

Informing the spatial management of Silky Shark (*Carcharhinus falciformis*) in the eastern Pacific Ocean



Brendan Talwar, Brice Semmens, Alexandre Aires-Da-Silva, Jenn Humberstone, Melanie Hutchinson, Jon Lopez, Carolina Minte-Vera, Dan Ovando, Salvador Siu, Lyall Bellquist

It is helpful to understand spatial population structure.



Figure 1. Conceptual model of north Pacific albacore spatial structure and movement patterns.

Minte-Vera², 2023

Conceptual models lead to better assessments.



Figure 1. Conceptual model of north Pacific albacore spatial structure and movement patterns.

Minte-Vera³, 2023



Spatial population structure is largely dictated by the movements of adults.

Chapman et al., 2015; Frisk et al., 2014

How much genetic and demographic mixing occurs across what spatial scales?

Fishery-dependent data

Local observations

Life history information

Population genetics

Tagging data

Spatially explicit 'classic stock delineation'

Units delineated according to spatial boundaries

[tagging, genetics, life history]

Spatially implicit 'Areas-as-fleets'

Units delineated according to selectivity [gear type, area of operation]

SILKY SHARK MANAGEMENT





Biomass has declined & fishing mortality has increased

SILKY SHARK CARCHARHINUS FALCIFORMIS

Leads shark bycatch in many shelf-edge and open-ocean fisheries

(Bonfil, 2008; Oliver et al., 2015)

'Coastal-pelagic', 'Semi-oceanic', 'Pelagic' Juvenile areas along the shelf edge, Adults more offshore

Neritic species (fine-scale population structure)

Silky Shark

Oceanic species (minimal population structure)

(Springer 1967; Branstetter, 1987, 1990; Bonfil, 2008)

Objective

Characterize Silky Shark spatial population structure in the (E)PO (i.e., explore stock structure)

Movement data {synthesis}
Population genetic data {review}
Spatial variation in life history parameters {new}





(Francis et al., 2023; Hutchinson et al., 2019, 2021; Kato & Carcallo, 1967; Ketchum et al., 2020; Lara-Lizardi et al., 2020; Musyl et al., 2011; Schaefer et al., 2019, 2021)



(Francis et al., 2023; Hutchinson et al., 2019, 2021; Kato & Carcallo, 1967; Ketchum et al., 2020; Lara-Lizardi et al., 2020; Musyl et al., 2011; Schaefer et al., 2019, 2021)



(Galvan-Tirado et al., 2013; Clarke et al., 2015; Kraft et al., 2018)



(Galvan-Tirado et al., 2013; Clarke et al., 2015; Kraft et al., 2018)





(Castillo-Olguin 2005; Rodriguez-Matus 2020)

Within-EPO: spatial variation in life history?

3,590 males with catch locations, measurements, and clasper calcification status





Clasper calcifisperiora (cification) (bifforaldb) mg (Totaldeng (bpatial group)



Explored 9 candidate spatial structures in estimating male TL₅₀

Highest support for the 'nearest genetic unit' spatial structure



Mexico TL₅₀ 156 cm TL

El Salvador TL₅₀ 148 cm TL

Ecuador TL₅₀ 172 cm TL



Does mating and pupping occur in each?

Mating habitats can be suggested by sex ratios.

Balanced sex ratios (1 male: 1 female) = potential mating habitat

Males and females mix in space – mating habitats widespread?



Longitude

Pups are all over – pupping areas widespread



Latitude

Observed catches

- Small-scale landings
- Longline
- Purse seine: Dolphin
- Purse seine: Unassociated tuna
- Purse seine: Floating object

Including offshore!

Ontogenetic shift from shelf-edge to oceanic habitats is not universal: Silky Sharks probably also pup offshore.





Spatially explicit 'classic stock delineation'

Units delineated according to spatial boundaries [tagging, genetics, life history]

Spatially implicit 'Areas-as-fleets'

Units delineated according to selectivity [gear type, area of operation]

(Waterhouse et al., 2014; Hurtado-Gerro et al., 2014; Punt 2019)







Small animals dominate

Large animals dominate

California Current Large Silkies

Summer: expand into higher latitudes Winter: condense towards equator

> Humboldt Current Large Silkies

Expect demographic mixing between genetic stocks in the boreal winter and austral winter Interested in developing similar conceptual models for other vulnerable sharks (e.g., hammerheads)



<u>Kraft et al 2018 – Pacific wide</u> *entire mitochondrial genome and thousands of nuclear loci

Pairwise comparisons between sites: nuclear DNA $F_{st} = 0.019 - 0.042$ mitochondrial DNA $F_{st} = 0.012 - 0.057$

<u>Rodriguez-Matus 2020 – EPO</u> *microsatellite data (nuclear)

Pairwise comparisons between sites: F_{st} range: 0.005 (within-Mexico comparisons) to 0.035 (central Mexico to El Salvador)



Flexed total length (cm)



















Longitude

