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SOME BIOLOGY ASPECTS OF OXEYE SCAD, SELAR BOOPS CAUGHT FROM BITUNG WATERS WITHIN MOLLUCAS SEA OF INDONESIA

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ABSTRACT

Trevally species of oxeye scad, *Selar boops* is one of small pelagic species which an economic values. The species was exploited throughout the year by small-scale fishermen in Bitung waters using handline. The aim of this study was to determine some biological aspects of *S. boops* in Bitung waters area through measuring their lengths and weights, sex ratios, stages of oocyte, gonadosomatic index and type of spawning. Samples of fish were randomly collected each month at Bitung Oceanic Fishing Port during February 2016 to January 2017. Total samples of fish were used in this study 1659 consisting 841 male, 754 female, and unidentified 64. Length at first maturity (L_m) was 16,3 cm (FL) and asymptotic length (L_∞) was 26,4 cm (FL). The overall length-weight relationships were $W=0,01136 * L^{3,1640}$ ($R^2=0,988$) with a positive allometric growth. The sex ratios were balanced between males and females. The average of gonadosomatic index was 1.02 while the lowest and highest values was 0.03 and 6.96 with partial spawning type. Identification and genetic confirmation of *S. boops* using DNA-COI analysis.

KEY WORDS

Biology aspects, DNA-COI, Selar boops, marine species.

Carangidae is estimated to have 25 total of genera and about 140 species in the world (Randal et al 1990). There are 3 genus of travellies in the world that has been declared valid namely *Selar crumenophthalmus*, *Selar boops* and *Selaroides leptolepis*. Trevally is a genus of the Carangidae which classified as small pelagic fish species that live in groups (Carpenter & Niem 2001) and migratory surrounding coastal waters at depths of 30-100 m (Froese & Pauly 2014) and distributed in the Indo-Pacific waters (Smith-Vaniz 1995). *S. boops* is species co-occurs with *Selar crumenophthalmus* on continental shelf waters where common between 20 and 100 m and it is most often found over soft bottom or seagrass bed areas (Gomelyuk 2009; Smith-Vaniz & Williams 2016), but it is also known to sometimes frequent coral and rocky reefs (Paxton et al 1989). This species forms large diurnal aggregations, dispersing at night to feed on planktonic and benthic invertebrates (crabs and shrimps) and small fishes (Smith-Vaniz 1986, Paxton et al 1989). The maximum size for this species is about 26 cm TL (Allen & Erdmann 2012).

Oxeye scad, *S. boops* locally name known as Tude Batu. This spesies is one of small pelagic resources in Bitung waters area and is one of the commercially important fishes in Indonesia. Yet, biological aspects data concerning the *S. boops* are lacking in Indonesian waters even in other countries. The research information about *S. boops* has been done by Hutubessy (2011), Isa et al (1996), and Gumanao et al (2016). There are virtually no data concerning the stages of maturity, spawning or other life history traits for this species and according the International Union for Conservation of Nature and Natural Resources (IUCN), this spesies was categorized red list of threatened species (Smith-Vaniz & Williams 2016). The purpose of the present study was to determine some aspects of the biology of *S. boops* in Bitung waters area, within Mallucas Sea of Indonesia. The results of this study as initial information related to some biologi aspects of *S. boops* and will provide valuable information to all stakeholders.

MATERIALS AND METHODS OF RESEARCH

Study site and sampling procedure. A total of 1659 oxeye scads (841 males, 754 females and 65 un-identification individuals) was randomly sampled from the commercial catch in the local fish in Bitung city, from February 2016 to January 2017. The samples were caught from Bitung waters area within Mollucas Sea of Indonesia. The samples were fresh captured fish. Measurements and dissections were quickly performed to prevent measurement bias and sample decay.

Research location. This research was conducted in Bitung Oceanic Fishing Port (BOFP), Bitung City, North Sulawesi Province, Indonesia. Geographically of BOFP located at 01°26'42"N - 125°12'24"E (Figure 1).

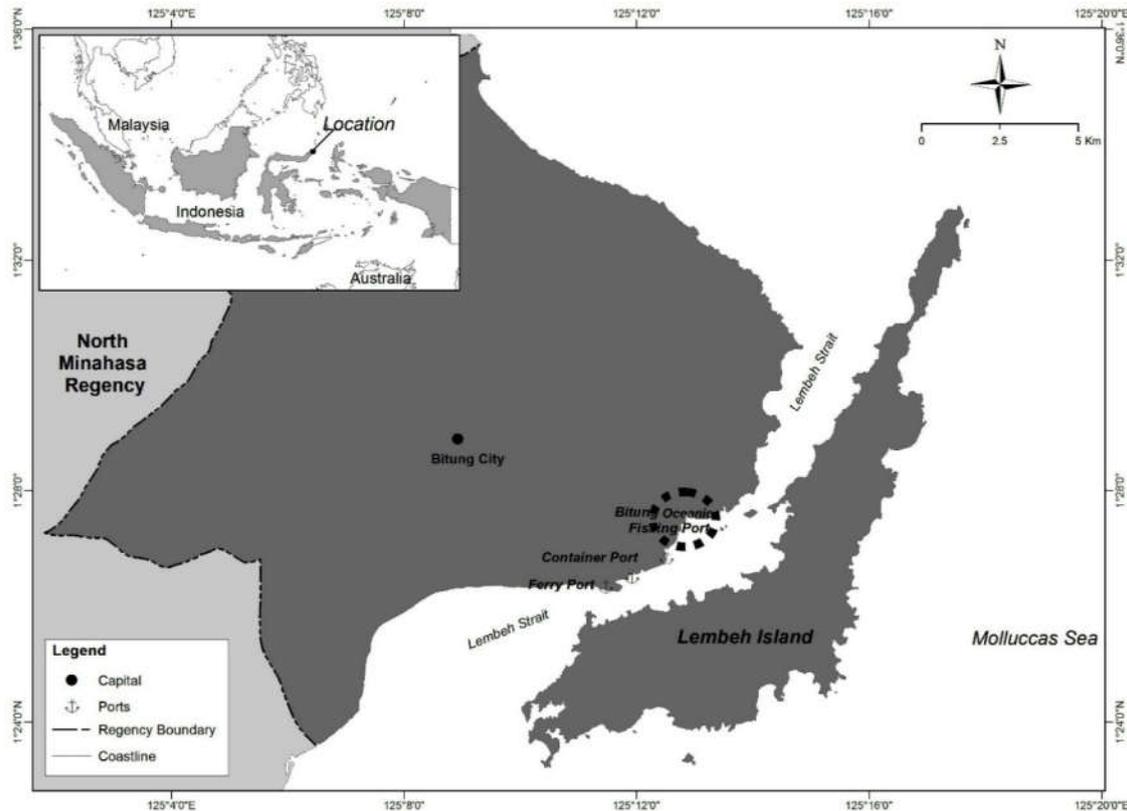


Figure 1 – Location sampling of Bitung Oceanic Fishing Port (01°26'42"N, 125°12'24"E)

Measurements. Each individual fish was measured for the fork length (FL) to the nearest 0.1 cm and body weight was measured using portable and battery power weighing scale of max. 500 g (0.1 g). The fish were later dissected from the abdominal region and their sexes were determined by visual examination of the gonads. The gonads were then removed carefully and their weights taken to the nearest 0.01 g with the portable electronic compact scale.

Species identification. The identification of *S. boops* was confirmed by DNA molecular analysis. DNA genome was collected from dorso-lateral tissue of the fish, preserved in acetone, and stored at -50°C prior to laboratory procedure. DNA extraction (Asahida et al 1996) was performed in 1.5 ml volume containing 600 µl TNESU 8 buffer extract (TNESU-Urea: 8 M urea; 10 mM Tris-HCl, pH 7.5; 125 mM NaCl; 10 mM EDTA; 1% SDS), and 300 µl Cl. The mix solution was incubated at 65°C for 2 h. The DNA was extracted with Phenol-Chloroform (1:1), 2 ethanol, 0.3 M NaCl, and TE Buffer (10 mM Tris-HCl, pH 8.0; 1 mM EDTA). Partial DNA region COI was amplified using PCR with primer (Ward et al 2005):

- FF2D: 5'-TTCTCCACCAACCACAARGAYATYGG-3'
- FR1D: 5'-CACCTCAGGGTGTCCGAARAAYCARAA-3'

PCR was performed at 10 µl volume containing dd H₂O 5.65 µl, 10X Fast Buffer 1 µl, dNTP mix 1 µl, each primer of 0.5 µl, SpeedSTAR *taq* polymerase 0,05 µl, and DNA template 0,5 µl based on protocol in Takara Inc., adjusted for Taq Enzyme SpeedSTAR HS DNA polymerase. PCR were carried out over 35 cycles with program setting: denaturation at 95°C for 30s, annealing at 55°C for 30s, and extension at 72°C for 30s. PCR product, after visualized in 1% agarose gel, was purified following Kit protocol of GE ExoSAP-IT. Sequencing was done by Firstbase Malaysia. The sequence was aligned (reverse complement, pairwise alignment, and consensus) using BioEdit (Hall 1999). Phylogenetic reconstruction of sequences DNA region COI were based on maximum-likelihood method with MEGA6 (Tamura et al 2013), bootstrap method with 1000 replicates and all parameters were set at default (Ikejima et al 2004).

Sex ratio. The sex of each specimen was identified by physical examination of the gonads. The proportion of the two sexes relative to one another was used to calculate the sex ratio. Based on gonad identification, Chi-square (χ^2) test was carried out on the observed male and female specimens to show the level or proportion of differentiation from the expected 1:1 ratio, if $\chi^2_{\text{test}} > \chi^2_{\text{table}}$, H_0 is accepted (Ayo-Olalusu 2014).

Oocyte development and maturity classification. To examine oocyte development and microscopically verify the precision of the macroscopic maturity scale, a subsample of 19 ovaries were fixed in 10% buffered formalin for histological analyses. Once fixed, a 4-mm thick transverse section was taken from the middle of each gonad, dehydrated through a series of alcohol and solvent solutions and infiltrated with paraffin wax on an automatic tissue processor (Sakura, Japan). A rotary microtome (HM 310, Thermo Fischer Scientific Inc., Germany) was used to cut 5-µm thick sections, which were stained with haematoxylin and eosin, cover-slipped with a mounting medium, and examined under a Nikon SMZ1000 light microscope with a Nikon DXM1200F digital camera.

Gonad stages identification. Macroscopic identification of female gonads was done based on five point maturity scales for partial spawners (Holden & Raitt 1974). The gonad developmental stages are categorised as immature (ovaries about 1/3 length of the body cavity), maturing (ovary about ½ length of the body cavity and ovary are pinkish without visible ova to the naked eyes), ripening (ovary takes about 2/3 length of the body cavity and ovary with granular appearance), ripe (ovary from 2/3 to the full length of the body cavity; ovary with conspicuous superficial blood vessels) and last stage as spent (ovary shrunken to about half-length of the body cavity and loose walls). Fresh gonads were usually removed from the fish within a few hours of capture, and their sex and stage of reproductive maturity determined using a macroscopic staging system. Gonads obtained from coastal fishers could usually be weighed fresh (0.01 g). Two or three transverse cuts were then made through each gonad to ensure proper fixation before placing them into a perforated cellophane bag and then into plastic drums containing 10% formalin in seawater.

Gonadosomatic index. The gonadosomatic index was calculated as a percentage of body mass. It is represented by the formula: $GSI = [\text{Gonad Weight} / \text{Total Tissue Weight (weight of fish)}] \times 100$. This was calculated for each individual and a monthly average for each sex was established. The GSI calculation were pooled based on the sex of fish regardless of the gonad maturity stage.

Isometric growth dimension. Isometric growth dimension was estimated using length-weight relationship (LWR). Cube law of length-weight relationship was transferred into log-liner following the quation (Froese 2006):

$$W = a * L^b$$

Where: 'W' stands for weight (g), 'L' stands for fork length (cm), 'a' is constant, and 'b' is isometric growth dimension. The constant 'a' and 'b' were derived by the method of linier least squares. To test 'b' values against the value of '3', Student's t-test was employed to predict any significant deviation (Snedecor & Cochran 1967).

The t-statistic was calculated as follows:

$$t = (b-3)/SEb$$

Where: SEb = standard error of 'b' = $SEb = \sqrt{(1/(n)) \times [(Sy/Sx)^2 - b^2]}$, Sx and Sy are the standard deviations of x and y, respectively. The t-value was compared with t-table value (n =1659) for degrees of freedom at 5% significance level.

Length at first capture (Lc) and Length at first maturity (Lm). We using spawning potential ratio (SPR) analysis provides an overview of the length distribution of fish, Lc and Lm of fish and the condition of *S. boops* in Bitung waters. In species where the males and females have distinct and different growth characteristics, the SPR Analysis focuses on the size structure and size of maturity of the females (Prince 2014). SPR is the proportion of spawning fish that are not caught in fishing policy (Walters & Martell 2004). SPR analysis was carried out by using assessment which is web-based in www.barefootecologist.com.

RESULTS AND DISCUSSION

Species identification. Only one sequence *S. boops* was used in phylogenetic analysis caught from Bitung waters area. Alignment of this sequence (690 bp) and based on estimates of evolutionary divergence identification resulted in 99% similarity to *S. boops* voucher ARO37 and *S. boops* voucher DBMF-M12 sequences from genBank (Table 1). The visualization of the morphology of *S. boops* from Bitung waters and phylogenetic are given in Figure 2 and 3.

Table 1 – Estimates of evolutionary divergence between sequence *S. boops* from Bitung with sequences from genBank (<https://blast.ncbi.nlm.nih.gov>)

Local name sequence	Description	Total score	Query cover	Query length	E value	Ident.	Accession
Tude Batu	<i>S. boops</i> voucher ARO 37	1188	95%	690	0.0	99%	KF009659.1
	<i>S. boops</i> voucher DBMF-M12	1171	94%	690	0.0	99%	HQ560953.1



Figure 2 – *S. boops* collected from Bitung waters area that used for species identification

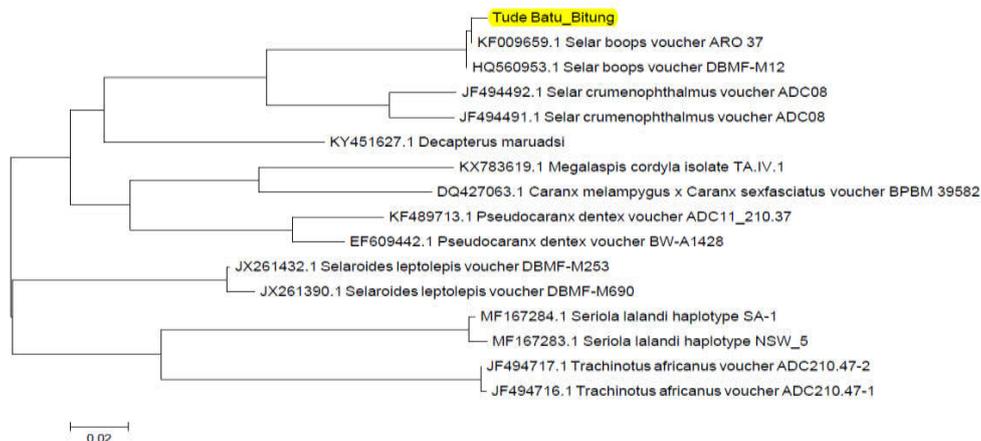


Figure 3 – Phylogenetic reconstruction (maximum likelihood-joining method) of sequences mt DNA region COI of sekuen sample from Bitung (Tude Batu_Bitung)

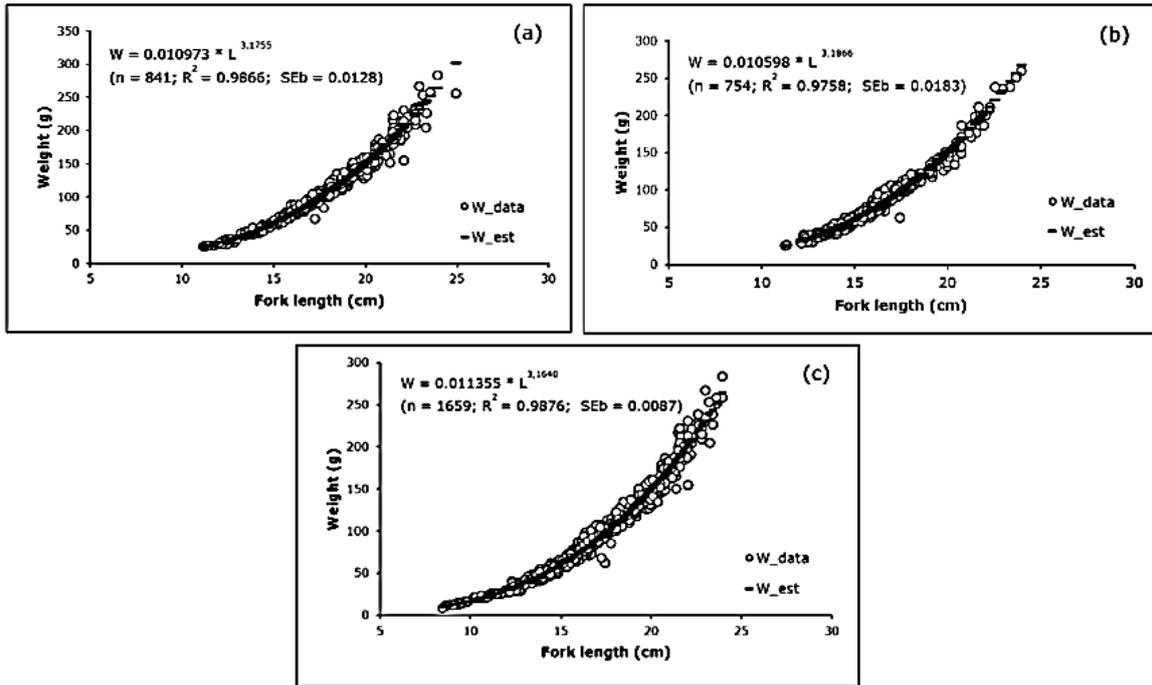


Figure 3 – Isometric growth dimension derived from length-weight equation of *S. boops*: (a) male; (b) female; (c) overall

Isometric growth dimension. A total of 841 and 754 randomly selected specimens of males and females were collected, with fork length and weight ranging from 11.2–25.0 cm and 25.0–282.0 g for males and 11.3–24.0 cm and 25.0–258.5 g for females. The scatter diagram showing the allometric relationship for male and female is given in Figure 4a and 4b. The LWR for male and female were established as $W = 0.010973 * L^{3.1755}$ ($R^2 = 0.9866$; $SEb = 0.0128$) and $W = 0.010598 * L^{3.1886}$ ($R^2 = 0.9758$; $SEb = 0.0183$), respectively. An overall LWR was established for all samples as $W = 0.011355 * L^{3.1640}$ ($R^2 = 0.9876$; $SEb = 0.0087$) (Figure 4c). The student's t-test of slope with $b = 3$ revealed the existence of positive allometric growth for both sexes (T-test > T-table) with confidence interval 'b' was $3.1504 < b < 3.2006$ for male and $3.1526 < b < 3.2254$ for female (Table 2).

Table 2 – Values for t-test for the slopes of regression equations with theoretical $b = 3$ of *S. boops* from landings at BOFP during the study

Sex	b (slope)	Interval of b	SEb	T-test	T-table
Male	3.1878	3.1504-3.2006	0.0128	13.74	1.96
Female	3.1886	3.1526-3.2245	0.0183	10.29	1.96
Overall	3.1640	3.1470-3.1810	0.0087	18.95	1.96

Sex ratio. Sex for *S. boops* with $N = 1659$ showed males dominated by 50.69% ($n = 841$) while females only 45.450% ($n = 754$) and not identified 3.86% ($n=64$). Chi square (χ^2) test revealed that the sex ratio was not significantly different deviate from the normal proportion ratio 1:1 (Tables 3) and number of male and female samples during the study is presented in Figure 4.

Table 3 – Chi square test for sex ratios of *S. boops* collected from BOFP during the study

Sex	Observation (O)	Expected (E)	O-E	(O-E) ²	(O-E) ² /E
Male	841	797.5	43.5	1,892.25	2.373
Female	754	797.5	-43.5	1,892.25	2.373
Total	1595				4.756

The Chi-square test was used to determine if a population contains equal proportion of males and females and it is a test of how well a model fits the observed data. Hypotesis: Ho = Male: Female is 1:1, Hi = Male: Female is not 1: 1. If n = 2, degree of freedom (df) = 2–1 = 1. From the critical values of the Chi-square distribution table at df = 1, $\chi^2_{0.05,1} = 3.841$. The calculated value 4.746 is more than table value, the null hypotesis Ho is accepted meaning proportion stable.

Table 4 – The estimated sex ratio values for *S. boops* collected from BOFP during this study

Month of observation	Male	Female	Un-identified	Proportion	
				Male	Female
February 2016	29	71	19	0.41	1
March 2016	82	42	40	1	0.51
April 2016	106	51	-	1	0.48
May 2016	76	80	-	0.95	1
June 2016	66	52	-	1	0.79
July 2016	71	61	-	1	0.86
August 2016	58	63	-	0.92	1
September 2016	98	78	-	1	0.80
October 2016	49	67	-	0.73	1
November 2016	53	72	1	0.74	1
December 2016	59	68	-	0.87	1
January 2017	94	49	4	1	0.52
Total	841	754	64		

Macroscopic identification at the gonad maturity stage: Macroscopic identification of the gonads was carried out for six months, from February 2016 to July 2016, to obtain an overview of gonadal development. The ovaries were orange in color, thin and ova were not visible to the naked eye during gonad stage one. Color, size and ova visibility gradually increased as the ovaries matured. Ovaries of gonad stage four appeared bright orange in color with conspicuous superficial blood vessels and ripe ova were visible to the naked eye and occupied almost the entire body cavity (Figure 5). The ranges of gonad weight in this study each stages was obtained that stage I total samples 324 range 0.02 – 0.19 g (mean value = 0.09 ± 0.04) and the maximum length was 16.9 cm, stage II total samples 128 ranges of gonad weight 0.09 – 0.43 g (mean value = 0.18 ± 0.07) and the maximum length was 17.4 cm, stage III total samples 167 ranges 0.18 – 1.99 g (mean value = 0.79 ± 0.46) and the maximum length was 20.1 cm, stage IV total samples 67 ranges 1.09 – 3.97 g (mean value = 2.64 ± 0.79) and the maximum length was 21.6 cm, and stage V total samples 68 ranges 3.00 – 15.81 g (means value = 6.30 ± 2.91) and the maximum length was 24.0 cm. Total sample of immature were 452 and matures were 302.

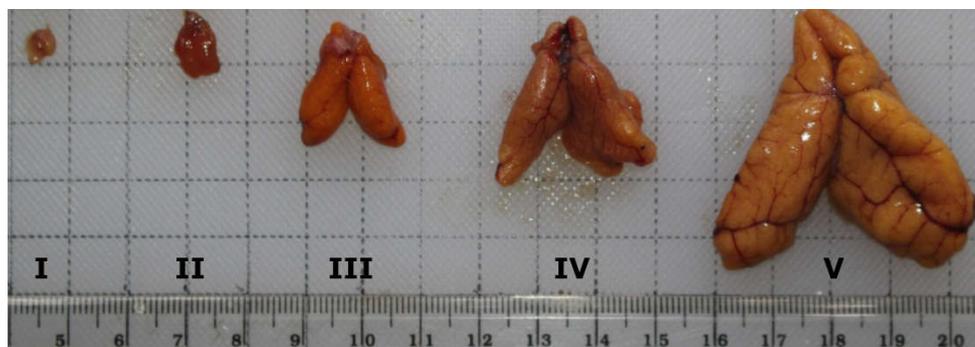
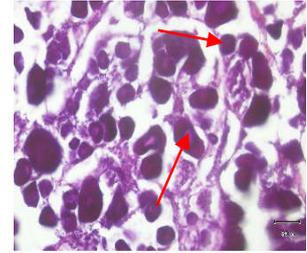


Figure 5 – Ovary size variation in *S. boops*: from right to the left, as the ovaries matured their size and development increased

Gonad stages identification. According to microscopic identification we obtained 5 stages variation size of oocytes development and this result explain that *S. boops* catch in Bitung waters is partial spawning type (Figure 6).

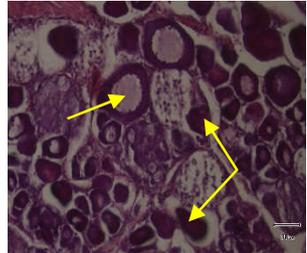
Stage I

Only oogonia (og) and primary growth oocytes (po) with large pink nuclei are present. Ovary wall is thin. Atresia may occur in the present and all of the following classes. Oocyte diameter ranges 10.56–37.84 μm (average 20.77 μm). Scale bar (Sb): 100 μm .



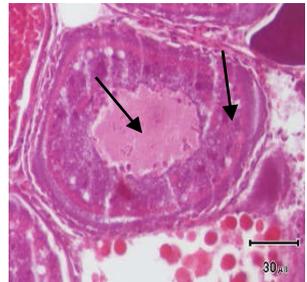
Stage II

Secondary growth oocyte, primary growth oocytes and oogonia are present. Ovary wall is thin in first time spawner, thick in repeat spawners. The ratio nucleus (nu) to oocyte area has decreased. Oocyte diameter ranges 25.52–58.08 μm (average 34.06 μm). Sb: 100 μm .



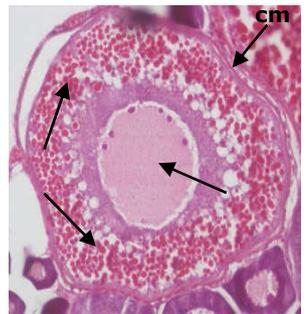
Stage III

Secondary growth oocytes containing yolk granules (yg), primary growth oocytes and oogonia are present. The oocyte diameter increases dramatically during III. By the end of class III the cytoplasm is half filled with yolk granules. Oocyte diameter ranges 27.28–132.90 μm (average 76.12 μm). Sb: 200 μm .



Stage IV

Secondary growth oocytes in the nuclear migration stage, primary growth oocytes and oogonia are present. By end of class IV oocytes are completely filled with yolk, cortical alveoli are pressed against cell membrane (cm) and nucleus begins migration towards micropyle. Oocyte diameter ranges 75.68–192.30 μm (average 117.66 μm). Sb: 200 μm .



Stage V

Final growth oocytes with hydrolyzed yolk granules (hyg), primary growth oocytes and oogonia are present. The nucleus has disintegrated and is no longer visible. Oocyte diameter ranges 177.80–239.40 μm (average 208.22 μm). Sb: 200 μm .

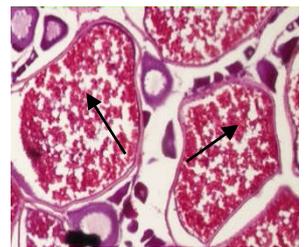


Figure 6 – Histological identification female gonads of *S. boops*

Gonadosomatic index. GSI value was fluctuated with range 0.46 – 1.89 and the highest and the lowest mean GSI value in October 2016 and September 2016. The average of GSI increased 1.05 on May 2016 from 0.63 on April 2016 and become decreased 0.70 on June 2016. After that, peak of GSI average increased 1.89 on October 2016 and become decreased 1.19 on November 2016 (Figure 7).

Length at first capture (Lc) and Length at first maturity (Lm). Based on SPR analysis, we can see the distribution values of *S. boops* caught during the research and most of the fish caught have a fork length of 16 cm (Figure 8). Length at first capture was 14.26 cm and length at first maturity was 16.30 cm. The comparison of length at first capture (Lc) and

length at first maturity (L_m) of *S. boops* shows in Figure 9. The value of SPR fishing rate relative to mortality (F/M) was 2.23 and asymptotic length (L_∞) was 26.40 cm.

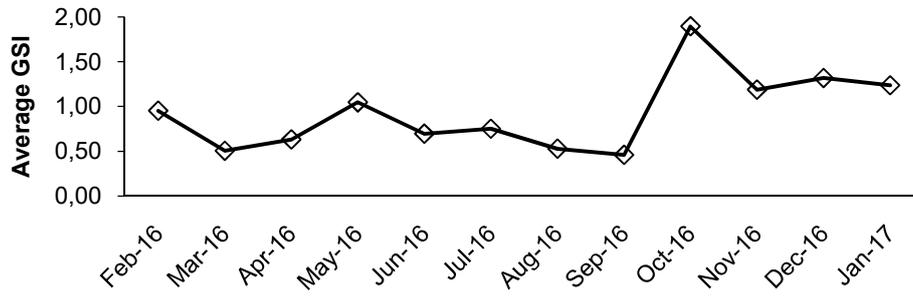


Figure 7 – Monthly average GSI of female *S. boops*

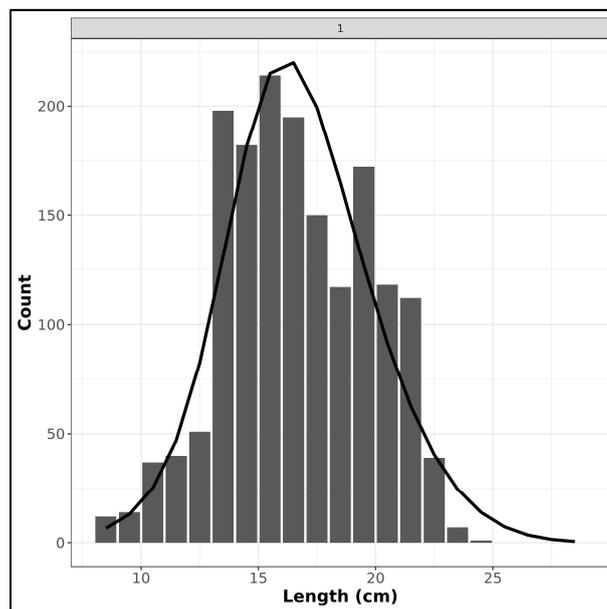


Figure 8 – Length distribution diagram of *S. boops*

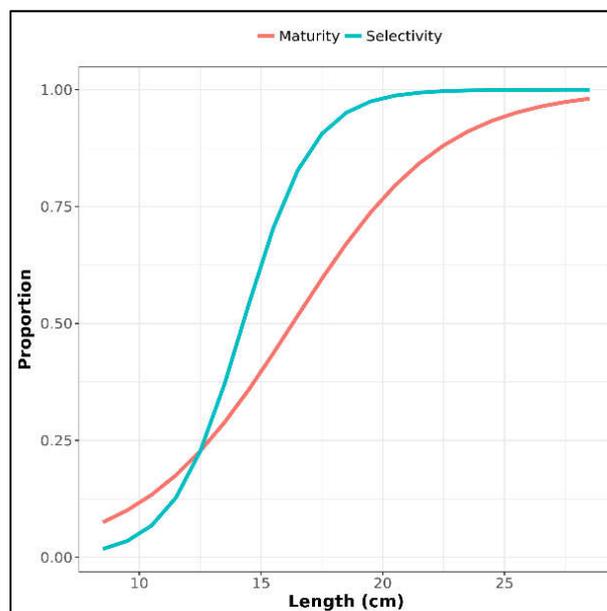


Figure 9 – Comparison of length at first capture (L_c) and Length at first maturity (L_m) of *S. boops*

Figure 9 above shows trends in which $L_c < L_m$, which means that the caught fishes are still immature. This condition was not good based on the fish biology aspect, because the generally fish were caught immature and suspected have not reproduced. If this condition is allowed to occur continuously it can be dangerous for *S. boops* stock in Bitung waters. One of indication that the utilization of fish resources doing well based on biologically if the average size of the caught fish is greater than the size of the first maturity ($L_c > L_m$), otherwise if $L_c < L_m$ can disturb the stability of the stock indeed cause over fishing of *S. boops* in Bitung waters. The value of F/M from the analysis shows that *S. boops* has F/M values of > 1 which means that the number of fishing mortality is higher than natural mortality due to fishing activity. Uncontrolled fishing can lead to a decrease in the average age and size of fish (Trippel 1995; Hutchings 2004; Allan & Castillo 2007). Walters & Martell (2004) state that fishing exploitation should be selective to the size of the fish, it's intended to avoid the occurrence of recruitment overfishing and growth overfishing.

This species is restricted to the Indo-West Pacific except for one confirmed record from Sesimbra Bay, Portugal. Elsewhere, this species has been recorded from the Andaman Islands to Vanuatu (Smith-Vaniz 1984), north to the Philippines, south to northern Australia (Paxton *et al* 1989), including Palau, New Caledonia and the Solomon Islands (Smith-Vaniz 1984). The result of genetic confirmation by phylogenetic tree (Figure 3), explains that the local type of trevallies namely Tude Batu which captured in waters around Bitung within Molluccas Sea is *S. boops*. Confirmation species using genetically is very important in the management of fish resources based on stock especially in Bitung waters area. By knowing valid species, sustainable management strategy can be developed. Phylogenetic trees have several uses such as summarizing the phylogeny of organisms by combining it with other data source analysis, studying co-speciation, calibrating the rate of molecular evolution, determining the age of estimate or genealogy, gene duplication analysis, estimating diversification rates, extinction, polymorphism, recombination, and population dynamics (Holder & Lewis 2003).

The findings of the overall growth this study $b = 3.164$ (FL) was similarly demonstrated by Isa *et al* (1996) and Gumanao *et al* (2016) which also it showed a positive allometry $b = 3.174$ (TL) and $b = 3.234$ (SL) respectively. Our result suggests that the *S. boops* in Bitung waters area grows faster in weight than length and the slope (b) value male slightly higher than female. This variation may be due to the difference in body forms, nutritional condition, and/or number of samples. Pauly (1984) suggested that "b" value less than 3 indicated that fish becomes more slender as it increases in length and with a value greater than 3 denotes stoutness indicating allometric growth. Differences in the value of b can give a picture of the differences in morphological characters of fish. The greater the value b the fish body is plumper. For example in this study the body shapes female of *S. boops* is plumper compared to male. LWR can be used as a character for taxonomic unit differentiation and changes in relationships with life events in life such as metamorphosis, growth and maturity (Thomas *et al* 2003), and as an important tool in exploiting appropriate fishing practices, as well as managing fish populations (Ahmed *et al* 2011; Valeria *et al* 2014).

Based on sex ratio this study was found that proportion of male slightly higher than female (1.12:1) and Chi-square analysis shown that balance proportion between male and female. The sex ratio male to female was high in six months: March (1:0.51), April (1:0.48), June (1:0.79), July (1:0.86), September (1:0.80) and January (1:0.52) while female to male ratio was high in six months too: February (0.41:1), May (0.95:1), August (0.92:1), October (0.73:1), November (0.74:1) and December (0.87:1). This conveys that sex ratio during this study was fluctuated, however males were observed to be dominant than females. Although the sex ratio in nature often occur deviations from ideal conditions, at least proportion of females should be more than males (2:1) for schooling fish to keep population conditions stable for sustainability resources (Ball & Rao 1984; Rahman *et al* 2013).

Gonadosomatic index (GSI) analysis for females of *S. boops* showed the lowest values in the beginning in February 2016 to September 2016 and progressively increased in October 2016 to January 2017 with the highest GSI recorded in October 2016. This indicated that the spawning periods of *S. boops* in Bitung waters area occurs throughout in October

2016 to January 2017 with peak season in October 2016. Based on GSI analysis it can be concluded that spawning season of *S. boops* in Bitung waters area occurs once in a year. This conveys that fishing exploitation have to attention the spawning season of *S. boops* for recruitment to keep stocks stable. Based on macroscopic and microscopic identification at the gonad maturity stage, the percentage between immature and mature were 59.95% and 40.05%, these result indicate that fish were caught generally immature. Sparre & Venema (1998) state that if there are too few old fish the stock is overfished and the fishing pressure on the stock should be reduced and if the fish are caught too young there is growth overfishing of the stock.

CONCLUSION

This study provides the first description of *S. boops* in Bitung water area in Indonesia. There were strong relationships between the lengths and weights of *S. boops* where there is high dependency of weight on length. Both of sexes were experiencing positive allometry growth. Spawning periods of *S. boops* throughout in October to January with peak season in October. Gonad development during spawning season is one factor that affects gonadosomatic index of fish which is also evident in this study. Sex ratios for *S. boops* were balanced with partial spawning type. The length at first caught (L_c) is smaller than the length at first maturity (L_m), therefore the some aspects of biology of *S. boops* related to legal-size are needed in fisheries management.

REFERENCES

1. Ahmed O. E., Mohammed E. A., Afra A. A., 2011 Length-weight relationship and condition factors of six fish species in Atbara River and Khashm El-girba reservoir, Sudan. *International Journal of Agriculture Sciences* 3(1):65-70.
2. Allan J. D., Castillo M. M., 2007 *Stream ecology: Structure and function of running waters*. Second edition. Springer. Netherland.
3. Allen G. R., Erdmann M. V., 2012 *Reef Fishes of the East Indies*. Tropical Reef Research, Perth, Australia.
4. Asahida T., Kobayashi T., Saitoh K., Nakayama I., 1996 Tissue Preservation and Total DNA Extraction from Fish Stored at Ambient Temperature Using Buffer Containing High Concentration of Urea. *Fisheries Science* 62 (5): 727-730.
5. Ayo-Alalusi C. I., 2014 Length weight relationship, condition factor and sex ratio of African Mud Catfish (*Clarias gariepinus*) reared in flow-through systems tanks. *Journal of Fisheries and Aquatic Science* 9(5): 430-434.
6. Ball D. V., Rao K. V., 1984 *Marine fisheries*. Tata McGraw-Hill Publishing Company, New Delhi, pp. 51-73.
7. Carpenter K. E., 2001 *The Living Marine Resources of the Western Central Pacific*. Volume 4. Bony fishes part 2 (Mugilidae to Carangidae). Rome, FAO: pp. 2659-2737.
8. Froese R., 2006 Cube law, condition factor and weight-length relationships: history, meta-analysis and recommendations. *Journal of Applied Ichthyology* 22:241-253.
9. Froese R., Pauly D., 2014. *Selar Crumenophthalmus* in Fish Base. August 2015 version. N.p.: FishBase, 2014. <http://www.fishbase.org>.
10. Gomelyuk, V.E. 2009. Fish assemblages composition and structure in three shallow habitats in north Australian tropical bay, Garig Gunak Barlu National Park, Northern Territory, Australia. *Journal of the Marine Biological Association of the United Kingdom* 89(3): 449-460.
11. Gumanao G. S., Cardoso M. M. S., Muller B., Bos A. R., 2016 Length-weight and length-length relationships of 139 Indo-Pacific fish species (Teleostei) from the Davao Gulf, Philippines. *Journal of Applied Ichthyology* 1-9.
12. Hutubessy G., 2011 Encircling gillnet selectivity for oxeye scad (*Selar boops* CUVIER, 1833) in the coast of Waai, Ambon Island. *Journal of Coastal Development* 14(2): 125-130.

13. Hall T. H., 1999 BioEdit: a user friendly biological sequence alignment editor and analysis program for Windows 95/98/NT. *Nucleic Acid Symposium Series* 4: 95-98.
14. Holder M., Lewis P. O., 2003 Phylogeny estimation: traditional and Bayesian approaches. *Nature Reviews Genetics* 4: 275-284.
15. Holden M. J, Raitt D. F. S., 1974 *Manual of fisheries science Part 2: Methods of resource investigation and their application.* Rome: FAO. <http://www.fao.org/docrep/003/f0752e/F0752E05.htm#ch5.2.2.2>.
16. Hutchings J. A., 2004 The cod that got away. *Nature* 428: 899-900.
17. Ikejima K., 2004 Molecular Phylogeneny and Possible Scenario of Ponyfish (Perciformes: Leiognathidae) Evolution. *Molluculer Phylogenet and Evolution.* 31: 904-909.
18. Isa M. M., Abdullah S., Yasin A. H., 1996 Population structure of small pelagic fish off the east coast of Peninsular Malaysia. *Fisheries Buletin No. 99:* 1-27. Department of Fisheries Malaysia. Ministry of Agriculture.
19. Paxton J. R., Hoese D. F., Allen G. R, Hanley J. E., 1989 *Pisces. Petromyzontidae to Carangidae.* Australian Government Publishing Service, Canberra.
20. Prince J., 2014 A technical report on an SPR size assessment of the blue swimmer crab fishery in Southeast Sulawesi. *Technical Report for IMACS, USAID,* 30 pp.
21. Pauly D., 1984 *Fish Population Dynamics in tropical Waters: A Manual for Use with Programmable Calculators.* ICLARM Studies and Reviews. No.8. 325 pp.
22. Rahman Y., Setyawati T. R., Yanti A. H., 2013. Characteristics of population Biawan fish (*Helostoma temminckii* Cuvier) in lake Kelubi, Tayan Hilir sub-district. *Jurnal Protobiont* 2(2): 80-86. [in Indonesian]
23. Randall J. E., Allen G. R., Steene R. C., 1990 *Fishes of the Great Barrier Reef and Coral Sea.* University of Hawaii Press, Honolulu. 506 pp.
24. Smith-Vaniz W. F., 1984. Carangidae. In: Fischer, W. and Bianchi, G. (eds), *FAO species identification sheets for fishery purposes. Western Indian Ocean fishing area 51. Vol. 1.,* Food and Agriculture Organization of the United Nations (FAO), Rome.
25. Smith-Vaniz W. F., 1986 Carangidae. In: M.M. Smith and P.C. Heemstra (eds), *Smith's Sea Fishes,* Springer-Verlag, Berlin, pp. 638-661.
26. Smith-Vaniz W. F., 1995 Carangidae. Jacks, branches, cojinúas, shoemakers, cooks, Casabes, mackerel, mackerel, humpback, amberjack, pilot fish p. 940-986 in W. Fischer, F. Krupp, W. Schneider, C. Sommer, KE Carpenter and V. Niem (eds.) *FAO Species Identification Guide for the Fines for Fisheries. Eastern Central Pacific.*Vol. 3. FAO, Rome.
27. Smith-Vaniz W. F., Williams I., 2016 Selar boops, Oxeye Scad. *The IUCN Red List of Threatened Species 2016:* e.T18158262A115368272 . 7pp
28. Snedecor G. W, Cochran W. G. 1967 *Statistical methods.* 6th edition. Oxford and IBH Publishing Co, New Dehli. 593 pp.
29. Sparre, P & S.C Venema. (1998). *Introduction to Tropical Fish Stock Assessment. Part I: Manual.* FAO Computerized Information Series (Fisheries). No. 12. FAO (Food and Agriculture Organization of the United Nations), Rome. 407p.
30. Tamura K., Stecher G., Peterson D., Filipski A., 2013 MEGA6: Molecular Evolutionary Genetics Analysis version 6.0. *Molecular Biology and Evolution* 30: 2725-2729.
31. Thomas J. S., Venus, Kurup B. M., 2003 Length-weight relationship of some deep-sea fish inhabiting continental slope beyond 250 m depth along West coast of India. *Naga, world fish center quart* 26: 17-21.
32. Trippel E. A., 1995. Age at immaturity as a stress indicator in fisheries. *BioScience* 45: 759-771.
33. Valeria R. B. F., Michelato M., da Cruz T. P., Furuya W. M., 2014 Length-weight relationships and prediction equations of body composition of farm raised *Astyanax aff. fasciatus* (Actinopterygii: Characiformes: Characidae). *Zoologia* 31(6): 521–524.
34. Ward R. D., Zemlak T. S., Innes B. H., Last P. R., 2005 DNA Barcoding Australia's fish species. *Philosophical Transaction of the Royal Society. B* 360, 1847-1857.
35. Walters C. J., 2004 *Fisheries ecology and management.* Princeton University, 448 pp.