

Phylogenetic and Taxonomic Studies of Paral ichthys (Paralichthyid): Evolutionary Insights from Molecular and Morphological Data

Murat Özkan

Istanbul University, Istanbul, Turkey.

Abstract

Paral ichthys forms a diverse genus among Paralichthyid flatfish which exhibit asymmetry and environmental importance in their development. The study of Paral ichthys evolution and species identities depends on taxonomy and phylogenetic research to discover the natural adaptations of its marine organisms. Researchers have contributed new advancements to Paral ichthys phylogenetic classification through a morphological trait analysis combined with molecular markers and genetic sequencing. This research explains how DNA barcoding methods alongside phylogenetic tree creation techniques help scientists define species limits and identify evolutionary relationships across the entire genus. This research evaluates how these discoveries contribute to evolutionary taxonomy excellence and affects the sustainable practices for Paral ichthys species management and conservation. Future research studying flatfish evolution and ecology as well their fisheries value must combine molecular data and morphological data simultaneously to achieve their objectives. Phylogenetic studies principally using mitochondrial and nuclear genes exposed the intricate evolutionary timeline for Paral ichthys which shows how natural selection split species between marine environments. The studies have positioned Paral ichthys among its family members of Paralichthyid while explaining evolutionary changes that led to benthic living adaptations. The integration of molecular data techniques with traditional morphological analyses continues to enhance Paral ichthys species classification which helps scientists better protect these commercially important fish species.

Keywords: Phylogenetic (PP), Taxonomic Studies (TS), Paral ichthys (PP), Evolutionary Insights (EI), Molecular (MM), Morphological Data (MD).

Citation: Murat, Ö. 2025. Phylogenetic and Taxonomic Studies of Paral ichthys (Paralichthyid): Evolutionary Insights from Molecular and Morphological Data. FishTaxa 35: 34-43.

Introduction

Paral ichthys is an important and specific family of flatfish which is also called sanddabs. These fishes are mostly found in various water reservoirs around the world. Here we are going to understand phylogenetic and Taxonomic Studies related to Paral ichthys. Usually, phylogenetic studies reveal a few important aspects such as molecular phylogeny, phylogenetic relationships, sister groups, and others. Taxonomic Studies are mostly used for the study of revised classification and species Delimitation of specific groups(Roje, 2014). Morphological studies are beneficial for the study of morphometric analysis, meristic characters, and osteological characters. The study of evolutionary Insights is related to the study of evolutionary history, adaptive radiations, and biogeographic patterns as well. Here we are going to discuss morphological studies first and then we will focus on taxonomic Studies and evolutionary Insights.

The important aspect of morphological studies is body morphology. These fishes are mostly flat, having oval-shaped bodies with small mouths and pointed snouts. These fishes have body sizes ranging from 10 to 60 cm. Morphological studies divide the body of these fishes into two parts, the anterior part consisting of the head and trunk and the exterior part consisting of the caudal fin and tail. If we discuss the skeletal system of these fishes, we may come to know these fishes have flat skulls and small mouths(Y Ph Kartavtsev et al., *Accepted: 14 August 2024; Published: 21 October 2024*

2007). The number of vertebrae in the vertebral column is 30 to 40. There are usually 10 to 12 pairs of ribs in these fishes. These fishes have four types of fins. Dorsal fins have almost 60 to 100 soft rays. The number of soft rays in anal fins ranges from 40 to 60. The pectoral and pelvic fins are usually smaller than the dorsal and anal fins. If we discuss the morphology of skin, we may come to know that there is the presence of scales on the body which are cycloid and cover the whole body except the head and fins. These scales have various functions such as protection, thermoregulation, flexibility, and others. Usually soon of these fishes is mostly smooth and slim which is quite helpful for easy and smooth swimming. The morphology of the eye is somehow different in these fishes. These fishes have a pair of eyes which is present on either side may be left or right(Timi et al., 2014; Young, 2023).

These fishes do have Gill rakers which are specialized for various functions. These gill rakers are useful for feeding, respiration, and other functions. Now we are going to discuss molecular data related to Paral ichthys. For analysis of molecular data, we may prefer mitochondrial DNA or nuclear DNA. If we use nuclear DNA for molecular data, we may focus on specific genes which may act as biomarkers(Roje, 2010). These specific genes are RAG1, RHOD gene, and TMO 4C4 gene. By studying these specific genes, we may get a better idea about the phylogenetic tree related to Paral ichthys. There are some other biotechnology techniques such as single nucleotide polymorphisms which are abbreviated as SNPs. Single nucleotide polymorphisms are used commonly these days because of the accuracy of results and detailed information. There are many other important applications of single nucleotide polymorphisms. One of its important applications is the high resolution of genetic data obtained by using single nucleotide polymorphisms(Chen, Bonillo, & Lecointre, 2003). The other important application of single nucleotide polymorphisms is that they can be used for largescale genotyping. Recent studies have convinced us that using single nucleotide polymorphisms for molecular data analysis is also cost-effective as compared to other analytical methods. Now we will enumerate important evolutionary Insights related to Paral ichthys. This Genus is much diversified in recent years because of various aspects. Geographic isolation and allopatric speciation played an important role in the evolution of new species of Paral ichthys. Geographic isolation means land has separated the same type of organisms but later on these organisms because of geographic isolation formed new forms of species(Roje, 2010). The important types of geographic isolation are continental isolation, Island isolation, mountain isolation, and riverine isolation. As a result of geographic isolation, many effects are to be considered here. The first major effect of Geographic isolation is that there is less gene flow between these populations. When these populations are separated by any factor, there will be less chance of gene flow from one population to another. The other important thing in population is gene drift. When sudden changes are seen in the genetics of a population over some time, this is called genetic drift which is possible because of geographic isolation. It has been seen that various species of this Genus evolved because of geographic isolation due to adaptation to the local environment(Bañón et al., 2005; Yagishita et al., 2009).

The other important aspect is allopatric speciation which is the result of the separation of populations by some physical barrier. Some important factors may contribute to allopatric speciation such as geographic barriers, climate change, and any kind of volcanic activity. Recent studies have shown that when there is more allopatric speciation, there is more chance of genetic diversity which may result in the evolution of many different types of species of the same Genus. If we discuss about phylogenetic relationship of this particular Genus, we may get to know that this Genus has having monophyletic lineage. This lineage means that all the kinds of species of this particular Genus have having same ancestor. There is also an important sister group of this Genus which is named Pseudorhombus(Redin & Kartavtsev, 2022). This is called so because of having the same characteristics as this Genus Paral ichthys. The phylogenetic tree has shown that there is a complex

relationship among various species of this Genus which is making it difficult to classify these fishes. Recent phylogenetic studies have shown that these fishes have shown transoceanic dispersal. It means that these fishes have dispersed the whole ocean basin because of various genetic adaptations. However, some species have shown regional endemism. It refers to the localization of any species in a specific region. These phylogenetic studies can be useful for taxonomic revision(Atta et al., 2022). As we know there is further addition of various Species in this Genus so there is a need to classify these fishes effectively in a new revised system. The other important application of these phylogenetic studies is that these can be used for sketching the conservative strategies for these fishes because we know that there are many conservative threats to these fishes in these days of increasing pollution(Villarreal et al., 2024).

Research Objective

The main objective of this research is to understand phylogenetic and taxonomic Studies of Paral ichthys on the basis of morphological studies and molecular data. These studies have convinced us that many other aspects of classification are related to morphology and phylogenetic relationships.

Literature Review

The Genus Paral ichthys comprises almost 15 species and is found mostly in tropical and subtropical water. When nuclear material was analyzed, there were a few important considerations related to Paral ichthys. The first and foremost outcome of these studies is the aspect of the monophyletic lineage of Paral ichthys (Yuri Phedorovich Kartavtsev et al., 2016). The word monophyletic lineage means they have the same common ancestor. This means that we have to admit that these species are mostly related to each other compared to other species of the genus (Wirtz, Davenport, & Chanet, 2023). We got an idea about the monophyletic lineage of Paral ichthys because of shared morphological traits. Recent studies have shown that there are some important Morphological characteristics such as the oval shape of the body, having a small mouth, and unique morphology of fins which are the same in all species of Genus Paral ichthys which convince us that they have monophyletic lineage. We also knew about the monophyletic lineage of Paral ichthys because of molecular signatures (Bruns, 2012). When we collected molecular data by analysis of mitochondrial DNA and nuclear DNA, we got to the point that this Genus has a monophyletic relationship. When we studied the phylogenetic tree of Paral ichthys, we came to know a few important phylogenetic relationships. These studies have shown that some species are similar to Paral ichthys so they are termed as the sister group of Paral ichthys (Cooper, 1996). If we know about the phylogenetic relationship of Paral ichthys, we will have some important implications. For example, the most important implication of phylogenetic studies is taxonomic stability. Because of phylogenetic studies, we classified Paral ichthys as a monophyletic group which helped us stabilize the taxonomic framework of Paral ichthys.

The other important implication of phylogenetic studies of Paral ichthys is that we may get Insights about the evolutionary history of Paral ichthys(Black, 2014). This kind of evolutionary history will help us understand diversification patterns, ways of adaptation, and biogeography of Genus Paral ichthys. The study of molecular phylogenetic will also help understand the relationship among different species of Genus Paral ichthys. These studies have shown that there are three important clades of Genus Paral ichthys. It is considered that Paral ichthys californium and Paral ichthys Woolman are sister species because they share sharing common ancestor(Olsson et al., 2019). If we know about the relationship of species of Genus Paral ichthys, we may have important taxonomic implications. For example, it will be useful for reevaluation of species boundaries. As we know in taxonomic classification, we need to differentiate one species from another species based on important characteristics. The other important implication of understanding the relationship among various species of Paral ichthys is that we may also be well known to cryptic species of this Genus Paral ichthys(Campbell, López, Satoh, Chen, & Miya, 2014). As we have studied molecular phylogenetic, so we got an idea about the divergence times of Genus Paral ichthys as well. These studies have shown that Paral ichthys evolved almost 30 to 40 million years ago during the time of the Eocene Oligocene transition. But this Genus is vastly diverged about 15 million years ago(Tan, Wang, Li, & Sha, 2024). These studies are more effective than studying fossils because there is a limited record of fossils. In recent years, there has been the use of molecular clock rates for estimating the divergence times of various Genera. However those proved to be uncertain, so molecular phylogenetic is the best method for determining the divergence times of Paral ichthys(Lavoué, Miya, Poulsen, Møller, & Nishida, 2008). Now we are going to discuss the morphological characteristics of Paral ichthys. These fishes are called flatfish groups as well because of their unique morphology. These fishes have a flat but oval-shaped body and the caudal fins have been seen as rounded and thick. It has been seen that some species belonging to Genus Paral ichthys may reach lengths up to 80 cm(Lavoué, Miya, Saitoh, Ishiguro, & Nishida, 2007). These fishes have all four types of fins named as caudal fin, anal fin, pectoral fin, and dorsal fin. If we talk about the texture of the skin of these fishes, they have mostly smooth texture but sometimes texture is rough on the blind side. Most of these fishes may show coloration but some species are well known for their dark stripes or blotches. These fishes have small heads, small mouths with curved jaws, and small but pointed teeth. These are gill rakers that are specialized for various purposes(Byrne, 2018). By studying morphological studies of Paral ichthys, we came to know important ecological adaptations of Genus Paral ichthys. One of these important ecological adaptations is behavioral adaptation. These fishes can show camouflage which helps them to hide from predators and also helps them to catch prey. The other important behavioral adaptation is that these fishes can show social behavior for example these fishes may communicate with each other with the help of their body language. In some cases, it has been seen that these fishes may aggregate in groups for communication. The other important ecological adaptation is physiological adaptation. One of the important physiological adaptations is osmoregulation(Munroe, 2014).

These species are fully advanced to the main concentration of water and salt in the body by different means. Recent studies have shown that these fishes are also able to tolerate a wide range of temperatures as well. Some other important Adaptations are related to the reproductive system of Paral ichthys. For example, most of the species of this Genus show spawning behavior in which there is the release of more than one gamete at a time. So there are more chances for fertilization(Shin, Jin, Sohn, & Lee, 2018). The other reproductive adaptation is that they form larvae which have suitable buoyancy and transparency for better survival in aquatic environments. These adaptations are also related to feeding habits in Paral ichthys. These fishes may show ambush predation in which they can catch prey by hiding themselves or by camouflage. These fishes may also use Sense of smell to find prey around them. These fishes mostly live near substrates which help them in finding and capturing prey so it is considered the most important habitat adaptation in Paral ichthys(Roje, 2014). Some species, they are having such structures that they use for dwelling soil so we can say that these fishes help in ecosystem engineering in the aquatic environment. There are important future perspectives of Paralichthys. One of these important future perspectives is that there is a need for an integrative taxonomy for Paral ichthys which can be proven effective and advanced these days. Secondly, there is still a need to study more about the ecological roles, social behaviors, and other behavioral patterns of these fishes. Thirdly there is a stringent need for the management of invasive species which are dangerous for the Paral ichthys population (Bañón et al., 2005).



Figure 1: Species Identification and Classification

Applications of Phylogenetic and Taxonomic Studies of *Paral ichthys* (Paralichthyid)

Species Identification and Classification

Phylogenetic and/or taxonomic analyses are used to classify Paral ichthys species in terms of scientific classification. During biodiversity and environmental evaluations, research teams use morphological examination and genetic testing to properly identify species (Figure 1). Understanding the biodiversity within Paralichthyid family is aided somewhat by knowing the species as well as classifying them as Paral ichthys (flounders). The distinction between different species is made on the basis of physical structure in relation with its molecular and genetics patterned.

However, according to conventional methods, the species of the Paral ichthys group can be categorized into a new species due to morphological body features comprising of body shape, fin structures and color patterns, as well as specific eye asymmetry patterns. However, mere physical characteristics of the species are not sufficient as several species and specific population groups differ by small morphological difference. Molecular methods such as taxonomic are used to solve such problems today, including 1) DNA barcoding (more detailed taxonomic work) with mitochondrial gene sequencing and genomic analysis. Analysis of cytochrome c oxidase I (COI) and ribosomal RNA genes enable researchers to make confident species distinctions for those with otherwise similar appearance. Phylogenetic tree construction using molecular data enables the definition of clear species taxonomic limits while providing evidence for ancestral relationship of species. The differences between cryptic species can still be found through their molecular markers, even though the different species can be indistinguishable from one another morphologically. There are two important reasons for accurate species classifications. They further the scientific learning and the set their protection measures to the unique genetic groups. Proper species classification and sustainable harvesting practices become more attainable, as well ecological studies become clearer on the different species' functions on marine ecosystem function.

Conservation and Management

Severe taxonomic research is required for the identification of separate populations and species of Paral ichthys for efficient conservation work. Species delineation and the variance in the genetic diversity levels provide information needed to establish conservation strategies for those endangered species while preserving these species' ecological balance. It is of critical importance for fisheries management to have accurate species identification for sustainable fishing that protects certain fish populations from overexploitation. Both phylogenetic and taxonomic studies as the primary basis for areas of conservation and management of the Paral ichthys species rely on taxonomic information together with a good understanding of species diversity, which are shaped by species diversity comprehension. The research provides critical information about flounder species' hereditary diversity and population networks and habitat sequences important for the development of effective conservation measures. This research leads to the researcher's recognition of isolated species populations which would allow them to detect endangered populations faced with overfishing and habitat destruction. Knowing the genetic diversity within as well as between populations reveals vulnerable genetic groups that are at risk of decrease in adaptative potential and genetic bottlenecks and inbreeding, to allow researchers to concentrate more on saving ecosystem health and biodiversity. The main instrument for sustainable fisheries management is species identification, and requires, for this reason, to be accurate. Accurate taxonomic data is required to establish best regulatory restrictions through regulatory bodies to protect Paral ichthys species from depletion during a fishery use. The only practical way to increase genetic biodiversity in Paral ichthys species is through breeding programs selecting fish for better disease tolerance and better growth abilities; knowledge of Paral ichthys species genetics provides a foundation for such programs. By conducting biological functions of Paral ichthys species in marine ecosystems, they become an essential resource for preserving the seafloor and estuary. Phylogenetic and taxonomic research is a basic tool for protecting Paral ichthys species populations as well as the marine ecosystem environments.

Understanding Evolutionary Relationships

Phylogenetic relationship studies help add genealogical pattern and evolutionary process of diversification and adaptive transformation for Paral ichthys species. Scientists are able to study these species and learn what evolutionary forces have produced the modern genus or 'family' of Paral ichthys, as well as environmental conditions and specialized habitats. Good knowledge regarding Paral ichthys species relationships is critical to phylogenetic studies to understand how this genus has evolved through time by adapting and diversifying. Because of its asymmetrical body form, Paral ichthys has exceptional conditions for investigating evolutionary processes associated with speciation and its habitat adaptation and eye realignment mechanisms. Conservation of mitochondrial and nuclear genes in a species allow scientists to reconstruct historically how the Paral ichthys genus evolved and to determine genetic differences and what resulted in the modern species distribution. Tests conducted on the temporal and patterns of speciation events in species confirm an evolutionary pathway of Paral ichthys species in order to respond rapidly to oceanic changes and density level changes from ecological niche gradual development. Adaptation of different species of the genus Paral ichthys to changes of depth related and underwater temperature ranges and methods of feeding are used for studies of key evolutionary features of the genus Paral ichthys and ways of their survival. The phylogenetic trees are analyzed in order to identify common ancestors and determine the correct position of this genus in the Paralichthyid family tree. Through the evolutionary viewpoint, the present-day genetic characteristics of Paral ichthys species are better understood as a result of genetic drift and gene flow as well as natural selection processes. An understanding of evolutionary relationships aids in understanding the adaptive radiation processes in Paral ichthys, as well as in understanding the biodiversity of a marine ecosystem.

Figure 2: Fisheries and Aquaculture

Fisheries and Aquaculture

For achieving sustainable fisheries operations and aquaculture management correct taxonomic identifications must be used (Figure 2). The understanding of Paral ichthys species diversity enables responsible managers to build management plans that promote sustainable fish population harvesting. The application of species genetic knowledge produces more effective breeding operation results by increasing productivity and strengthening disease resistance in aquaculture. The sustainable management and production of flounders can be achieved with proper flounder identification methods and information about Paral ichthys evolutionary patterns in fisheries and aquaculture. Several species identification methods together with phylogenetic studies permit scientists to differentiate all species so appropriate fishing restrictions can be established separately for maintaining fishing outputs alongside wild species conservation. After obtaining genetic and population data of Paral ichthys species collections fisheries managers can create more effective conservation plans to protect endangered species. The genetic structure of Paral ichthys species maintains conditions for selective breeding programs to enhance growth speed together with disease immunity while also optimizing performance under farming conditions. Genetic information provides protection to farm-raised populations by preventing genetic population decline as well as inbreeding problems. Advanced aquaculture systems become possible through ecological preference understanding because this information enables better production environments through enhanced water management and dietary plans. Failure to understand phylogenetics makes aquaculture operations unable to determine appropriate species fits for optimal production locations. Taxonomic discoveries result in increased sustainability of both Paral ichthys wild-caught fishes and aquaculture operations due to their ability to promote enduring success of these flounders in their respective environments. These scientific approaches enable biodiversity maintenance through their support for both natural and farmbased Paral ichthys population health.

Ecological and Environmental Studies

Research conducted on taxonomy helps scientists understand the functional position of Paral ichthys species in marine environments. Studies about species distribution and ecological interaction patterns and habitat selection patterns support both environmental monitoring and marine conservation policy development. These fish adapt uniquely to benthic life which makes them excellent biological indicators to study the relationship between ecosystems and environmental transition. Studies about flounders as Paral ichthys species contribute essential insights to ecological and environmental research which helps understand their marine ecosystem function and behaviors with other organisms. The environmental health indicators Paral ichthys provides stem from their benthic lifestyle and their specialized feeding mechanisms as well as their asymmetrical body

structure and unique physiological and behavioral traits. Phylogenetic and taxonomic research identifies the habitats where Paral ichthys species reside while showing their adaptation to such areas as sandy and muddy seafloors and continental shelves and estuaries. The tracking of population trends by researchers depends on biological patterns knowledge for scientific monitoring because it measures impacts from habitat degradation and climate change and pollution on species populations. Calculating changes in population sizes of Paral ichthys species across their habitats serves as an efficient method for measuring ocean biodiversity alterations resulting from temperature and salinity or sediment changes that stem from climate variations. The scientific study of taxonomy reveals Paral ichthys ecological relationships through two key observations: first by examining how the fish feed on benthic invertebrates and second by analyzing their relationship with broader species as prey for marine mammals and large fish species. Phylogenetic data allows ecological research teams to identify which Paral ichthys species manage benthic ecological nutrient cycles and transfer energy between marine habitats. Better protected areas result from effective marine protected area management which utilizes knowledge about species' ecological preferences and protects the species' vital habitats. Studying both phylogenetic and taxonomic characteristics of Paral ichthys species allows scientists to protect marine ecosystems while promoting sustainable management practices during present environmental challenges. Phenetic alongside systematic research functions as a vital methodology to protect the genetic value of Paral ichthys species. Scientific research of population genetics helps detect genetic weakness and adaptable regions that support species conservation throughout environmental challenges. Phylogenetic analysis and taxonomy require present-day methods for protecting genetic resources of the Paral ichthys species because these research fields provide long-term guidance for their sustainable development. The findings of phylogenetic genetic analyses assist scientists to detect multiple genetic groups and endangered genetic populations stemming from habitat destruction and fishing activities together with climate change events. Species survival requires genetic diversity because population adjustments to environmental changes become easier through this species variety and this diversity protects health while offering disease resistance. Research in taxonomy shows the existence of genetically separated groups which poses additional population protection requirements alongside gene flow improvements throughout endangered groups. Preservation of genetic resources in aquaculture applications prevents population inbreeding risks while preserving culturing stock health. Multiple genetic types within Paral ichthys aquaculture populations protect industry from diseases and result in better productivity and stock health. Through joining genetic marker analysis with genomic study researchers can identify distinct populations containing traits for resistance against diseases and improved rapid growth possibilities for future breeding projects. Marine ecosystems require the conservation of genetic resources because different population losses create progressive negative consequences on species to ecosystem interactions. The Paral ichthys species research for genetic diversity protection uses phylogenetic and taxonomic methods to conserve biodiversity and maintain ecosystem health which delivers access to genetic resources for future populations. Studies focusing on Paral ichthys at the molecular level demonstrate promising opportunities for biomedical research along with biotechnology despite current research being scant. The distinct biological features together with evolutionary adaptations of flatfish provide the opportunity to discover developmental processes and repair methods because of their unique eye migration pattern. These fish species provide genomic data to facilitate biotechnological programs for disease protections alongside adaptation technology development. Study of Paral ichthys species taxonomy brings important value for the advancement of drug discovery and genetic research at the same time as biotechnology industry development. Scientists can find therapeutic compounds in biochemical substances through their knowledge of Paral ichthys species genomic sequences and their evolutionary characteristics. The ability of Paral ichthys to survive environmental stress along with specific metabolic processes allows scientists to

develop biomedical compounds that might change medical practices. The discovered compounds derived from Paral ichthys species studies provide the groundwork for developing new pharmaceutical agents and treatment methods that primarily benefit the fields of cancer therapy with neurodegenerative diseases and metabolic disorder control. The immune response mechanisms of Paral ichthys species represent a valuable resource for medical vaccine production as well as anti-infection therapeutic advancement. Phylogenetic genetic studies help researchers develop molecular markers and genetically modified organisms (GMOs) of aquaculture value while also enabling them to assess environments and conduct bioremediation. The study of Paral ichthys species and their closely related marine fish allows scientists to track genes responsible for growth rate and disease resistance and stress tolerance traits because these traits hold vital importance for aquaculture and biotechnological sector developments. The phylogenetic evaluation of Paral ichthys species bridges evolutionary science with functioning methodologies that promote drug development and biotechnology and sustainable practices and genetic research. Studies of Paral ichthys phylogeny and taxonomy yield practical benefits across conservation needs together with sustainable fishery management and sustainable ecological investigation and biotechnological development opportunities.

Conclusion

We have overviewed different studies that were related to the phylogeny and taxonomy of Paral ichthys. We have also overviewed those important studies that are related to evolutionary Insights from molecular and morphological data of Paral ichthys. The taxonomic and phylogenetic investigations of Paral ichthys generated numerous breakthroughs about its evolutionary background along with species richness together with adaptive characteristics that evolved within the Paralichthyid family. Research combining morphological traits with genetic sequencing data has become essential for solving difficult phylogenetic relationships while confirming species divisions among Paral ichthys species. Enhanced understanding of the genus helped authorities acquire better expertise for protecting Paral ichthys species among their vital ecological roles and economic value. Vietnam uses a thermal barrier zone to prevent the spread of large-scale fires. Researchers intensely study measurements of taxonomic dimensions and evolutionary patterns of Paral ichthys genus because Paralichthyid features this family. Scientific authorities classify members of the genus as flounders reveals fundamental morphological details because their symmetry appears initially but asymmetry develops through eye movement to one body side during their development process. This transformation, alongside other traits like body shape and fin structures, plays a crucial role in the identification and classification of various species within the genus. Scientific classification depends on multiple data sources which include morphological observations together with DNA sequencing techniques and molecular analytical methods.

References

- Atta, C. J., Yuan, H., Li, C., Arcila, D., Betancur-R, R., Hughes, L. C., . . . Tornabene, L. (2022). Exon-capture data and locus screening provide new insights into the phylogeny of flatfishes (Pleuronectoidei). *Molecular Phylogenetics and Evolution*, 166, 107315.
- Bañón, R., Gómez Pardo, B., Machordom, A., Foresti, F., Porto-Foresti, F., Azevedo, M. F., . . . Martínez, P. (2005). Phylogenetic analysis of flatfish (Order Pleuronectiformes) based on mitochondrial 16s rDNA sequences.
- Black, C. R. (2014). Geometric morphometric analysis of skeletal shape variation across the pleuronectiformes.
- Bruns, N. (2012). Morphological and genetic similarity among three species of halibut (Paralichthys spp.) in Baja California, Mexico.
- Byrne, L. (2018). A phylogenetic assessment of flatfish (Order Pleuronectiformes) intrarelationships based on molecular evidence. Université d'Ottawa/University of Ottawa,
- Campbell, M. A., López, J. A., Satoh, T. P., Chen, W.-J., & Miya, M. (2014). Mitochondrial genomic investigation of flatfish monophyly. *Gene*, 551(2), 176-182.

- Chen, W.-J., Bonillo, C., & Lecointre, G. (2003). Repeatability of clades as a criterion of reliability: a case study for molecular phylogeny of Acanthomorpha (Teleostei) with larger number of taxa. *Molecular Phylogenetics and Evolution*, *26*(2), 262-288.
- Cooper, J. A. (1996). *Monophyly and intrarelationships of the family Pleuronectidae (Pleuronectiformes), with a revised classification*: University of Ottawa (Canada).
- Kartavtsev, Y. P., Park, T.-J., Vinnikov, K. A., Ivankov, V., Sharina, S., & Lee, J.-S. (2007). Cytochrome b (Cyt-b) gene sequence analysis in six flatfish species (Teleostei, Pleuronectidae), with phylogenetic and taxonomic insights. *Marine Biology*, *152*, 757-773.
- Kartavtsev, Y. P., Sharina, S. N., Saitoh, K., Imoto, J. M., Hanzawa, N., & Redin, A. D. (2016). Phylogenetic relationships of Russian far eastern flatfish (Pleuronectiformes, Pleuronectidae) based on two mitochondrial gene sequences, Co-1 and Cyt-b, with inferences in order phylogeny using complete mitogenome data. *Mitochondrial DNA Part A*, 27(1), 667-678.
- Lavoué, S., Miya, M., Poulsen, J. Y., Møller, P. R., & Nishida, M. (2008). Monophyly, phylogenetic position and interfamilial relationships of the Alepocephaliformes (Teleostei) based on whole mitogenome sequences. *Molecular Phylogenetics and Evolution*, 47(3), 1111-1121.
- Lavoué, S., Miya, M., Saitoh, K., Ishiguro, N. B., & Nishida, M. (2007). Phylogenetic relationships among anchovies, sardines, herrings and their relatives (Clupeiformes), inferred from whole mitogenome sequences. *Molecular Phylogenetics and Evolution*, 43(3), 1096-1105.
- Munroe, T. A. (2014). Systematic diversity of the Pleuronectiformes. Flatfishes: biology and exploitation, 13-51.
- Olsson, D., Marquez, A., Tellechea, J. S., Carvalho, P. H., Pereira, A. N., & Norbis, W. (2019). Genetic and morphometric analyzes of Paralichthys species confirm the presence of P. brasiliensis in the Uruguayan waters. *Neotropical Biodiversity*, 5(1), 30-35.
- Redin, A. D., & Kartavtsev, Y. P. (2022). The Mitogenome Structure of Righteye Flounders (Pleuronectidae): Molecular Phylogeny and Systematics of the Family in East Asia. *Diversity*, 14(10), 805.
- Roje, D. M. (2010). Incorporating molecular phylogenetics with larval morphology while mitigating the effects of substitution saturation on phylogeny estimation: A new hypothesis of relationships for the flatfish family Pleuronectidae (Percomorpha: Pleuronectiformes). *Molecular Phylogenetics and Evolution*, 56(2), 586-600.
- Roje, D. M. (2014). *Molecules, morphology and monophyly: resolving pleuronectiform phylogeny and investigating why it has been so difficult to do.*
- Shin, S. P., Jin, C. N., Sohn, H. C., & Lee, J. (2018). Parvicapsula curvatura n. sp. in cultured olive flounder Paralichthys olivaceus and phylogenetic characteristics of the genus Parvicapsula. *Diseases of Aquatic Organisms*, 130(3), 199-207.
- Tan, S., Wang, W., Li, J., & Sha, Z. (2024). Comprehensive analysis of 111 Pleuronectiformes mitochondrial genomes: insights into structure, conservation, variation and evolution. *bioRxiv*, 2024.2009. 2022.614327.
- Timi, J. T., Paoletti, M., Cimmaruta, R., Lanfranchi, A. L., Alarcos, A. J., Garbin, L., . . . Mattiucci, S. (2014). Molecular identification, morphological characterization and new insights into the ecology of larval Pseudoterranova cattani in fishes from the Argentine coast with its differentiation from the Antarctic species, P. decipiens sp. E (Nematoda: Anisakidae). *Veterinary Parasitology*, 199(1-2), 59-72.
- Villarreal, F., Burguener, G. F., Sosa, E. J., Stocchi, N., Somoza, G. M., Turjanski, A. G., . . . Mechaly, A. S. (2024). Genome sequencing and analysis of black flounder (Paralichthys orbignyanus) reveals new insights into Pleuronectiformes genomic size and structure. *BMC genomics, 25*(1), 297.
- Wirtz, P., Davenport, J., & Chanet, B. (2023). Further investigations on fincrawling in flatfishes (Teleostei: Pleuronectiformes): phylogenetic implications. *Cahiers de Biologie Marine*, 64, 219-227.
- Yagishita, N., Miya, M., Yamanoue, Y., Shirai, S. M., Nakayama, K., Suzuki, N., . . . Nakabo, T. (2009). Mitogenomic evaluation of the unique facial nerve pattern as a phylogenetic marker within the percifom fishes (Teleostei: Percomorpha). *Molecular Phylogenetics and Evolution*, 53(1), 258-266.
- Young, H. (2023). Evaluating the efficacy of a marine reserve in enhancing the population growth and size of endangered fish species. *FishTaxa*, 27, 25-35.