

# An initial attempt at a sex-disaggregated assessment for the South African hake resource, fitting directly to age-length keys

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December 2008

# INTRODUCTION

Geromont *et al.* (1995) estimated a female proportion in the south coast longline catches of 83%. Furthermore, there are very clear sex-specific differences in somatic growth for both *M. paradoxus* and *M. capensis*, in fact more so than between species (see Fig. 1). Routine application of age-length keys to obtain catch-at-age proportions is conducted without attention to sex-specific differences, but sex-differential growth means that larger sized of males are not well represented in the catch. This could confound estimates based on catch-at-age data developed from a sex-aggregated age-length key, which might consequently underrepresent the number of older hake present (and therefore affect the estimates of natural mortality).

This document presents a first attempt at modelling the sexes separately, which requires fitting directly to age-length keys (ALKs) and length frequencies (e.g. Punt *et al.* 2006).

# THE MODEL

The model is sex-specific; however, apart from sex-specific age-length keys, all the data available are sex-aggregated. Some assumptions have therefore to be made.

As it is not possible to estimate a sex-specific selectivity-at-age, it is rather assumed that the selectivityat-length is the same for both males and females. The model therefore needs to keep track of the numbers in each length class. Rather than move to a length-based model, the option being pursued is to keep the model age-structured and convert numbers-at-age into numbers-at-length using a length-at-age distribution (assumed to remain constant over time).

The catches cannot be disaggregated by sex, and there seems no reason to suppose that there is any preferential targeting by sex; thus, we assume that the annual fishing mortality generated by each fleet is the same on both males and females.

Note: for ease of reading, the 'species' superscript has been omitted below.

The numbers-at-age are converted to numbers-at-length as follows:

$$N_{y,a,l}^{g} = N_{y,a}^{g} P_{a,l}^{g}$$
(1)

where  $P_{a,l}^{g}$  is the proportion of fish of age *a* and sex *g* that fall in the length group *l* (i.e.,  $\sum_{l} P_{a,l}^{g} = 1$  for all ages *a*). The matrix *A* is calculated under the assumption that length-at-age is normally distributed about a mean given by the von Bertalanffy equation, i.e.:

$$L_a \sim N \Big[ L_{\infty} \Big( 1 - e^{-\kappa (a - t_0)} \Big); \theta_a^2 \Big]$$
<sup>(2)</sup>

where  $\theta_a$  is the standard deviation of length-at-age *a*, which is modelled as a function of the expected length at age *a*, i.e.:

$$\theta_a = \beta \left[ L_{\infty} \left( 1 - e^{-\kappa (a - t_0)} \right) \right]^{\gamma}$$
(3)

Population dynamics:

$$N_{y+1,a+1,l}^{g} = \left(N_{y,a,l}^{g} e^{-M_{a}/2} - \sum_{f} C_{y,l}^{g,f}\right) e^{-M_{a}/2} \quad \text{for } 0 \le a \le m-2$$
(4)

$$N_{y+1,m,l}^{g} = \left(N_{y,m-1,l}^{g} e^{-M_{m-1}/2} - \sum_{f} C_{y,l}^{g,f}\right) e^{-M_{m-1}/2} + \left(N_{y,m,l}^{g} e^{-M_{m}/2} - \sum_{f} C_{y,l}^{g,f}\right) e^{-M_{m}/2}$$
(5)

where

 $N_{y,a,l}^{g}$  is the number of fish of sex g, age a and length l at the start of year y,

 $M_a$  denotes the natural mortality rate on fish of age *a* (assumed – for the moment - to be the same for males and females),

 $C_{y,l}^{g,f}$  is the estimated number of hake of sex g and length l caught in year y by fleet f, and

*m* is the maximum age considered (taken to be a plus-group).

A Beverton-Holt stock-recruitment relationship is assumed, with the recruitment ( $R_y^g$ ) dependent on the female component of the spawning biomass and assuming a 50:50 sex-split at recruitment.

$$R_{y}^{g} = 0.5 \frac{\alpha B_{y}^{sp, females}}{\beta + B_{y}^{sp, females}} e^{(\varsigma_{y} - \sigma_{R}^{2}/2)}$$
(6)

where

 $\alpha$  and  $\beta$  are spawning biomass-recruitment relationship parameters,

 $\varsigma_y$  reflects fluctuation about the expected recruitment for year y, which is assumed to be normally distributed with standard deviation  $\sigma_R$ ;

 $B_{y}^{sp,g}$  is the spawning biomass of sex g at the start of year y, computed as:

$$B_{y}^{sp,g} = \sum_{a} \sum_{l} f_{a} w_{l}^{g} N_{y,a,l}^{g}$$
(7)

where

 $w_l^g$  is the mass of fish of sex g and length l and  $f_a$  is the proportion of fish of age a that are mature.

Catch

$$C_{y,a,l}^{g,f} = N_{y,a,l}^{g} \ e^{-M_{a}/2} \ S_{y,l}^{f} \ F_{y}^{f}$$
(8)

where

 $S_{y,l}^{f}$  is the commercial selectivity,

 $F_y^f$  is the fished proportion of a fully selected length class, for fleet *f*, assumed to be the same for males and females.

Note: S and F are assumed to be independent of g.

The estimated sex-aggregated catches-at-length to be compared with observations of length frequencies are:

$$C_{y,l}^{f} = \sum_{g} \sum_{a} C_{y,a,l}^{g,f}$$

$$\tag{9}$$

The (known) annual catch-by-mass of fleet *f* is given by:

$$C_{y,l}^{f} = \sum_{g} \sum_{a} \sum_{l} w_{l}^{g} C_{y,a,l}^{g,f}$$
(10)

So that:

$$F_{y}^{f} = C_{y}^{f} / \left( \sum_{g} \sum_{a} \sum_{l} N_{y,a,l}^{g} e^{-M_{a}/2} S_{y,l}^{f} \right)$$
(11)

#### The likelihood function

1.

The model is fitted to CPUE and survey abundance indices, commercial and survey catch-at-length data, as well as to the stock-recruitment curve to estimate model parameters, as in the baseline assessment. The contributions by each of these to the negative log-likelihood are not repeated here.

For years for which ALKs are available, the baseline assessment is also fitted to commercial and survey catch-at-age data. Here however, the model is fitted to the data underlying the ALKs directly, so that catch-at-length are used throughout. The ALKs are the only data that are available in sex-disaggregated form. The contribution of the ALKs to the negative log-likelihood is as follows.

Under the assumption that fish are sampled randomly with respect to age within each length-class, the contribution to the negative log-likelihood for the ALK data (ignoring constants) is:

$$-\ln L^{ALK} = -\sum_{i} \sum_{l} \sum_{a} \left[ A^{obs}_{i,l,a} \ln(\rho_{i,l,a}) - A^{obs}_{i,l,a} \ln(A^{obs}_{i,l,a}) \right]$$
(12)

were

 $A_{i,a,l}^{obs}$  is the observed number of fish of age *a* that fall in the length class *l*, for ALK *i* (a specific

combination of survey (or commercial fleet), year, species and gender)

 $\boldsymbol{\rho}_{i,a,l}$  is the model estimate of  $A_{i,a,l}^{obs}$  , computed as:

$$\rho_{i,a,l} = W_{i,l} \frac{C_{i,l} A_{a,l}}{\sum_{a} C_{i,l} A_{a,l}}$$
(13)

where  $W_{i,l}$  is the number of fish in length class *l* that were aged for ALK *i*.

# DISCUSSION

With the addition of the length dimension, the model now runs extremely slowly and results have not yet been obtained.

### REFERENCES

- Geromont HF, Butterworth DS, Japp D and Leslie RW. 1995. Preliminary assessment of longline experiments: south coast hake. Unpublished document, Marine and Coastal Management, South Africa. WG/11/95/D:H:28. 12pp.
- Punt AE, Smith DC, Tuck GN and Methot RD. 2006. Including discard data in fisheries stock assessments: two case studies from south-eastern Australia. Fisheries Research 79: 239-250.

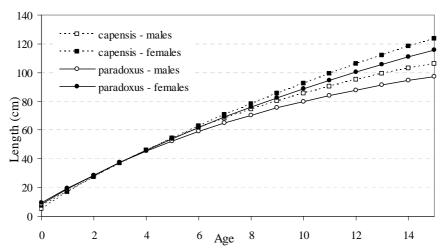


Fig. 1: Estimated mean length-at-age from the von Bertalanffi equation for males and females *M. capensis* and *M. paradoxus*