



Content of the document for data description

1. Methodology

1.1 Sampling methodology, frequency, and size

We collected the chub mackerel samples since 2016. In 2017-2019, more than 2000 individuals were measured in the lab. The size range of fork length was 129~388 mm (Table 1).

Since in the December, 2017, we began to collect the samples for their otolith identification of the chub mackerel population in the CA areas (Table 2). The random samples were collected by the fishermen in two different fishing boats every month, then taken to the lab for further analysis. First, we measured the fork length (FK, mm), total length (mm) and wet weight (g) of each fish. Second, we took out a pair of sagittal otoliths and ground the right otolith. Finally, the age of every otolith was identified by otolith ring method under the microscope.

Table1 Data, frequency, and size on the samples of the chub mackerel in the CA areas

Data	Frequency	Size range (FK, mm) of fishes
August to November, 2016	3 months	129~378
May to November, 2017	9 months, every month	130~345
April to October, 2018	7 months, every month	178~388
April to November, 2019	8 months, every month	154~358

Table2 Data, frequency, and size on the samples for otolith identification of the chub mackerel in the CA areas

Data	Frequency	Size range (FK, mm) of fishes
December, 2017	1 month	178~244
April to October, 2018	7 months, every month	195~367
April to November, 2019	8 months, every month	154~339

1.2 Methodology of ALK-development

The forward age-length key (ALK) was first developed by Fridriksson (1934). The method works on the premise that given a random sample of N fish for which only lengths have been measured and a subsample of n fish whose lengths and ages have been measured, the probability $P(i|j)$ that a fish is age i given that it belongs to length bin j is the same for both samples. This probability can be estimated from the age-length sample as:

$$\hat{P}(i|j) = \hat{q}_{ij} = n_{ij}/n_j \quad (1)$$

where \hat{q}_{ij} is the estimated probabilities of age given length that populate the cells of the forward ALK. n_{ij} is number of fish of age i and length bin j in the age-length sample, n_j is total number of fish belonging to the j th length bin of the age-length sample.

The probabilities of age given length from the forward ALK are then simply multiplied by the marginal probabilities $\hat{P}(j) = y_j/N$ to obtain an estimate of age composition from the forward key, \hat{A} . This can be expressed using matrix algebra as follows:

$$\hat{A} = QY/N \quad (2)$$

where Q is the I by J matrix with elements \hat{q}_{ij} . Equation (2) can be shown to give maximum likelihood estimates; it is presented in the form above to emphasize the logic of the approach.

1.3 Criteria if multiple ALKs are used in different regions

The forward age-length key:

(1) The age-length and the length frequency samples must originate from the same statistical population, i.e. within a length class, the underlying age composition must be the same for the two samples. In other words, the two samples must be drawn from the same available population. This implies that:

(a) A forward key developed from one year cannot be applied to another year.

(b) A forward key developed from one area cannot be applied to another area if the two areas are characterized by differences in age composition.

(2) A forward key developed from one gear can be used to age catch from a different gear even if the two gears have different size selectivities, so long as the two selectivity curves within a length bin are parallel. With narrow length bins, selectivity is almost constant, hence, the requirement of parallel selectivity curves is met.

The inverse age-length key:

(1) The number of length bins (J) must be greater than or equal to the number of age classes (I) in order to obtain a unique solution (in some cases, a plus group will need to be implemented).

(2) The age-length and the length frequency sample do not need to have been collected in the same year. They can be collected from two populations with different age compositions as long as size at age does not differ between the two populations.

(3) The Hoenig and Heisey (1987) method is the superior method for applying inverse keys when there is a single length frequency and a single age-length sample as it allows for uncertainty in both the length frequency sample and the age-length sample.

The combined forward-inverse age-length (FIAL) key:

(1) The number of length bins (J) must be greater than or equal to the number of age classes (I) in order to obtain a unique solution.

(2) Size-at-age is assumed constant among samples.

(3) The estimator is valid even if length stratification is used.

1.4 Methodology of estimating catch-at-age from ALK

$$N_{i,j} = \frac{TC_j \times P_{i,j}}{BW_i} \quad (3)$$

where i is the i th class interval of KnL; j is the j th cell of year; P is the proportion of samples' weight; TC is the catch; \widehat{BW} is the fitted body weight by Length-weight relationships (Froese, 1998), the formula is:

$$BW_1 = a \times KnL^b \quad (4)$$

We have obtained the relationship between age and length through ALK, therefore, therefore,

the relationship between age and weight can be obtained.

$$BW_2 = f(age) \quad (5)$$

So, the yearly catch-at-age can be estimated by

$$N_{i,j} = \frac{TC_j \times P_{i,j}}{BW_2} \quad (6)$$

2 Results to be shown

2.1 Sample sizes of length measurements and age determination

We have collected 2310 individuals of chub mackerel to measure the biological parameter from 2016 to 2019. Every month, 2~3 samples including 50~100 individuals were collected during the fishing boats and then be transferred to lab to further measured (Table 3).

Table3 Sample sizes of length measurements

Year	month	samples	individuals
2016	August to November	8	254
2017	May to November	15	842
2018	April to October	14	345
2019	April to November	16	869

Table4 Sample sizes of age determination

Year	month	samples	individuals
2017	December	2	40
2018	April to October	14	260
2019	April to November	16	469

2.2 Length and age distribution

Using of catch samples from fishing boats, the biological parameters of chub mackerel were measured. In 2017 the mean fork length on the chub mackerel was 227.3mm, and then increased to 266.1mm in 2018 and 239.4 mm in 2019. On the whole, it found that the mean fork length became a slight increasing trend of body length in 2016-2019.

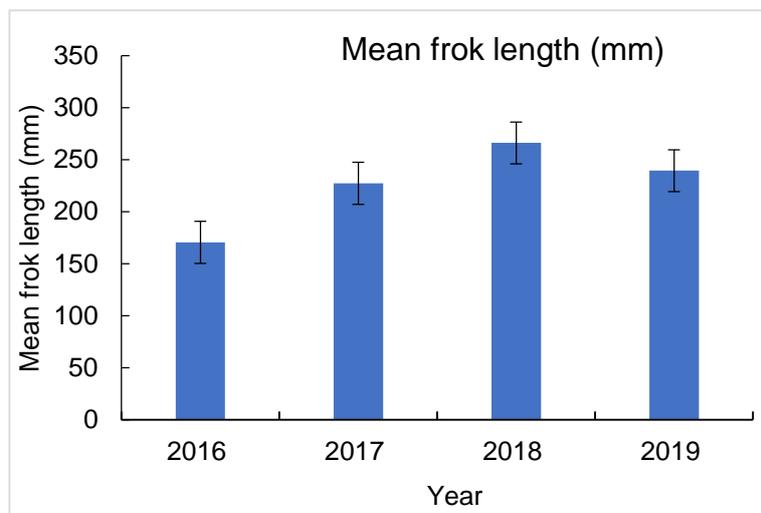


Fig.1 the mean fork length on the chub mackerel in CA areas in 2016-2019

We analyzed that the distribution of different fork length groups and ages distribution in 2018 and 2019. In the 0+ age group, mainly fork length range was less than 200mm, which mean that more than 98% of the individuals less than 200 mm were 0+-year-old fish. The individuals which fork length ranged from 200-260mm were mainly 1+ year old. The individuals which fork length ranged from 240-320mm were mainly 2+ year old. The individuals which fork length ranged from 280-360mm were mainly 3+ year old. In 2019, it was found that the individuals which fork length ranged from 300-360mm were also distributed 4+ year old.

Table5 Fork length and age distribution of chub mackerel in the North Pacific Ocean in 2018

Fork length (mm)	Age group (percentage, %)				
	0+	1+	2+	3+	4+
160-180	66.7				
180-200	33.3	15.9			
200-220		38.1			
220-240		28.6			
240-260		15.9	7.1		
260-280		1.6	23.8		
280-300			16.7		
300-320			42.9	22.2	
320-340			9.5	50	
340-360				22.2	
360-380				5.6	

Table6 Fork length and age distribution of chub mackerel in the North Pacific Ocean in 2019

Fork length (mm)	Age group (percentage, %)				
	0+	1+	2+	3+	4+
160-180	66.7				
180-200	28.8				
200-220	3.0	11.2			
220-240	1.5	61.2	20.0		
240-260		27.8	60.0		
260-280			0		
280-300			20.0	11.1	
300-320				55.6	24.5
320-340				33.3	50.0
340-360					25.5
360-380					

2.3 ALK (if used)

Next SC meeting we will discuss the ALK method to estimate catch-at-age on the chub mackerel.

2.4 Length-Weight relationship

Body size is a basic biological characteristic in fish populations and can reflect individual physiology as well as changing environment conditions. Length-Weight relationship is an important characteristic to describe this change. For figures 2-4, The R^2 of length-weight of *Scomber japonicus* were larger than 0.94. The represent chub mackerel has a significant relationship between length and weight. In accordance with the samples in 2019, the relationship between with fork length and biomass of chub mackerel was $y=0.00000405 x^{3.36}$ ($R^2 = 0.9583$), where y is the weight (g) of chub mackerel individual, x is the fork length (mm)

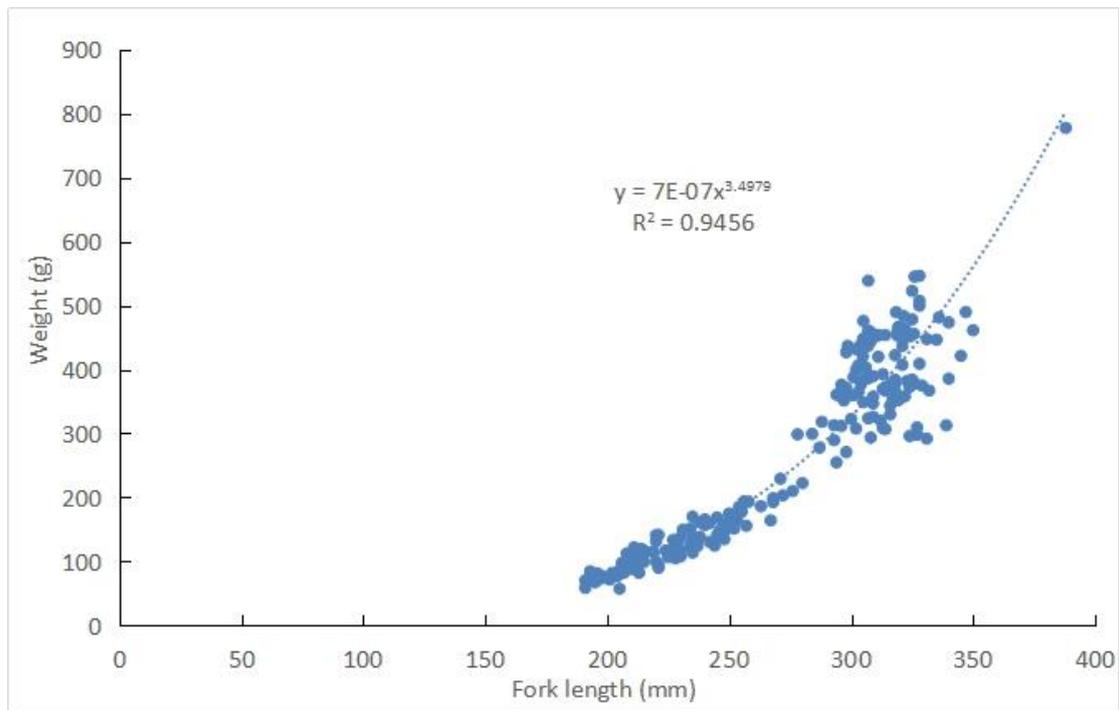


Figure2 Length-Weight relationship of China_chub mackerel in 2018 (n=345)

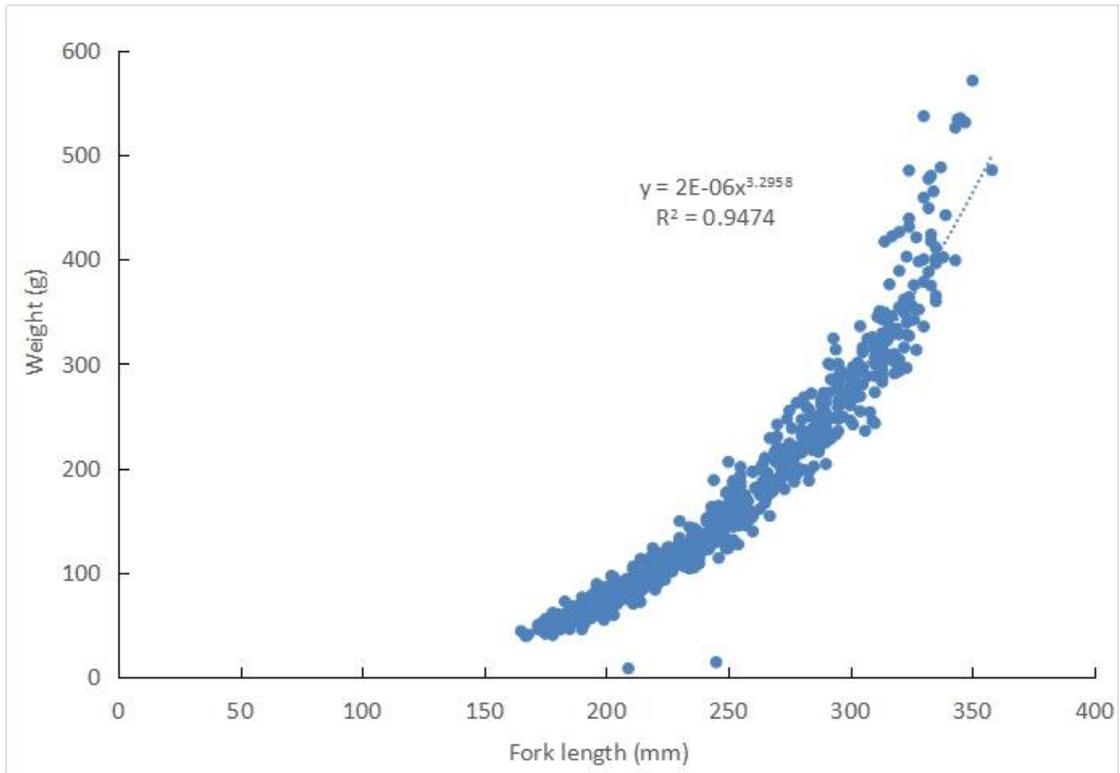


Figure3 Length-Weight relationship of China_chub mackerel in 2019 (n=869)

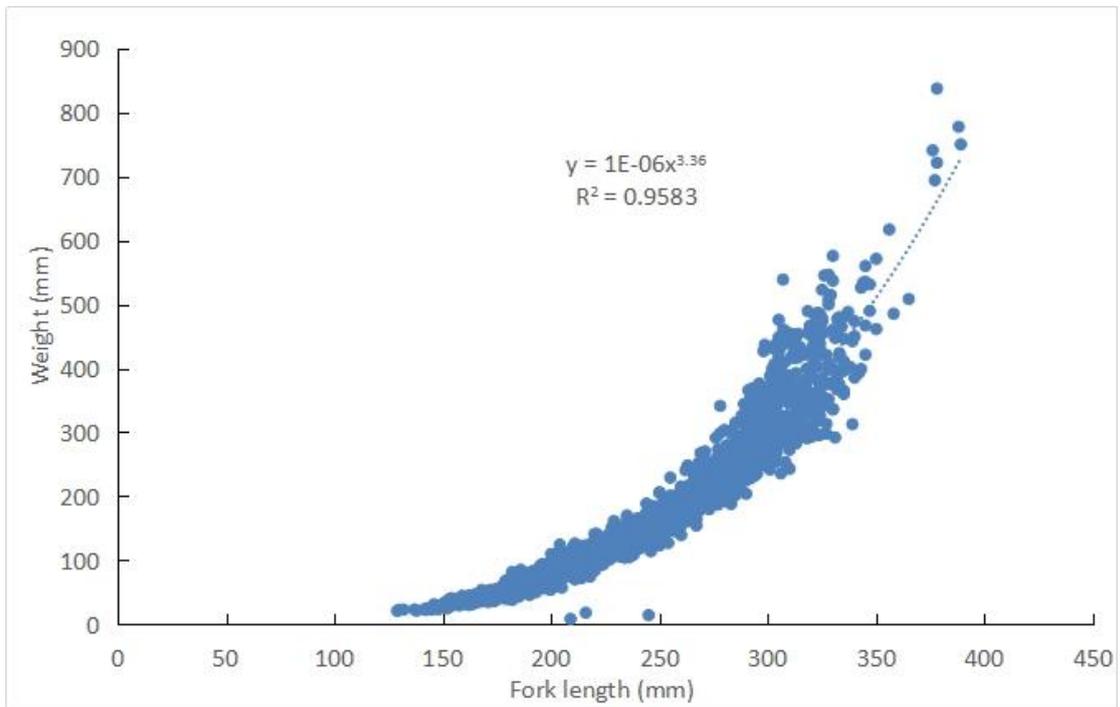


Figure4 Length-Weight relationship of China_chub mackerel in 2016-2019 (n=2213)

2.5 Catch-at-age

Using otolith age identification technology, we analyzed the Catch-at-age composition in chub mackerel catches (Fig.3). The catch-at-age 0⁺, 3⁺ and 4⁺ in 2019 both exceed those in 2018. In these two years, the catch-at-age 5⁺ to 7⁺ both have not caught in the CA areas.

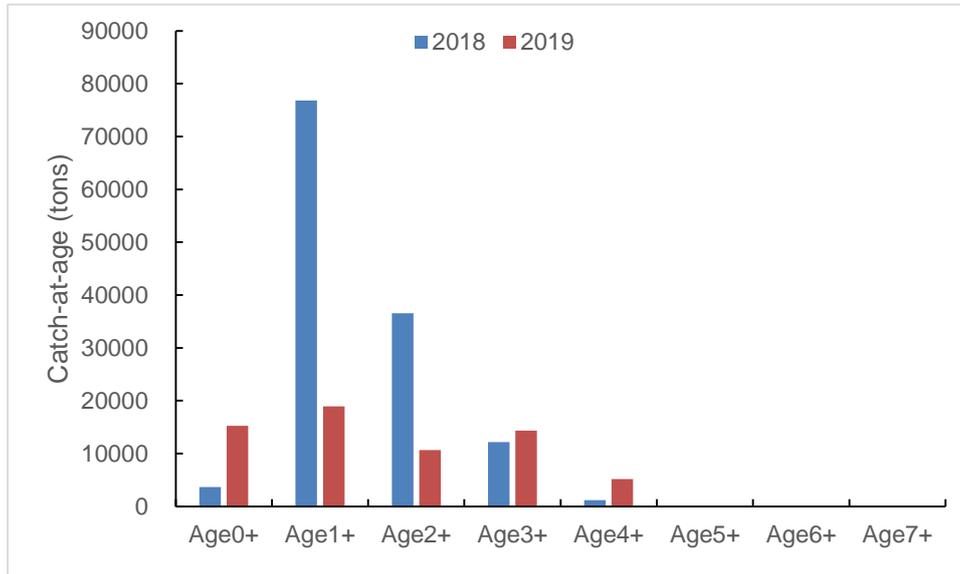


Fig.5 Catch-at-age on the chub mackerel in the CA areas in 2018-2019