

What speciesmake upthe Nike fish assemblages at the macrotidal estuary in Gorontalo Bay, Indonesia?

Femy M. Sahami^{1*}, Rene Charles Kepel², Abdul Hafidz Olii¹, Silvester Benny Pratasik²

Corresponding author: Femy M. Sahami (femysahami@ung.ac.id)

Abstract

Background:No study has documented the species composition of Nikefish(fam:Gobiidae)schools. The aim of this study is to document the species composition of the Nike-fish schooling.

Methods: All samples were collected randomly from fisher's catch during the fishing season on 5th-11thOctober 2018 at macrotidal area in Leato. Then, all specimens were identified morphologically by melanophore pattern differences. Subsequently, all identified-samples by melanophores pattern differences were sent to the genetic laboratory for identification. **Results:** The morphological resultsshow there are five individuals with a different melanophores pattern. On the contrary, the genetic results only showfour species from those five individuals. They are *Sicyopterus pugnans*, *S. cynocephalus*, *Belobranchus segura*, and *Bunaka gyrinoides*.

Conclusions: Our findings show that there are only four species that composethe Nike fish schooling in Gorontao Bay. They are *Sicyopterus pugnans*, *Sicyopterus cynocephalus*, *Belobranchussegura*, and *Bunakagyrinoides*

Keywords: Nike-fish, gorontalo, melanophores pattern, genetic, morphology

Style Definition: Heading 4
Style Definition: Header: Font:
Style Definition: Footer: Font:

Style Definition: Comment Text:

Font:

¹Faculty of Fisheries and Marine Science, Gorontalo State University, Jl. Jendral Sudirman. No. 6, Gorontalo City, 96128, Gorontalo Province, Indonesia;

²Faculty of Fisheries and Marine Science, Sam Ratulangi University, Jl. KampusUnsrat Bahu, Manado City, 95115, North Sulawesi Province, Indonesia.

Introduction

Estuaries area crucial habitat for biota and small fish, in particular juveniles of commercially relevant species. They are considered as the most productive and dynamic ecosystem in the world (Cantera et al., 2001; Lahjie et al., 2019; McHugh, 1967; Sreekanth et al., 2017). They also perform the most crucial role in the population dynamic for a lot of invertebrate and fish species. These ecosystems also significantly contribute to providing some ecological services such as nursery ground, feeding ground and breeding habitats for both freshwater and marine species (Beck et al., 2001; McLusky and Elliott, 2004; Sun et al., 2019). The most well-known species that occupy the seas and estuary area in Gorontalo Bay is Nikefish.

Nike (pronounced nee-K)is a local name for transparent juvenile of unknown fish. These fish areapproximately2–4 cm in length;they appear seasonally and fished at estuary waters around the Gorontalo Bay. These juvenile fish has been fished and marketed traditionally for a long time. They are preferable for consumption by the local people than other fisheries products. As a consequence, fishing activity has increased over time to supply local demand for Nike (Wolok et al., 2019).

However, the impact of fishing activities is unknown. A recent paper concerningNikeonly reports the seasonal appearance during the fishing season (Pasisingi & Abdullah, 2018), total length and morphometric measurements (Zakaria, 2018), nutrition content (Liputo et al., 2013), and mercury contamination of these fish(Salam et al., 2016). To our knowledge, no studies have documented the species diversity that composed the schooling of Nike. Although, Yamasaki et al., (2011) have reported that species in juvenile form can be determined by its melanophores pattern and genetic determination.

The objective of the present study is to address this lack of knowledge by identifying the fish species that composed a Nike fish schooling. This information is very urgent and required for fisheries management. Therefore, we aimed to identify the species that composed the schooling of Nike fish in Gorontalo Bay by melanophores pattern and genetic identification.

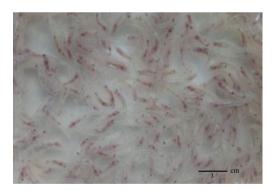


Figure 1.Nike fish assemblages.

Methods

This study was conducted in October 2018 at Leato (0°30'0.58"N, 123°3'55.42"E), Gorontalo Bay, Indonesia (Figure2). Approximately 100 g of the Nike-fish were collected randomly from the fishermen's catch at fishing grounds during the catch-season (on October 5th–11th). All samples were transported using a cool-box

to the lab for measurement. Immediately after collection, all samples were identified visually by (Yamasaki et al., 2011). and the specimens with different melanophores pattern were separated according to their melanophores display. We assummed that those separated individuals are different on species.

Then, we selected one individual at each group and labeled as N1, N2, N3, N4, N5, for genetic identification. Images of the selected samples were captured using Canon EOS 100d with 58 mm pro Digital Wide Converter 0.45X Lens and subsequently converted to black and white using CorelDraw Graphic Suite 2019.

After selection, all of the individual with different melanophores were preserved with ethanol 70% in a separate bottle and sent to the Genetics Laboratory at Manokwari for genetic identification by Sanger sequencing. The DNA cytochrome oxidase subunit I (CO1) of the sample was isolated with a GeneaidTM DNA Isolation Kit. Editing, proofreading, and the phylogenetic tree was generated with MEGA 5.0 software.



Figure 2. Study site. The red dot indicates the position of fishing ground where the samples were collected from fishermen.

Results

Five unspecified individuals of Nike-fish that have identified mofphologically by melanophores differences showing in Figure 3. N1 revealed as *Sicyopterus pugnans*; N2 as *Sicyopterus cynocephalus*, N3 and N5 as *Belobranchus segura*, and N4 as *Bunaka gyrinoides*. The specimens with melanophores differences of the each group showing in figure 4.

Melanophores pattern

Nike-fish schools consisted of various species with the same body-shape, but different melanophore displays. Moreover, from 100 g (~145 individuals) of the total specimens that have identified, only found five individuals with a different melanophores pattern were identified (Figure 3).

Comment [JT1]: By whom? Please describe separation of the fish into different groups and how/why you assumed them to be different species

Formatted: Default Paragraph Fon Font: Times New Roman

Comment [JT2]: Please state:
•Which gene was sequenced
•The software used to generate Figure

Comment [JT3]: This section contai insufficient detail – you must at least st all of the species identified

Comment [JT4]: Please list all of the species identified in this section, and stawhich pattern they have

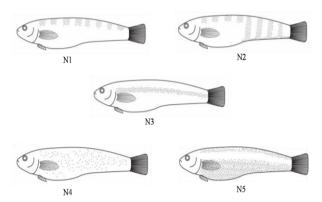


Figure 3. Nike fish with different melanophore patterns.

Genetic identification

Figure 3shows the genetic identification among the individuals (species). The outcomes of genetic identification for N3 and N5 shows that both samples are the same species: *Belobranchus segura*.

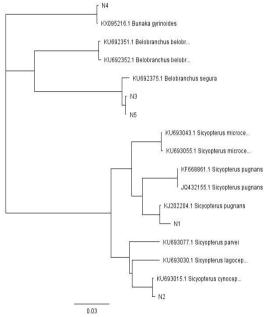


Figure 4. Phylogenetic tree of individuals with different melanophores patterns.

Discussion

Although the melanophore patterns N3 and N5 are different, their genetics are identical, meaning they are the same species (*Belobranchus segura*). This dissimilarity might be affected by the changes of melanophore during the development of the larvae. Valade et al., (2009) report that such melanophores chang on *Sicyopterus langocephalus* during the larvae stage. These changes could represent a problem for morphological identification. We can not count the species by morphological differences. Therefore, for the next examination we strongly recommended determining the species composition of the Nike fish schools by genetic rather than morphological identification because for that reason.

Conclusion

Our findings show that there are four species that compose Nike fish schooling. They are Sicyopterus pugnans, Sicyopterus cynocephalus, Belobranchus segura, and Bunaka gyrinoides.

Data availability

Underlying data

Group N1, *Sicyopterus pugnans* isolate N1_LEATO_1 cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial. GenBank accession number MN065178.

Group N2, Sicyopterus cynocephalus isolate N2_LEATO_1 cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial. GenBank accession number MN069305.

Group N3, Belobranchussegura isolate N3_LEATO_1 cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial. GenBank accession number MN069306.

Group N4, Bunakagyrinoides isolate N4_LEATO_1 cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial. GenBank accession number MN069307.

Group N5, Belobranchussegura isolate N5_LEATO_1 cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial. GenBank accession number MN069308.

Grant information

The authors declare that no grants were involved in supporting this work

Acknowledgments

The authors would like to thank La Nane, Sitty Ainsyah Habibie, and Nuralim Pasisingi for technical writing and support during this research.

References

Baldwin CC, Mounts JH, Smith DG, and Weigt LA, 2008. Genetic identification and color descriptions of early life-history stages of Belizean Phaeoptyx and Astrapogon (Teleostei: Apogonidae) with comments on identification of adult Phaeoptyx. Zootaxa. 1–22.

Beck, M.W., Heck, K.L., Able, K.W., Childers, D.L., Eggleston, D.B., Gillanders, B.M., Halpern, B., Hays, C.G., Hoshino, K., Minello, T.J., Orth, R.J., Sheridan, P.F., Weinstein, M.P., 2001. A better understanding of the habitats that serve as nurseries for marine species and the factors that create site-specific variability in nursery quality will improve conservation and management of these areas. BioScience 51, 633–641.

Boutin, B. P., & Targett, T. E. (2019). Density, Growth, Production, and Feeding Dynamics of Juvenile Weakfish (Cynoscionregalis) in Delaware Bay and Salt Marsh Tributaries: Spatiotemporal Comparison of Nursery Habitat Quality. *Estuaries and Coasts*, 42(1), 274-291.

Formatted: Line spacing: single

- Cantera, J. R., & Blanco, J. F. (2001). The estuary ecosystem of Buenaventura bay, Colombia. In *Coastal marine ecosystems of Latin America* (pp. 265-280). Springer, Berlin, Heidelberg.
- Lahjie, A. M., Nouval, B., Lahjie, A. A., Ruslim, Y., & Kristiningrum, R. (2019). Economic valuation from direct use of mangrove forest restoration in Balikpapan Bay, East Kalimantan, Indonesia. *F1000Research*, 8.
- Larmuseau, M. H., Huyse, T., Vancampenhout, K., Van Houdt, J. K., &Volckaert, F. A. (2010). High molecular diversity in the rhodopsin gene in closely related goby fishes: A role for visual pigments in adaptive speciation? Molecular phylogenetics and evolution, 55(2), 689-698.
- Liputo, S. A., Berhimpon, S., & Fatimah, F. (2013). Analisa Nilai Gizi Serta KomponenAsam Amino dan Asam Lemak dari Nugget IkanNike (Awaousmelanocephalus) DenganPenambahan Tempe. *CHEMISTRY PROGRESS*, 6(1).
- McHugh, J.L., 1967. Estuarine nekton. In: Lauff, G.H. (Ed.), Estuaries, vol. 83. American Association for the Advancement of Science Special Publication, Washington, DC, pp. 581–620.
- McLusky, D.S., Elliott, M., 2004. The Estuarine Ecosystem. Ecology, Threats and Management, third ed. Oxford University Press, 214 pp.
- Pasisingi, N., & Abdullah, S. (2018). Pola kemunculanikan nikeNike (Gobiidae) di PerairanTeluk Gorontalo, Indonesia. *DEPIK JurnalIlmu-IlmuPerairan*, *Pesisir dan Perikanan*, 7(2), 111-118.
- Salam, A., Sahami, F. M., &Panigoro, C. (2016). Nike (Awaousmelanocephalus) Fishery and Mercury Contamination in the Estuary of BoneBolango River. *Omni-Akuatika*, 12(2).
- Sanger, F., Nicklen, S., & Coulson, A. R. (1977). DNA sequencing with chain-terminating inhibitors. Proceedings of the national academy of sciences, 74(12), 5463-5467.
- Sreekanth, G. B., Lekshmi, N. M., & Singh, N. P. (2017). Temporal patterns in fish community structure: environmental perturbations from a well-mixed tropical estuary. *Proceedings of the National Academy of Sciences, India Section B: Biological Sciences*, 87(1), 135-145.
- Wolok, T., Fachrussyah, Z. C., &Yantu, I. (2019). Technical And Economic Analysis Of Catching Equipment Totaluo In Nike Fishing (AwaousMelanocephalus) In Gorontalo City. Jambura Science of Management, 1(2), 65–71.
- Yamasaki, N., Kondo, M., Maeda, K., &Tachihara, K. (2011). Reproductive biology of three amphidromous gobies, Sicyopterus japonicus, Awaousmelanocephalus, and Stenogobius sp., on Okinawa Island/Biologie de la reproduction de trois gobies amphidromes de l'iled'Okinawa: Sicyopterus japonicus, Awaousmelanocephalus et Stenogobius sp. *Cybium, International Journal of Ichthyology*, *35*(4), 345-360.
- Zakaria, Z. (2018). AnalisisMorfometrik Schooling IkanNike di PerairanLautPesisir Kota Gorontalo. *Jambura Journal of Educational Chemistry*, 13(1), 77-80.

Formatted: Font: Arial, 8,5 pt, Foncolor: Custom Color(RGB(34;34;34)



What speciesmake upthe Nike fish assemblages at the macrotidal estuary in Gorontalo Bay, Indonesia?

Femy M. Sahami^{1*}, Rene Charles Kepel², Abdul Hafidz Olii¹, Silvester Benny Pratasik²

Corresponding author: Femy M. Sahami (femysahami@ung.ac.id)

Abstract

Background:No study has documented the species composition of Nikefish(fam:Gobiidae)schools. The aim of this study is to document the species composition of the Nike-fish schooling.

Methods: All samples were collected randomly from fisher's catch during the fishing season on 5th-11thOctober 2018 at macrotidal area in Leato. Then, all specimens were identified morphologically by melanophore pattern differences. Subsequently, all identified-samples by melanophores pattern differences were sent to the genetic laboratory for identification. **Results:** The morphological resultsshow there are five individuals with a different melanophores pattern. On the contrary, the genetic results only showfour species from those five individuals. They are *Sicyopterus pugnans*, *S. cynocephalus*, *Belobranchus segura*, and *Bunaka gyrinoides*.

Conclusions: Our findings show that there are only four species that composethe Nike fish schooling in Gorontao Bay. They are *Sicyopterus pugnans*, *Sicyopterus cynocephalus*, *Belobranchussegura*, and *Bunakagyrinoides*

Keywords: Nike-fish, gorontalo, melanophores pattern, genetic, morphology

Style Definition: Heading 4
Style Definition: Header: Font:
Style Definition: Footer: Font:

Style Definition: Comment Text:

Font:

¹Faculty of Fisheries and Marine Science, Gorontalo State University, Jl. Jendral Sudirman. No. 6, Gorontalo City, 96128, Gorontalo Province, Indonesia;

²Faculty of Fisheries and Marine Science, Sam Ratulangi University, Jl. KampusUnsrat Bahu, Manado City, 95115, North Sulawesi Province, Indonesia.

Introduction

Estuaries area crucial habitat for biota and small fish, in particular juveniles of commercially relevant species. They are considered as the most productive and dynamic ecosystem in the world (Cantera et al., 2001; Lahjie et al., 2019; McHugh, 1967; Sreekanth et al., 2017). They also perform the most crucial role in the population dynamic for a lot of invertebrate and fish species. These ecosystems also significantly contribute to providing some ecological services such as nursery ground, feeding ground and breeding habitats for both freshwater and marine species (Beck et al., 2001; McLusky and Elliott, 2004; Sun et al., 2019). The most well-known species that occupy the seas and estuary area in Gorontalo Bay is Nikefish.

Nike (pronounced nee-K)is a local name for transparent juvenile of unknown fish. These fish areapproximately2–4 cm in length;they appear seasonally and fished at estuary waters around the Gorontalo Bay. These juvenile fish has been fished and marketed traditionally for a long time. They are preferable for consumption by the local people than other fisheries products. As a consequence, fishing activity has increased over time to supply local demand for Nike (Wolok et al., 2019).

However, the impact of fishing activities is unknown. A recent paper concerningNikeonly reports the seasonal appearance during the fishing season (Pasisingi & Abdullah, 2018), total length and morphometric measurements (Zakaria, 2018), nutrition content (Liputo et al., 2013), and mercury contamination of these fish(Salam et al., 2016). To our knowledge, no studies have documented the species diversity that composed the schooling of Nike. Although, Yamasaki et al., (2011) have reported that species in juvenile form can be determined by its melanophores pattern and genetic determination.

The objective of the present study is to address this lack of knowledge by identifying the fish species that composed a Nike fish schooling. This information is very urgent and required for fisheries management. Therefore, we aimed to identify the species that composed the schooling of Nike fish in Gorontalo Bay by melanophores pattern and genetic identification.

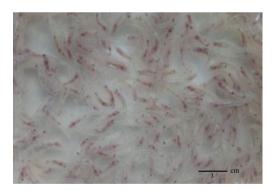


Figure 1.Nike fish assemblages.

Methods

This study was conducted in October 2018 at Leato (0°30'0.58"N, 123°3'55.42"E), Gorontalo Bay, Indonesia (Figure2). Approximately 100 g of the Nike-fish were collected randomly from the fishermen's catch at fishing grounds during the catch-season (on October 5th–11th). All samples were transported using a cool-box

to the lab for measurement. Immediately after collection, all samples were identified visually by (Yamasaki et al., 2011). and the specimens with different melanophores pattern were separated according to their melanophores display. We assummed that those separated individuals are different on species.

Then, we selected one individual at each group and labeled as N1, N2, N3, N4, N5, for genetic identification. Images of the selected samples were captured using Canon EOS 100d with 58 mm pro Digital Wide Converter 0.45X Lens and subsequently converted to black and white using CorelDraw Graphic Suite 2019.

After selection, all of the individual with different melanophores were preserved with ethanol 70% in a separate bottle and sent to the Genetics Laboratory at Manokwari for genetic identification by Sanger sequencing. The DNA cytochrome oxidase subunit I (CO1) of the sample was isolated with a GeneaidTM DNA Isolation Kit. Editing, proofreading, and the phylogenetic tree was generated with MEGA 5.0 software.



Figure 2. Study site. The red dot indicates the position of fishing ground where the samples were collected from fishermen.

Results

Five unspecified individuals of Nike-fish that have identified mofphologically by melanophores differences showing in Figure 3. N1 revealed as *Sicyopterus pugnans*; N2 as *Sicyopterus cynocephalus*, N3 and N5 as *Belobranchus segura*, and N4 as *Bunaka gyrinoides*. The specimens with melanophores differences of the each group showing in figure 4.

Melanophores pattern

Nike-fish schools consisted of various species with the same body-shape, but different melanophore displays. Moreover, from 100 g (~145 individuals) of the total specimens that have identified, only found five individuals with a different melanophores pattern were identified (Figure 3).

Comment [JT1]: By whom? Please describe separation of the fish into different groups and how/why you assumed them to be different species

Formatted: Default Paragraph Fon Font: Times New Roman

Comment [JT2]: Please state:
•Which gene was sequenced
•The software used to generate Figure

Comment [JT3]: This section contai insufficient detail – you must at least st all of the species identified

Comment [JT4]: Please list all of the species identified in this section, and stawhich pattern they have

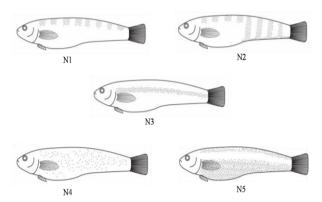


Figure 3. Nike fish with different melanophore patterns.

Genetic identification

Figure 3shows the genetic identification among the individuals (species). The outcomes of genetic identification for N3 and N5 shows that both samples are the same species: *Belobranchus segura*.

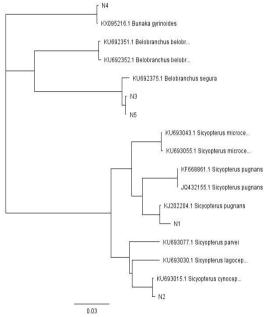


Figure 4. Phylogenetic tree of individuals with different melanophores patterns.

Discussion

Although the melanophore patterns N3 and N5 are different, their genetics are identical, meaning they are the same species (*Belobranchus segura*). This dissimilarity might be affected by the changes of melanophore during the development of the larvae. Valade et al., (2009) report that such melanophores chang on *Sicyopterus langocephalus* during the larvae stage. These changes could represent a problem for morphological identification. We can not count the species by morphological differences. Therefore, for the next examination we strongly recommended determining the species composition of the Nike fish schools by genetic rather than morphological identification because for that reason.

Conclusion

Our findings show that there are four species that compose Nike fish schooling. They are Sicyopterus pugnans, Sicyopterus cynocephalus, Belobranchus segura, and Bunaka gyrinoides.

Data availability

Underlying data

Group N1, *Sicyopterus pugnans* isolate N1_LEATO_1 cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial. GenBank accession number MN065178.

Group N2, Sicyopterus cynocephalus isolate N2_LEATO_1 cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial. GenBank accession number MN069305.

Group N3, Belobranchussegura isolate N3_LEATO_1 cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial. GenBank accession number MN069306.

Group N4, Bunakagyrinoides isolate N4_LEATO_1 cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial. GenBank accession number MN069307.

Group N5, Belobranchussegura isolate N5_LEATO_1 cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial. GenBank accession number MN069308.

Grant information

The authors declare that no grants were involved in supporting this work

Acknowledgments

The authors would like to thank La Nane, Sitty Ainsyah Habibie, and Nuralim Pasisingi for technical writing and support during this research.

References

Baldwin CC, Mounts JH, Smith DG, and Weigt LA, 2008. Genetic identification and color descriptions of early life-history stages of Belizean Phaeoptyx and Astrapogon (Teleostei: Apogonidae) with comments on identification of adult Phaeoptyx. Zootaxa. 1–22.

Beck, M.W., Heck, K.L., Able, K.W., Childers, D.L., Eggleston, D.B., Gillanders, B.M., Halpern, B., Hays, C.G., Hoshino, K., Minello, T.J., Orth, R.J., Sheridan, P.F., Weinstein, M.P., 2001. A better understanding of the habitats that serve as nurseries for marine species and the factors that create site-specific variability in nursery quality will improve conservation and management of these areas. BioScience 51, 633–641.

Boutin, B. P., & Targett, T. E. (2019). Density, Growth, Production, and Feeding Dynamics of Juvenile Weakfish (Cynoscionregalis) in Delaware Bay and Salt Marsh Tributaries: Spatiotemporal Comparison of Nursery Habitat Quality. *Estuaries and Coasts*, 42(1), 274-291.

Formatted: Line spacing: single

- Cantera, J. R., & Blanco, J. F. (2001). The estuary ecosystem of Buenaventura bay, Colombia. In *Coastal marine ecosystems of Latin America* (pp. 265-280). Springer, Berlin, Heidelberg.
- Lahjie, A. M., Nouval, B., Lahjie, A. A., Ruslim, Y., & Kristiningrum, R. (2019). Economic valuation from direct use of mangrove forest restoration in Balikpapan Bay, East Kalimantan, Indonesia. *F1000Research*, 8.
- Larmuseau, M. H., Huyse, T., Vancampenhout, K., Van Houdt, J. K., &Volckaert, F. A. (2010). High molecular diversity in the rhodopsin gene in closely related goby fishes: A role for visual pigments in adaptive speciation? Molecular phylogenetics and evolution, 55(2), 689-698.
- Liputo, S. A., Berhimpon, S., & Fatimah, F. (2013). Analisa Nilai Gizi Serta KomponenAsam Amino dan Asam Lemak dari Nugget IkanNike (Awaousmelanocephalus) DenganPenambahan Tempe. *CHEMISTRY PROGRESS*, 6(1).
- McHugh, J.L., 1967. Estuarine nekton. In: Lauff, G.H. (Ed.), Estuaries, vol. 83. American Association for the Advancement of Science Special Publication, Washington, DC, pp. 581–620.
- McLusky, D.S., Elliott, M., 2004. The Estuarine Ecosystem. Ecology, Threats and Management, third ed. Oxford University Press, 214 pp.
- Pasisingi, N., & Abdullah, S. (2018). Pola kemunculanikan nikeNike (Gobiidae) di PerairanTeluk Gorontalo, Indonesia. *DEPIK JurnalIlmu-IlmuPerairan*, *Pesisir dan Perikanan*, 7(2), 111-118.
- Salam, A., Sahami, F. M., &Panigoro, C. (2016). Nike (Awaousmelanocephalus) Fishery and Mercury Contamination in the Estuary of BoneBolango River. *Omni-Akuatika*, 12(2).
- Sanger, F., Nicklen, S., & Coulson, A. R. (1977). DNA sequencing with chain-terminating inhibitors. Proceedings of the national academy of sciences, 74(12), 5463-5467.
- Sreekanth, G. B., Lekshmi, N. M., & Singh, N. P. (2017). Temporal patterns in fish community structure: environmental perturbations from a well-mixed tropical estuary. *Proceedings of the National Academy of Sciences, India Section B: Biological Sciences*, 87(1), 135-145.
- Wolok, T., Fachrussyah, Z. C., &Yantu, I. (2019). Technical And Economic Analysis Of Catching Equipment Totaluo In Nike Fishing (AwaousMelanocephalus) In Gorontalo City. Jambura Science of Management, 1(2), 65–71.
- Yamasaki, N., Kondo, M., Maeda, K., &Tachihara, K. (2011). Reproductive biology of three amphidromous gobies, Sicyopterus japonicus, Awaousmelanocephalus, and Stenogobius sp., on Okinawa Island/Biologie de la reproduction de trois gobies amphidromes de l'iled'Okinawa: Sicyopterus japonicus, Awaousmelanocephalus et Stenogobius sp. *Cybium, International Journal of Ichthyology*, *35*(4), 345-360.
- Zakaria, Z. (2018). AnalisisMorfometrik Schooling IkanNike di PerairanLautPesisir Kota Gorontalo. *Jambura Journal of Educational Chemistry*, 13(1), 77-80.

Formatted: Font: Arial, 8,5 pt, Foncolor: Custom Color(RGB(34;34;34)