.1_ <i>S. oualaniensis</i> (South China Sea)	0.029									
3	EU660577.1_ <i>S. oualaniensis</i> (Eastern Pacific)	0.029	0.000								
4	MF411106.1_ <i>S. oualaniensis</i> (South China Sea)	0.031	0.005	0.005							
5	MF411105.1_ <i>S. oualaniensis</i> (South China Sea)	0.031	0.005	0.005	0.003						
6	MF411100.1_ <i>S. oualaniensis</i> (South China Sea)	0.031	0.005	0.005	0.003	0.003					
7	MF411094.1_ <i>S. oualaniensis</i> (South China Sea)	0.031	0.007	0.007	0.005	0.005	0.005				
8	MF411108.1_ <i>S. oualaniensis</i> (South China Sea)	0.032	0.007	0.007	0.005	0.005	0.005	0.007			
9	DQ885836.1_ <i>S. oualaniensis</i> (Mid Pacific)	0.029	0.003	0.003	0.002	0.002	0.002	0.003	0.003		
10	BP_ <i>S. oualaniensis</i> (Indonesia- North Sulawesi)	0.007	0.032	0.032	0.034	0.034	0.034	0.034	0.036	0.032	
11	MF980608.1_ <i>Sthenoteuthis pteroptus</i> (Spain)	0.119	0.114	0.114	0.116	0.116	0.112	0.116	0.114	0.114	0.116

119 The difference is demonstrated in specimen grouping (Figure 3) as well in which there are 4 different groups formed, and  
 120 only one compared *S. oulaniensis* of six NCBI records is in the same group as *S. oulaniensis* specimen on study. It reflects  
 121 that this species has sufficient genetic diversity. Group 1 comprises 5 specimens from South China Sea, one from Mid  
 122 Pacific, and one from Eastern Pacific. Group 2 comprises specimen from South China Sea and Eastern Pacific, while  
 123 group 3 comprised the specimen from Sulawesi Sea and 1 NCBI of South China Sea specimen indicating that the species  
 124 on study has the closest similarity to one of the South China Sea specimen. Group 4 is an outgroup that is taken from  
 125 different species of Spain specimen, *Sthenoteuthis pteropus*. This grouping reflects the kinship of compared *S.*  
 126 *oulaniensis* specimens from different areas.



127 **Figure 3.** Phylogenetic tree of *S. oulaniensis*  
 128

129 This evidence is supported by the closest kinship to the deep sea squid *S. oulaniensis* specimens from South China waters  
 130 of NCBI record and they could originate from the same ancestor. Besides, this comparison also indicates that there are  
 131 intraspecific genetic variations in *S. oulaniensis* (Table 1 and Figure 3). Genetic diversity could result from population  
 132 size, in which the larger population size within species is, the higher genetic diversity will be. This condition is also  
 133 supported by Hague & Routmant (2016). It could also be caused by different environmental conditions with localities that  
 134 restrict the organism distribution due to its tolerance limit to the environments.

135 **Morphological description**

136 *Sthenoteuthis oulaniensis* has the following morphometric characteristics: Head length is 0.28 of mantle length (ML), tentacle length is  
 137 1.158 ML, fin width is 0.3 ML, and fin length is 0.4 ML. The fourth arm is the shortest and the second arm is the longest among the  
 138 squid arms (Figure 4). These morphometric characteristics could become typical characteristics in *S. oulaniensis* from  
 139 Sulawesi Sea. Hence, the present study is in agreement with Staaf et al (2010) and Xua et al (2020) who found the  
 140 existence of a morphologically distinct form that is highly genetically divergent in *S. oulaniensis* that separates the dwarf  
 141 and medium-sized species and exhibits a distinct biogeographic break at equatorial waters of Eastern Pacific Oceans.  
 142

**Commented [JB6]:** Was this based on measurement of one individual, or several? If several, please state that these lengths are based on averages. If from on individual, please state this here.

**Commented [C7R6]:** This description is based on 600 individual measurements



Figure 4. Tentacular club (left) and Squid *Sthenoteuthis oulaniensis*.

*S. oulaniensis* is known as purpleback flying squid, a tropical Indo-Pacific species that occurs in the Pacific from southern Japan to southern Queensland and from just south of Baja California to northern Chile. *S. oulaniensis* belongs to family Ommastrephidae widely distributed in the tropical and subtropical areas of the Pacific Ocean and Indian Ocean, and this species is important for resource exploitation in South China Sea as one of the major target species of large-scale light falling-net fishing (Yu et al 2019; Zhao et al 2021). This species is known to do diurnal vertical migration between the surface at night to the deeper layer during the day, and thus, the species is caught by local fishermen of Manado Bay using the flashlight artificial bait near the surface water. Fishing operations used a traditional outboard-motored outrigger boat and fishing activities relied on hand-line facilitated with flashlight artificial bait that was lowered down to 20-25 m depth to get the squid to bite.

Cephalopods are sensitive to water temperature (Li et al 2020) that becomes one of the major environmental factors affecting squid activities, including aggregation, breeding, and emigration (Klemas 2013). Deep sea squids *Sthenoteuthis oulaniensis* are distributed from 0-4,500 M depth with a peak at 0-500 M depth, water temperature of 25 – 32°C, salinity of 34-35 PSU (Jerap and Roper 2010). These seawater temperature ranges cover that of Sulawesi Sea that enables to support the occurrence of this squid species based on 10 years of Sulawesi Sea water temperature data (<https://www.seatemperature.org/asia/indonesia/north-sulawesi>). Besides sea surface temperature (SST), sea surface height anomaly (SSHA) at -0.05–0.05 m and chlorophyll-a concentration higher than 0.18 µg/L are required to gain higher catch (Zhao et al 2021).

*S. oulaniensis* are sexually dimorphic and the females tend to grow larger than the males in most cases (Chembian and Mathew 2014). Purpleback squid *S. oulaniensis* has a short lifecycle, a rapid growth rate, and high fecundity (Zhang et al 2013). According to Roper et al (2010), this species is highly active predator with major prey groups of fish, cephalopods, crustaceans, and others. This species is known to be one of the fastest-growing squid species with daily length increase of about 1.0 mm in the dwarf and middle-sized forms and 3.8 mm in the giant form is about 3.8 mm. This species can move at high speeds through the water, easily manoeuver and quickly respond to their environment changes. Under distress of external factors, such as predation, this species can reach a high speed and glide above the surface over ten meters. They also found that the cruising speed of an adult squid is between 3 to 10 km per hour, their burst speed achieves greater speeds of up 35 km per hour, and it can be important for sudden changes in movement and escape behavior. They can occur into shoals from 2 individuals to 800 individuals, and when geographical distributions overlap they can exist with

173 other members of similar-sized family Ommastrephidae schools, *Dosidicus gigas* and *Ommastrephes bartramii* that are  
174 distributed from the Indo-Pacific to Indian Ocean as well (Liu et al 2016). Ommastrephidae squids are known as voracious  
175 and adaptable predators of a broad range of prey including small crustaceans and fishes at early life stages and shift to  
176 micronekton, larger fishes, and cephalopods (including cannibalism) (Gong et al. 2020). These species are highly  
177 migratory, undertake diel vertical migrations of several hundred meters and seasonal migrations between the shelf and  
178 open ocean (Stewart et al 2013). Thus, they can work as important linkages between both neritic and oceanic food webs  
179 (Arkhipkin 2013; Alegre et al 2014).

180 In North Sulawesi, this squid species fishing is done by artisanal fishermen using jigging with artificial flashing bait,  
181 and the fishing operation has been done in Sulawesi Sea. In the present study, the squids *S. oulaniensis* caught in  
182 Sulawesi Sea had a size range of 68 mm to 119 mm ML, with mean mantle length of 89.833±0.9000 mm, reflecting that  
183 the squid catches are dominated by large individuals. According to the classification of Nesis (1993), this size range  
184 belong to dwarf form with body size of 90-100 mm ML for mature males and 90-120 mm ML in mature females, and  
185 occurs in the equatorial waters, and lacks of dorsal mantle photophore patch. The occurrence of *S. oulaniensis* in Sulawesi  
186 waters could result from the influence of environmental factors. Similar finding is also reported that changes in marine  
187 environments affect cephalopod fishery resources at different spatial and temporal scales (Zhang et al 2012). Furthermore,  
188 the distribution of purpleback flying squid *S. oulaniensis* in this area could not be separated from the impact of the  
189 Kuroshio from the Pacific Northern Equatorial current toward the east coast of Luzon, Philippines that forms the  
190 southward-flowing Mindanao Current (Qiu and Lukas 1996) and transfers the upper ocean waters from the Pacific to the  
191 Indian Ocean through the Indonesian Seas (Taufiqurrahman et al 2020). As the water mass enters the Indonesian Seas, the  
192 warm and relatively salty characteristics of the Pacific water tend to disappear due to strong vertical tidal mixing. The  
193 mixing is believed to affect the carbon, oxygen, and nutrient (nitrate, phosphates) concentrations in the eastern Indonesian  
194 Seas and subsequently in the Indian Ocean (Ayers et al. 2014). This event makes Indonesian marine waters fertile enough  
195 to support the high occurrence of marine animals in the area. The Indonesian Throughflow is the only ocean connector  
196 pathway in the equator (Sprintall et al 2014), and it has an important purpose in the transport of mass and heat from the  
197 Pacific into the Indian Ocean (Feng et al 2018). The throughflow brings the eggs and larvae, along with the rich detritus of  
198 the sea that is swept up from the offshore deep basins. It is in agreement with Dell et al (2011) that the biophysical  
199 environment plays an essential role in controlling the distribution and abundance of pelagic predators in the ocean. Cheng  
200 et al (2018) who studied the swordtip squid *Uroteuthis edulis* found that complex oceanographic conditions might affect  
201 their population in the Southern East China Sea due to seasonal changes in the Kuroshio Current and Mainland China  
202 Coastal Current during the northeasterly monsoon and southwesterly monsoon seasons.

203 This evidence is supported by the closest kinship to the deep sea squid *S. oulaniensis* specimens from South China  
204 waters of NCBI record and they could originate from the same ancestor despite in genetic diversity. Genetic diversity  
205 could result from population size, in which the larger population size within species is, the higher genetic diversity will be.  
206 This finding has reconfirmed the distribution of *S. oulaniensis* reported in previous studies. The present study has provided  
207 information on the occurrence of one of the importantly economic squid resources in Indonesian waters, especially the  
208 Sulawesi Sea, that can be used for national economic development. As a center of biodiversity, we strongly believe that  
209 there are more squid species living in these waters that need to be described, so that there are more taxonomic works need  
210 to be done on species richness of squids in Sulawesi Sea waters as well for future development. Besides, other biological  
211 studies of the species need to be done as well in order to have better understanding on this species that future management  
212 and conservation efforts could be well prepared.

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215 facilities. Our high appreciation is also addressed to Ms. Isroja Paransa who has involved in this study, but passed away in  
216 the middle of study.

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