The cuttlefish Sepia officinalis L. (Mollusca: Cephalopoda) in the English Channel and the Bay of Biscay - is there a single panmictic population?

Katja Wolfram¹, Felix C. Mark^{1,2}, Hans O. Pörtner¹

¹Alfred Wegener Institute for Polar and Marine Research, Bremerhaven, Germany ²Zoology Dept, Cambridge University, United Kingdom



Introduction

Population substructure of the common European cuttlefish Sepia officinalis (Mollusca: Cephalopoda) has never been investigated in its northern distribution range from the Bay of Biscay through the English Channel into the North Sea.

For the southern distribution range, cuttlefish population substructure has been reported to be weak in the semi-enclosed Adriatic Sea [1] and extensive around the Iberian Peninsula [2] (Fig 1).

So far, the existence of separated Sepia officinalis populations has been postulated by numerous authors due to differences in reproductive behaviour [3, 4], migration behaviour [5, 6, 7] and in vitro oxygen binding properties of the respiratory pigment haemocyanin [8]. Yet, population structure has never been investigated on the genetic level, which was the aim of this study.



Microsatellite analysis for the first time indicates the existence of a panmictic Sepia officinalis population in the northern distribution

The level of genetic substructure of the English Channel, Bay of Biscay and North Sea cuttlefish (measured as FST) is comparable to the weak population subdivision reported for the panmictic Adriatic cuttlefish [1] (Figure 4). The genetic divergence between cuttlefish from the northern distribution range and specimens off Southern Portugal, however, is comparable to the genetic variability between Iberian cuttlefish on either side of the Strait of Gibraltar, which are thought to be genetically distinct subpopulations [2].

This was also confirmed by neighbour-joining trees (Figure 2) and principal component analysis (PCA, Figure 3): Neighbour-joining trees exhibited a much larger genetic distance (standard genetic distance of Nei 1972) between Southern Portugal cuttlefish and northern cuttlefish than between either of the northern sampling areas. PCA revealed a clear discrimination of individuals from South Portugal and



Material and Methods

To examine genetic variation among the sampling sites, lengths of seven DNA microsatellites were analysed for 220 individuals from the Bay of Biscay (n = 58), the English Channel (n = 63), the southern North Sea (n = 25) and Southern Portugal (n = 74). Statistical data analysis was carried out and genotypes at the seven loci were compared within the individuals from the northern distribution range and with those from Southern Portugal.



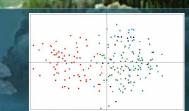
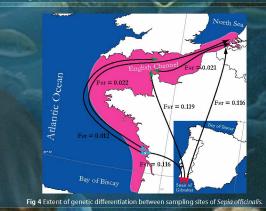


Fig 3 Principal component analysis of genotypes in 220 Sepia officinalis specimens from the North Sea (blac's), the English Channel (green), the Bay of Biscay (blue) and Southern Portugal (set).



Conclusion

Ability of organisms to adapt to climate also depends on the genetic variation or polymorphisms present amongst populations of a species.

The cuttlefish Sepia officinalis in the Bay of Biscay, the English For example, across a latitudinal gradient, populations with Channel and adjacent waters should be regarded as a appropriate mutations to survive higher temperatures may panmictic population displaying only minor subdivision. exist, leading to selective sweeps within the panmictic population.