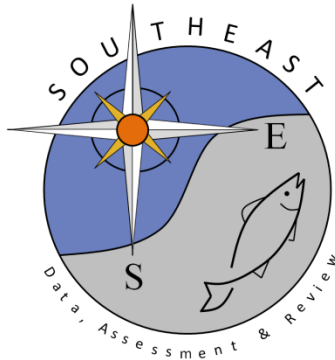


Updated Estimates of Batch Fecundity vs. Total Length, Total Weight, and  
Calendar Age for South Atlantic Red Snapper in Support of the SEDAR 73  
Operational Assessment

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SEDAR 73-WP07

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SEDAR73-WP07  
MARMAP/SEAMAP-SA Reef Fish Survey Technical Report 2020-11

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## **Background**

Red Snapper, *Lutjanus campechanus*, is a large, long-lived, member of the family Lutjanidae, with a maximum reported age of 51 yr (SEDAR41, 2017). Red Snappers are distributed in marine waters throughout the Gulf of Mexico south to the Yucatan Peninsula and in United States (U.S.) Atlantic waters north to North Carolina (Nelson and Manooch, 1982; Manooch and Potts, 1997). Along the southeastern U.S., adult Red Snapper are associated with structured habitats such as coral reefs, wrecks, gas and oil platforms, rocky outcroppings, and live-bottom (Moseley, 1966; Nelson and Manooch, 1982; Barans and Henry, 1984; Sedberry and Van Dolah, 1984). Red Snapper are gonochorists with indeterminate fecundity (Woods, 2003; Brulé et al., 2010) that spawn during April through October in the Atlantic waters of the southeastern U.S. (White and Palmer, 2004; Lowerre-Barbieri et al. 2015).

The objective of this report is to present updated batch fecundity (BF) equations for the SEDAR73 Operational Assessment. Additional specimens (n=28) were processed by staff of the SouthEast Reef Fish Survey and the data were added to the dataset analyzed for SEDAR41. The dataset from SEDAR41 included data from Lowerre-Barbieri et al. (2015).

## **Methods**

### *Survey Design and Gear*

(see Smart et al. 2015 for full description)

### Sampling area

- Cape Hatteras, NC, to St. Lucie Inlet, FL

### Sampling season

- May through September
  - Limited earlier and later sampling in some years

### Survey Design

- Simple random sample survey design
  - Annually, randomly selected stations from a chevron video trap universe of confirmed live-bottom and/or hard-bottom habitat stations
  - No two stations are randomly selected that are closer than 200 m from each other
    - Minimum distance is typically closer to 400 m
- Video traps deployed on suspected live-bottom and/or hard-bottom in a given year (reconnaissance) are evaluated based on catch and/or video or photographic evidence of bottom type for inclusion in the universe in subsequent years
  - If added to the known habitat universe, data from the reconnaissance deployment is included in index development

### Sampling Gear – Chevron Video Traps (video camera(s) added in 2010)

(see Collins 1990 and MARMAP 2009 for more detailed descriptions)

- Arrowhead shaped, with a total interior volume of 0.91 m<sup>3</sup>
- Constructed of 35 x 35 mm square mesh plastic-coated wire with a single entrance funnel (“horse neck”)
- Baited with a combination of whole or cut clupeids (*Brevoortia* or *Alosa* spp., family Clupeidae), most often *Brevoortia* spp.

- o Four whole clupeids on each of four stringers suspended within the video trap
- o Approximately 8 clupeids placed loose in the video trap
- Soak time of approximately 90 minutes
- Daylight hours

#### Oceanographic Data

- Hydrographic data collected via CTD during soaking of a “set” (typically 6 video traps, but may be less) of chevron video traps deployed at the same time and same reef patch
  - o Bottom temperature (°C) is defined as the temperature of the deepest recording within 5 m of the bottom

#### *Data Filtering/Inclusion*

##### Projects coordinated by MARMAP/SERFS

- o P05/T59/T60 – MARMAP/SEAMAP-SA/SEFIS
- o P50 – Port Sampling (Fishery-Dependent)

##### Gear

- o 043 - Snapper reel
- o 061 - SBLL
- o 324 - Chevron trap

#### *Specimen processing*

Fish were kept on ice until they were processed on the vessel or in the laboratory, generally within 24 h of capture. Maximum total length ( $\pm 1$  mm), total weight ( $\pm 1$  g), and gonad weight ( $\pm 1$  g;  $\pm 0.1$  g for smaller gonads) were measured and usually both sagittal otoliths were removed and stored dry. A sample of gonad tissue was removed and fixed for histological processing and an additional sample of ovarian tissue was fixed to estimate batch fecundity if there was macroscopic evidence of oocyte hydration but no ovulation.

#### *Fecundity*

Histological samples were processed and examined to assess reproductive phase and search for evidence that ovulation had begun (i.e., postovulatory follicle complexes (POC)). All specimens with new POCs were not used for the estimation of batch fecundity. Batch fecundity was estimated gravimetrically by MARMAP/SERFS and Lowerre-Barbieri et al. (2015) using the hydrated oocyte method (see Hunter et al., 1985; Murua et al., 2003). The protocols for processing Red Snapper ovarian tissue were similar, except for the methods of fixing tissue and separating oocytes. Lowerre-Barbieri et al. (2015) hydraulically separated the oocytes prior to fixation in 2% neutrally-buffered formalin and then weighed subsamples, whereas MARMAP/SERFS fixed the tissue in 10% seawater-buffered formalin, weighed subsamples, transferred subsamples to 5% seawater-buffered formalin, and then separated oocytes. MARMAP/SERFS investigators obtained 2 or 3 subsamples per specimen that weighed 75-175 mg, versus 2 subsamples weighing 100 mg in Lowerre-Barbieri et al. (2015).

The count of hydrated oocytes in each subsample was extrapolated to gonad weight to compute batch fecundity. For MARMAP/SERFS subsamples, it was necessary to develop a regression equation to

convert fresh gonad weight to preserved (i.e., fixed) gonad weight because only a portion of most large gonads was preserved to reduce the amount of formalin utilized. For Red Snapper, the equation to estimate preserved weight was: preserved wt (g) = fresh wt (g) \* 0.886218 – 1.61585, with the range of fresh gonad wt = 4 to 275 g, adj.  $r^2=0.99$ ,  $n=15$ .

The relationship between batch fecundity vs. maximum total length, total weight, and calendar age was estimated using simple linear and non-linear (power function;  $BF = bX^2$ ) regression analyses. In the most recent Vermilion Snapper assessment, the power function was recommended (Buble and Wyanski, 2017) because fecundity tends to exhibit a non-linear relationship with fish length due to body cavity volume increasing in a non-linear manner with fish length (Wooten, 1979). Two power function models (with and without intercept) were utilized, the latter to relax forcing the intercept through the origin. Simple linear regression analyses were performed with EXCEL software and non-linear analyses were performed with Statistical Analysis System (SAS) software, vers. 9.4 (SAS Institute Inc, 1989).

### *Ageing*

Otoliths were embedded and sectioned following standard protocols and assigned increment counts and edge codes by two readers independently (Smart et al., 2015). Calendar ages were determined from consensus ages via the rule that if the edge code was 3 or 4 and month of capture was January through July, then calendar age was increment count + 1. For all other edge codes and months, the calendar age was increment count.

### **Results**

A total of 97 Red Snapper were collected during 1999-2019 to assess batch fecundity, including 28 specimens collected by MARMAP/SERFS during 2007-2019 that were added to the dataset ( $n=69$ ) analyzed for SEDAR41. The dataset from SEDAR41 was composed of data from Lowerre-Barbieri et al. (2015;  $n=44$ ) and 25 specimens processed by MARMAP/SERFS. The specimens processed by Lowerre-Barbieri et al. (2015) were collected with hooked gear (longlines, rod and reel) during April through August in 2012 off the upper half of Florida's east coast, whereas the 53 specimens from MARMAP/SERFS were collected throughout the sampling area (28.3 to 34.9 °N), nearly all (94%) with chevron traps.

The 97 specimens ranged in length from 360 to 958 mm TL and in calendar age from 2 to 16 yr. Although the fits of the linear regression equations were good (Table 1; Adj.  $R^2 = 0.50-0.62$ ), the power function model was applied to the data because batch fecundity exhibited a non-linear relationship with total length (TL) and total weight (TW), and to a lesser degree with calendar age (Fig. 1). The power function with no intercept parameter (2-parameter) tended to overestimate batch fecundity at smaller sizes (length and weight) due to the model forcing the intercept through the origin (Fig. 2); therefore the 3-parameter model is recommended equations for predicting batch fecundity versus TL and TW (Table 2). The drawback to the 3-parameter model is that the predicted values are negative at  $TL < 378$  mm and  $TW < 728$  g (Tables 3 and 4). For smaller fish, it is recommended that the mean observed batch fecundity be used at sizes  $< 400$  mm TL (= 55523 eggs) or  $< 1000$  g TW (=61808 eggs). The 2-parameter power function is the recommended equation for predicting batch fecundity versus calendar age (Tables 2 and 5) because the 3-parameter function underestimated batch fecundity at younger ages (Age 2) and potentially at the oldest ages (Fig. 3)

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Table 1. Simple linear regression equations ( $BF = a + bX$ ) for Red Snapper batch fecundity (BF) versus total length (TL), Total weight (g), and calendar age. Red Snapper ( $n=97$ ) were collected in 1999-2019 primarily during fishery-independent sampling off the Atlantic coast of the southeastern United States by Florida's Fish & Wildlife Research Institute and the SouthEast Reef Fish Survey at S. Carolina Department of Natural Resources.

X	Range of X	a	SE <sub>a</sub>	b	SE <sub>b</sub>	n	Adj. R <sup>2</sup>
TL (mm)	360-958	-1833719	210224	4479.011	355.3476	97	0.62
Total wt (g)	630-11,830	-180964	84244.44	257.0033	18.73116	97	0.66
Cal. Age (yr)	2-16	2.934978	0.253587	2.14E-06	2.21E-07	94	0.50

Table 2. Recommended power function equations relating batch fecundity (BF) to total length (TL), total weight, and calendar age for Red Snapper off the Atlantic coast of the southeastern United States.

Independent Variable (X)	Range	a	SE <sub>a</sub>	b	SE <sub>b</sub>	Z	SE <sub>Z</sub>	n	Model	Notes
TL (mm)	360-958	-271137	246741	0.0235	0.1126	2.7404	0.7030	97	BF = a+(b*X <sup>Z</sup> )	Due to negative predicted value at TL < 378, let BF for females < 400 mm TL = mean obs. BF for TL < 400 (= 55523).
Total Weight (g)	630-11,830	-198789	211224	317.0	704.7	0.9775	0.2380	97	BF = a+(b*X <sup>Z</sup> )	Due to negative predicted value at WW < 728, let BF for females < 1000 g WW = mean obs. BF for WW < 1000 (= 61808).
Calendar Age (yr)	2-16	n/a	n/a	137936.0	37942.3	1.1382	0.1301	94	BF = b*X <sup>Z</sup>	

Table 3. Predicted batch fecundity versus total length (TL; mm) for Red Snapper off the Atlantic coast of the southeastern United States using the equation in Table 2.

TL (mm)	Batch Fecundity (# of eggs)		
360	-33256	710	1258739
370	-14708	720	1318514
380	4734	730	1379752
390	25088	740	1442467
400	46370	750	1506675
410	68598	760	1572390
420	91791	770	1639627
430	115964	780	1708401
440	141137	790	1778727
450	167324	800	1850620
460	194545	810	1924094
470	222815	820	1999163
480	252152	830	2075843
490	282572	840	2154148
500	314091	850	2234092
510	346727	860	2315690
520	380497	870	2398956
530	415415	880	2483905
540	451499	890	2570550
550	488765	900	2658906
560	527230	910	2748988
570	566908	920	2840809
580	607817	930	2934383
590	649971	940	3029725
600	693388	950	3126849
610	738083	960	3225769
620	784071		
630	831368		
640	879991		
650	929953		
660	981272		
670	1033961		
680	1088038		
690	1143516		
700	1200411		

Table 4. Predicted batch fecundity versus total weight (g) for Red Snapper off the Atlantic coast of the southeastern United States using the equation in Table 2.

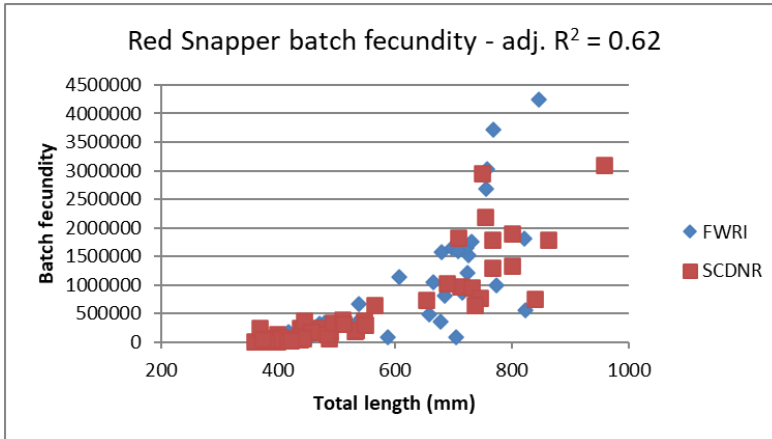
<b>Total Weight (g)</b>	<b>Batch Fecundity (# of eggs)</b>
500	-60972
1000	72579
1500	204567
2000	335549
2500	465788
3000	595439
3500	724602
4000	853349
4500	981734
5000	1109798
5500	1237573
6000	1365086
6500	1492361
7000	1619415
7500	1746265
8000	1872925
8500	1999406
9000	2125720
9500	2251877
10000	2377884
10500	2503749
11000	2629479
11500	2755081
12000	2880560

Table 5. Predicted batch fecundity versus calendar age (yr) for Red Snapper off the Atlantic coast of the southeastern United States using the equation in Table 2.

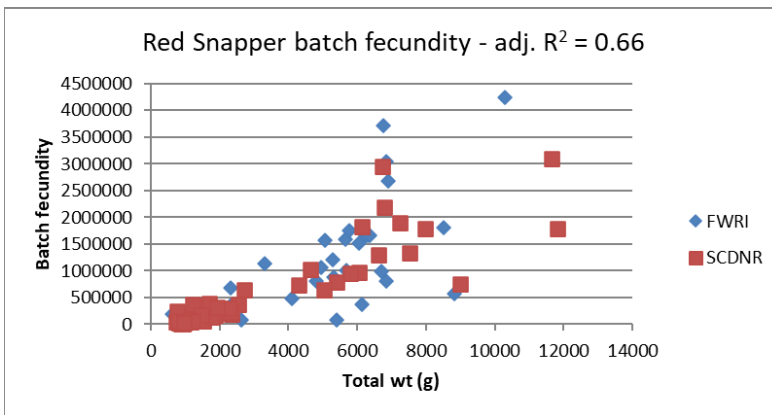
<b>Calendar Age (yr)</b>	<b>Batch Fecundity (# of eggs)</b>
2	303606
3	481656
4	668255
5	861480
6	1060155
7	1263480
8	1470871
9	1681886
10	1896172
11	2113444
12	2333467
13	2556042
14	2780997
15	3008186
16	3237479

Figure 1. Simple linear regression of batch fecundity versus A) total length, B) total weight, and C) calendar age in Red Snapper (n=97) collected in 1999-2019 primarily during fishery-independent sampling off the Atlantic coast of the southeastern United States by Florida's Fish & Wildlife Research Institute (FWRI) and the SouthEast Reef Fish Survey at S. Carolina Dept. of Natural Resources (SCDNR).

A)



B)



C)

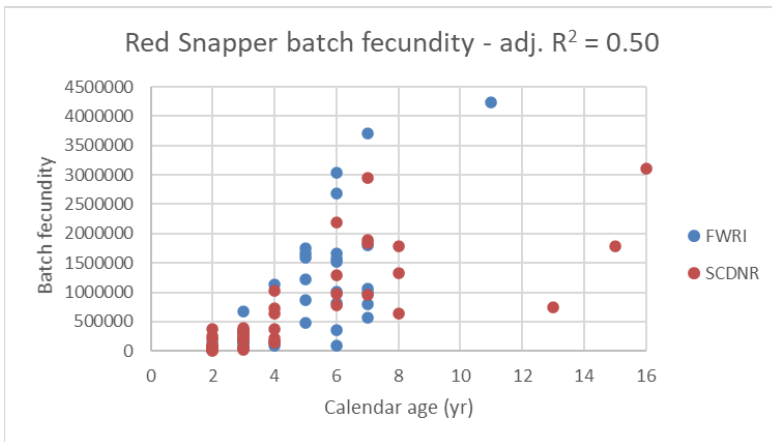
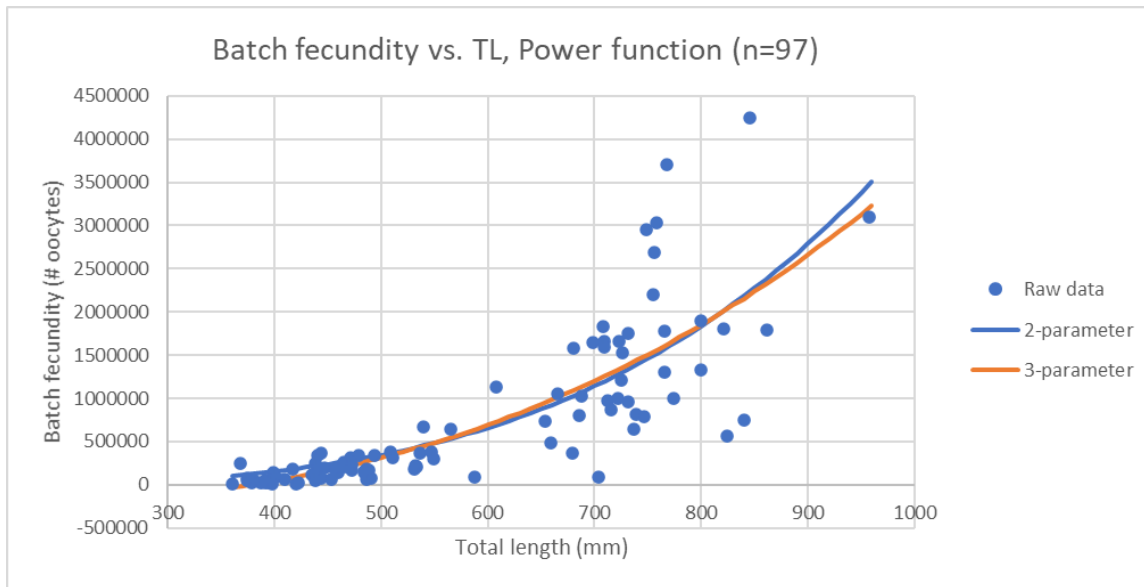


Figure 2. Non-linear regression analysis of batch fecundity (BF) versus A) total length and B) total weight in Red Snapper using the power function ( $BF = bX^2$ ). The 3-parameter model included an estimate of the intercept ( $BF = a + bX^2$ ). The Red Snapper (n=97) were collected in 1999-2019 primarily during fishery-independent sampling off the Atlantic coast of the southeastern United States by Florida's Fish & Wildlife Research Institute (FWRI) and the SouthEast Reef Fish Survey at S. Carolina Department of Natural Resources (SCDNR).

A)



B)

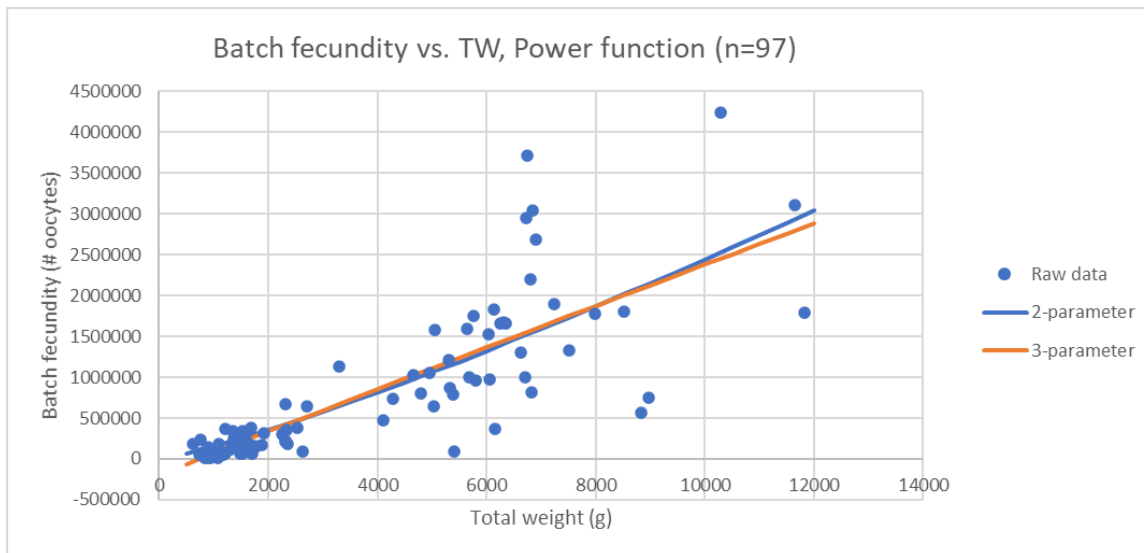


Figure 3. Non-linear regression analysis of batch fecundity (BF) versus calendar age in Red Snapper using the power function ( $BF = bX^2$ ). The 3-parameter model included an estimate of the intercept ( $BF = a + bX^2$ ). The Red Snapper (n=94) were collected in 1999-2019 primarily during fishery-independent sampling off the Atlantic coast of the southeastern United States by Florida's Fish & Wildlife Research Institute (FWRI) and the SouthEast Reef Fish Survey at S. Carolina Department of Natural Resources (SCDNR).

