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Life cycle of Chokka squid Loligo reynaudii in South African waters Lipinski, Marek; van der Vyver, Frikkie; Shaw, Paul W; Sauer, W. H. H.

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1	LIFE CYCLE OF CHOKKA SQUID, LOLIGO REYNAUDII
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11	
12	Abstract
13	
14	This short note summarizes past and present knowledge about life cycle of chokka squid
15	(Loligo reynaudii D'Orb., 1848). In the past (until about 2010) chokka stock was considered
16	simple and uniform, with one paralarval pool, drift of palararvae westwards, one main
17	nursery area and one main long spawning migration of adult squid eastwards, back to the
18	main spawning grounds. New findings revised this life cycle. Although genetically the stock
19	is uniform, but morphologically it comprises three main geographic groups. It is proposed
20	that their differences originate from many different palaralval "events", and that short (not
21	long) migrations dominate the life cycle.
22	
23	Keywords: life cycle, Loligo reynaudii, squid.
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25	

26 Introduction

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28 The life cycle of the chokka squid (Loligo reynaudii D'Orb., 1848) has been debated in a 29 number of publications (e.g. Augustyn 1989; Augustyn et al. 1992, 1994; Oylott et al. 2006, 30 2007; Sauer et al. 2013; van der Vyver et al. 2015). The first three accounts came up with a 31 simple biological scheme. This scheme, set out in Figure 1 and Figure 2, was based primarily 32 on biological analyses during research conducted on both commercial vessels and the 33 Fisheries Research Vessel "Africana" from 1983 onwards. 34 In the distribution terms, main spawning areas were detected and mapped (e.g. Sauer et al. 35 1992) inshore between Plettenberg Bay and Port Alfred. Other spawning grounds, although 36 detected and mentioned in various publications, were considered minor. Paralarvae were 37 mainly detected around spawning grounds, but were present along the whole south coast 38 (Augustyn et al. 1994). Paralarvae detected along the west coast were identified (Vecchione 39 & Lipinski 1995) as Afrololigo mercatoris (Adam, 1941). 40 Juveniles of 20-80 mm ML were mainly detected between Plettenberg Bay and Cape 41 Agulhas, although they were present year round along the whole south coast. Adult squid 42 which were usually detected offshore hunting in small schools all over Agulhas Bank, were 43 thought to return in their bulk to main spawning grounds, and the whole cycle will repeat 44 itself (Augustyn et al. 1994). 45 Figure 2 fills some details of the scheme illustrated on Figure 1. According to this scheme, 46 most of the squid spawns in the east, all paralarvae drift to the common paralarval pool (from 47 which some hypothetically drift to the west coast and grow, but most of these get lost). One 48 stock of squid, recruited from this paralarval pool, feeds and grows on the Agulhas Bank. 49 Some part of it spawns locally inshore (short migration), some migrate to the west coast 50 waters, but most return eastward to spawn.

52	More thorough analysis of the existing data (summarized in Augustyn et al. 1994) and then		
53	additional analyses of old and new data (e.g. Olyott et al. 2006, 2007) supplemented this		
54	established view. Most important points of departure were as follows:		
55	1. There is unquantified spawning of <i>Loligo reynaudii</i> in deep waters (deeper than 70 m;		
56	Augustyn et al. 1994; Roberts & Sauer 1994);		
57	2. Juveniles 20-80 mm are much wider distributed along the south coast, highest		
58	densities of them are detected between Algoa Bay and Cape St. Francis, slightly		
59	offshore in relation to their spawning grounds (Augustyn et al. 1994);		
60	3. Migrations of adults between Tsitsikamma and Port Alfred indeed take place up to		
61	200 km, they mainly in west to east direction but are complicated and interpretation of		
62	emerging patterns is difficult. Each spawning concentration is very dynamic;		
63	exchange there may be 0.2 of its biomass per day, or more (Lipinski et al. 1998; Sauer		
64	et al. 2000).		
65	Results and Discussion		
66	New evidence		
67	More recently a combination of ecological, morphological, environmental and genetic		
68	research has questioned our understanding of the life cycle of chokka, calling for more		
69	complicated structure than first envisaged (Shaw et al. 2010, Sauer et al. 2013; van der Vyver		
70	et al. 2015), however, the published accounts of these findings stopped short of providing a		
71	new life cycle scheme of chokka squid, which is the aim of the present note.		
72			

73 The following new facts and interpretations were become available:

74	1.	Spawning in the deep was confirmed, mapped and quantified as having 18% share in
75		total spawning. Ecological experiments have proven the viability of this spawning as
76		producing healthy hatchlings (Oosthuizen & Roberts 2009; Roberts et al. 2012).
77	2.	Simulation experiments pointed out to complicated distribution of paralarvae and
78		possible substantial losses during their drift (Roberts & Mullon 2010).
79	3.	Scarcity, but constant presence of chokka between St. Helena Bay and Kunene River
80		was confirmed (Lipinski unpublished results of R/V Dr Fridtjof Nansen cruises).
81	4.	Separate but viable sub-population of chokka in the southern Angola (up to 500 km
82		from Kunene) is the object of some artisanal fisheries. Mature squid were noted there,
83		but nothing is known about egg beds and paralarvae (van der Vyver 2015; Sauer
84		unpublished results).
85	5.	As the result of genetic and morphometric studies it was found that there is little
86		genetic diversity even between most distant sub-populations (Angolan vs. Port
87		Alfred). However, morphometric diversity was significant between south coast of SA,
88		western Agulhas and west coast of SA, and Angola (van der Vyver 2015; Fig. 3). This
89		regional patterns of morphological divergence observed, occurred against a backdrop
90		of high gene flow, which was interpreted as the influence of environmental
91		heterogeneity and not genetic drift/isolation as the primary driver of the phenotypic
92		differences. The observed phenotypic heterogeneity probably reflects the interplay
93		between genetic adaptation and short term plasticity, which may vary throughout the
94		geographic range of the study, and be a start of more profound morphological
95		differences (e.g. in beaks or statoliths) and then stable genetic differences. The
96		existence of the three morphological domains (Eastern and Central Agulhas, Western
97		Agulhas and West Coast, and southern Angola) calls for further revision of the

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existing life cycle on a geographic and temporal background, especially when more biological data will became known about Angolan population.

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102 Life cycle as known today (2016) is presented on the Figures 3-4 as follows.

103 Loligo reynaudii forms mobile, large metapopulation. Most northern (Angolan) part of this 104 metapopulation is not genetically isolated from other, southerly components, but differs 105 morphologically. Since mixing with nearest abundant group (St. Helena – Western Agulhas) 106 is minimal due to scarcity of individuals over nearly 1800 km of coastline, this Angolan sub-107 population is likely to be a recent extension of the species range northwards, and has its own 108 breeding and paralarval transport regime. Morphological differences between west coast plus 109 Western Agulhas, and Eastern Agulhas plus Tsitsikamma - Port Alfred are maintained 110 throughout two different paralarval pools, which are further divided into the smaller groups. 111 Deep water spawning, on the other hand helps to maintain relative homogeneity of this part 112 of the meta-population (exclusively South African), as is migration in space and time (i.e. 113 subsequent generations in different areas, as changing environment will dictate). Migration of 114 adult squid is generally short (around 200 km) and may proceed in all directions, including 115 inshore – offshore (Sauer et al. 2000). Hypothetical long migration may exist on a small 116 scale, although it was never documented. There also may be some adult squid which does not 117 migrate at all, but again this has not been documented.

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This life cycle scheme differs sharply with the first simple proposal. The latter superficially agrees well with the genetic results of van Vyver et al. 2015, but not with morphological part of their study. Observed morphological differences have to be rooted in the early development (on a paralarval stage). This in turn may be related with timing of hatching

124 subsequent small movements of juveniles on their nursery grounds. It is hoped that proposed

scheme (Figs. 3-4) reflects well this biological reality.

126

127 Funding and Ethical Considerations

128

129 This note does not provide information about new data – all data were already collected

130 under various programs and acknowledged in publications cited in this note. This note is

about new idea concerning the life cycle of squid and the only cost is the time spent by

132 authors during writing it – no specific funding was obtained. This note does not contain any

133 studies with animals performed by any of the authors. There is no conflict of any interests

134 whatsoever.

135

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196 Figure Legends

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Fig. 1. The scheme of geographic distribution and movement of paralarvae and adults of
chokka squid (*Loligo reynaudii*), as understood in the early years of research.

200

201 Fig. 2. The life cycle of chokka squid, according to distributional scheme illustrated on Fig. 202 1. There are two clusters of spawning sites: main off the Eastern Cape, and accessory off the 203 Western Agulhas coast. There is only one paralarval pool, fed along similar routes throughout 204 the year but mainly in November-December. Thick arrows indicate main circulation in the 205 life cycle scheme; thinner arrows indicate supplementary processes. Broken lines indicate 206 paralarval movements. Size of rectangular boxes represent approximate strength of each 207 migration event. Lost paralarvae were, as many larvae of other species, a result of being 208 carried away from coast by the Agulhas Current and its offshoots. It is unclear if any 209 paralarvae reach the west coast (marked by question marks).

210

Fig.3. A revised representation of the geographic distribution of chokka:.(A) west coast of
southern Africa; (B) south coast of South Africa.

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Fig. 4. New life cycle scheme of chokka. There are three blocks of information: Angola, where very little data is available; Eastern Agulhas and Eastern Cape spawning grounds; and Central and Western Agulhas spawning grounds. Last two also include deep water spawning grounds. Main departure from the previous scheme is a partition of one large paralarval pool into separate paralarval "events" which are different in space, time, or both. Also, possible loss of paralarvae was documented from both Eastern Agulhas and Western Agulhas (Roberts and Mullon 2010). Thick arrows indicate main circulation in the life cycle scheme; thinner

- 221 arrows indicate supplementary processes. Broken lines indicate paralarval movements. Size
- 222 of rectangular boxes represent approximate strength of each migration event. It is unclear if
- any paralarvae reach the west coast (marked by question marks).











